

PROGRAM IN TRANSLATIONAL AND MOLECULAR MEDICINE DIMET

UNIVERSITY OF MILANO-BICOCCA SCHOOL OF MEDICINE AND SCHOOL OF SCIENCE

The multidrug-resistance transporter Abcc3 protects NK cells from chemotherapy in a murine model of malignant glioma

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Chapter 1

General introduction

Glioblastoma, the deadly tumor

Glioblastoma (GBM), a grade IV astrocytoma, is the highest-grade and the most common form of malignant brain tumor. It is one of the most lethal form of human cancer: without treatment, the median survival is approximately 3 months with a rapid development of clinical symptoms ¹. Clinically, patients with GBM may present with headaches, focal neurologic deficits, confusion, memory loss, personality changes or seizures depend on the location and size of tumor ².

Each year in western world 3-4 new cases of glioblastomas are diagnosed every 100,000 inhabitants. Although brain tumors account for <2% of all primary neoplasms, high mortality rates mean they rank high as leading cause of cancer-related death ³. Most patients die within one year from the diagnosis, and only 5% survive more than 5 years despite aggressive therapies (CBTRUS, 2011). Despite several progresses, the current standard treatment for GBM patients still consists of maximal surgical resection followed by radiotherapy plus concomitant and adjuvant temozolomide (TMZ) ^{4,5}. Many features help to explain the refractory nature of GBM to treatments including capacity for infiltration, and presence of the blood–brain barrier, a significant obstacle to chemotherapy. Recurrences are almost inevitable remaining an incurable disease with a survival range of 12–15 months ⁴.

GBM is usually described into primary or secondary form, indistinguishable histologically but differ in genetic and epigenetic profiles ⁶ (figure 1).

Primary GBM is the most common form (about 95%) and arises *de novo* from glial cells as a combination of genetic and epigenetic alterations. Typically has a clinical history < 6 months and is most common in older patients. It displays deletion in cell cycle-related genes p16^{INK4A} and p19/p14^{ARF}, mutations in the promoter of the telomerase reverse transcriptase (TERT), loss of chromosome 10 or PTEN mutation and gene amplification of the epidermal growth factor receptor (EGFR), usually occurring in conjunction with the variant III (EGFRvIII) activating mutation.

Secondary GBM, less frequent and studied, develops over months or years from preexisting low-grade astrocytomas and predominantly affects younger patients ⁷. It is characterized by mutations in TP53, amplifications of cyclin-dependent kinase 4 (CDK4) or loss of retinoblastoma (Rb) and overexpression of platelet-derived growth factor (PDGF). IDH1 mutations were found in less than 5% of primary GBM and in greater than 80% of secondary GBM, characteristically associated with improved survival.

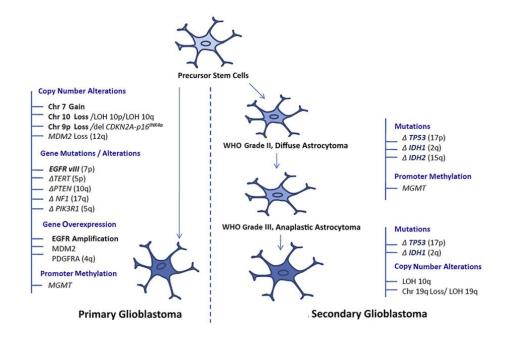


Figure 1. Primary and secondary GBM development. The most frequent and relevant molecular abnormalities of primary versus secondary GBM ⁸.

The recent characterization of the genome ^{9,10} and transcriptome ^{11–13} of GBM results in the generation of molecularly defined subtypes (i.e., proneural, neural, classical and mesenchymal). Molecular subclasses are typically associated with different genetic alterations and expression profiles that can better predict survival and treatment response. Thus, GBM subtypes suggest the possibility to appropriately stratify patients and to treat them with more innovative and personalized cares.

It is reported, for example, that the proneural (PN) signature can be considered the less aggressive form with a more favorable prognosis ¹². On the contrary, the mesenchymal (MES) subtype is considered the most aggressive profile carrying not only an angiogenic but also an

inflammatory signature that makes MES GBM more responsive to immune-based therapies ^{14,15}.

Due to the infiltrative nature of GBM, with diffuse dissemination of tumor cells beyond the tumor mass and general resistance to therapy, there have been intensified research efforts studying new targeted therapies.

Novel therapeutic approaches created expectations for increased survival, including anti-angiogenic therapy and immunotherapy, however heterogeneity favors escape strategies.

In the last years, research focused on targets as EGFR, vascular endothelium growth factor (VEGF) and platelet-derived growth factor (PDGF) receptors to inhibit mechanisms of proliferation and angiogenesis. Bevacizumab is a recombinant humanized monoclonal antibody that neutralizes the activity of VEGF and is normally used as treatment for recurrent GBM. Recently, results of a phase III study of bevacizumab added to standard treatments in newly diagnosed GBM showed a significantly increased in progression free survival ^{16,17}.

A growing body of evidence indicate encouraging results with immunotherapeutic approaches to fight GBM ^{18,19}. Immunotherapy with dendritic cells (DCs), loaded with whole tumor lysate or glioma-associated antigen peptides, revealed tolerable and safe with promising increased survival ^{20–22}. Mitchell et al. suggested in a recent clinical study that pre-conditioning the vaccine site with a potent recall antigen as tetanus toxoid can significantly improve the lymph node homing and the efficacy of DC immunotherapy ²³.

EGFRvIII is a constitutively activated and immunogenic mutation not expressed in normal tissues but widely expressed in GBM and for that the most important target for different immunotherapeutic approaches. A phase II peptide-based immunotherapy trial was conducted and demonstrated safety with evidence of immune response and increased patient survival ²⁴. A phase III randomized study of Rindopepimut/GM-CSF (ACT IV) in patients affected by newly diagnosed GBM is ongoing but the recruitment is concluded. The EGFRvIII epitope is also proposed as target for chimeric antigen receptor (CAR) T cell therapy in murine models demonstrating a robust antitumor activity ^{25,26}. Different clinical trials exploring the safety and effectiveness of CARs against HER2, IL-13α2 and EGFRvIII just started ²⁷.

Finally, immunotherapy with immune checkpoint inhibitors for cytotoxic T lymphocyte associated antigen 4 (CTLA-4) and programmed cell death 1 (PD-1) have provided substantial improvements in melanoma and lung cancer, and there is evidence of a potential benefit for GBM patients ²⁸. Tumor progression involves multiple immunosuppressive mechanisms, making combination of immunotherapeutic agents that target different pathways could be a promising approach.

Glioblastoma & Immune System

The central nervous system (CNS) has long been considered an immunological privileged site not affected by apparent immune responses. Initially, CNS immune privilege was supported by (i) restrictions imposed by the blood-brain barrier (BBB), structured to restrict the transport of molecules and cells from the circulation to the

parenchyma, (ii) the lack of draining lymphatics and (iii) the dearth of antigen-presenting cells (APCs).

Over the last 20 years, accumulating data re-evaluated this idea suggesting that the CSN is neither isolated nor passive in its interactions with the immune system ²⁹. Indeed, peripheral immune cells can cross the intact BBB and infiltrate the parenchyma to perform routine immune surveillance functions against infections and cancer ³⁰. BBB represents only a relative barrier to lymphocyte trafficking, especially in pathological states where it is disrupted, often in the case of GBM. Microglia are the resident immunocompetent cells with the role of APCs participating in the regulation of innate and adaptive immune responses. Moreover, under a variety of inflammatory conditions, peripheral DCs are quickly recruited to the site of the brain lesion ³¹. Although conventional lymphatics are missing in the CNS, antigenloaded DCs have access to secondary lymphoid organs, where T-cell priming takes place, through cerebrospinal fluid and perivascular spaces (figure 2). Furthermore, subcutaneous vaccination with genetically modified cytokine-secreting tumor cells or systemic immunotherapy with DCs induce an efficient immune response against intracranial tumors, confirming that the CNS is not truly immunologically privileged but a site for immunological responses and a candidate for immunotherapeutic approaches ^{32–34}.

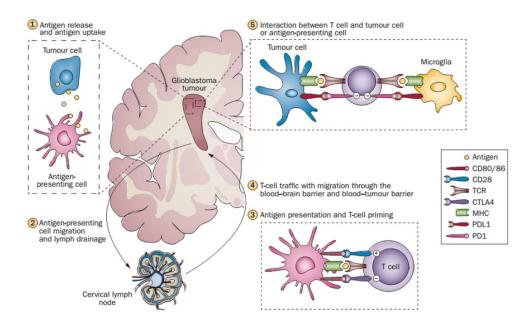


Figure 2. Immune response cycle of GBM. Antigens released from dying tumor cells are up-taken by APCs and presented to T cells that are then primed and activated in lymph nodes. The cytotoxic T cells migrate to tumor site where they infiltrate and specifically recognized tumor cells. During this cycle, various ligand-receptor interactions between APCs and T cells and between tumor cells and T cells provide signals to stimulate or inhibit the immune response ²⁸.

Very recently Louveau and colleagues discovered the presence of functional lymphatic vessel in mouse CNS ³⁵. In searching for some immune cell gateway, the scientists observed that meninges, that cover the brain and contain blood vessels and cerebrospinal fluid are associated with structures showing a vessel-like patterns and expressing markers of the lymphatic system. These vessels carried fluids and immune cells from the cerebrospinal fluid into cervical lymph nodes. They also identified a potentially similar structure in human dura, however additional studies are necessary to demonstrate and

characterize the existence and the location and organization of meningeal lymphatic vessels in the human CNS.

Clinical observations suggests that high densities of tumor-infiltrating lymphocytes (TILs) correlate with improved clinical outcome in melanoma, ovarian, colon, breast, lung, and other types of solid tumors. The majority of these studies observed a robust infiltration of CD3⁺T cells, cytotoxic T lymphocytes (CTLs) and CD45RO⁺ memory T cells that are associated with an improved survival ³⁶.

The characterization of TILs needs to be properly extended in malignant glioma. Indeed, contradictory reports exist regarding the presence and the prognostic factor of immune infiltration in GBM patients ³⁷. Donson et al. showed as gene expression microarray profiles of high-grade astrocytomas, including GBM, from long-term survivors have increased expression of immune-related genes. Histology validated results revealing an increased immune cell infiltration significantly associated with longer survival and a better Karnofsky performance status ³⁸. Similarly, a recent study described a significant correlation between increased CD3⁺ and CD8⁺ T cell infiltration in tumor bed and improved patient survival, even in presence of CD4⁺CD25⁺FOXP3⁺ regulatory T (Treg) cells and surface expression of inhibitory molecules ³⁹. Yang et al. reported natural killer (NK) cell infiltration in GBM suggesting their important role in antitumor immune responses ⁴⁰.

On the contrary, some reports indicated a predominant and variable infiltration of CD3⁺ T cells, often represented by Treg cells, in GBM specimens with a negative or not significant correlation with patient prognosis ^{41–43}. The difficulty to sustain an anti-tumor T cell mediated

cytotoxic response is principally due to the strong immunosuppressive microenvironment of GBM. Indeed, Fecci et al. found TILs with poor effector functions and an up-regulation of inhibitory receptors, such as CD95, PD-1, PD-L1, CTLA-4, LAG3 and TIM3, indicating immune exhaustion ⁴⁴.

Tumors cells responds with numerous strategies to counteract the immune system. It has largely reported that GBM cells secrete high levels of transforming growth factor beta (TGF-β), responsible for decreased expression of the activating receptor NKG2D on effector cells ⁴⁵, IL-10, prostaglandin E2 (PGE₂), VEGF and galectins ¹⁵. Moreover, glioma cells promote trafficking of suppressive Tregs, myeloid-derived suppressor cells (MDSCs) and M2 macrophages by production of chemoattractants as indoleamine 2,3- dioxygenase (IDO), CCL22 and CCL2 ^{46,47}. Down-regulation of major histocompatibility complex (MHC) and B7 family proteins ^{48,49}, increase of PD-L1 and HLA-E expression ^{50,51} and secretion of NKG2D ligands MICA/B ⁵² contribute to the immunosuppressive feature of GBM environment.

These mechanisms of immune suppression and the dynamic nature of interactions between T lymphocytes and tumor cells develop a mechanism of immune escape as a negative consequence of an antitumor immune response (figure 3).

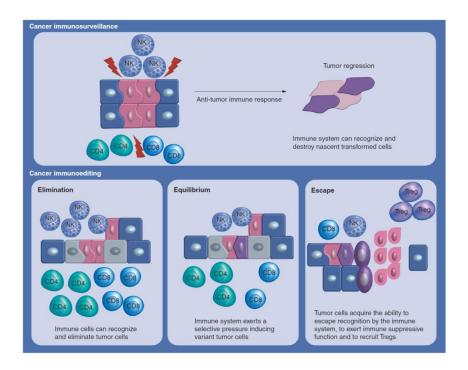


Figure 3. The three Es of cancer immunoediting: elimination, equilibrium, and escape. Immunosurveillance hypothesis describes the ability of the immune system to recognize and eliminate tumor cells. Immunoediting is a process triggered after the encounter between immune components and tumor cells. Transformed tumors cells may be recognized and eliminated by different cell types of the immune system. If elimination is unsuccessful, the immune system and the cancer can reach an equilibrium phase. Equilibrium involves the continuous elimination of tumor cells and the production of resistant tumor variants by immune selection pressure. This continuous sculpting can lead to escape, in which mutated cancer cells become able to inhibit the immune system and cancer can growth uncontrolled ⁵³.

Several different immunotherapeutic approaches are proposed to potentiate immune system against glioma cells with encouraging clinical results ^{18,27}. However, the mechanism of cancer immunoediting

used by GBM to evade immune clearance ⁵⁴ during immunotherapy suggests that this cancer requires multimodal therapeutic strategies for an effective treatment. Additionally, given the heterogeneity and the complexity of mechanisms involved in GBM pathogenesis, a more detailed characterization of tumor immune microenvironment is necessary.

Chemotherapy: a secret ally

Conventional anticancer drugs were firstly selected for their ability to arrest cell cycle progression leading to cell death. A growing body of evidence suggests that chemotherapeutic agents, when used at clinically useful doses, have an impact on therapeutic outcome beyond their cytotoxic properties on neoplastic cells. Indeed, the majority of these agents, initially considered immunologically silent or immunosuppressive, are able to stimulate the innate and adaptive immune system and evoke long-term protective memory T cell responses facilitating tumor eradication ⁵⁵. It has been shown that tumor cells, pretreated with chemotherapy, can evoke a durable adaptive immune response upon transfer into immunocompetent mice and an efficient rejection of tumor rechallenge ⁵⁶.

It is reported that chemotherapy is able to enhance antitumor immune response by (i) directly inducing tumor cell death with consequent release of tumor-derived antigens and immunostimulatory danger signals, (ii) by modifying tumor microenvironment and (iii) by modulating functions of immune cells, including DCs, MDSCs, tumor-associated macrophages (TAMs), CTLs and Tregs.

Certain chemotherapeutic agents confer an immunogenic cell death, which stimulates an immune response against dead-cell antigens, in particular when they derive from cancer cells. The immunogenic cell death involves changes occurring in a defined temporal sequence, in the composition of the cell surface as well as the release of soluble mediators called damage-associated molecular patterns (DAMPs). DAMPs stimulate inflammasome-dependent response, recruit specific cells of the innate and adaptive immune system to the tumor bed and trigger a therapeutically relevant immune response against malignant cells (figure 4).

Pre-apoptotic exposure of calreticulin (CRT) on dying tumor cell surfaces promote phagocytosis by DCs and the up-regulation of their costimulatory molecules and antigen-processing machinery components ⁵⁷. It is reported that the inhibition of CRT exposure in tumors negatively affects the efficacy of chemotherapy in immunocompetent mice, suggesting that this stress-dependent immunogenic signal is indispensable for the elicitation of anticancer immune responses ^{58,59}. The exposure of CRT is accompanied by changes in the endoplasmic reticulum (ER) ultrastructure and by signs of an ER stress response ⁶⁰. Moreover, the transcriptional up-regulation of heat-shock proteins (HSPs) is another important mechanism induced in response to cellular stress. HSPs are considered potently immunoadjuvants able to determine a strong immunogenicity of stressed tumor cells, facilitating cross-presentation of tumor antigens, DC maturation and also activating NK cell immune responses ⁶¹. Thus, a second line of immunogenic factors that stimulate antigen processing is required in addition to these signals. In late apoptosis the highmobility group box 1 (HMGB1), a non-histone chromatin-binding nuclear protein, is released and acts as a pro-inflammatory signaling factor. The release of HMGB1 and its interaction with the toll-like receptor 4 (TLR4) on the DC surface are mandatory for the optimal presentation of tumor antigens to T lymphocytes ⁶² and stimulate the synthesis of pro-inflammatory cytokines leading to NK and T cell activation ⁶³. Furthermore, many chemotherapeutic agents induce production and release of other alarmin molecules as uric acid, adenosine triphosphate (ATP) and uridine triphosphate (UTP) that bind the purinergic receptor P2X7 at the surface of DCs, resulting in the activation of the NLRP3 inflammasome and therefore in the secretion of interleukin-1β (IL-1β). Several reports indicated as IL-1β is required for the efficient priming and expansion of CD4⁺ and CD8⁺ T lymphocytes and elicitation of antitumor immune response ⁶⁴.

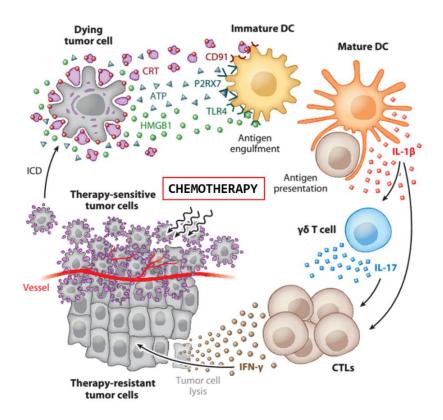


Figure 4. Immunogenic cell death (ICD) induced by chemotherapy.

Stressed-cancer cells undergo to ICD releasing DAMPs that facilitate the recruitment and activation of DCs into the tumor bed and the conditions for optimal antigen presentation to T cells. These processes result in a potent IFN- γ mediated immune response involving both $\gamma\delta T$ cells and CTLs, which eventually can lead to the eradication of chemotherapy-resistant tumor cells. Adapted from 65 .

In addition, chemotherapy can also render cancer cells more susceptible to CTL and NK cell killing inducing changes in their immunogenic properties. It is reported the capacity of some anti-neoplastic agents to increase the permeability of tumor cells to granzymes facilitating the CTL-mediated lysis even if they do not express the antigen recognized by CTLs ⁶⁶. Similarly, van der Most et al. showed as the alkylating

agent cyclophosphamide can sensitize tumor cells to TRAIL-dependent CD8⁺ T cell anti-tumor responses ⁶⁷. Anti-cancer agents inducing DNA damage promote functional down-regulation of inhibitory molecules and up-regulation of stimulatory DNAM-1 and NKG2D ligands on the surface of cancer cells, resulting in augmented NK and CD8⁺ T cell cytotoxicity and IFN-γ production ^{68–70}.

Tumor microenvironment contains immune-suppressive cytokines, immune checkpoints and regulatory subsets of cells, such as TAM, MDSCs and Tregs, that can contribute to a failure of immune-based approaches in clinics. Chemotherapy has the ability to modify tumor microenvironment converting it into a permissive state for the activation of a local and systemic antitumor immunity 71 . Indeed, current data have indicated as chemotherapeutic drugs induce an alteration of cytokine and chemokine profile in the tumor microenvironment decreasing amounts of immunosuppressive IL-4, IL-10 and TGF- β , and promoting production of pro-inflammatory mediators such as IL-8, TNF- α , IL-2, IL-7, IL-15, IL-21 and IFNs 72 .

In response to anti-neoplastic agents, the composition of tumor immune infiltrate changes with a crucial effect on therapy outcomes. Thus, an increased number of IFN- γ producing effector cells over immunosuppressive cells within the tumor microenvironment after chemotherapy predict favorable therapeutic responses in different types of tumors 73 . Indeed, many conventional chemotherapeutic compounds can lead to a transient suppression of functions or direct depletion of intra-tumoral and peripheral inhibitory cells restoring a potent and durable cytotoxic antitumor immune response and improving patient

survival ^{74,75}. Accumulating evidence suggest that chemotherapy treatment, most of all a metronomic administration, results in an enhanced serum levels of IFN-γ and IL-2 with consequent CD8⁺ T cell response in both murine models and patients with different solid tumors ^{76,77}. Similarly, some types of anticancer drugs elicit NK cell recruitment to tumors and regulate their cytotoxic activity in several models of cancer ^{68,74,78}.

These evidences suggest that many anticancer agents have the ability to evoke an anti-tumor immune response and may cooperate with immunostimulatory approaches, as immunotherapy, to develop combined protocols. The rationale for this dual approach is that chemotherapy has the potential to induce sufficient tumor cell death and counteract the immunosuppressive factors in tumor microenvironment synergizing immune-based therapy and harnessing the maximum antitumor immune response.

Temozolomide (TMZ)

TMZ is the most widely used and effective chemotherapeutic drug for GBM patients. A major benefit of TMZ is its ability to cross the bloodbrain barrier with almost 100% bioavailability. TMZ is a prodrug that spontaneously hydrolyzes at physiological pH to the active metabolites 5-(3-methyltriazen-1-yl)-imidazo-4-carboximide (MTIC) and methylhydrazine. TMZ exerts its antitumor effects by methylating guanine at the N⁷ and O⁶ positions and adenine at the N³ position. These adducts create a DNA damage in tumor cells that results in apoptosis and cytotoxicity ⁷⁹.

TMZ was first evaluated as a single treatm

ent for recurrent malignant glioma, where it showed a good antitumor activity and acceptable side effects. A successive pilot phase 2 trial demonstrated the feasibility of the concomitant administration of TMZ with fractionated radiotherapy, followed by six cycles of adjuvant TMZ for newly diagnosed GBM patients with a median survival benefit of 2.5 months compared to radiations alone ⁴.

The most important prognostic factor for survival and TMZ response is the methylation status of O⁶-methylguanine-DNA methyltransferase (MGMT), an alkyl guanine transferase that repairs lesions created by TMZ and other alkylating agents. It has been well documented that the level of MGMT activity is correlated with tumor cell resistance to chemotherapeutic drugs ⁸⁰.

Although TMZ is well-tolerated and has an overall beneficial impact on patient survival, it has also been well-described to induce immunosuppression, most often described as various forms of lymphopenia and abnormal levels of white blood cells ⁷⁹. Although bone marrow cells are particularly vulnerable to TMZ due to their high proliferation rate, severe myelosuppression and bone marrow failure are rare adverse effects. It has been suggested that TMZ selectively targets monocytes but not dendritic cells that derived from them ⁸¹.

Several studies pointed out that alkylating agents have immune inhibitory effects via selective toxicity to proliferating lymphocytes and inhibition of differentiation of immune effectors ^{82,83} highlighting the drug-mediated toxicity of chemotherapy to immune effector cells. However, lymphopenia caused by TMZ is preferential associated to depletion of highly potent immunosuppressive cells as Tregs and

MDSCs in blood and within the tumor microenvironment 84. This lymphopenic state has been positively correlated with clinical outcome due to relieve suppression of antitumor effector cells 85,86 Treg reduction was associated with the possibility to exploit a "timewindow" of lymphopenia in order to amplify immune responses. Recently, concomitant use of immunotherapy and TMZ regimen has been proposed: a lymphopenic state can induce homeostatic lymphocyte proliferation and enhance antitumor immune responses ^{87,88}. Thus, the efficacy of antitumor vaccines can be enhance by vaccination immediately after lymphodepleting chemotherapy ⁸⁹. Growing amount of evidence show as systemic TMZ does not inhibit antitumor immunity and has a robust efficacy when used in combination with recombinant IFN-β or DC vaccination in different animal models 90-92. Clinically, TMZ appears to be beneficial when combined with peptide vaccines in GBM and metastatic melanoma patients ^{93,94}. Moreover, prolonged survival has been observed in patients with GBM who receive chemotherapy after DC vaccination, compared with patients who receive vaccination only 95. Significant clinical benefits were observed also in a group of GBM patients treated with DC vaccines and subsequent chemotherapy suggesting a synergism between two approaches

TMZ contribution to tumor microenvironment is mediated by down-regulation of the expression of TGF- β , the most immunosuppressive factor in GBM, ⁹⁶ and by reduced CCL2 secretion, an important chemokine involved in Treg migration, by glioma cells ⁴⁶. TMZ has also demonstrated significant anti-angiogenic effects in experimental gliomas ⁹⁷. It has been recently described the ability of TMZ to modify

the chemokine network, in particular CXC chemokines with a role in glioma cell biology ⁹⁸. Furthermore, Hong et al. showed that TMZ can induce the intratumoral expression of different chemokines resulting in the recruitment of immune cells with antitumor activity ⁹⁹. Finally, it has been reported also the ability of TMZ to induce on tumor cells the expression of stress-associated NKG2D ligands on TMZ-resistant glioma cells, potentially rendering them vulnerable to lymphocyte recognition and lysis ¹⁰⁰.

ABC transporters

ATP-binding cassette (ABC) transporters are one of the largest protein families evolutionarily conserved present in all living organism from prokaryotes to mammals.

ABC transporters are large, membrane-bound proteins that utilize the energy of ATP hydrolysis to carry out the translocation of various substrates across membrane ¹⁰¹.

All ABC transporters consist of two distinct domains: the transmembrane domain (TMD) and the nucleotide-binding domain (NBD). The sequence and architecture of TMDs is variable, reflecting the chemical diversity of substrates that can be translocated. The NBD or ABC-domain is located in the cytoplasm and has a highly conserved sequence. ABC proteins are organized either as full transporters containing two TMDs and two NBDs, or as half transporters. The latter must form either homodimers or heterodimers to constitute a functional transporter.

It has been established that these pumps recognize a very wide range of substrates and their major physiological function is to transport exogenous and endogenous hydrophobic compounds through the membrane ¹⁰².

The ABC transporters, expressed in numerous cell types under physiological conditions, are also targets of therapeutic interventions in medicine, including cancer drug-resistance, lipid and other metabolic disorders.

Multidrug-resistance

Multidrug-resistance (MDR) is defined as the ability to resist to several drugs that may have different structures and mechanism of action. MDR is one of the major reason for the failure in the treatment of cancer. Chemotherapeutic drugs commonly enter cells by passive diffusion throughout the membrane. The MDR phenotype in tumors is usually associated with the overexpression of certain ABC transporters, resulting in increased drug efflux and reduced intracellular drug accumulation ¹⁰³.

It is known that exist at least 48 members of ABC transporters associated to MDR belonging to 7 different subfamilies that are divided on the basis of sequence and structural homology in humans. The best clinically characterized MDR transporters belong to three major groups: ABCB, ABCC and ABCG families ¹⁰⁴.

ABCB and ABCG preferentially extrude large hydrophobic, positively charged molecules, while the members of the ABCC family can extrude both hydrophobic uncharged molecules and water-soluble anionic compounds in physiological conditions.

The most extensively studied gene in ABCB family is ABCB1, also known as MDR1 or P-glycoprotein (Pgp1). Pgp1 was the first described

membrane transporter characterized by its ability to confer multidrugresistance in cancer cells to cytotoxic drugs such as methotrexate, anthracyclines, vinca alkaloids and taxanes ¹⁰⁵.

The ABCC family, also known as multidrug resistance-associated protein family (MRPs), contains the greatest number of known drug transporters that confer resistance to a similar, but not identical, spectrum of drugs of Pgp1 ¹⁰⁶. It has been established that there are 13 genes belonging to the ABCC family and only three of these genes encode proteins are not involved in drug resistance. Thus, ABCC7 transporter is the cystic fibrosis transmembrane conductance regulator (CFTR), the only member that functions as a channel for chloride ion efflux. ABCC8 and ABCC9 are cell surface receptors that regulate potassium channels modulating insulin secretion ¹⁰⁷.

The most characterized member in ABCG family is ABCG2, an half transporter with a single NBD and TMD, also known as the breast cancer resistance protein (BCRP) found for the first time in cell lines resistant to anthracycline and mitoxantrone. It is expressed in many types of solid tumors, conferring resistance to most of topoisomerase I or II inhibitors such as topotecan, irinotecan and doxorubicin ¹⁰⁸.

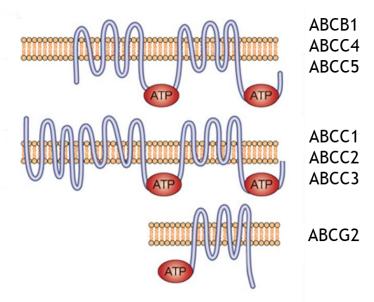


Figure 5. Structure of the best clinically characterized MDR transporters. ABCB1 and ABCC4/5 transporters have 12 transmembrane domains and two ATP-binding sites. The structures of ABCC1/2/3 are similar but contain also an additional domain that is composed of five transmembrane segments at the amino-terminal end, giving them a total of 17 transmembrane domains. ABCG2 contains 6 transmembrane domains and one ATP-binding region ¹⁰⁹.

Other Functions

Since ABC transporters have an important role in tumor biology, they are large studied and associated to some of the cancer hallmarks as apoptosis, proliferation and migration or invasion.

Several reports provide evidences that ABC proteins might promote cell survival independently of cytotoxic drug efflux ^{110,111}. For example, the increased expression of Pgp1 has been reported to delay apoptosis in response to a wide range of harmful stimuli in normal and cancer cells. Indeed, apoptotic cascade needs intracellular acidification and the over-expression of multidrug-resistance proteins, associated with

intracellular alkalinization, inhibits the progression of death ^{112,113}. The downregulation of ABC transporter expression or the inhibition of their activity may result in cell cycle arrest and alteration of the apoptotic cascade with reduced expression of anti-apoptotic proteins ^{114,115}.

Moreover, ABC transporters seem to be involved in proliferative mechanisms. Kuss et al. described as a down-regulation of ABCC1 expression not only increases tumor cell death but also reduces the mitotic index indicating a lower cell turnover in vitro and in vivo ¹¹⁶. Similarly, other ABC pumps, normally associated to MDR, have a role in contributing directly to the proliferation of tumor and normal cells ^{117,118}

Lastly, several ABC transporters play some roles in normal and tumor cell migration. The physiological expression of ABC proteins promotes migration in different type of cells, such as DCs or endothelial cells ^{119,120}. It is reported that the reduction of ABC protein levels reduced migration in vitro and in vivo in different cancer cell lines promoting their invasive phenotype ^{121,122}.

ABC transporters and Immune system

In the last years, the expression of ABC proteins has been reported in numerous cell types under physiological conditions. Peripheral blood cells have also been found to express several ABC transporters with an important role in the resistance to anti-inflammatory, anti-viral, and anti-cancer drugs ^{123–125}. Although the relative expression of this type of proteins in lymphocytes are controversial ^{126,127}, it is accepted that ABC transporters can be protagonists of immune processes influencing

the differentiation and maturation of immune cells and promoting the migration of immune effectors to sites of inflammation ^{123,128}.

Indeed, ABC proteins can transport inflammatory mediators such as platelet activating factor, leukotrienes and prostaglandins ^{129–131} exerting a pro-inflammatory effect and facilitating the migration of immune cells to lymph nodes ¹³².

It is well known the role of ABC transporters, in particular Pgp1, ABCC1 and ABCC4, in DC migration and differentiation ^{119,133}. In addition, several member of ABCB family are responsible for intracellular peptide transport, including a role in major histocompatibility complex (MHC) class I antigen presentation ¹³⁴.

A controversial point remains whether ABC transporters are involved in the efflux of inflammatory cytokines and chemokines. Several studies proposed an active role of ABC transporters in extrude molecules as TNF- α , IFN- γ , perforins and interleukins ^{135–137}. However, the molecular weight of some of these molecules exceeds size of the most common substrates of these transporters. Cytokine secretion may result as a secondary and indirect effect in response to efflux of other substrates.

Several studies reported that Pgp1, ABCC1 and ABCC2 transporter activity is related to cytotoxicity in NK cells and its inhibition was accompanied by suppression of IFN-γ secretion ^{123,135}. On the contrary, there are evident discrepancies about the expression and function of ABCC1 and Pgp1 in T cells: some studies suggested an increased expression upon T cell activation while others indicated a decrease ^{138,139}.

Since ABC transporters have an important role during chemotherapy treatment due to MDR acquisition, their function should be investigated intensively in immune cells to assess their potential roles in basal drugresistance. For now, the greatest interest for ABC transporters is for drug therapy of leukemias: Pgp1, BCRP and several MRPs have already been found to confer resistance against anti-leukaemic drugs and indicated as prognostic factors ^{140,141}.

A substantial variability in ABC expression levels of each transporter is reported in different lymphocyte subsets, ¹²⁶. Expression of ABC transporter in CD8⁺ and CD4⁺ T cells are controversial and related to activation and differentiation status ¹⁴². In Tregs a low expression of ABC transporters and an extreme sensitivity to chemotherapy ^{123,143} support the absence of a specific drug efflux. Conversely, a number of studied revealed that NK cells appear more chemo-resistant than other lymphocytes principally due to their relatively high levels of Pgp1 and ABCC1 ^{128,144,145}. Moreover, it has been suggested that the highly aggressive clinical course and strong refractory to chemotherapy of patients with NK lymphomas could be related to activity of Pgp1 and other ABC transporters ¹⁴⁶.

ABCC3, a marker of drug-resistance

ABCC3, also known as MRP3, is a member of the large C family involved in active extrusion of substrates out of the cells. ABCC3 is expressed in many polarized cells and localized in the basolateral membrane. It was first localized in human and rat hepatocytes where it can mediate the efflux of organic anions into sinusoidal blood ^{147,148}. It

is expressed in the adrenal glands and the intra-hepatic bile ducts, colon, kidney, pancreas, spleen and some lymphoid cells ^{123,149}.

ABCC3 is also an organic anion transporter and prefers glucuronate conjugates as transported substrates. Like other MRPs, it was largely characterized for its role in detoxification through export of numerous toxic molecules. This detoxification is reported also in numerous tumors where it contributes to drug-resistance actively extruding a broad spectrum of anticancer drugs ^{107,150}. Overexpression and activation of ABCC3 leads to the reduction of intracellular drug levels and consequent drug insensitivity. Indeed, ABCC3 is proposed as a marker of MDR and predictor for clinical outcome and response to chemotherapy in several cancer types ^{151–154}. Moreover, it has been shown to be upregulated and related to tumor grading in pancreatic cancer and malignant gliomas ^{155,156}.

Its expression correlated with decreased sensitivity to different anticancer drugs such as vincristine, etoposide, and cisplatin ^{157,158}, and also to methotrexate and doxorubicin ^{148,159,160}.

Giannoudis et al. suggested that ABCC3 efflux activity could have a role in imatinib treatment failure in patients with chronic myeloid leukemia. ABCC3 pharmacological inhibition showed a better cellular retention of imatinib demonstrating as this drug is actively efflux by ABCC3 transporter. Moreover, patients with high ABCC3 expression at diagnosis were less responsive to imatinib treatment supporting this hypothesis ¹⁶¹.

Although ABCC3 expression and efflux activity is not broadly studied in lymphocytes and other immune cells, increasing evidences supported its role in lymphoid leukemias ¹⁶². Mature leukocytes display basal

constitutive MRP-related transport activity regardless of cell lineage suggesting a potential role of ABCC3 in chemo-protection of immune cells.

Unraveling the specific role of ABCC3 and other transporters in MDR phenotype of the immune system may therefore reveal new links for future immune-chemotherapeutic approaches.

Scope of the thesis

Glioblastoma (GBM) is one of the most lethal human cancer. Standard treatment of GBM includes three steps: surgical resection of the tumor mass, radiotherapy and chemotherapy with temozolomide (TMZ). Despite this multimodal and aggressive strategy, GBM continues to have a dismal prognosis with almost inevitable recurrences. There is the need to find long-term strategies to target residual tumor cells infiltrated in the adjacent areas of brain and responsible for tumor perpetuation.

Cancer immunotherapy is now considered a potent weapon aims to harness the body's own immune system to fight tumor cells. Recently, the number of immunotherapy approaches to fight GBM is growing with encouraging results^{18,19}. Moreover, data show that several anticancer agents, traditionally viewed as immunosuppressive, are now proposed as immune-adjuvants able to enforce anti-tumor immune response when combined with immunotherapy¹⁶³.

In our institute, we are currently carrying out a clinical trial based on immunotherapy with dendritic cells (DCs) loaded with whole tumor lysate. Patients with first diagnosis of GBM after surgery and leukapheresis receive standard treatment with radio-chemotherapy (RT-TMZ) and TMZ as adjuvant during DC vaccines. Flow cytometry analyses on peripheral blood lymphocytes indicated a significant activation of an NK, but not CD8⁺ T, cell response correlated with a prolonged survival of patients. In light of these data, we moved from clinical observations to a murine malignant glioma model to evaluate a potential direct effect of TMZ on immune system.

We have found that NK cells, but not CD8⁺ T lymphocytes, are able to up-regulate Abcc3, a multidrug resistant ATP transporter, as we extensively demonstrated in "The multidrug-resistance transporter Abcc3 protects NK cells from chemotherapy in a murine model of malignant glioma" OncoImmunology, Oct 29 2015 (DOI 10.1080/2162402X.2015.1108513).

In this study, our principal goal was to analyze the potential ability of TMZ to modulate local and systemic immune cells involved in antitumor response and influence their activity in GL261 glioma-bearing mice.

The first step was to characterize the direct impact of TMZ on frequency, cytotoxic ability, migration and consequent homing of NK and CD8⁺ T cells.

Nine days after GL261 intracranial injection, mice were treated with TMZ for five days. Trafficking of NK1.1⁺ CD3⁻ NK and of CD8⁺ T cells was investigated by flow cytometry in blood, their homing ability into the brain was evaluated as tumor infiltration by flow cytometry and immunofluorescence.

We also investigated the role of TMZ in modulating tumor microenvironment and providing conditions that can favor NK cell homing and activation. We observed in fact that TMZ treatment is able to exert a local modulation of the immunogenicity and to modify the chemokine-cytokine pattern increasing infiltration of NK cells with a stronger anti-tumor effector activity.

The second step was to evaluate the direct molecular mechanisms induced by TMZ on NK cells, important protagonists in anti-tumor

immune response of our patients, focusing on their ability to react to the cytotoxic effects of chemotherapy. By microarray analysis we identified three important gene signature related to migration, inhibition of apoptosis and multidrug resistance. Notably a high expression of Abcc3 was found physiologically expressed in NK cells and upregulated during TMZ treatment.

The third goal was to specifically characterize the functional role of Abcc3 expressed by NK cells and its involvement in resistance to TMZ-induced apoptosis. NK cells expressing Abcc3 do not undergo apoptosis and show an activation of Akt, a key protein for immune cell survival. Pharmacological inhibition of Abcc3 in NK cells caused an increase of apoptosis demonstrating the crucial role of Abcc3 in resistance of NK cells to TMZ.

Results obtained will guide us to implement the knowledge of combined therapy modalities taking into account our clinical trials where GBM patients received chemo-immunotherapy.

Based on these pre-clinical observations we plan to verify the expression of Abcc3 in NK cells of patients before, during and after chemo-immunotherapy in order to define its potential role in the immune response correlated with a greater survival.

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Chapter 2

The multidrug-resistance transporter Abcc3 protects N	X (cell	S
from chemotherapy in a murine model of malignant gl	ioi	ma	

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Abstract

Abcc3, a member of the ATP-binding cassette transporter superfamily, plays a role in multidrug-resistance. Here, we found that Abcc3 is highly expressed in blood-derived NK cells but not in CD8⁺ T cells. In GL261 glioma-bearing mice treated with the alkylating agent temozolomide (TMZ) for five days, an early increased frequency of NK cells was observed. We also found that Abcc3 is strongly up-regulated and functionally active in NK cells from mice treated with TMZ compared to controls. We demonstrate that Abcc3 is critical for NK cell survival during TMZ administration; more importantly, Akt, involved in lymphocyte survival, is phosphorylated only in NK cells expressing Abcc3. The resistance of NK cells to chemotherapy was accompanied by increased migration and homing in the brain at early time points. Cytotoxicity, evaluated by IFN-y production and specific lytic activity against GL261 cells, increased peripherally in the later phases, after conclusion of TMZ treatment. Intra-tumor increase of the NK effector subset as well as in IFN-y, granzymes and perforin-1 expression, were found early and persisted over time, correlating with a profound modulation on glioma microenvironment induced by TMZ.

Our findings reveal an important involvement of Abcc3 in NK cell resistance to chemotherapy and have important clinical implications for patients treated with chemo-immunotherapy.

Introduction

Current therapeutic options for glioblastoma (GBM) patients include chemotherapy with the alkylating agent temozolomide (TMZ). 1,2 Recent data have suggested that some chemotherapeutic agents, previously viewed as immunosuppressive, possess immunemodulatory effects^{3,4} and influence the vaccine-induced immune response affecting the quality and efficacy of the T cell response⁵ or enhancing the immunogenicity of dying tumor cells.^{3,4} TMZ leads to transient lymphodepletion⁶ and may interfere with regulatory T cell (Treg) trafficking to the tumor⁷, thereby creating a "time-window" for improved efficacy of vaccinations; moreover, dendritic cell (DC) immunotherapy may increase TMZ sensitivity.8 Preclinical evidence has implicated the inhibition of glioma growth by NK cells^{9,10} and recently, we reported a significant, positive correlation of NK cell response and survival of patients affected by recurrent GBM treated with DCs loaded with autologous tumor lysates. 11 Treg depletion by TMZ could relieve the suppression of NK cells restoring the innate antitumor response. 12

Previous attempts have been made to decipher the mechanisms through which NK cells are more radio- and chemotherapy resistant than other lymphoid cells. It has been observed that NK cells express high levels of P-glycoprotein 1 (P-gp1), a transmembrane transporter encoded by the multidrug-resistance 1 (MDR1) gene, as well as MRP1 (ABCC1) and MRP2 (ABCC2).¹³ Discrepancies were found in terms of the expression and function in T cells of multidrug-resistance proteins, specifically P-gp1 and ABCC1.^{14,15}

In a clinical trial currently active at our institution (DENDR1 - EUDRACT N° 2008-005035-15), 24 patients with first diagnosis of GBM have been treated with DC loaded with autologous tumor lysate together with standard radiotherapy and chemotherapy with TMZ. Peripheral blood lymphocytes (PBLs) from patients were analyzed by flow cytometry for immunotherapy follow-up. Their ratio of vaccine/baseline frequencies (V/B ratio) was correlated with the progression-free survival (PFS) of each patient. The increased V/B ratio of NK cells but not CD8⁺ T cells was significantly associated with prolonged PFS (Pellegatta et al, manuscript in preparation).

To investigate the specific contribution of TMZ-based chemotherapy to differential responses of NK and T cells, we used the GL261 preclinical model of glioma.

We found that blood-derived NK cells (but not CD8⁺ T cells) are resistant to and activated by TMZ. Multidrug-resistance is primarily associated with Abcc3 expression (a member of the MRP family), which was up-regulated and functionally active in NK cells during TMZ treatment. Furthermore, NK cells displayed migratory and cytotoxic activities that were positively influenced by TMZ.

Results

Local and systemic NK cell frequency is positively influenced by TMZ

Nine days after intracranial implantation of GL261 gliomas, immune competent glioma-bearing mice were treated with intraperitoneal injections (i.p.) of 5 mg/kg TMZ or DMSO for 5 days (Figure 1A). To characterize the effect of TMZ on the immune system, PBLs and tumorinfiltrating lymphocytes (TILs) were harvested at different time points, and immune cell populations quantified using flow cytometry. TMZ induced rapid and reversible lymphopenia: CD8+ T cells decreased significantly at 48 hours (hrs), after two administrations of chemotherapy (p < 0.0001 vs controls) and quickly increased at 72 hrs (p < 0.01 vs 48hrs; Figure 1B). On the contrary, peripheral blood NK cells increased significantly at early time point, doubled 72 hrs after the first TMZ administration and remained higher than controls throughout the entire treatment (Figure 1C). To assess a possible delayed effect of TMZ on immune cells, we performed similar evaluations at day 19, 5 days after ending chemotherapy. We did not observe a significant difference between CD8+ T cells in blood of TMZ-treated mice compared to controls (Figure 1B) while NK cells were still increased in blood of TMZ- compared to vehicle-treated mice (Figure 1C). In nonglioma-bearing mice TMZ induced a modulation of CD8+ T lymphocytes and NK cells similar to TMZ-treated tumor bearing mice (Figure S1).

Tumor infiltrating immune cells were isolated from fresh gliomas by Percoll gradient and quantified by flow cytometry as NK1.1⁺CD3⁻ and

CD8⁺ CD3⁺: at 48 hrs CD8⁺ T cells decreased in TMZ-treated mice compared with controls (Figure 1D); at 240 hrs, no differences were found between TMZ-treated mice and controls (p = 0.3; Figure 1D). NK cells in the tumor showed a similar pattern of systemic NK cells, increasing significantly at 72 hrs after the first TMZ administration and continued to be significantly elevated at 240 hrs (p < 0.005 at both time points, Figure 1E).

Hematoxilin & eosin staining confirmed the evidence of a vigorous immune cell infiltration in response to TMZ compared with the vehicle at 72 hrs. In situ immunofluorescence established the predominant infiltration of NK cells located into the tumor mass (Figure 1F). Quantitative determination of TILs from the same groups of treatment was obtained by flow cytometry (Figure 1G).

In another set of experiments, immune cells from spleen, cervical lymph nodes and bone marrow were analyzed: the results did not indicate a significant influence of TMZ on lymphocytes in these organs (not shown).

These results show that trafficking and NK cell homing to the tumor are positively influenced by TMZ administration.

Expression of genes involved in drug-resistance and chemotaxis is up-regulated in NK cells from TMZ-treated mice

To further characterize the molecular effects of TMZ, we compared gene expression profiles of NK cells obtained by magnetic sorting of PBLs from TMZ- and vehicle-treated glioma-bearing mice (n = 50/group) 72 hrs after treatment onset. We used the GeneChip Mouse Gene 2.0 ST Array and identified differentially expressed genes (DEGs)

using a \geq 2 fold-change (FC) threshold for transcript comparisons. A robust difference was observed between the transcriptome levels of the two NK cell groups (Table S1) and 211 DEGs passed the FC cut-off.

Based on Gene Ontology annotations, the transcripts were grouped because of their involvement in multidrug-resistance, anti-apoptosis and migration. We focused the validation experiments on genes indicating the relationship of NK cells with drug-resistance. In particular, three up-regulated genes were related to ABC drug transporters: Abcc3, Abca9 and $Abca6^{16}$. Other genes were related to the inhibition of apoptosis (CD5L and Naip1) and cell survival (Ednrb, Gata6 and Fgfr1), an indication of the predisposition of NK cells to resist the cytotoxic effect of TMZ^{17-20} .

In addition, data from genes regulating cytoskeleton organization, microtubule-based movement, actin polymerization and chemotaxis (*Ccr1*, *Efnb2*, *Alox15*, *Lbp* and *Lrp1*) supported the idea that NK cells from TMZ-treated mice migrated more than NK cells from controls^{21,22}. Notably, at this time point, down-regulated genes were related to NK cell-mediated cytotoxicity (*GzmD*, *GzmE*, *GzmG* and *GmzC*) and to secretory pathway or inflammatory response (*Scgb1a1* and *Elane*)²³.

Gene expression profiling was also performed on CD8⁺T cells purified from the same mice. No significant differences could be observed between CD8⁺ T cells sorted from TMZ-treated mice and those of controls (Table S1).

Overall, these findings suggest that TMZ influences the activity of NK cells by activating pathways relevant in the acquisition of chemoresistance.

NK cells respond to chemotherapy by over-expressing Abcc3

The validation of ABC transporter over-expression was performed by real time PCR. The analysis revealed that two of the three ABC transporters, Abcc3 and Abca6, were significantly up-regulated in peripheral NK cells from TMZ-treated mice compared to controls (3.46 \pm 0.01-fold; p < 0.0001, and 2.75 \pm 0.045-fold; p < 0.001, respectively; Figure 2A and B).

To investigate *in vitro* the expression of Abca6 and Abcc3 during TMZ administration, we treated PBLs from naïve mice with 1 μ M TMZ or DMSO at different time points. The dosage was determined according to TMZ concentrations measured in the plasma of patients treated with the "standard" schedule.²⁴ The up-regulation of Abca6 expression was observed only after 6 and 8 hrs of TMZ treatment (2.0 \pm 0.1–fold and 3.3 \pm 0.1–fold, respectively vs DMSO-treated PBLs; p = 0.01, p < 0.005; Figure 2C). On the contrary, the up-regulation of Abcc3 expression was detectable after 4 hrs (p < 0.05) and increased over the time during TMZ treatment, suggesting a direct effect of chemotherapy on its expression (Figure 2D).

The remarkable increase of Abcc3 expression on the surface of NK cells was confirmed *in vivo*. At flow cytometry, NK cells (but not CD8⁺ T cells) displayed a high basal expression of Abcc3 (31.2 \pm 0.8% Abcc3⁺ NK cells vs 2.0 \pm 0.6% Abcc3⁺ CD8⁺ T cells; p < 0.00001). Moreover, NK cells from TMZ-treated mice exhibited a significant up-regulation of Abcc3 compared to controls during chemotherapy (31.2 \pm 0.8% vehicle NK cells vs 59.8 \pm 1.1% TMZ-treated NK cells, p < 0.0001; Figure 2E, left). No significant difference in Abcc3 expression was

found in CD8⁺ T cells from TMZ-treated mice compared to controls $(2.0 \pm 0.6\% \text{ vehicles vs } 1.5 \pm 0.5\% \text{ TMZ-treated mice, Figure 2E, right)}.$

These results show that Abcc3 is differentially expressed in NK cells compared to CD8⁺ T cells and is increased in NK cells from TMZ-treated glioma-bearing mice compared with controls.

Abcc3 expressed in NK cells is functionally active

To investigate whether Abcc3 expression is related to a greater ABC transporter activity and a drug-resistant phenotype, we used a flow cytometry assay to measure the efflux activity of the three clinically most important ABC transporter families involved in cancer multidrugresistance. The assay is based on determining the fluorescence intensities of cells after a short incubation with a fluorescent substrate in the presence or absence (control) of specific ABC transporter inhibitors. Inhibition of active ABC transporters results in increased fluorescence intensity due to the accumulation of the substrate. PBLs from naïve mice were treated in vitro with 1 µM TMZ or DMSO for 4 hrs. NK and CD8⁺ T cells were gated on PBLs, and the multidrugresistance activity factor (MAF) was calculated. Cells exhibiting drugresistance have increased fluorescence and a MAF greater than 25%. TMZ-NK cells showed greater fluorescence in the presence of a multidrug-resistance protein inhibitor for MRP, of which Abcc3 is a key member resulting in MAF = 73.4% (Figure 2G).

TMZ-treated NK cells were also tested for the MDR and BCRP inhibitors included in the assay, confirming a high efflux activity. MRP exhibited the strongest efflux activity (p<0.001; Figure 2G). Results obtained with a lower dose of TMZ suggested that the efflux activity

was dose-dependent in NK cells (Figure S2). No evidence of a resistant phenotype was found in TMZ-treated CD8⁺ T cells, showing a MAF < 25% for all ABC transporter families (Figure 2H).

These findings highlight the rapid activation of ABC multidrugresistance transporters in NK cells but not in CD8⁺ T lymphocytes during TMZ treatment, supporting the NK cell ability to react to the cytotoxic effects of chemotherapy.

Abcc3 is critical for survival and expansion of NK cells during TMZ administration

To determine whether Abcc3 is required for NK cell survival, we treated PBLs from naïve mice with 1 µM TMZ or DMSO for 2, 6 and 15 hrs. Apoptotic cells were measured by flow cytometry on gated NK and CD8⁺ T cell populations using Annexin V and Propidium Iodide (PI) staining (Figure 3 A-C).

Cells in early apoptosis (EA) were Annexin V positive and PI negative, cells in late apoptosis (LA) or dead were Annexin V and PI positive. NK cells showed a low percentage of early apoptotic cells that slightly increased after 6 and 15 hrs (p < 0.01; Figure 3A). Abcc3 negative NK cells showed a higher percentage of apoptotic cells compared to Abcc3 positive NK cells in response to TMZ (Figure 3B). CD8⁺ T cells showed a remarkable increase of early and late apoptosis after 2 hrs and later (p < 0.005, p < 0.001; Figure 3C).

Because the pathways involving Akt activation could promote lymphocyte survival 25,26 , we investigated Akt activation by analyzing phosphorylation in PBLs in response to chemotherapy. NK cells and CD8⁺T cells were purified from naïve mice and treated with 1 μ M TMZ

or DMSO *in vitro* at two different time points. Western blots showed a basal phosphorylation of Akt in NK cells with a time-dependent increase of pAkt with TMZ treatment (Figure 3D). On the contrary, in CD8⁺ T cells, no significant difference in Akt activation was detected in TMZ-treated cells (Figure 3E). We confirmed these results by analyzing Akt activation in NK and CD8⁺ T cells from glioma-bearing mice treated with TMZ or vehicle. We sacrificed mice after the third TMZ (or DMSO) administration every 5 minutes for 30 minutes, and Akt activation in PBLs was measured by a flow cytometry phosphospecific staining (Miltenyi Biotec). Akt phosphorylation was only detected in NK cells from TMZ-treated mice expressing Abcc3 (p=0.01 vs vehicle-treated mice; Figure 3E). Abcc3⁺CD8⁺ T cells from TMZ-treated and control mice did not exhibit Akt phosphorylation (2.9 ± 1.2% TMZ-treated vs 3.3 ± 0.9% control mice; Figure 3E).

Inactivation of Abcc3 function by its specific inhibitor MK571 induced a significant increase of NK cell apoptosis in PBLs treated with 1 μ M TMZ for 4hrs (Figure 3F).

Together, these results confirm the important role of Abcc3 in survival and response to cytotoxic effects of chemotherapy.

NK cell migration and maturation are positively influenced by TMZ

To validate the signature related to migration and chemotaxis in NK cells from TMZ-treated glioma-bearing mice, we measured NK cell migration using the transwell system. NK cells purified from PBLs of TMZ-treated glioma-bearing mice and controls were evaluated for their ability to migrate toward conditioned medium derived from GL261

cells treated with 150 μ M TMZ or DMSO for 24 hrs. We observed a 1.8-fold increase in the migration of NK cells from TMZ-treated mice toward the supernatant from TMZ-stimulated GL261 compared to the vehicle supernatant (p<0.005). Migration of NK cells from treated mice increased by 2.3- and 30-fold compared with NK cells from control mice (p<0.0001, Figure 4A).

We then analyzed the expression of CD49b and CD49d integrin subunits involved in cellular adhesion and leukocyte tissue infiltration. CD49b+ CD49d+ double positive NK cells increased in blood of TMZ-treated mice compared to controls (57.1 \pm 3.4% vs 39.7 \pm 2.2%, respectively, p<0.01), supporting their greater homing ability into tumors (Figure 4B).

We also characterized the effect of TMZ on the maturation status of NK cells by evaluating the surface density of CD11b and CD27, two markers associated with the four-stage developmental program in mice and humans. In blood, immature CD11blowCD27low NK cells significantly decreased after three days of chemotherapy (p < 0.001 vs controls). Interestingly, TMZ led to a significant enrichment of the CD11blowCD27high and CD11bhighCD27high NK cell subsets with migratory potential and a simultaneous decrease of the CD11bhigh CD27low cytotoxic subset (p < 0.001 vs controls; Figure 4C). The increase of the CD11bhigh CD27low cytotoxic subset was observed five days after the end of TMZ administration (p=0.001; Figure 4D).

On the contrary, in gliomas of TMZ-treated mice there was a strong accumulation of the CD11b^{high}CD27^{low}NK effector subset 72 hrs after the first TMZ administration (p < 0.005, Figure 4E). This increase persisted at later time points (p < 0.005, Figure 4F).

These data showed a direct effect of TMZ on the progressive maturation of NK cells, with consequent influence on their migratory and cytotoxic phenotype in blood and gliomas.

Local and systemic NK cell cytotoxicity is triggered by TMZ

We further aimed to verify the cytotoxic ability of NK cells during and after completion of TMZ treatment.

The significant accumulation of the effector subset in blood of TMZ-treated mice suggested an activation of systemic cytotoxicity against tumors. In parallel, IFN- γ production in blood-derived NK cells from TMZ-treated mice increased compared to controls (7.5 \pm 1.0% vs 16.9 \pm 1.2%; p < 0.005; Figure 5A).

To test the cytotoxic specificity of NK cells, PBL from naïve mice, and TMZ- or vehicle-treated glioma-bearing mice were stimulated with autologous irradiated tumor cells. NK cells from TMZ-treated mice, purified by magnetic sorting, exhibited a greater lytic activity against GL261 cells than NK from vehicle or naïve mice (p < 0.0001, Figure 5B).

The accumulation of the NK cytotoxic subset in gliomas was associated to increased expression of *Perforin 1 (Prf1)*, *Granzyme B (GzmB)* and $Ifn-\gamma$ (by 9.3± 0.04-fold, 8.1± 0.02-fold and 4.10 ± 0.03-fold, respectively; p<0.0001) in TMZ-treated mice (Figure 5C). This upregulation persisted on day 19, five days after the end of TMZ treatment (*Prf1*: 3.97± 0.07-fold, *GzmB*: 2.2 ± 0.06-fold and $Ifn-\gamma$: 2.4 ± 0.02-fold, p<0.0001 vs controls; Figure 5D).

Thus, TMZ is able to modulate NK cell function increasing their effector activity.

Glioma microenvironment is converted by TMZ into a site permissive for an efficient effector immune response

To investigate whether the glioma microenvironment could be modulated by TMZ, we looked for expression levels of galectin-1, -3 and -9, that suppress NK immune surveillance^{30–32} and found that they were expressed at high levels in tumors of vehicle-treated mice and significantly less in those of TMZ-treated mice (Figure 6A).

A decreased expression of H2-Q1 (Hla-e), a non-classical Major Histocompatibility Complex class I (Mhc I) molecule normally implicated in immune escape mechanism and inhibition of NK cellmediated lysis, was also found in gliomas from TMZ-treated mice (Figure 6B). We also investigated the effects of TMZ administration on intra-tumor expression of chemokines and cytokines involved in modulating the infiltration of immune cells. CCL3 (which expression is related to NK cell accumulation 10), TNF- α (which is not only involved in anti-tumor immune response but also responsible for decreased GL261 proliferation³³), IL-7 (which plays a role in promoting NK cell survival and inhibiting apoptosis³⁴), and IL-27 (which is an important stimulator of NK cell effector function³⁵), all increased in gliomas from TMZ-treated mice compared to controls (Figure 6C). In contrast, CXCL10, which role in glioma progression is contradictory^{36–38}, and CCL5, which expression was recently most related with CD8A levels³⁹, and previously described as immunosuppressor by us and others ^{10,40}, significantly decreased.

Finally, we investigated *in vitro* the effect of TMZ on glioma immunogenicity. GL261 cells were treated with 50 and 150 μ M TMZ or DMSO at different time points. TGF- β 1 and TGF- β 2 concentrations

in the supernatant from TMZ-treated GL261 cells significantly decreased, as evaluated by ELISA (Figure 6D). Nkg2d ligand (Nkg2dI), involved in Nkg2d-mediated NK cell recognition of tumor cells and weakly expressed in GL261 cells, was up-regulated in a time- and dose-dependent manner after TMZ treatment (Figure S3). Similarly, we found a time- and dose-dependent increase of the Rae-1 - β , - ϵ and - δ , ligands for the Nkg2d receptor (Figure 6E); while B7-h3, a NK cell inhibitory molecule highly expressed in GL261 cells, was significantly decreased (Figure S3).

These results indicate that TMZ modulate glioma microenvironment into a site favoring NK cell infiltration and anti-tumor cytotoxicity.

Discussion

Our results show, for the first time, that NK cells in peripheral blood are resistant to chemotherapy due to expression of Abcc3, which was slightly or not expressed by CD8⁺ T cells. Abcc3 was up-regulated and active in NK cells from glioma-bearing mice during TMZ treatment, whereas CD8⁺ T cells did not exhibit a drug-resistant phenotype. We also confirmed the ability of NK, but not CD8⁺ T cells, to react to cytotoxic effects of chemotherapy by measuring their apoptosis *in vitro*, which was low or almost absent in NK cells during TMZ treatment.

ABC transporters promote cell survival independently of cytotoxic drug efflux, as shown in a study where the inhibition of endogenous expression of ABC transporters resulted in reduced expression of Bcl-2 protein levels and activation of the apoptotic cascade. 41 Based on these observations, ABC transporters can be responsible for the drugresistance phenotype through direct drug efflux and by other intrinsic pathways, including the phosphorylation of Akt, a key regulatory molecule involved in cell survival, that is activated in response to TMZ contributing to chemo-resistance.⁴² This is supported by data presented here showing that Abcc3-expressing NK cells from TMZ-treated glioma-bearing mice exhibit significant Akt phosphorylation, a protective mechanism against cell death. ³⁶ Notably, Akt activation has been correlated with increased expression and activity of some ABC transporters^{44–46} and there is evidence that Akt and PI3K/Akt pathway are responsible for cytotoxic and killing ability of NK cells. 47-50 Our data showing that Akt is phosphorylated only in Abcc3+ NK cells support the involvement of Abcc3 in the cytotoxicity of NK cells.

We also found that TMZ led to an enrichment of NK cells with migratory function, as observed by investigating the 4-stage model of NK cell maturation.²⁸ While different studies have shown that ABC transporters play a role in the migration of cancer and normal cells, including immune cells^{51,52}, a direct action of TMZ on NK cell maturation/migration has not been previously reported. The increase of migratory subsets is concomitant with a consistent modulation of the glioma microenvironment of TMZ-treated mice, where we found a significant increase of chemokines that are important for NK cell recruitment.^{53,54}

Intriguingly, the expression of galectin-1 and other galectins, that are potent suppressors of antitumor immune surveillance, decreased in glioma-bearing mice treated with TMZ. Recently, galectin-1 was found responsible for the inhibition of NK cell function and viability: galectin-1 deficient gliomas could be eradicated by infiltrating NK cells with cytotoxic function.³⁰

The evidence that the "cytotoxicity signature" after three chemotherapy treatments resulted down-modulated in the periphery and up-regulated in the gliomas of TMZ-treated mice, supports the contention that TMZ can modulate glioma microenvironment facilitating NK cell infiltration and cytotoxic function.

It has long been known that NK cells can control and reinforce antitumor immune responses mediated by dendritic cells.⁵⁷ NK cells and DCs work in synergy, taking advantage of one another.^{58,59} The reciprocal interaction of NK cells and DCs and the ability of NK cells to recognize and kill tumor cells represent an important rationale for their monitoring during immunotherapeutic approaches with DCs in correlation with clinical outcomes. Our results obtained on a group of recurrent GBM patients treated with DCs loaded with autologous tumor lysate showed a significant increase of NK cells producing IFN-□ in patients with prolonged survival.¹¹ We are now evaluating the combination of DC immunotherapy with TMZ in patients with first diagnosis GBM. The NK cell response significantly correlates with survival, whereas the CD8⁺ T cell response does not appear to influence clinical outcomes. Patients with evidence of an NK cell response showed a significant up-regulation of ABCC3 in PBLs in comparison with non-responders (manuscript in preparation).

In agreement with these results, we observed that trafficking and NK cell homing increased in glioma-bearing mice during TMZ treatment. Notably, we also found that murine NK cells can efficiently overcome the drug-mediated toxicity of chemotherapy by expressing multidrugresistance genes. It has been previously described that NK cells are resistant to chemotherapy. ^{56,60} After chemotherapy treatment, NK cells are the first lymphoid cells to recover and may represent the principal lymphocytes for the initial months after treatment, suggesting a more rapid reaction to cytotoxic effect of drugs than other immune cells.⁴⁹ Murine and human NK cells were found to express high levels of multidrug transporters, which could confer protection against chemotherapeutic agents.⁶¹ TMZ administration, similar to other chemotherapeutics, can cause the development of moderate to severe lymphopenia and myelosuppression^{62,63}, indicating that immune inhibitory effects take place through selective toxicity on proliferating lymphocytes and inhibition of immune effector differentiation. 63,64 However, there is evidence that following a transient chemotherapyinduced lymphopenia, lymphocytes undergo homeostatic proliferation that enhances anti-tumor, vaccine-induced immune responses^{6,65,66} with positive clinical outcome correlation.⁶⁷ Discrepancies have been reported about the effect of TMZ on NK cells. It has been shown that TMZ significantly decrease the absolute number of CD3⁻ CD56⁺ effector cells in blood of GBM patients.⁵⁷ However, other studies have reported that circulating NK cells are relatively resistant to chemotherapy, with their frequency and absolute number, as well as their effector functions, unaffected by TMZ.^{55,56}

In conclusion, our data indicate that chemotherapy is able to modulate tumor microenvironment and reinforce tumor infiltration of NK effector cells and that can contribute to the adjuvant effect of chemotherapy. The different sensitivity of NK and T cells to TMZ, however, may disrupt their interactions, like that of 2B4 (CD244) and CD48⁶⁹⁷⁰, relevant to generate a T cell memory and possibly amplify anti-tumor responses. Confirmation of this hypothesis, driven by the results we reported here, may imply a careful re-evaluation of chemotherapy/immunotherapy schedules.

Materials & Methods

Cell culture

GL261 cells were cultured as neurospheres in a stem cell growth medium Technologies), DMEM-F12 Glutamax, B-27 (Life penicillin/streptomycin, human recombinant epidermal growth factor (EGF; 20 ng/mL) and human recombinant fibroblast growth factor-2 (FGF-2; 20 ng/mL; Peprotech). PBLs were cultured in RPMI 1640 containing 10% fetal bovine serum (FBS), β-mercaptoethanol, sodium acids. pyruvate, nonessential amino L-glutamine, and penicillin/streptomycin (all from EuroClone). Human recombinant interleukin-2 (hIL-2; 10 U/ml; Roche) was added to the medium.

In vivo experiments

C57BL/6N 6-week-old female mice (Charles River Laboratories) received intracranial injections of 10⁵ GL261 cells using specific stereotactic coordinates into the nucleus caudatum (0.7 mm posterior, 3 mm left lateral and 3.5 mm deep, with respect to the bregma). Mice were divided in 2 groups, treated 9 days after glioma implantation with intraperitoneal injections (i.p.) of 5 mg/kg TMZ or vehicle (DMSO) on days 1-5 and sacrificed at different time points. The animals were monitored every day and euthanized when suffering, in accordance with the current directives of the Campus animal IFOM-IEO house facility and the Minister of Health. Animal experiments were performed in accordance to the Italian Principle of Laboratory Animal Care (D.Lgs. 26/2014) and European Communities Council Directives (86/609/EEC and 2010/63/UE).

Isolation of local and systemic lymphocytes

PBLs were isolated using Lympholyte-M (Cedarlane Labs) according to the manufacturer's instructions. An indirect magnetic labeling system was used to immune-isolate CD8⁺ T and NK cells (NK and CD8⁺ T Cell Isolation Kits, Miltenyi Biotec) resulting in a 97±1.5% and 93.2±2.9% pure CD8⁺ T and NK cell population, respectively, as evaluated by flow cytometry. TILs were isolated with the Tumor Dissociation Kit and GentleMACS (Miltenyi Biotec) according to the manufacturer's instruction. A Percoll-density gradient centrifugation (30% - 40% - 80% - 100% isotonic Percoll, 400xg, 15 min at 20°C) was used to separate lymphocytes from the tumor single cell suspension. Immune cells were recovered from the 40-80 gradient interphase.

Flow cytometry

Cells were stained in a cold staining buffer at 4°C in the dark. The following antibodies were used: CD45, CD8, CD4, CD3, CD11b, CD27, NK1.1, NKp46, CD49b, CD49d, IFN-γ and pAkt (Miltenyi Biotec), Abcc3 (Abcam), B7-h3 (Biolegend) and Nkg2dl (eBioscience). DAPI was added to exclude dead cells. Flow cytometry acquisition was performed on a MACSQuant instrument, and data were analyzed with MACSQuantify Software (Miltenyi Biotec).

Microarray analyses and Real Time-PCR

Total RNA from purified NK and CD8⁺ T cells was extracted with TRIzol reagent (Life Technologies) using RNeasy Mini Kit and RNase-Free DNase Set (Qiagen). Microarray analyses were performed after three TMZ or DMSO administrations. Mouse Gene 2.0 ST Array GeneChip (Affymetrix), which includes 35,240 mouse transcripts, was used following standard procedures. Differentially expressed genes were identified using a fold-change threshold \geq 2 for all transcript comparisons. The functional annotation of genes that passed the FC and expression signal cut-offs was performed using the Gene Ontology (GO) Biological

Process category. Fast SYBR Green chemistry (Life Technologies) was used for real-time PCR expression analyses. Relative mRNA levels were measured using a ViiA7 Real Time-PCR System (Life Technologies) and calculated using the $\Delta\Delta$ Ct method normalizing to the housekeeping Gapdh, Actin and β 2M levels. The primer sequences (Primm S.r.l.) are reported in Table S2.

Migration assay

Migration was assessed in vitro using 8μm Transwell migration chambers. Purified NK cells (4x10⁵/transwell) were placed in the upper chamber and evaluated for their ability to transmigrate toward the lower chamber. Chemoattractant in the lower chamber was represented by medium from GL261 cells previously treated for 24 hours with DMSO or 150μM TMZ. After 12h, migrating cells were stained with crystal violet solubilized in 10% acetic acid.

Western blot analysis and proteome profiler array

Briefly, cells were washed with cold PBS and lysed in a buffer supplemented with protease and phosphatase inhibitors. Membranes with transferred proteins were incubated with the primary antibody anti-pAKT (Ser473) (1:1000, Cell Signaling) or anti-vinculin (1:10000). The primary antibody incubation was followed by incubation with peroxidase conjugated to the secondary antibody (anti-rat, 1:10000). A chemiluminescence reaction using the ECL Plus kit (GE Healthcare) was detected using G:BOX iChemi system (Syngene). Tumor relative levels of cytokines and chemokines were measured using the Mouse Cytokine Array Panel A kit (R&D Systems) following the manufacturer's instructions. Images of the blots were acquired with G:BOX Chemi system (Syngene) and quantitative analyses were performed using ImageJ

software. The 40 cytokines and chemokines of interest were normalized to the corresponding positive controls.

Apoptosis Assay

Resistance of NK and CD8 $^+$ T cells to the cytotoxic effects of TMZ was evaluated with Annexin V (Biolegend) and propidium (PI). Early and later apoptosis were distinguished with Annexin V positivity and Annexin V-PI double positivity, respectively. The selective MRP inhibitor MK571 at a concentration of 25 μ M was used to test the Abcc3 role in chemoresistance. Briefly, naïve lymphocytes were treated with 1 μ M TMZ or DMSO for 4 h in vitro. MK571 was added to the medium 30 minutes before or after the pharmacological treatment and apoptosis was evaluated.

ABC transporter activity

The eFluxx-ID® Green Multidrug-Resistance Assay (Enzo Life Sciences) was used to detect the multidrug-resistant phenotype of NK and CD8 $^+$ T cells by monitoring the efflux activity of the three major multidrug-resistance proteins: MDR1, MRP and BCRP. Following the manufactures' instructions, specific inhibitors (Verapamil, MK571 and Novobiocin) were used to define the resistance activity factor (MAF) in PBLs from naïve mice treated with 1 μ M TMZ or DMSO for 4 h in vitro. MAF values >25 are indicative of multidrug-resistance positive phenotype.

IFN-γ production and LDH assay

PBLs were cultured in a 24-well tissue culture plate in 1 mL of completed RPMI supplemented with 10 U/mL IL-2. Unspecific stimulation was performed with 50 ng/mL phorbol myristate acetate (PMA), 1 μ g/mL Ionomycin and 10 μ g/mL Brefeldin A for a total of 4 hrs. Lymphocytes were stained for IFN- γ (BD Cytofix/Cytoperm Fixation/Permeabilization Solution Kit, BD Bioscience) according to the manufacturer's instructions.

A non-radioactive cytotoxic assay (Cytotoxicity detection kit^{plus} LDH, Roche) was performed to test PBL capacity of TMZ-treated, control and naïve mice to recognize and lyse GL261 cells, according to manufactures' instructions. PBLs were pre-stimulated for 5 days in the presence of 20 Gy-irradiated GL261 cells and tested 5 days later for GL261 cell specific cytotoxicity.

Histology and immunofluorescence

Double immunofluorescence was performed on 5μm paraffin-embedded sections. Paraffin was removed with xylene and the sections were rehydrated in graded alcohol. Tumor sections were incubated with anti-CD8 (1:10, BD Bioscience) and anti-NK1.1 (1:10, Miltenyi) antibodies overnight at 4°C. After a counterstained with DAPI, sections were examined using a Nikon confocal microscope and analyses were performed on three to five independent fields per tumor using the 40X objective. Tumor sections were also stained with hematoxylin and eosin to assess the volume of tumor growth and acquired using the Aperio ScanScope slide scanner (Leica).

Statistical analysis

Statistical comparison were performed using two-tailed Student's T-test. Results were considered significant at p < 0.05.

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Figures

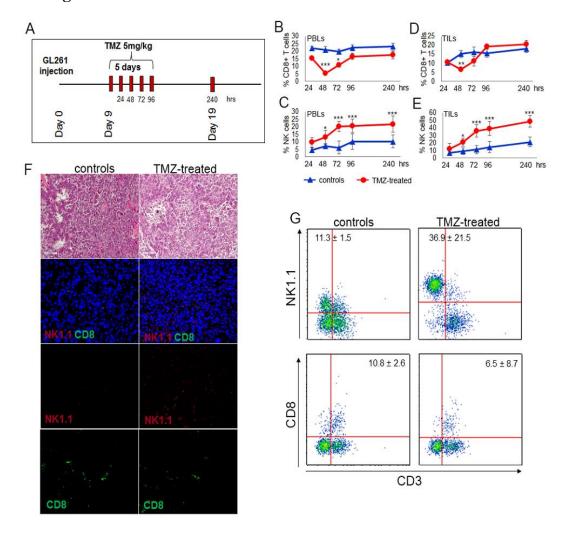


Figure 1. TMZ treatment influences local and peripheral immune cell frequency.

(A) Experimental schema of in vivo treatment. C57BL6 were i.c. injected with GL261 cells and treated for five days with i.p. injection of 5mg/kg TMZ or vehicle (DMSO) 9 days after tumor implantation. On days 9-13 and 19 after tumor implantation (24-96 and 240 hrs after TMZ treatment), n=5 mice per group/each time point were sacrificed for immune

monitoring. (B) Peripheral percentages of CD8⁺T cells (CD8⁺CD3⁺): 22.2 \pm 1.2% vs 15.5 \pm 0.2% at 24 hrs; 21.1 \pm 2.0% vs 5.5 \pm 1.0% at 48hrs; 19.8 \pm 1.4% vs 10.8 \pm 1.3% at 72hrs; 22.4 \pm 2.1 vs 16.6 \pm 2.1% at 96 hrs; 22.6 \pm 0.4% vs 17.3 \pm 1.4% at 240 hrs, controls vs TMZ-treated mice, respectively; *p < 0.01; ***p<0.0001. (C) Percentages of blood NK cells (NK1.1+CD3-): $4.8 \pm 1.2\%$ vs $9.8 \pm 3.2\%$ at 24 hrs; $7.2 \pm 1.2\%$ vs $13.3 \pm 1.2\%$ 1.2% at 48 hrs; $6.3 \pm 1.6\%$ vs $20.2 \pm 1.9\%$ at 72 hrs; $10.2 \pm 2.1\%$ vs 20.6 \pm 2.3% at 96 hrs, 7.2 \pm 2.1% vs 21.8 \pm 2.3% at 240 hrs, controls vs TMZtreated mice respectively; *p < 0.01; ***p<0.0001. (**D**) Tumor-infiltrating CD8⁺ T cells: $10.3 \pm 1.2\%$ vs $10.5 \pm 0.2\%$ at 24 hrs; $15.2 \pm 2.1\%$ vs 6.5 $\pm 1.0\%$ at 48 hrs; $16.3 \pm 2.7\%$ vs $11.2 \pm 2.6\%$ at 72 hrs; $15.7 \pm 2.2\%$ vs 19.3 \pm 1.1% at 96 hrs, 18.2 \pm 2.2% vs 20.4 \pm 2.3% at 240 hrs, controls vs TMZtreated mice respectively; **p < 0.001. (E) Tumor-infiltrating NK cells during and after TMZ administration: $7.2 \pm 12.5\%$ vs $12.5 \pm 7.4\%$ at 24 hrs; $9.1 \pm 4.5\%$ vs $21.1 \pm 5.3\%$ at 48 hrs; $11.3 \pm 5.6\%$ vs $36.2 \pm 8.1\%$ at 72 hrs; $14.5 \pm 7.6\%$ vs $39.2 \pm 9.2\%$ at 96 hrs; $21.4 \pm 6.5\%$ vs $48.9 \pm 7.7\%$ at 240 hrs, controls vs treated mice respectively; *p < 0.01; ***p<0.0001. (F) Characterization of immune infiltration in gliomas by IF at 72 hrs after beginning of TMZ treatment. (G) Quantitative determination of TILs by flow cytometry obtained from the same groups of gliomas used for IF studies.

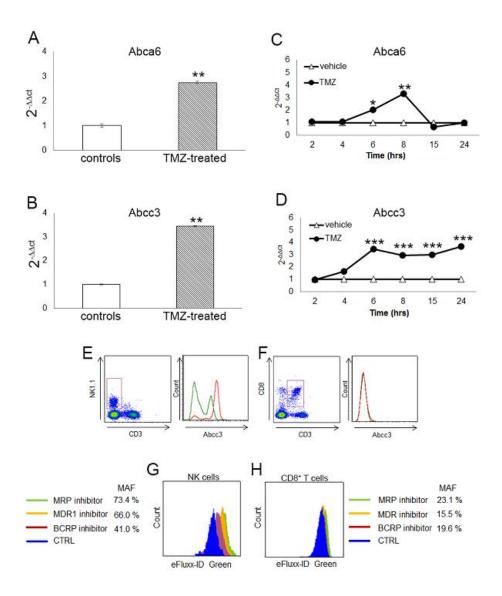


Figure 2. Abcc3 is responsible for NK cell chemo-resistance.

(**A and B**) Relative expression of Abca6 and Abcc3 transporters in blood NK cells at 72 hrs, **p < 0.001. (**C and D**) Time course of Abca6 and Abcc3 expression in PBLs from naïve mice treated in vitro with $1\mu M$ TMZ

or DMSO; *p = 0.01, **p < 0.005 and ***p < 0.0001. (**E and F**) Abcc3 levels in NK and CD8⁺ T cells from glioma-bearing mice after 3 treatments of TMZ (red line) or DMSO (green line). (**G and H**) MDR activity assay in NK and CD8⁺ T cells from PBLs of naïve mice treated in vitro with 1 μ M TMZ or DMSO for 4hrs. MAF values > 25 are indicative of multidrugresistant phenotype.

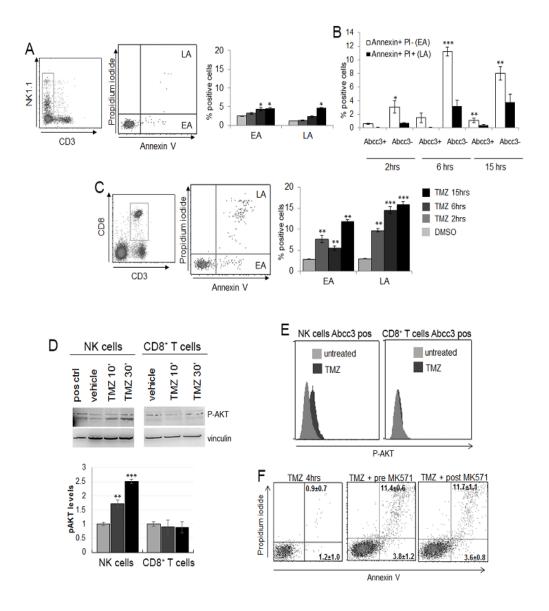


Figure 3. NK cells exhibit resistance to TMZ-induced apoptosis.

(**A and C**) Early and late apoptosis (EA and LA, respectively) induced by TMZ treatment in NK and CD8⁺ T cells from PBLs of naïve mice. (**B**) EA and LA of Abcc3⁺ and Abcc3⁻ NK cells treated in vitro with 1 μM TMZ.

(**D**) WB analysis and densitometric quantification of pAkt in blood immune-separated NK and CD8⁺ T cells of naïve mice treated with 1 μ M TMZ or DMSO. (**E**) Intracellular staining of pAkt in blood-derived Abcc3⁺ NK and CD8⁺ T cells from glioma-bearing mice at 72hrs. (**F**) Representative dot plots showing apoptosis in NK cells treated for 4 hrs in vitro with 1 μ M TMZ and 25 μ M Abcc3 inhibitor added to the medium 30 minutes before (pre MK571) or after (post MK571) the pharmacological treatment. *p < 0.01, **p < 0.001 and ***p < 0.0001.

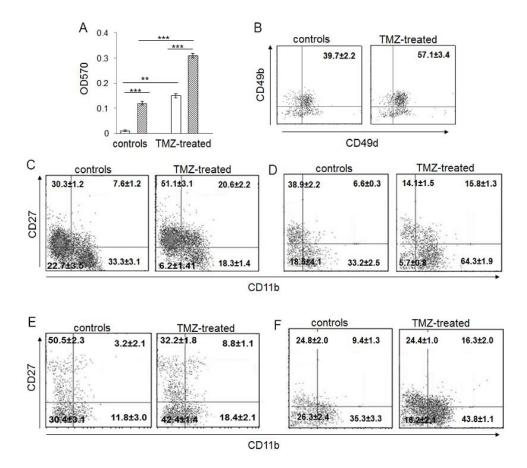


Figure 4. The developmental stage of NK cells from TMZ-treated mice was influenced by TMZ. (A) Migration of NK cells from PBLs of TMZ-treated and control mice at 72hrs toward medium from DMSO-treated (white bars) and TMZ-treated (striped bars) GL261 cells. **p < 0.005; ***p < 0.0001. (B) CD49b and CD49d integrin surface expression in blood-derived NK cells from TMZ-treated and control mice isolated at 72 hrs; p < 0.001. (C and D) Representative dot plots of the NK cell four-developmental stages basing on surface expression of CD27 and CD11b in blood of TMZ- and vehicle-treated mice on days 12 (C) and 19 (D); p < 0.01. (E and F) Representative dot plots showing CD27 and CD11b expression in tumor-infiltrating NK cells isolated on day 12 (E) and 19 (F) from gliomas, p < 0.005.

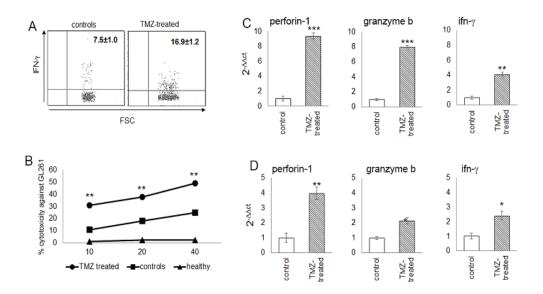


Figure 5. Cytotoxicity of glioma-infiltrating NK cells is promoted by TMZ treatment.

(A) IFN- γ production in blood NK cells (n = 4/group); p < 0.005. (B) Cytotoxic specificity of peripheral NK cells from naïve mice, TMZ- and vehicle-treated glioma-bearing mice (n = 10/group) against GL261 cells in vitro. Effector: target ratio (E:T) 10:1, 20:1, 40:1. p < 0.001. (C and D) Relative expression of *Prf1*, *GzmB* and *Ifn-\gamma* in TILs at early (C) and later time point (D). *p < 0.01; **p < 0.001; ***p < 0.0001.

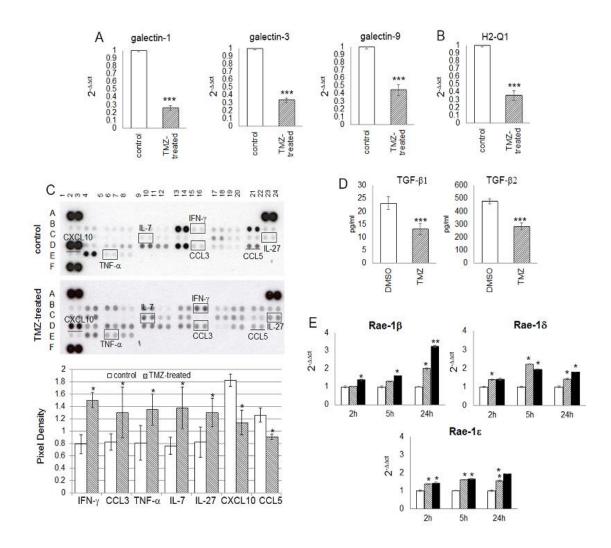


Figure 6. TMZ modifies glioma microenvironment favoring an NK cell antitumor activity.

(**A and B**) Relative expression of the most immunosuppressive molecules normally expressed in glioma cells: galectin-1, -3, -9 (3.85 \pm 0.03-, 2.94 \pm 0.02- and 2.22 \pm 0.03-fold, respectively vs controls) and H2-q1 (2.81 \pm 0.08-fold vs controls). (**C**) Cytokine and chemokine protein array blots with pixel density quantification of lysates from gliomas of control and TMZ-treated mice. Blots are representative of experiments performed on

a pool of gliomas from controls or TMZ-treated mice. Cytokines and chemokines which expression changed in both experiments were considered: IFN- γ , CCL3, TNF- α , IL-7 and IL-27 were 1.9 \pm 0.6-, 1.6 \pm 0.4-, 1.7 \pm 0.2-, 1.8 \pm 0.4-, 1.6 \pm 0.2-fold higher, CXCL10 and CCL5 were 0.6 \pm 0.7- and 0.7 \pm 0.4-fold lower in TMZ-treated compared to controls; * p \leq 0.01. (**D**) TGF- β 1 and - β 2 levels in medium of GL261 cells treated in vitro with DMSO or TMZ: from 23.2 \pm 2.5 pg/ml and 478.4 \pm 19.3 pg/ml in vehicle to 13.2 \pm 2.2 pg/ml and 283.8 \pm 25.9 pg/ml in TMZ-treated cells. (**E**) Relative expression of Nkg2d ligands in GL261 cells treated in vitro with 50 and 150 μ M TMZ (striped and black bars, respectively) and DMSO used as vehicle (white bars) at different time points. *p < 0.01; ***p < 0.001; ***p < 0.0001.

Supplementary

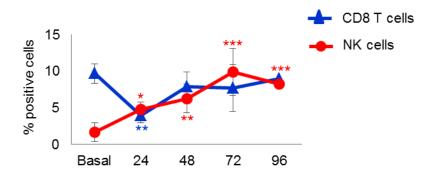


Figure S1. TMZ enhances the NK cell frequency in non-gliomabearing mice.

A group of non-glioma-bearing mice were treated with TMZ. PBLs were harvested 24, 48, 72 and 96 hrs after treatment. TMZ induced a modulation of CD8 $^+$ T cells and NK cells similar to TMZ-treated tumor bearing mice (Figure 1B and D). Specifically, NK cell frequency increased significantly at early time points (* p < 0.01; ** p < 0.005; p*** p<0.001), CD8 $^+$ T cells decrease was rapid but reversible (** p < 0.005).

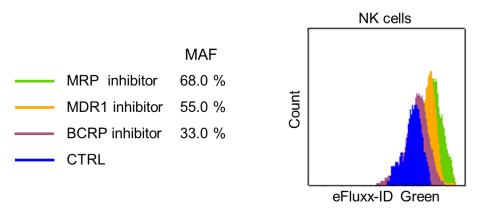


Figure S2. Peripheral blood NK cells show a dose-dependent drugresistance activity.

Efflux activity of ABC transporters assessed by flow cytometry with a fluorescent dye in presence or absence of specific ABC inhibitors to calculate the multidrug-resistance activity factor values (MAF). NK cells from naïve mice were treated in vitro with 0,5 μ M TMZ or DMSO for 4 hrs. MAF values > 25 are indicative of multidrug-resistant phenotype.

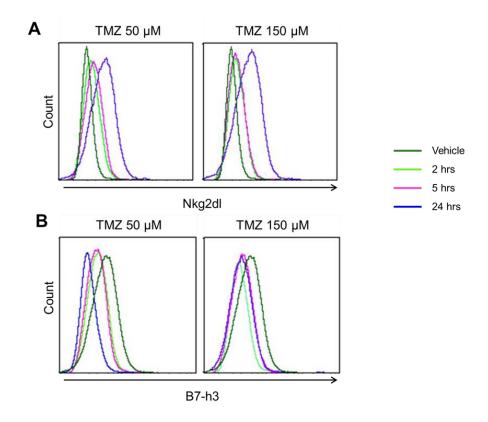


Figure S3. TMZ influences the GL261 cell immunogenicity.

Representative dot plots of GL261 cells treated in vitro with 50 and 150 μ M TMZ or DMSO (vehicle) at different time points. (**A**) Nkg2dl rapidly increased (4.1± 0.2% vs 9.4 ± 1.1% and 10.6 ± 0.9% at 2 hrs; 3.2 ± 1.0% vs 13.6 ± 1.0% and 17.6 ± 1.3% at 5 hrs;3.9 ± 0.6% vs 20.2 ± 2.3% and 27.7 ± 1.3% at 24hrs, controls vs. 50 and 150 μ M TMZ-treatedmice respectively; *p < 0.01). (**B**) B7-H3 significantly decreased (23.9± 0.2% vs 18.4 ± 1.5% and 6.1 ± 1.6% at 2 hrs; 23.2 ± 1.0% vs 10.7 ± 1.0% and 6.0 ± 1.5% at 5 hrs;23.9 ± 0.6% vs 4.8 ± 0.3% and 6.4 ± 0.9% at 24hrs, controls vs. 50 and 150 μ M TMZ-treated mice respectively; *p < 0.01). Flow cytometry analyses were obtained from 3 different experiments.

Table S1.

Probe Set ID	NK CTRL	NK TMZ	CD8 CTRL	CD8 TMZ	logFC NK	logFC CD8	Gene Symbol	Gene Bank
17362539	7,72202	3,242859	2,866335	2,828359	-4,479161	-0,037976	Scgb1a1	NM_011681
17306991	5,994791	2,021417	2,445533	2,329827	-3,973374	-0,115706	Gzmd	NM_010372
17306983	6,139857	2,709189	2,941385	3,022386	-3,430668	0,081001	Gzme	NM_010373
17489363	6,428623	3,300454	2,478789	2,344999	-3,128169	-0,13379	Hamp	NM_032541
17466530	8,309629	5,47775	3,184305	3,266585	-2,831879	0,08228	Rny1	NR_004419
17306999	5,211462	2,416042	2,940237	2,979622	-2,79542	0,039385	Gzmg	NM_010375
17235011	7,567307	4,952219	3,304193	3,500477	-2,615088	0,196284	Elane	NM_015779
17283634	6,251194	3,755193	3,030057	3,156768	-2,496001	0,126711	Serpina1c	NM_009245
17438886	9,429761	7,098767	4,373413	3,481985	-2,330994	-0,891428	Alb	NM_009654
17307025	4,977774	2,666936	2,911402	3,095509	-2,310838	0,184107	Gzmc	NM_010371
17351831	6,627881	4,440394	7,413254	6,670288	-2,187487	-0,742966	Snord58b	NR_028552
17305520	7,387096	5,22775	3,074114	2,902435	-2,159346	-0,171679	Ear1	NM_007894
17477832	8,77969	6,654347	4,169685	5,017468	-2,125343	0,847783	Ftl1	NM_010240
17312032	7,696645	5,612951	3,031745	2,946713	-2,083694	-0,085032	Ndrg1	NM_008681
17285706	10,43502	8,39452	5,278312	5,203849	-2,0405	-0,074463	Hist1h4m	NM_001195421
17264036	9,386904	7,403713	7,042769	6,920182	-1,983191	-0,122587	Ubb	ENSMUST00000019649
17250742	8,66349	6,7166	7,953413	7,965178	-1,94689	0,011765	Snord49a	NR_028550
17547723	7,702611	5,854451	5,840998	6,089122	-1,84816	0,248124	Ywhaq	NM_011739
17548348	9,131264	7,313562	6,380653	6,421731	-1,817702	0,041078	Actg1	NM_009609
17218680	9,861839	8,075171	8,931215	8,56097	-1,786668	-0,370245	Snord47	NR_028543
17314679	7,01271	5,22708	5,930602	6,067333	-1,78563	0,136731	Tuba1b	NM_011654
17236800	7,39912	5,62707	4,930883	4,31751	-1,77205	-0,613373	Dcn	NM_007833
17232731	7,586909	5,842309	6,427805	6,052614	-1,7446	-0,375191	Rnu3a	NR_002842
17339392	8,181211	6,505793	4,807532	4,453646	-1,675418	-0,353886	Myl12a	ENSMUST00000024846
17547608	8,508842	6,837495	5,176863	5,225361	-1,671347	0,048498	Actg1	NM_009609
17471778	7,76548	6,138906	2,566724	2,469116	-1,626574	-0,097608	Klra3	NM_010648
17246936	7,113566	5,522482	3,727224	3,828931	-1,591084	0,101707	Rasl10a	NM_145216
17225173	10,7736	9,208171	10,10508	9,887005	-1,565429	-0,218075	Snord82	NR_002851
17254877	7,119788	5,560782	3,753373	3,839166	-1,559006	0,085793	Мро	ENSMUST00000020779
17324398	7,262779	5,705933	7,378842	7,178941	-1,556846	-0,199901	Snora81	NR_034048
17357659	8,12729	6,61431	7,682272	8,015916	-1,51298	0,333644	Ms4a4b	NM_021718
17547620	8,859197	7,348626	6,197084	6,349689	-1,510571	0,152605	Ftl1	NM_010240
17471757	10,1179	8,609843	2,713829	3,135477	-1,508057	0,421648	Klra9	NM_010651
17295032	7,459177	5,975341	3,641421	3,915324	-1,483836	0,273903	Arsb	NM_009712
17356556	8,938833	7,455145	8,78065	8,328193	-1,483688	-0,452457	Malat1	NR_002847
17305986	10,50259	9,055317	10,01262	10,27451	-1,447273	0,26189	Rpph1	NR_002142
17329023	8,605267	7,160343	10,82889	10,93757	-1,444924	0,10868	Iglv1	M94350
17471768	9,013568	7,593667	3,737961	3,716724	-1,419901	-0,021237	Klra10	NM_008459
17349552	8,164172	6,808004	7,231226	7,424065	-1,356168	0,192839	Snora74a	NR_002905
17306975	6,341461	4,988468	2,87951	2,807576	-1,352993	-0,071934	Ctsg	NM_007800
17547869	9,344457	7,997941	7,318279	7,238977	-1,346516	-0,079302	Ywhaz	AK139259
17306937	7,291237	5,946251	2,916593	2,985589	-1,344986	0,068996	Cma1	NM_010780
17228102	8,13068	6,802405	5,947146	5,78095	-1,328275	-0,166196	Arpc5	NM_026369
17304510	7,301034	5,990347	7,204192	6,906984	-1,310687	-0,297208	Snord69	NR_028531
17308731	9,176225	7,885982	8,533373	8,526903	-1,290243	-0,00647	Tpt1	NM_009429
17318020	7,034488	5,746267	9,09833	9,27974	-1,288221	0,18141	Ly6d	NM_010742

17265126	NINA 01016A	Fun1	0.222620	-1,255397	6.461220	6 22060	7 5 6 1 4 2 2	0.016020	17221275
17258664 8,899.17 7,6708.25 5,1028.48 5,358.63 -1,226.652 0,2557.82 Fasi 17358271 9,989287 8,824363 9,68526 9,14133 -1,164924 -0,54393 Cycs E 17254283 7,990671 4,455153 4,639244 -1,141065 0,184091 Ccl4 1731059 7,339288 6,215622 2,914825 2,850724 -1,123666 -0,064101 Kira1 17301932 7,296147 6,183538 3,109515 3,151845 -1,112609 0,04233 Esd 17321086 8,713758 7,60283 6,662064 6,585664 -1,110928 -0,0704 Grg5 17262174 7,99739 6,859603 7,351913 7,409043 -1,107936 0,05713 Grb211 17230664 7,834934 6,736828 10,45232 9,548893 -1,098106 -0,903427 Myb 17300264 7,903318 6,810907 7,05496 7,324223 -1,092351 -0,381273 Snora34 17330567 6,903749 5,834694 5,866147 6,026212 -1,069055 0,160065 Ube2d2a 17250740 10,88953 9,834885 10,23115 10,32846 -1,054645 0,09731 Snord49b 17211416 7,314589 6,268205 2,887611 2,48839 -1,046384 -0,039252 Krbc1a 17209668 6,89051 5,851244 3,465873 3,27207 -1,039266 0,193803 Mcpt8 17210707 7,418639 6,389576 4,594179 4,386136 -1,029608 0,208043 Kira6 17300733 10,94476 9,924092 5,745749 6,068019 -1,020668 0,32227 Gzmb 17215079 6,185431 5,16647 5,207719 5,148394 -1,018961 -0,059925 83gm7 17312679 8,37664 7,36377 7,314499 7,190969 -1,01287 -0,12248 Apol7e 1735148 7,941168 6,930844 7,214105 7,279716 -1,010324 0,065611 Snord22 1723116 6,516153 5,521481 7,197991 7,295009 -0,994672 0,095618 Gstp2 1731364 8,558916 7,580473 9,948925 9,481834 -0,097925 3,00000 3,000000 3,0000000 3,0000000000	NM_010164	,		· ·					
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17262174	NM_016903		-			-		<u> </u>	
17239664	NM_010318	_		· ·	· ·	-		-	
17302054	NM_008143					-			
17321241	NM_001198914	'		-	-	-			
17353673	NR_028481	Snora31		-1,092351		7,705496		<u> </u>	17302054
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17211416	NM_019912	Ube2d2a	0,160065	-1,069055	6,026212	5,866147	5,834694	6,903749	17353673
17306968	NR_028526	Snord49b	0,09731	-1,054645	10,32846	10,23115	9,834885	10,88953	17250740
17229187	NM_183322	Khdc1a	-0,039252	-1,046384	2,848359	2,887611	6,268205	7,314589	17211416
17471707	NM_008572	Mcpt8	-0,193803	-1,039266	3,27207	3,465873	5,851244	6,89051	17306968
17307033	NM_008510	Xcl1	-0,122188	-1,032787	5,516995	5,639183	9,605013	10,6378	17229187
17215097	NM_008464	Klra6	-0,208043	-1,029063	4,386136	4,594179	6,389576	7,418639	17471707
17312679	NM_013542	Gzmb	0,32227	-1,020668	6,068019	5,745749	9,924092	10,94476	17307033
17357148	NM_145222	B3gnt7	-0,059325	-1,018961	5,148394	5,207719	5,16647	6,185431	17215097
17231116	NM_001134802	Apol7e	-0,12348	-1,01287	7,190969	7,314449	7,36377	8,37664	17312679
17231116	NR 004445	Snord22	0,065611	-1,010324	7,279716	7,214105	6,930844	7,941168	17357148
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17231477 9,1174 8,305917 3,192828 3,214253 -0,811483 0,021425 Myct1	ENSMUST00000110293	Serpinb6b	0,097902	-0,837692	5,486094	5,388192	8,197952	9,035644	17286354
	NM_031395	Sytl3	-0,505582	-0,823827	7,7073	8,212882	9,544233	10,36806	17340532
17330751 8,75443 7,944432 9,089464 8,598955 -0,809998 -0,490509 Cd96	NM_026793	Myct1	0,021425	-0,811483	3,214253	3,192828	8,305917	9,1174	17231477
	NM_032465	Cd96	-0,490509	-0,809998	8,598955	9,089464	7,944432	8,75443	17330751
17219435 6,612782 5,807183 9,426852 9,5171 -0,805599 0,090248 Slamf6	NM_030710						5,807183		
17332317 6,368155 5,564988 6,241686 6,450589 -0,803167 0,208903 Pigp	NM_001159619								
	NM_026793 NM_032465 NM_030710	Myct1 Cd96 Slamf6	0,021425 -0,490509 0,090248	-0,811483 -0,809998 -0,805599	3,214253 8,598955 9,5171	3,192828 9,089464 9,426852	8,305917 7,944432 5,807183	9,1174 8,75443 6,612782	17231477 17330751 17219435

17327613	8,441427	7,639778	7,565238	7,518968	-0,801649	-0,04627	Nlrc3	NM_001081280
17343426	9,015028	8,213956	8,712572	8,520144	-0,801072	-0,192428	Rasal3	NM_178785
17262600	8,569902	7,772296	10,95838	10,53023	-0,797606	-0,42815	Tcf7	ENSMUST00000072425
17229391	8,958097	8,160543	9,765763	9,347804	-0,797554	-0,417959	Uck2	NM_030724
17296286	10,53042	9,736224	5,335327	5,123867	-0,794196	-0,21146	Gzma	NM_010370
17212252	9,681068	8,889685	5,093229	5,293192	-0,791383	0,199963	Il18rap	ENSMUST00000027237
17308800	7,871169	7,083346	7,71402	7,596732	-0,787823	-0,117288	Dnajc15	NM_025384
17425276	9,582831	8,797203	6,357544	6,544458	-0,785628	0,186914	Amd1	NM_009665
17216976	7,15169	6,366389	8,953716	9,296986	-0,785301	0,34327	Faim3	ENSMUST00000038829
17342027	8,325926	7,540999	7,258554	7,163834	-0,784927	-0,09472	Snora78	NR_028515
17236531	9,51541	8,738447	8,522184	8,313869	-0,776963	-0,208315	Lta4h	NM_008517
17471744	9,6378	8,862185	4,052271	4,167145	-0,775615	0,114874	Klra14	NM_008461
17243458	9,718661	8,944473	9,732668	9,571295	-0,774188	-0,161373	Rpl23a	NM_207523
17219011	6,447505	5,677557	7,381426	6,774175	-0,769948	-0,607251	Pou2f1	NM_011137
17268820	9,263456	8,497277	9,60421	9,592226	-0,766179	-0,011984	Ikzf3	NM_011771
17281419	8,160708	7,396818	8,305159	8,238655	-0,76389	-0,066504	Fkbp3	NM_013902
17289986	7,534341	6,774198	4,048173	4,140341	-0,760143	0,092168	Cdc20b	XM_138861
17241181	10,00853	9,252814	4,06883	4,06882	-0,755716	0	Adamts14	NM_001081127
17227025	8,63718	7,888536	7,874898	7,820453	-0,748644	-0,054445	Lax1	NM_001159649
17340629	9,74552	8,99758	8,46234	8,285722	-0,74794	-0,176618	Mir692-1	NR_030465
17351384	9,285126	8,537655	8,262767	8,408219	-0,747471	0,145452	Sec11c	NM_025468
17356173	8,65758	7,910756	8,37185	8,335217	-0,746824	-0,036633	Ptprcap	ENSMUST00000061086
17335330	9,559889	8,816904	9,749522	9,945054	-0,742985	0,195532	Rpl10a	NM_011287
17268445	10,23153	9,492514	5,510755	5,70501	-0,739016	0,194255	Tbx21	NM_019507
17226622	7,168702	6,434578	8,563266	8,657395	-0,734124	0,094129	Cd55	NM_010016
17290682	11,02663	10,29295	10,26991	10,21478	-0,73368	-0,05513	Ywhaq	NM_011739
17217566	8,743455	8,016308	7,829902	7,54802	-0,727147	-0,281882	Ptpn7	NM_177081
17308280	9,241116	8,521441	8,624108	8,770431	-0,719675	0,146323	Ррр3сс	NM_008915
17329345	3,740905	3,025362	2,788297	2,784504	-0,715543	-0,003793	Crygs	NM_009967
17348833	5,346639	4,632923	3,660481	3,628634	-0,713716	-0,031847	Ttr	NM_013697
17231229	7,278599	6,566498	6,328564	6,240336	-0,712101	-0,088228	Hsd11b1	ENSMUST00000016338
17229178	9,667757	8,957817	8,188601	7,813097	-0,70994	-0,375504	Atp1b1	NM_009721
17232394	5,073312	4,363488	9,812919	8,944581	-0,709824	-0,868338	Themis	ENSMUST00000056097
17323828	8,027083	7,319416	9,449989	9,728542	-0,707667	0,278553	B3gnt5	NM_001159407
17383127	7,6682	6,968411	6,989335	7,271997	-0,699789	0,282662	Snora43	NR_028572
17284247	7,976331	7,276897	8,597324	8,410034	-0,699434	-0,18729	Gpr132	NM_019925
17309644	8,015487	7,327188	8,120798	8,666903	-0,688299	0,546105	Gpr18	NM_182806
17211867	8,066287	7,378091	8,145213	7,481831	-0,688196	-0,663382	Zap70	NM_009539
17262102	8,574433	7,886278	9,694916	9,317152	-0,688155	-0,377764	Itk	NM_010583
17337110	9,637356	8,949302	8,988371	9,101721	-0,688054	0,11335	H2-Q5	NR_051981
17507462	7,133271	6,446615	7,377557	7,209774	-0,686656	-0,167783	Ankrd10	NR_030779
17335628	6,7878	6,101765	7,864737	7,407816	-0,686035	-0,456921	Dnahc8	NM_013811
17345926	7,694689	7,010238	9,67535	9,05047	-0,684451	-0,62488	Satb1	ENSMUST00000144331
17330544	7,453719	6,769765	8,367424	8,487717	-0,683954	0,120293	Sidt1	NM_001159419
17256823	7,597329	6,916411	4,491824	4,06953	-0,680918	-0,422294	Arl4d	NM_025404
17217247	6,422698	5,747318	7,675901	7,838312	-0,67538	0,162411	Pik3c2b	NM_001099276
17266362	9,867433	9,195583	9,996347	10,00752	-0,67185	0,011173	Rpl23a	NM_207523
17335550	8,462671	7,791561	8,043688	7,703098	-0,67111	-0,34059	Tbc1d22b	NM_198647
17336476	7,195491	6,525324	9,606166	9,856952	-0,670167	0,250786	H2-Ob	NM_010389
17292117	7,507554	6,837662	5,226664	5,06265	-0,669892	-0,164014	Gfod1	NM_001033399
17219397	9,409743	8,747982	8,108435	8,425161	-0,661761	0,316726	Cd48	ENSMUST00000068584
17319401	6,992645	6,332792	6,944668	6,644951	-0,659853	-0,299717	Snord83b	NR_028282
17318100	10,88581	10,22758	9,696198	9,64756	-0,65823	-0,048638	Ly6c2	NM_001099217
17248064	6,053524	5,395494	6,173096	6,231428	-0,65803	0,058332	Mtif2	NM_133767
	-,	-,	-, 5050	-,	-,-5000	-,		

17248894	7,56391	6,91022	7,977855	7,748202	-0,65369	-0,229653	Snord95	NR_028564
17212724	10,62824	9,975967	9,806267	9,843521	-0,652273	0,037254	Stat4	NM_011487
17215820	8,188369	7,538746	5,001016	4,968524	-0,649623	-0,032492	Gpc1	NM_016696
17249980	10,05675	9,408862	7,838535	7,730333	-0,647888	-0,108202	Igtp	NM_018738
17283871	6,381108	5,733329	10,0225	9,14764	-0,647779	-0,87486	Bcl11b	ENSMUST00000066060
17331173	7,873653	7,2305	6,054777	6,184859	-0,643153	0,130082	St3gal6	ENSMUST00000137035
17258118	8,615083	7,972767	6,738274	6,69655	-0,642316	-0,041724	Rab37	NM 001163753
17306785	8,420643	7,778876	7,792899	7,73721	-0,641767	-0,055689	Nedd8	NM 008683
17313050	9,564374	8,924964	9,949554	10,26131	-0,63941	0,311756	Apobec3	NM 001160415
17220475	7,509834	6,87098	9,259194	8,817899	-0,638854	-0,441295	Dusp10	ENSMUST00000048655
17309580	6,074012	5,43732	6,551866	6,678033	-0,636692	0,126167	Dock9	NM 001081039
17548405	8,764709	8,131117	7,932292	7,820493	-0,633592	-0,111799	Myl12a	ENSMUST00000123686
17278268	6,947299	6,318162	6,245343	6,50313	-0,629137	0,257787	Serpina3f	NM 001168294
17229767	8,692522	8,06673	8,438124	8,665705	-0,625792	0,227581	Ly9	NM 008534
17214053	6,552958	5,928838	7,361716	7,348809	-0,62412	-0,012907	Xrcc5	NM 009533
17323473	8,297984	7,674548	7,720813	7,557796	-0,623436	-0,163017	Crkl	NM 007764
17286375	9,415531	8,792261	4,403741	4,416109	-0,62327	0,012368	Serpinb9b	ENSMUST00000006392
17251518	7,858993	7,238272	8,915357	8,997569	-0,620721	0,082212	Vamp2	NM 009497
17298791	10,30448	9,686543	10,31118	10,28728	-0,617937	-0,0239	Rpl23a	NM 207523
17336649	5,579643	4,964034	7,517755	6,389677	-0,615609	-1,128078	Prrt1	ENSMUST00000015620
17226043	9,665359	9,053116	9,322617	9,368759	-0,612243	0,046142	Bcl2	NM 009741
17232853	8,603196	7,991085	7,621121	7,669891	-0,612111	0,04877	Mical1	NM 138315
17223574	8,005469	7,396096	8,322408	8,05392	-0,609373	-0,268488	Sumo1	NM 009460
17214983	8,228064	7,622594	7,870612	7,599275	-0,60547	-0,271337	Cab39	NM 133781
17294974	8,579552	7,977632	8,450578	8,327712	-0,60192	-0,122866	Papd4	NM_133905
17266967	11,24631	10,64597	5,718506	7,141956	-0,60034	1,42345	Ccl3	NM 011337
17306690	6,931663	6,333368	5,60846	5,38979	-0,598295	-0,21867	Dcaf11	ENSMUST00000133256
17237937	9,804963	9,208323	10,07581	10,09739	-0,59664	0,02158	B4gaInt1	NM 008080
17366278	8,679462	8,084273	9,162449	9,135664	-0,595189	-0,02138	Sp100	ENSMUST00000141998
17334147	8,354542	7,760951	9,794441	9,650993	-0,593591	-0,143448	Mir5125	NR 039586
17303472	9,876981	9,284418	9,61978	9,204252	-0,593591	-0,143448	Ptma	NM 008972
17348054	7,463392	6,870926	7,23273	7,429677	-0,592466	0,196947	Mettl4	NM 176917
17219542	7,403392	6,746457	7,964671	7,715325	-0,592400	-0,249346		NM 026234
						-	Pigm Hist1h2bc	_
17285851 17217619	11,86335	11,27338	8,265327	8,410759	-0,58997	0,145432	Phlda3	NM_023422
17217619	7,122962 5,564948	6,533694 4,976171	4,994402 7,482156	5,072389 6,778565	-0,589268 -0,588777	-0,703591	Rcsd1	NM_013750 AK030934
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17257085	8,504443	7,920609	7,905697	7,050145	-0,583834	-0,855552	Hexim1	NM_138753
17342182	7,245903 5,607321	6,664486	5,909534	5,763075	-0,581417	-0,146459 0,10091	Baiap3	NM_001163270
17228788		5,026563	7,424744	7,525654	-0,580758		Mir1927	NR_035448
17231287	5,103554	4,522939	6,738204	6,651904	-0,580615	-0,0863	Cd46	ENSMUST00000162650
17324394	7,261841	6,685762	8,435038	8,076769	-0,576079	-0,358269	Snord2	NR_030705
17308556	9,055775	8,483037	5,86439	5,770777	-0,572738	-0,093613	Cysltr2	NM_133720
17293362	10,86562	10,29385	7,145607	6,9282	-0,57177	-0,217407	Ctla2a	NM_007796
17357306	7,347357	6,775671	6,608177	6,832348	-0,571686	0,224171	B3gat3	NM_024256
17224620	6,768267	6,198813	5,639653	5,926612	-0,569454	0,286959	Chpf	NM_001001565
17347210	7,166876	6,601146	6,018254	5,717389	-0,56573	-0,300865	Memo1	NM_133771
17218653	7,267068	6,70248	7,448176	7,384833	-0,564588	-0,063343	Gas5	NR_002840
17294222	8,51128	7,946997	9,256173	8,571746	-0,564283	-0,684427	Rpl9	ENSMUST00000080545
17345740	7,342778	6,778806	7,456551	7,398767	-0,563972	-0,057784	Foxp4	NM_001110824
17288896	10,56974	10,00764	9,081757	9,02475	-0,5621	-0,057007	Actg1	NM_009609
17229782	9,601611	9,040549	8,03731	8,42053	-0,561062	0,38322	Slamf7	NM_144539
17286365	9,762955	9,204165	6,711953	6,944495	-0,55879	0,232542	Serpinb9	ENSMUST00000063191
17312829	10,91237	10,35781	9,37781	9,346898	-0,55456	-0,030912	Lgals1	NM_008495
17284360	10,12793	9,574241	11,49501	11,71039	-0,553689	0,21538	Ighm	AB067787

17265030	8,777009	8,223427	8,799217	8,777853	-0,553582	-0,021364	Acap1	NM_153788
17222913	8,523819	7,976355	6,471075	6,595484	-0,547464	0,124409	Nab1	ENSMUST00000069792
17336190	6,969216	6,421898	6,079786	6,078209	-0,547318	-0,001577	Ndufa7	NM_023202
17278056	8,655529	8,108963	6,976704	7,079153	-0,546566	0,102449	Rin3	NM_177620
17289181	9,885352	9,340539	6,678686	7,168914	-0,544813	0,490228	Arsb	NM 009712
17225360	7,866826	7,323525	7,354815	6,904544	-0,543301	-0,450271	Arl4c	NM 177305
17329636	10,5496	10,00633	4,519377	4,065839	-0,54327	-0,453538	Gp5	NM 008148
17233694	10,6716	10,12976	6,68435	6,681859	-0,54184	-0,002491	Prf1	NM 011073
17353143	6,841847	6,302286	6,779686	6,590427	-0,539561	-0,189259	Rprd1a	NM 144861
17306272	7,103533	6,56557	7,005251	7,037005	-0,537963	0,031754	Snord8	NR 028542
17234339	8,580913	8,044294	7,66598	7,83583	-0,536619	0,16985	Adora2a	NM 009630
17275524	7,111043	6,574971	7,746321	7,52486	-0,536072	-0,221461	Srp54c	NM 001100110
17356194	7,388518	6,8529	7,512085	7,639639	-0,535618	0,127554	Clcf1	ENSMUST00000046506
17328958	7,394947	6,859927	4,261745	4,061539	-0,53502	-0,200206	Gp1bb	NM 001001999
17326336	7,189328	6,655676	5,506394	5,218909	-0,533652	-0,287485	Мрр5	NM 019579
17260364	8,331414	7,800178	7,768806	7,593963	-0,533032	-0,174843	Purb	NM 011221
17216563	7,337948	6,807835	7,708800	7,193344	-0,531230	-0,174843	Mki67ip	NM 026472
	8,440359			· ·				_
17303916		7,912599	7,858069	7,637767	-0,52776	-0,220302	Camk2g	NM_178597
17223793	6,712744	6,185215	4,588263	4,669118	-0,527529	0,080855	Fzd5	NM_022721
17297678	8,587286	8,059834	8,513708	7,980078	-0,527452	-0,53363	Cnot1	NM_178078
17315312	7,677176	7,150307	4,044717	4,359251	-0,526869	0,314534	Soat2	ENSMUST00000023806
17337065	6,816663	6,29005	6,570727	6,380967	-0,526613	-0,18976	H2-Q2	ENSMUST00000074806
17319009	10,84138	10,31536	8,721974	8,647979	-0,52602	-0,073995	II2rb	NM_008368
17320705	9,633808	9,107987	6,715457	6,907989	-0,525821	0,192532	Yaf2	NR_028315
17243051	10,27437	9,7493	8,798029	8,924009	-0,52507	0,12598	Mob3a	NM_172457
17242931	7,670016	7,145276	8,482405	8,466669	-0,52474	-0,015736	Tcf3	NM_001164147
17261417	7,354632	6,830637	4,281227	4,151361	-0,523995	-0,129866	Chac2	NM_026527
17212229	9,042225	8,518259	6,757498	7,141586	-0,523966	0,384088	Il18r1	NM_008365
17237915	7,683266	7,160321	7,512674	7,463905	-0,522945	-0,048769	Agap2	NM_001033263
17284795	7,671269	7,149072	7,534875	7,488251	-0,522197	-0,046624	Sp4	NM_009239
17255626	7,164841	6,643003	8,287263	7,724408	-0,521838	-0,562855	Skap1	NM_001033186
17280184	7,900993	7,379378	7,173523	7,285791	-0,521615	0,112268	Zfp125	AJ005350
17274658	7,954896	7,43544	6,310131	6,29651	-0,519456	-0,013621	Tssc1	NM_201357
17292941	7,599355	7,083307	7,799572	7,684948	-0,516048	-0,114624	Fam193b	NM_145382
17236339	8,570982	8,056081	7,569371	7,686084	-0,514901	0,116713	Gnptab	NM_001004164
17348850	7,117806	6,603748	7,42284	7,270076	-0,514058	-0,152764	Rnf138	NM_207623
17284037	9,858269	9,345045	9,022526	8,830379	-0,513224	-0,192147	Brp44l	NM 018819
17217145	8,085062	7,573195	8,398483	8,361729	-0,511867	-0,036754	Elk4	NM 007923
17239702	8,274086	7,762582	7,35342	7,107204	-0,511504	-0,246216	Tbpl1	NM 011603
17280842	7,299767	6,788553	6,336206	6,370578	-0,511214	0,034372	Ifrd1	ENSMUST00000001672
17337122	10,86945	10,35939	10,6478	10,65594	-0,51006	0,00814	H2-Q6	NM 207648
17289094	5,65484	5,144958	5,18576	5,323533	-0,509882	0,137773	Ankrd34b	ENSMUST00000168871
17343449	7,703459	7,193591	7,304299	7,122027	-0,509868	-0,182272	Pglyrp2	NM 021319
17293273	8,442746	7,934557	6,665953	6,375139	-0,508189	-0,290814	Isca1	NM 026921
17309340	7,865636	7,359308	5,659228	5,526604	-0,506328	-0,132624	Spry2	NM 011897
17230304	5,966102	5,461136	7,050429	6,985984	-0,504966	-0,064445	Smyd3	ENSMUST00000128302
17231859	10,70357	10,1993	10,53372	10,34626	-0,504900	-0,004443	Ifngr1	NM_010511
17302251	6,655107	6,151074	6,592021	6,633879		0,041858		NM_010311 NM_026474
					-0,504033		Sugt1	_
17284327	5,988863	5,487208	6,405217	5,919301	-0,501655	-0,485916	Ighg	AB097847
17240645	7,610886	7,109783	6,143065	6,523561	-0,501103	0,380496	Prdm1	NM_007548
17354212	6,555723	6,096356	6,955244	6,410246	-0,459367	-0,544998	Sumo2	NM_133354
17483301	6,555723	6,096356	6,955244	6,410246	-0,459367	-0,544998	Sumo2	NM_133354
17300173	5,167911	4,723652	9,845593	9,202727	-0,444259	-0,642866	Trav12-2	X04330
17349514	7,9982	7,5809	6,231769	7,173673	-0,4173	0,941904	Egr1	NM_007913

17217846	3,599054	3,190356	8,247457	6,893147	-0,408698	-1,35431	Mir181b-1	NR_029820
17330998	5,543056	5,150113	7,887892	7,242175	-0,392943	-0,645717	Cep97	NM_028815
17219407	10,61739	10,24315	7,882639	7,112572	-0,37424	-0,770067	Slamf1	NM_013730
17547965	5,862079	5,548983	6,617781	6,070514	-0,313096	-0,547267	Cbr1	NM_007620
17246803	6,90621	6,607695	6,20713	7,316542	-0,298515	1,109412	Osm	ENSMUST00000075221
17225169	8,387702	8,09376	9,488194	8,766941	-0,293942	-0,721253	Snora75	NR 028478
17215697	4,98603	4,710763	6,885127	5,980136	-0,275267	-0,904991	Ramp1	NM 001168392
17228353	7,584553	7,312157	8,530529	7,851785	-0,272396	-0,678744	Mr1	NM 008209
17216753	7,310708	7,048003	7,472921	6,965268	-0,262705	-0,507653	Mgat5	NM 145128
17330805	4,371064	4,13599	8,840948	7,944216	-0,235074	-0,896732	Trat1	NM 198297
17229658	11,33652	11,10233	7,533551	8,608191	-0,23419	1,07464	Fcer1g	NM 010185
17230111	8,455375	8,257897	4,345897	5,736881	-0,197478	1,390984	Ifi205	NM 172648
17222825	9,240271	9,04487	8,963826	8,402598	-0,195401	-0,561228	Obfc2a	NM 028696
17354219	5,162158	4,989333	3,493811	3,857747	-0,172825	0,363936	Trim36	NM 178872
17227797	11,51649	11,3647	6,526243	7,409307	-0,172823	0,883064	Rgs18	ENSMUST00000027603
		, , , , , , , , , , , , , , , , , , ,				-		ENSMUST00000172388
17227780	6,86869	6,727451	7,286811	7,835445	-0,141239	0,548634	Rgs1	
17213462	6,473576	6,347395	8,528661	7,98141	-0,126181	-0,547251	Cd28	NM_007642
17360842	8,921074	8,80429	8,246848	8,340968	-0,116784	0,09412	Rps12	NM_011295
17289037	7,305953	7,201729	9,245312	8,311029	-0,104224	-0,934283	Ssbp2	NM_024272
17356583	6,609526	6,528538	7,458825	7,658788	-0,080988	0,199963	Syvn1	NM_028769
17295038	8,448845	8,378065	4,964139	5,573284	-0,07078	0,609145	Scamp1	ENSMUST00000022197
17313433	5,233052	5,168207	7,865443	7,261367	-0,064845	-0,604076	Xrcc6	ENSMUST00000100399
17300360	7,971485	7,914164	9,144283	8,977753	-0,057321	-0,16653	Pabpn1	NM_019402
17214685	4,339372	4,296576	6,619501	5,902818	-0,042796	-0,716683	Acsl3	NM_028817
17289792	3,060398	3,020478	8,447563	7,356251	-0,03992	-1,091312	Mir1904	NR_035442
17321263	6,800642	6,81018	7,478119	6,448468	0,009538	-1,029651	Adcy6	NM_007405
17260137	6,715715	6,747581	9,148178	8,215382	0,031866	-0,932796	Kremen1	NM_032396
17290695	5,639749	5,682642	7,909864	7,998283	0,042893	0,088419	Gpr137b	NM_031999
17253912	7,120241	7,164826	7,733023	7,594581	0,044585	-0,138442	Rhot1	NM_001163354
17220407								
17228497	2,782127	2,828932	7,964214	6,536144	0,046805	-1,42807	Tdrd5	NM_001134741
17353676	2,782127 4,489139	2,828932 4,544283	7,964214 4,165155	6,536144 3,689264	0,046805 0,055144	-1,42807 -0,475891	Tdrd5 Nrg2	NM_001134741 ENSMUST00000115713
	, -			-				
17353676	4,489139	4,544283	4,165155	3,689264	0,055144	-0,475891	Nrg2	ENSMUST00000115713
17353676 17297374	4,489139 5,642543	4,544283 5,705226	4,165155 7,109665	3,689264 6,36235	0,055144 0,062683	-0,475891 -0,747315	Nrg2 Nkiras1	ENSMUST00000115713 NM_023526
17353676 17297374 17213548	4,489139 5,642543 4,067666	4,544283 5,705226 4,134457	4,165155 7,109665 6,286251	3,689264 6,36235 6,871441	0,055144 0,062683 0,066791	-0,475891 -0,747315 0,58519	Nrg2 Nkiras1 Nrp2	ENSMUST00000115713 NM_023526 NM_001077406
17353676 17297374 17213548 17374488	4,489139 5,642543 4,067666 11,25486	4,544283 5,705226 4,134457 11,32608	4,165155 7,109665 6,286251 8,187112	3,689264 6,36235 6,871441 10,18821	0,055144 0,062683 0,066791 0,07122	-0,475891 -0,747315 0,58519 2,001098	Nrg2 Nkiras1 Nrp2 Thbs1	ENSMUST00000115713 NM_023526 NM_001077406 NM_011580
17353676 17297374 17213548 17374488 17312700 17288145	4,489139 5,642543 4,067666 11,25486 7,76849 5,896847	4,544283 5,705226 4,134457 11,32608 7,842728 5,982243	4,165155 7,109665 6,286251 8,187112 7,889555 5,876764	3,689264 6,36235 6,871441 10,18821 8,438079 6,039476	0,055144 0,062683 0,066791 0,07122 0,074238 0,085396	-0,475891 -0,747315 0,58519 2,001098 0,548524 0,162712	Nrg2 Nkiras1 Nrp2 Thbs1 Ncf4 Habp4	ENSMUST00000115713 NM_023526 NM_001077406 NM_011580 NM_008677 NM_019986
17353676 17297374 17213548 17374488 17312700 17288145 17332084	4,489139 5,642543 4,067666 11,25486 7,76849 5,896847 6,933	4,544283 5,705226 4,134457 11,32608 7,842728 5,982243 7,036024	4,165155 7,109665 6,286251 8,187112 7,889555 5,876764 7,462723	3,689264 6,36235 6,871441 10,18821 8,438079 6,039476 7,457321	0,055144 0,062683 0,066791 0,07122 0,074238 0,085396 0,103024	-0,475891 -0,747315 0,58519 2,001098 0,548524 0,162712 -0,005402	Nrg2 Nkiras1 Nrp2 Thbs1 Ncf4 Habp4 Gcfc1	ENSMUST00000115713 NM_023526 NM_001077406 NM_011580 NM_008677 NM_019986 ENSMUST00000118522
17353676 17297374 17213548 17374488 17312700 17288145	4,489139 5,642543 4,067666 11,25486 7,76849 5,896847	4,544283 5,705226 4,134457 11,32608 7,842728 5,982243 7,036024 5,282709	4,165155 7,109665 6,286251 8,187112 7,889555 5,876764	3,689264 6,36235 6,871441 10,18821 8,438079 6,039476 7,457321 5,742994	0,055144 0,062683 0,066791 0,07122 0,074238 0,085396 0,103024 0,128911	-0,475891 -0,747315 0,58519 2,001098 0,548524 0,162712	Nrg2 Nkiras1 Nrp2 Thbs1 Ncf4 Habp4	ENSMUST00000115713 NM_023526 NM_001077406 NM_011580 NM_008677 NM_019986
17353676 17297374 17213548 17374488 17312700 17288145 17332084 17291953 17218861	4,489139 5,642543 4,067666 11,25486 7,76849 5,896847 6,933 5,153798 11,14262	4,544283 5,705226 4,134457 11,32608 7,842728 5,982243 7,036024 5,282709 11,3079	4,165155 7,109665 6,286251 8,187112 7,889555 5,876764 7,462723 5,598669 6,46808	3,689264 6,36235 6,871441 10,18821 8,438079 6,039476 7,457321 5,742994 9,011907	0,055144 0,062683 0,066791 0,07122 0,074238 0,085396 0,103024 0,128911 0,16528	-0,475891 -0,747315 0,58519 2,001098 0,548524 0,162712 -0,005402 0,144325 2,543827	Nrg2 Nkiras1 Nrp2 Thbs1 Ncf4 Habp4 Gcfc1 Muted F5	ENSMUST00000115713 NM_023526 NM_001077406 NM_011580 NM_008677 NM_019986 ENSMUST00000118522 NM_139063 NM_007976
17353676 17297374 17213548 17374488 17312700 17288145 17332084 17291953 17218861 17250178	4,489139 5,642543 4,067666 11,25486 7,76849 5,896847 6,933 5,153798 11,14262 6,567119	4,544283 5,705226 4,134457 11,32608 7,842728 5,982243 7,036024 5,282709 11,3079 6,739287	4,165155 7,109665 6,286251 8,187112 7,889555 5,876764 7,462723 5,598669 6,46808 7,416512	3,689264 6,36235 6,871441 10,18821 8,438079 6,039476 7,457321 5,742994 9,011907 6,655526	0,055144 0,062683 0,066791 0,07122 0,074238 0,085396 0,103024 0,128911 0,16528 0,172168	-0,475891 -0,747315 0,58519 2,001098 0,548524 0,162712 -0,005402 0,144325 2,543827 -0,760986	Nrg2 Nkiras1 Nrp2 Thbs1 Ncf4 Habp4 Gcfc1 Muted F5 Hist3h2a	ENSMUST00000115713 NM_023526 NM_001077406 NM_011580 NM_008677 NM_019986 ENSMUST00000118522 NM_139063 NM_007976 NM_178218
17353676 17297374 17213548 17374488 17312700 17288145 17332084 17291953 17218861 17250178 17212355	4,489139 5,642543 4,067666 11,25486 7,76849 5,896847 6,933 5,153798 11,14262 6,567119 7,147372	4,544283 5,705226 4,134457 11,32608 7,842728 5,982243 7,036024 5,282709 11,3079 6,739287 7,342451	4,165155 7,109665 6,286251 8,187112 7,889555 5,876764 7,462723 5,598669 6,46808 7,416512 8,072539	3,689264 6,36235 6,871441 10,18821 8,438079 6,039476 7,457321 5,742994 9,011907 6,655526 7,059476	0,055144 0,062683 0,066791 0,07122 0,074238 0,085396 0,103024 0,128911 0,16528 0,172168 0,195079	-0,475891 -0,747315 0,58519 2,001098 0,548524 0,162712 -0,005402 0,144325 2,543827 -0,760986 -1,013063	Nrg2 Nkiras1 Nrp2 Thbs1 Ncf4 Habp4 Gcfc1 Muted F5 Hist3h2a Nck2	ENSMUST0000115713 NM_023526 NM_001077406 NM_011580 NM_08677 NM_019986 ENSMUST00000118522 NM_139063 NM_007976 NM_178218 NM_010879
17353676 17297374 17213548 17374488 17312700 17288145 17332084 17291953 17218861 17250178 17212355 17300251	4,489139 5,642543 4,067666 11,25486 7,76849 5,896847 6,933 5,153798 11,14262 6,567119 7,147372 7,847435	4,544283 5,705226 4,134457 11,32608 7,842728 5,982243 7,036024 5,282709 11,3079 6,739287 7,342451 8,072048	4,165155 7,109665 6,286251 8,187112 7,889555 5,876764 7,462723 5,598669 6,46808 7,416512 8,072539 6,149077	3,689264 6,36235 6,871441 10,18821 8,438079 6,039476 7,457321 5,742994 9,011907 6,655526 7,059476 6,75549	0,055144 0,062683 0,066791 0,07122 0,074238 0,085396 0,103024 0,128911 0,16528 0,172168 0,195079 0,224613	-0,475891 -0,747315 0,58519 2,001098 0,548524 0,162712 -0,005402 0,144325 2,543827 -0,760986 -1,013063 0,606413	Nrg2 Nkiras1 Nrp2 Thbs1 Ncf4 Habp4 Gcfc1 Muted F5 Hist3h2a Nck2 Abhd4	ENSMUST0000115713 NM_023526 NM_001077406 NM_011580 NM_08677 NM_019986 ENSMUST00000118522 NM_139063 NM_007976 NM_178218 NM_010879 NM_134076
17353676 17297374 17213548 17374488 17312700 17288145 17332084 17291953 17218861 17250178 17212355 17300251 17357947	4,489139 5,642543 4,067666 11,25486 7,76849 5,896847 6,933 5,153798 11,14262 6,567119 7,147372 7,847435 5,34164	4,544283 5,705226 4,134457 11,32608 7,842728 5,982243 7,036024 5,282709 11,3079 6,739287 7,342451 8,072048 5,585937	4,165155 7,109665 6,286251 8,187112 7,889555 5,876764 7,462723 5,598669 6,46808 7,416512 8,072539 6,149077 5,893805	3,689264 6,36235 6,871441 10,18821 8,438079 6,039476 7,457321 5,742994 9,011907 6,655526 7,059476 6,75549 7,185633	0,055144 0,062683 0,066791 0,07122 0,074238 0,085396 0,103024 0,128911 0,16528 0,172168 0,195079 0,224613 0,244297	-0,475891 -0,747315 0,58519 2,001098 0,548524 0,162712 -0,005402 0,144325 2,543827 -0,760986 -1,013063 0,606413 1,291828	Nrg2 Nkiras1 Nrp2 Thbs1 Ncf4 Habp4 Gcfc1 Muted F5 Hist3h2a Nck2 Abhd4 Olfr1502	ENSMUST0000115713 NM_023526 NM_001077406 NM_011580 NM_01986 ENSMUST00000118522 NM_139063 NM_007976 NM_178218 NM_010879 NM_134076 NM_146797
17353676 17297374 17213548 17374488 17312700 17288145 17332084 17291953 17218861 17250178 17212355 17300251 17357947 17280125	4,489139 5,642543 4,067666 11,25486 7,76849 5,896847 6,933 5,153798 11,14262 6,567119 7,147372 7,847435 5,34164 8,232164	4,544283 5,705226 4,134457 11,32608 7,842728 5,982243 7,036024 5,282709 11,3079 6,739287 7,342451 8,072048 5,585937 8,524548	4,165155 7,109665 6,286251 8,187112 7,889555 5,876764 7,462723 5,598669 6,46808 7,416512 8,072539 6,149077 5,893805 6,924429	3,689264 6,36235 6,871441 10,18821 8,438079 6,039476 7,457321 5,742994 9,011907 6,655526 7,059476 6,75549 7,185633 7,566913	0,055144 0,062683 0,066791 0,07122 0,074238 0,085396 0,103024 0,128911 0,16528 0,172168 0,195079 0,224613 0,244297 0,292384	-0,475891 -0,747315 0,58519 2,001098 0,548524 0,162712 -0,005402 0,144325 2,543827 -0,760986 -1,013063 0,606413 1,291828 0,642484	Nrg2 Nkiras1 Nrp2 Thbs1 Ncf4 Habp4 Gcfc1 Muted F5 Hist3h2a Nck2 Abhd4 Olfr1502 Pqlc3	ENSMUST0000115713 NM_023526 NM_001077406 NM_011580 NM_008677 NM_019986 ENSMUST00000118522 NM_139063 NM_007976 NM_178218 NM_010879 NM_134076 NM_146797 NM_001161111
17353676 17297374 17213548 17374488 17312700 17288145 17332084 17291953 17218861 17250178 17212355 17300251 17357947 17280125 17300189	4,489139 5,642543 4,067666 11,25486 7,76849 5,896847 6,933 5,153798 11,14262 6,567119 7,147372 7,847435 5,34164 8,232164 1,933763	4,544283 5,705226 4,134457 11,32608 7,842728 5,982243 7,036024 5,282709 11,3079 6,739287 7,342451 8,072048 5,585937 8,524548 2,267333	4,165155 7,109665 6,286251 8,187112 7,889555 5,876764 7,462723 5,598669 6,46808 7,416512 8,072539 6,149077 5,893805 6,924429 7,829066	3,689264 6,36235 6,871441 10,18821 8,438079 6,039476 7,457321 5,742994 9,011907 6,655526 7,059476 6,75549 7,185633 7,566913 7,166917	0,055144 0,062683 0,066791 0,07122 0,074238 0,085396 0,103024 0,128911 0,16528 0,172168 0,195079 0,224613 0,244297 0,292384 0,33357	-0,475891 -0,747315 0,58519 2,001098 0,548524 0,162712 -0,005402 0,144325 2,543827 -0,760986 -1,013063 0,606413 1,291828 0,642484 -0,662149	Nrg2 Nkiras1 Nrp2 Thbs1 Ncf4 Habp4 Gcfc1 Muted F5 Hist3h2a Nck2 Abhd4 Olfr1502 Pqlc3 Trav3-4	ENSMUST0000115713 NM_023526 NM_001077406 NM_011580 NM_08677 NM_019986 ENSMUST00000118522 NM_139063 NM_007976 NM_178218 NM_010879 NM_134076 NM_146797 NM_001161111 Z86035
17353676 17297374 17213548 17374488 17312700 17288145 17332084 17291953 17218861 17250178 17212355 17300251 17387947 17280125 17300189 17314460	4,489139 5,642543 4,067666 11,25486 7,76849 5,896847 6,933 5,153798 11,14262 6,567119 7,147372 7,847435 5,34164 8,232164 1,933763 7,833005	4,544283 5,705226 4,134457 11,32608 7,842728 5,982243 7,036024 5,282709 11,3079 6,739287 7,342451 8,072048 5,585937 8,524548 2,267333 8,176063	4,165155 7,109665 6,286251 8,187112 7,889555 5,876764 7,462723 5,598669 6,46808 7,416512 8,072539 6,149077 5,893805 6,924429 7,829066 7,187557	3,689264 6,36235 6,871441 10,18821 8,438079 6,039476 7,457321 5,742994 9,011907 6,655526 7,059476 6,75549 7,185633 7,566913 7,166917 7,708617	0,055144 0,062683 0,066791 0,07122 0,074238 0,085396 0,103024 0,128911 0,16528 0,172168 0,195079 0,224613 0,244297 0,292384 0,33357 0,343058	-0,475891 -0,747315 0,58519 2,001098 0,548524 0,162712 -0,005402 0,144325 2,543827 -0,760986 -1,013063 0,606413 1,291828 0,642484 -0,662149 0,52106	Nrg2 Nkiras1 Nrp2 Thbs1 Ncf4 Habp4 Gcfc1 Muted F5 Hist3h2a Nck2 Abhd4 Olfr1502 Pqlc3 Trav3-4 Irak4	ENSMUST0000115713 NM_023526 NM_001077406 NM_011580 NM_008677 NM_019986 ENSMUST00000118522 NM_139063 NM_007976 NM_178218 NM_010879 NM_134076 NM_146797 NM_146797 NM_001161111 Z86035 NM_029926
17353676 17297374 17213548 17374488 17312700 17288145 17332084 17291953 17218861 17250178 17210255 17300251 17380125 17300189 17314460 17262178	4,489139 5,642543 4,067666 11,25486 7,76849 5,896847 6,933 5,153798 11,14262 6,567119 7,147372 7,847435 5,34164 8,232164 1,933763 7,833005 6,269192	4,544283 5,705226 4,134457 11,32608 7,842728 5,982243 7,036024 5,282709 11,3079 6,739287 7,342451 8,072048 5,585937 8,524548 2,267333 8,176063 6,631094	4,165155 7,109665 6,286251 8,187112 7,889555 5,876764 7,462723 5,598669 6,46808 7,416512 8,072539 6,149077 5,893805 6,924429 7,829066 7,187557 6,452964	3,689264 6,36235 6,871441 10,18821 8,438079 6,039476 7,457321 5,742994 9,011907 6,655526 7,059476 6,75549 7,185633 7,566913 7,166917 7,708617 6,440458	0,055144 0,062683 0,066791 0,07122 0,074238 0,085396 0,103024 0,128911 0,16528 0,172168 0,195079 0,224613 0,244297 0,292384 0,33357 0,343058 0,361902	-0,475891 -0,747315 0,58519 2,001098 0,548524 0,162712 -0,005402 0,144325 2,543827 -0,760986 -1,013063 0,606413 1,291828 0,642484 -0,662149 0,52106 -0,012506	Nrg2 Nkiras1 Nrp2 Thbs1 Ncf4 Habp4 Gcfc1 Muted F5 Hist3h2a Abhd4 Olfr1502 Pqlc3 Trav3-4 Irak4 Trim41	ENSMUST00000115713 NM_023526 NM_001077406 NM_011580 NM_019986 ENSMUST00000118522 NM_139063 NM_007976 NM_178218 NM_010879 NM_134076 NM_146797 NM_01161111 Z86035 NM_029926 NM_145377
17353676 17297374 17213548 17374488 17312700 17288145 17332084 17291953 17218861 17250178 17218255 17300251 17357947 17280125 17300189 17314460 17262178 17262178	4,489139 5,642543 4,067666 11,25486 7,76849 5,896847 6,933 5,153798 11,14262 6,567119 7,147372 7,847435 5,34164 8,232164 1,933763 7,833005 6,269192 6,986139	4,544283 5,705226 4,134457 11,32608 7,842728 5,982243 7,036024 5,282709 11,3079 6,739287 7,342451 8,524548 2,267333 8,176063 6,631094 7,358756	4,165155 7,109665 6,286251 8,187112 7,889555 5,876764 7,462723 5,598669 6,46808 7,416512 8,072539 6,149077 5,893805 6,924429 7,829066 7,187557 6,452964 7,614451	3,689264 6,36235 6,871441 10,18821 8,438079 6,039476 7,457321 5,742994 9,011907 6,655526 7,059476 6,75549 7,185633 7,566913 7,708617 6,440458 8,147987	0,055144 0,062683 0,066791 0,07122 0,074238 0,085396 0,103024 0,128911 0,16528 0,172168 0,195079 0,224613 0,244297 0,292384 0,33357 0,343058 0,361902 0,372617	-0,475891 -0,747315 0,58519 2,001098 0,548524 0,162712 -0,005402 0,144325 2,543827 -0,760986 -1,013063 0,60413 1,291828 0,642484 -0,662149 0,52106 -0,012506 0,533536	Nrg2 Nkiras1 Nrp2 Thbs1 Ncf4 Habp4 Gcfc1 Muted F5 Hist3h2a Abhd4 Olfr1502 Pqlc3 Trav3-4 Irak4 Trim41 Zfp330	ENSMUST00000115713 NM_023526 NM_001077406 NM_011580 NM_08677 NM_019986 ENSMUST00000118522 NM_139063 NM_00776 NM_178218 NM_010879 NM_134076 NM_146797 NM_01161111 Z86035 NM_029926 NM_145377 AK139907
17353676 17297374 17213548 17374488 17312700 17288145 17332084 17291953 17218861 17250178 17212355 17300251 17357947 17280125 17300189 17314460 17262178 17225472 17329026	4,489139 5,642543 4,067666 11,25486 7,76849 5,896847 6,933 5,153798 11,14262 6,567119 7,147372 7,847435 5,34164 8,232164 1,933763 7,833005 6,269192 6,986139 1,681696	4,544283 5,705226 4,134457 11,32608 7,842728 5,982243 7,036024 5,282709 11,3079 6,739287 7,342451 8,072048 5,585937 8,524548 2,267333 8,176063 6,631094 7,358756 2,062349	4,165155 7,109665 6,286251 8,187112 7,889555 5,876764 7,462723 5,598669 6,46808 7,416512 8,072539 6,149077 5,893805 6,924429 7,829066 7,187557 6,452964 7,614451 2,484488	3,689264 6,36235 6,871441 10,18821 8,438079 6,039476 7,457321 5,742994 9,011907 6,655526 7,059476 6,75549 7,185633 7,566913 7,708617 6,440458 8,147987 2,368513	0,055144 0,062683 0,066791 0,07122 0,074238 0,085396 0,103024 0,128911 0,16528 0,172168 0,195079 0,224613 0,244297 0,292384 0,33357 0,343058 0,361902 0,372617 0,380653	-0,475891 -0,747315 0,58519 2,001098 0,548524 0,162712 -0,005402 0,144325 2,543827 -0,760986 -1,013063 0,606413 1,291828 0,642484 -0,662149 0,52106 -0,012506 0,533536 -0,115975	Nrg2 Nkiras1 Nrp2 Thbs1 Ncf4 Habp4 Gcfc1 Muted F5 Hist3h2a Nck2 Abhd4 Olfr1502 Pqlc3 Trav3-4 Irak4 Trim41 Zfp330 Olfr164	ENSMUST00000115713 NM_023526 NM_001077406 NM_011580 NM_01986 ENSMUST00000118522 NM_139063 NM_007976 NM_178218 NM_010879 NM_134076 NM_146797 NM_01161111 Z86035 NM_029926 NM_145377 AK139907 ENSMUST00000056727
17353676 17297374 17213548 17374488 17312700 17288145 17332084 17291953 17218861 17250178 17212355 17300251 17357947 17280125 17300189 17314460 17262178 17225472 17329026 17344387	4,489139 5,642543 4,067666 11,25486 7,76849 5,896847 6,933 5,153798 11,14262 6,567119 7,447372 7,847435 5,34164 8,232164 1,933763 7,833005 6,269192 6,986139 1,681696 5,255955	4,544283 5,705226 4,134457 11,32608 7,842728 5,982243 7,036024 5,282709 11,3079 6,739287 7,342451 8,072048 2,267333 8,176063 6,631094 7,358756 2,062349 5,647359	4,165155 7,109665 6,286251 8,187112 7,889555 5,876764 7,462723 5,598669 6,46808 7,416512 8,072539 6,149077 5,893805 6,924429 7,829066 7,187557 6,452964 7,614451 2,484488 7,687472	3,689264 6,36235 6,871441 10,18821 8,438079 6,039476 7,457321 5,742994 9,011907 6,655526 7,059476 6,75549 7,185633 7,566913 7,166917 7,708617 6,440458 8,147987 2,368513 7,07621	0,055144 0,062683 0,066791 0,07122 0,074238 0,085396 0,103024 0,128911 0,16528 0,172168 0,195079 0,224613 0,244297 0,292384 0,33357 0,343058 0,361902 0,372617 0,380653 0,391404	-0,475891 -0,747315 0,58519 2,001098 0,548524 0,162712 -0,005402 0,144325 2,543827 -0,760986 -1,013063 0,606413 1,291828 0,62149 0,52106 -0,012506 0,533536 -0,115975 -0,611262	Nrg2 Nkiras1 Nrp2 Thbs1 Ncf4 Habp4 Gcfc1 Muted F5 Hist3h2a Nck2 Abhd4 Olfr1502 Pqlc3 Trav3-4 Irak4 Trim41 Zfp330 Olfr164 Gtf2h4	ENSMUST00000115713 NM_023526 NM_001077406 NM_011580 NM_01986 ENSMUST00000118522 NM_139063 NM_007976 NM_178218 NM_010879 NM_134076 NM_146797 NM_01161111 Z86035 NM_029926 NM_145377 AK139907 ENSMUST00000056727 ENSMUST00000001565
17353676 17297374 17213548 17374488 17312700 17288145 17332084 17291953 17218861 17250178 17212355 17300251 17300189 17314460 17262178 17225472 17329026 17344387 17343897	4,489139 5,642543 4,067666 11,25486 7,76849 5,896847 6,933 5,153798 11,14262 6,567119 7,147372 7,847435 5,34164 8,232164 1,933763 7,833005 6,269192 6,986139 1,681696 5,255955 6,468628	4,544283 5,705226 4,134457 11,32608 7,842728 5,982243 7,036024 5,282709 11,3079 6,739287 7,342451 8,072048 2,267333 8,176063 6,631094 7,358756 2,062349 5,647359 6,881639	4,165155 7,109665 6,286251 8,187112 7,889555 5,876764 7,462723 5,598669 6,46808 7,416512 8,072539 6,149077 5,893805 6,24429 7,829066 7,187557 6,452964 7,614451 2,484488 7,687472 7,309086	3,689264 6,36235 6,871441 10,18821 8,438079 6,039476 7,457321 5,742994 9,011907 6,655526 7,059476 6,75549 7,185633 7,566913 7,166917 7,708617 6,440458 8,147987 2,368513 7,07621 6,618672	0,055144 0,062683 0,066791 0,07122 0,074238 0,085396 0,103024 0,128911 0,16528 0,172168 0,195079 0,224613 0,24247 0,33357 0,343058 0,361902 0,372617 0,380653 0,391404 0,413011	-0,475891 -0,747315 0,58519 2,001098 0,548524 0,162712 -0,005402 0,144325 2,543827 -0,760986 -1,013063 0,606413 1,291828 0,642484 -0,662149 0,52106 -0,012506 0,533536 -0,115975 -0,611262 -0,690414	Nrg2 Nkiras1 Nrp2 Thbs1 Ncf4 Habp4 Gcfc1 Muted F5 Hist3h2a Nck2 Abhd4 Olfr1502 Pqlc3 Trav3-4 Irak4 Trim41 Zfp330 Olfr164 Gtf2h4 Ppt2	ENSMUST00000115713 NM_023526 NM_001077406 NM_011580 NM_01986 ENSMUST00000118522 NM_139063 NM_007976 NM_178218 NM_010879 NM_134076 NM_146797 NM_01161111 Z86035 NM_029926 NM_145377 AK139907 ENSMUST0000001565 ENSMUST000000169067
17353676 17297374 17213548 17374488 17312700 17288145 17332084 17291953 17218861 17250178 17213355 17300251 17357947 17280125 17300189 17314460 17262178 17225472 17329026 17344387 17343897 17292634	4,489139 5,642543 4,067666 11,25486 7,76849 5,896847 6,933 5,153798 11,14262 6,567119 7,147372 7,847435 5,34164 8,232164 1,933763 7,833005 6,269192 6,986139 1,681696 5,255955 6,468628 6,804196	4,544283 5,705226 4,134457 11,32608 7,842728 5,982243 7,036024 5,282709 11,3079 6,739287 7,342451 8,072048 2,267333 8,176063 6,631094 7,358756 2,062349 5,647359 6,881639 7,224471	4,165155 7,109665 6,286251 8,187112 7,889555 5,876764 7,462723 5,598669 6,46808 7,416512 8,072539 6,149077 5,893805 6,924429 7,829066 7,187557 6,452964 7,614451 2,484488 7,687472 7,309086 6,585979	3,689264 6,36235 6,871441 10,18821 8,438079 6,039476 7,457321 5,742994 9,011907 6,655526 7,059476 6,75549 7,185633 7,566913 7,166917 7,708617 6,440458 8,147987 2,368513 7,07621 6,618672 7,177952	0,055144 0,062683 0,066791 0,07122 0,074238 0,085396 0,103024 0,128911 0,16528 0,172168 0,195079 0,224613 0,244297 0,292384 0,33357 0,343058 0,361902 0,372617 0,380653 0,391404 0,413011 0,420275	-0,475891 -0,747315 0,58519 2,001098 0,548524 0,162712 -0,005402 0,144325 2,543827 -0,760986 -1,013063 0,606413 1,291828 0,622149 0,52106 -0,012506 0,533536 -0,115975 -0,611262 -0,690414 0,591973	Nrg2 Nkiras1 Nrp2 Thbs1 Ncf4 Habp4 Gcfc1 Muted F5 Hist3h2a Nck2 Abhd4 Olfr1502 Pqlc3 Trav3-4 Irak4 Trim41 Zfp330 Olfr164 Gtf2h4 Ppt2 Nfil3	ENSMUSTO0000115713 NM_023526 NM_001077406 NM_011580 NM_011580 NM_008677 NM_019986 ENSMUST00000118522 NM_139063 NM_007976 NM_178218 NM_010879 NM_134076 NM_146797 NM_001161111 Z86035 NM_029926 NM_145377 AK139907 ENSMUST000000169565 ENSMUST000000169067 ENSMUST00000011655
17353676 17297374 17213548 17374488 17312700 17288145 17332084 17291953 17218861 17250178 1721355 17300251 17357947 17280125 17300189 17314460 17262178 17225472 17329026 17344387 17343897 17292634 17218366	4,489139 5,642543 4,067666 11,25486 7,76849 5,896847 6,933 5,153798 11,14262 6,567119 7,147372 7,847435 5,34164 8,232164 1,933763 7,833005 6,269192 6,986139 1,681696 5,255955 6,468628 6,804196 7,210876	4,544283 5,705226 4,134457 11,32608 7,842728 5,982243 7,036024 5,282709 11,3079 6,739287 7,342451 8,072048 5,585937 8,524548 2,267333 8,176063 6,631094 7,358756 2,062349 5,647359 6,881639 7,224471 7,640054	4,165155 7,109665 6,286251 8,187112 7,889555 5,876764 7,462723 5,598669 6,46808 7,416512 8,072539 6,149077 5,893805 6,924429 7,829066 7,187557 6,452964 7,614451 2,484488 7,687472 7,309086 6,585979 8,214086	3,689264 6,36235 6,871441 10,18821 8,438079 6,039476 7,457321 5,742994 9,011907 6,655526 7,059476 6,75549 7,185633 7,566913 7,708617 6,440458 8,147987 2,368513 7,07621 6,618672 7,177952 7,580537	0,055144 0,062683 0,066791 0,07122 0,074238 0,085396 0,103024 0,128911 0,16528 0,172168 0,195079 0,224613 0,244297 0,292384 0,33357 0,343058 0,361902 0,372617 0,380653 0,391404 0,413011 0,420275 0,429178	-0,475891 -0,747315 0,58519 2,001098 0,548524 0,162712 -0,005402 0,144325 2,543827 -0,760986 -1,013063 0,606413 1,291828 0,642484 -0,662149 0,52106 -0,012506 0,533536 -0,115975 -0,611262 -0,690414 0,591973 -0,633549	Nrg2 Nkiras1 Nrp2 Thbs1 Ncf4 Habp4 Gcfc1 Muted F5 Hist3h2a Nck2 Abhd4 Olfr1502 Pqlc3 Trav3-4 Irak4 Trim41 Zfp330 Olfr164 Gtf2h4 Ppt2 Nfil3 Stx6	ENSMUST0000115713 NM_023526 NM_001077406 NM_011580 NM_011580 NM_098677 NM_019986 ENSMUST00000118522 NM_139063 NM_007976 NM_178218 NM_010879 NM_134076 NM_146797 NM_010161111 Z86035 NM_029926 NM_145377 AK139907 ENSMUST00000056727 ENSMUST0000001565 ENSMUST0000001605 NM_021433
17353676 17297374 17213548 17374488 17312700 17288145 17332084 17291953 17218861 17250178 17212355 17300251 17357947 17280125 1730189 17314460 17262178 17225472 17329026 17344387 17243866 17319607	4,489139 5,642543 4,067666 11,25486 7,76849 5,896847 6,933 5,153798 11,14262 6,567119 7,147372 7,847435 5,34164 8,232164 1,933763 7,833005 6,269192 6,986139 1,681696 5,255955 6,468628 6,804196 7,210876 7,622211	4,544283 5,705226 4,134457 11,32608 7,842728 5,982243 7,036024 5,282709 11,3079 6,739287 7,342451 8,072048 5,585937 8,5245483 2,267383 8,176063 6,631094 7,358756 2,062349 5,647359 6,881639 7,224471 7,640054 8,063448	4,165155 7,109665 6,286251 8,187112 7,889555 5,876764 7,462723 5,598669 6,46808 7,416512 8,072539 6,149077 5,893805 6,924429 7,829666 7,187557 6,452964 7,614451 2,484488 7,687472 7,309086 6,585979 8,214086 6,758389	3,689264 6,36235 6,871441 10,18821 8,438079 6,039476 7,457321 5,742994 9,011907 6,655526 7,059476 6,75549 7,185633 7,566913 7,166917 7,708617 6,440458 8,147987 2,368513 7,07621 6,618672 7,177952 7,580537 7,300317	0,055144 0,062683 0,066791 0,07122 0,074238 0,085396 0,103024 0,128911 0,16528 0,172168 0,195079 0,224613 0,244297 0,292384 0,33357 0,343058 0,361902 0,372617 0,380653 0,391404 0,413011 0,420275 0,429178 0,441237	-0,475891 -0,747315 0,58519 2,001098 0,548524 0,162712 -0,005402 0,144325 2,543827 -0,760986 -1,013063 0,606413 1,291828 0,642484 -0,662149 0,52106 -0,012506 0,533536 -0,115975 -0,611262 -0,690414 0,591973 -0,633549 0,541928	Nrg2 Nkiras1 Nrp2 Thbs1 Ncf4 Habp4 Gcfc1 Muted F5 Hist3h2a Nck2 Abhd4 Olfr1502 Pq1c3 Trav3-4 Irak4 Trim41 Zfp330 Olfr164 Gtf2h4 Ppt2 Nfil3 Stx6 Naga	ENSMUST0000115713 NM_023526 NM_001077406 NM_011580 NM_011580 NM_098677 NM_019986 ENSMUST00000118522 NM_139063 NM_007976 NM_178218 NM_010879 NM_134076 NM_146797 NM_001161111 Z86035 NM_029926 NM_145377 AK139907 ENSMUST00000056727 ENSMUST0000001565 ENSMUST000000169067 ENSMUST00000071065 NM_021433 NM_008669
17353676 17297374 17213548 17374488 17312700 17288145 17332084 17291953 17218861 17250178 1721355 17300251 17357947 17280125 17300189 17314460 17262178 17225472 17329026 17344387 17343897 17292634 17218366	4,489139 5,642543 4,067666 11,25486 7,76849 5,896847 6,933 5,153798 11,14262 6,567119 7,147372 7,847435 5,34164 8,232164 1,933763 7,833005 6,269192 6,986139 1,681696 5,255955 6,468628 6,804196 7,210876	4,544283 5,705226 4,134457 11,32608 7,842728 5,982243 7,036024 5,282709 11,3079 6,739287 7,342451 8,072048 5,585937 8,524548 2,267333 8,176063 6,631094 7,358756 2,062349 5,647359 6,881639 7,224471 7,640054	4,165155 7,109665 6,286251 8,187112 7,889555 5,876764 7,462723 5,598669 6,46808 7,416512 8,072539 6,149077 5,893805 6,924429 7,829066 7,187557 6,452964 7,614451 2,484488 7,687472 7,309086 6,585979 8,214086	3,689264 6,36235 6,871441 10,18821 8,438079 6,039476 7,457321 5,742994 9,011907 6,655526 7,059476 6,75549 7,185633 7,566913 7,708617 6,440458 8,147987 2,368513 7,07621 6,618672 7,177952 7,580537	0,055144 0,062683 0,066791 0,07122 0,074238 0,085396 0,103024 0,128911 0,16528 0,172168 0,195079 0,224613 0,244297 0,292384 0,33357 0,343058 0,361902 0,372617 0,380653 0,391404 0,413011 0,420275 0,429178	-0,475891 -0,747315 0,58519 2,001098 0,548524 0,162712 -0,005402 0,144325 2,543827 -0,760986 -1,013063 0,606413 1,291828 0,642484 -0,662149 0,52106 -0,012506 0,533536 -0,115975 -0,611262 -0,690414 0,591973 -0,633549	Nrg2 Nkiras1 Nrp2 Thbs1 Ncf4 Habp4 Gcfc1 Muted F5 Hist3h2a Nck2 Abhd4 Olfr1502 Pqlc3 Trav3-4 Irak4 Trim41 Zfp330 Olfr164 Gtf2h4 Ppt2 Nfil3 Stx6	ENSMUST0000115713 NM_023526 NM_001077406 NM_011580 NM_011580 NM_098677 NM_019986 ENSMUST00000118522 NM_139063 NM_007976 NM_178218 NM_010879 NM_134076 NM_146797 NM_010161111 Z86035 NM_029926 NM_145377 AK139907 ENSMUST00000056727 ENSMUST0000001565 ENSMUST0000001605 NM_021433

17281013									
17326510 9,258003 9,748003 4,678825 6,492658 0,488 1,813833 Pros1 NM_011173 17296128 6,049206 6,54127 6,06002 7,071895 0,492010 0,755875 Gapt NM_017771 17329433 7,165043 7,6678781 7,661284 6,973663 0,488738 -0,687621 Bcl6 ENSMUSTO0000023151 17334419 8,05252 8,553817 6,752113 7,047427 0,501297 0,295814 Sept1 NM_01379 17254341 6,761891 7,21333 7,400024 7,217378 0,05152 -0,186646 Flotz NM_008028 1,7270888 7,797262 8,300146 6,875265 7,076592 0,502884 0,201327 Ern1 ENSMUST00000001059 17227038 3,109274 9,640001 8,559824 8,666 0,590227 0,28833 Glud1 NM_00139 1,7284084 4,345088 4,856655 8,669348 7,572428 0,511537 1,09992 Tubel NM_002806 1,7234938 8,003888 8,46246 7,467991 7,67975 0,511573 0,180049 Irm2x ENSMUST000000027425 1,7232951 7,109286 7,621434 7,799578 7,739091 0,512057 -0,060487 Sec63 NM_153055 1,7212779 6,243887 6,793185 4,656544 6,6593 0,512087 0,068487 Sec63 NM_153055 1,7212779 1,6221819 6,74617 7,073561 6,6593 0,502085 0,433940 Chaf1b NM_002808 1,7240907 7,148594 7,669847 8,091006 7,397471 0,570853 0,093535 Majle NM_002808 1,7236074 7,73585 6,455940 6,6693 0,502085 0,433940 Chaf1b NM_002808 1,723001 6,02445 6,552396 6,552346 6,6693 0,502085 0,433940 Chaf1b NM_002808 1,723001 6,02445 6,552396 6,55534 7,502523 0,52301 0,016993 4,044866 Reep4 NM_0016961 1,7230747 1,73585 8,038515 7,767850 0,500018 0,016993 0,00000002732 1,721090 6,02445 6,552396 6,555396 6,6693 0,503056 0,433940 0,4024114 Wdr7 NM_00016981 1,723056 0,500016 0,50006	17289103	6,449024	6,902662	8,602953	7,944226	0,453638	-0,658727	Serinc5	NM_172588
17296128	17231066	5,945387	6,42391	7,232356	6,687385	0,478523	-0,544971	Dtl	NM_029766
17329433 7,165043 7,663781 7,661284 6,973662 0,498738 -0,687621 Bcl6 ENSMUST00000023151 17334419 8,05252 5,553617 6,752113 7,047427 0,501297 -0,295514 Sept	17326510	9,258803	9,746803	4,678825	6,492658	0,488	1,813833	Pros1	NM_011173
17324419	17296128	6,049269	6,54127	6,30602	7,071895	0,492001	0,765875	Gapt	NM 177713
17324419 8,05252 8,55817 6,75113 7,047427 0,501297 0,299314 Sept1 NM_008028 1725343 6,716819 7,218339 7,440024 7,217378 0,50152 -0,186646 Flot2 NM_008028 17270388 7,797562 8,300146 6,875265 7,075592 0,502884 0,201327 Em1 ENSMUSTO0000001059 17327038 9,130974 9,640001 8,55582 8,8566 0,599027 0,297018 Min 1672 NM_001599 17328038 8,802176 9,312473 8,938039 9,226872 0,510297 0,288833 Glud1 NM_001833 17228634 4,345088 4,856625 8,669348 7,572428 0,511537 -1,09692 Tube1 NM_028006 17212998 8,0030888 8,542461 7,467901 7,64795 0,511537 -1,09692 Tube1 NM_028006 172129395 7,019286 7,62134 7,799785 0,511537 0,180049 Tube2 ENSNUSTO0000027425 17212677 6,243887 6,759185 4,086674 4,605269 0,515298 0,568959 5,85955 NM_028787 17212677 6,243887 6,759185 4,086674 4,605269 0,515298 0,568959 5,85955 NM_028787 17212677 7,073561 6,538659 0,518051 -0,534902 Chaf1b NM_02808 17239056 7,70744 7,597256 6,425904 6,6693 0,520853 -0,939355 Maple9 NM_018561 17350563 7,70744 7,597256 6,425904 6,6693 0,520853 -0,939355 Maple9 NM_018561 17350553 7,07644 7,597256 6,425904 6,6693 0,520865 0,523904 0,46886 Repat N1_4003885 17293031 6,029445 6,552446 5,68673 5,702523 0,523901 0,016793 Akrlet NM_001869 17241667 7,735385 8,018515 7,781590 0,500814 0,46886 Repat N1_4003859 1725096 8,216847 8,74993 8,30149 0,533314 0,026497 0,46894 0,46941 0,	17329433	7,165043	7,663781	7,661284	6,973663	0,498738	-0,687621	Bcl6	ENSMUST00000023151
17253431	17334419			6.752113				Sepx1	NM 013759
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1732038 9,130974 9,640001 8,559582 8,8566 0,509027 0,297018 1fnar2 NM 010509 17298055 8,02176 9,312473 8,938039 9,226872 0,510297 0,288833 Glud1 NM 00813 17230264 4,454088 4,865625 8,669348 7,572428 0,511537 1,080049 1tm2c ENSMUST00000027425 17232951 7,109286 7,621343 7,799578 7,739091 0,512057 0,060487 Sec63 NM 1,50057 17216727 6,244887 6,759185 4,05674 4,605269 0,515298 0,568595 Sc2515 NM 0,28787 17237331 6,228119 6,74617 7,073561 6,58659 0,518051 0,534902 Chaf1b NM 0,28083 17240957 7,448994 7,66847 8,91006 7,397471 0,520853 0,69335 Mapk9 NM 0,16961 17356563 7,0764 7,597256 6,425904 6,6693 0,520856 0,243396 Sk25a45 ENSMUST0000025732 17301757 7,214467 7,735385 8,018515 7,871659 0,520918 0,046856 Reep4 NM 1,80588 1735122 6,764033 7,294382 7,520123 7,478109 0,530494 0,042014 Wdr7 NM 0,01014981 17241062 7,14453 7,674551 7,674									_
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1722179	17233629	10,51824	11,0534	10,74282	10,89393	0,53516	0,15111		NM_001146120
17298312	17272785	9,435004	9,972488	7,731281	8,14407	0,537484	0,412789	Lgals3bp	ENSMUST00000043722
17297227	17222179	5,384631	5,922555	7,069615	6,700094	0,537924	-0,369521	Lman2l	ENSMUST00000125304
17347475 6,579448 7,120022 6,85364 6,190247 0,540574 -0,663317 Hnrpll NM_144802 17307588 6,86782 7,412256 7,040932 7,089822 0,544436 0,04889 Fdft1 NM_010191 17219546 5,077194 5,623463 6,268832 6,785285 0,546269 0,516453 Slamf9 ENSMUST00000027830 17350517 6,285724 6,832091 6,848124 6,692298 0,546367 -0,155826 Hsd17b4 NM_008996 17247533 9,079891 9,628637 9,443911 9,41308 0,548746 -0,030831 Rab1 NM_008996 17308132 7,169549 7,718704 8,83729 8,725042 0,549155 -0,112248 Sic25a37 NM_026331 17255402 7,48515 8,034584 7,738726 7,710865 0,549434 -0,027861 Spop NM_025287 17317296 6,656888 7,20871 6,531343 6,694017 0,551822 0,162674 Tme65 ENSMUST00000072113	17298312	6,117821	6,657413	6,170248	6,119926	0,539592	-0,050322	Glt8d1	NM_029626
17307588	17297227	6,853437	7,393343	7,702649	7,707954	0,539906	0,005305	Slc4a7	NM_001033270
17219546 5,077194 5,623463 6,268832 6,785285 0,546269 0,516453 Slamf9 ENSMUST00000027830 17350517 6,285724 6,832091 6,848124 6,692298 0,546367 -0,155826 Hsd17b4 NM_008292 17247533 9,079891 9,628637 9,443911 9,41308 0,548746 -0,030831 Rab1 NM_008996 17308132 7,169549 7,718704 8,83729 8,725042 0,549155 -0,112248 Slc25a37 NM_026331 17255402 7,48515 8,034584 7,738726 7,710865 0,549434 -0,027861 Spop NM_025287 17347571 6,20408 6,754478 6,795957 6,560254 0,550398 -0,235703 Map4k3 NM_001081357 17317296 6,6565888 7,0871 6,531433 6,694017 0,551899 0,269912 Plxnc1 NM_01018797 17256345 7,387208 7,941107 8,280304 8,081751 0,553899 0,134982 Hiatl1 ENSMUST0000007213 <tr< td=""><td>17347475</td><td>6,579448</td><td>7,120022</td><td>6,853564</td><td>6,190247</td><td>0,540574</td><td>-0,663317</td><td>Hnrpll</td><td>NM_144802</td></tr<>	17347475	6,579448	7,120022	6,853564	6,190247	0,540574	-0,663317	Hnrpll	NM_144802
17350517 6,285724 6,832091 6,848124 6,692298 0,546367 -0,155826 Hsd17b4 NM_008292 17247533 9,079891 9,628637 9,443911 9,41308 0,548746 -0,030831 Rab1 NM_008996 17308132 7,169549 7,718704 8,83729 8,725042 0,549155 -0,112248 Slc25a37 NM_026331 17255402 7,48515 8,034584 7,738726 7,710865 0,549434 -0,027861 Spop NM_025287 17347571 6,620408 6,754478 6,795957 6,560254 0,550398 -0,235703 Map4k3 NM_00181357 17317296 6,656888 7,20871 6,531343 6,694017 0,551822 0,162674 Tmem65 ENSMUST00000072113 17244439 7,007654 7,560632 7,696064 7,553899 -0,189553 Eif1 NM_0118797 17256345 7,387208 7,941107 8,280304 8,081751 0,553899 -0,138553 Hiatl1 ENSMUST000000155487 17253235 <td>17307588</td> <td>6,86782</td> <td>7,412256</td> <td>7,040932</td> <td>7,089822</td> <td>0,544436</td> <td>0,04889</td> <td>Fdft1</td> <td>NM_010191</td>	17307588	6,86782	7,412256	7,040932	7,089822	0,544436	0,04889	Fdft1	NM_010191
17247533	17219546	5,077194	5,623463	6,268832	6,785285	0,546269	0,516453	Slamf9	ENSMUST00000027830
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17255402 7,48515 8,034584 7,738726 7,710865 0,549434 -0,027861 Spop NM_025287 17347571 6,20408 6,754478 6,795957 6,560254 0,550398 -0,235703 Map4k3 NM_001081357 17317296 6,656888 7,20871 6,531343 6,694017 0,551822 0,162674 Tmem65 ENSMUST00000072113 17244439 7,007654 7,560632 7,696054 7,965966 0,552978 0,269912 Plxnc1 NM_018797 17256345 7,387208 7,941107 8,280304 8,081751 0,553899 -0,198553 Eif1 NM_011508 17293757 7,636679 8,190578 7,367367 7,502349 0,553899 0,134982 Hiatl1 ENSMUST00000155487 17253235 8,582083 9,139152 9,168846 9,389271 0,557069 0,220425 Ssh2 NM_177710 17230256 8,095563 8,652717 8,205182 8,29938 0,557154 0,094198 Ads ENSMUST00000016105	17247533	9,079891	9,628637	9,443911	9,41308	0,548746	-0,030831	Rab1	NM_008996
17347571 6,20408 6,754478 6,795957 6,560254 0,550398 -0,235703 Map4k3 NM_001081357 17317296 6,656888 7,20871 6,531343 6,694017 0,551822 0,162674 Tmem65 ENSMUST00000072113 17244439 7,007654 7,560632 7,696054 7,965966 0,552978 0,269912 Plxnc1 NM_018797 17256345 7,387208 7,941107 8,280304 8,081751 0,553899 -0,198553 Eif1 NM_011508 17293757 7,636679 8,190578 7,367367 7,502349 0,553899 0,134982 Hiatl1 ENSMUST00000155487 17253235 8,582083 9,139152 9,168846 9,389271 0,557069 0,220425 Ssh2 NM_177710 17230256 8,095563 8,652717 8,205182 8,29938 0,557154 0,094198 Adss ENSMUST00000016105 17301722 8,124253 8,683105 8,155062 8,36490 0,558469 0,215167 Slc25a30 NM_0026232	17308132	7,169549	7,718704	8,83729	8,725042	0,549155	-0,112248	Slc25a37	NM_026331
17347571 6,20408 6,754478 6,795957 6,560254 0,550398 -0,235703 Map4k3 NM_001081357 17317296 6,656888 7,20871 6,531343 6,694017 0,551822 0,162674 Tmem65 ENSMUST00000072113 17244439 7,007654 7,560632 7,696054 7,965966 0,552978 0,269912 Plxnc1 NM_018797 17256345 7,387208 7,941107 8,280304 8,081751 0,553899 -0,198553 Eif1 NM_011508 17293757 7,636679 8,190578 7,367367 7,502349 0,553899 0,134982 Hiatl1 ENSMUST00000155487 17253235 8,582083 9,139152 9,168846 9,389271 0,557069 0,220425 Ssh2 NM_177710 17230256 8,095563 8,652717 8,205182 8,29938 0,557154 0,094198 Ads ENSMUST0000016105 17301722 8,124253 8,683105 8,155062 8,36497 0,558469 0,215167 Slc25a30 NM_0026232 <	17255402	7,48515	8,034584	7,738726	7,710865	0,549434	-0,027861	Spop	NM 025287
17244439 7,007654 7,560632 7,696054 7,965966 0,552978 0,269912 Plxnc1 NM_018797 17256345 7,387208 7,941107 8,280304 8,081751 0,553899 -0,198553 Eif1 NM_011508 17293757 7,636679 8,190578 7,367367 7,502349 0,553899 0,134982 Hiatl1 ENSMUST00000155487 17253235 8,582083 9,139152 9,168846 9,389271 0,557069 0,220425 Ssh2 NM_177710 17230256 8,095563 8,652717 8,205182 8,29938 0,557154 0,094198 Adss ENSMUST00000016105 17308714 7,069029 7,627498 7,17178 7,386947 0,558469 0,215167 Slc25a30 NM_026232 17301722 8,124253 8,683105 8,155062 8,36409 0,558852 0,209028 Bin3 NM_021328 17228860 8,883883 9,444352 9,288135 9,294095 0,560469 0,00596 Aph1a NM_146104 <td< td=""><td>17347571</td><td>6,20408</td><td>6,754478</td><td>6,795957</td><td>6,560254</td><td>0,550398</td><td>-0,235703</td><td></td><td>NM 001081357</td></td<>	17347571	6,20408	6,754478	6,795957	6,560254	0,550398	-0,235703		NM 001081357
17244439 7,007654 7,560632 7,696054 7,965966 0,552978 0,269912 Plxnc1 NM_018797 17256345 7,387208 7,941107 8,280304 8,081751 0,553899 -0,198553 Eif1 NM_011508 17293757 7,636679 8,190578 7,367367 7,502349 0,553899 0,134982 Hiatl1 ENSMUST00000155487 17253235 8,582083 9,139152 9,168846 9,389271 0,557069 0,220425 Ssh2 NM_177710 17230256 8,095563 8,652717 8,205182 8,29938 0,557154 0,094198 Adss ENSMUST00000016105 17308714 7,069029 7,627498 7,17178 7,386947 0,558469 0,215167 Slc25a30 NM_026232 17301722 8,124253 8,683105 8,155062 8,36409 0,558852 0,209028 Bin3 NM_021328 17228860 8,883883 9,444352 9,288135 9,294095 0,560469 0,00596 Aph1a NM_146104 <td< td=""><td>17317296</td><td>6,656888</td><td>7,20871</td><td>6,531343</td><td>6,694017</td><td>0,551822</td><td>0,162674</td><td>Tmem65</td><td>ENSMUST00000072113</td></td<>	17317296	6,656888	7,20871	6,531343	6,694017	0,551822	0,162674	Tmem65	ENSMUST00000072113
17256345 7,387208 7,941107 8,280304 8,081751 0,553899 -0,198553 Eif1 NM_011508 17293757 7,636679 8,190578 7,367367 7,502349 0,553899 0,134982 Hiatl1 ENSMUST00000155487 17253235 8,582083 9,139152 9,168846 9,389271 0,557069 0,220425 Ssh2 NM_177710 17230256 8,095563 8,652717 8,205182 8,29938 0,557154 0,094198 Adss ENSMUST00000016105 17308714 7,069029 7,627498 7,17178 7,386947 0,558469 0,215167 Slc25a30 NM_026232 17301722 8,124253 8,683105 8,155062 8,36409 0,558852 0,209028 Bin3 NM_026232 17340332 7,602308 8,161225 7,995773 8,164858 0,558917 0,169085 Klraq1 NM_028658 17228060 8,883883 9,444352 9,288135 9,294095 0,560469 0,00596 Aph1a NM_146104 <td< td=""><td>17244439</td><td></td><td></td><td></td><td></td><td></td><td></td><td>Plxnc1</td><td></td></td<>	17244439							Plxnc1	
17293757 7,636679 8,190578 7,367367 7,502349 0,553899 0,134982 Hiatl1 ENSMUST00000155487 17253235 8,582083 9,139152 9,168846 9,389271 0,557069 0,220425 Ssh2 NM_177710 17230256 8,095563 8,652717 8,205182 8,29938 0,557154 0,094198 Adss ENSMUST00000016105 17308714 7,069029 7,627498 7,17178 7,386947 0,558469 0,215167 Slc25a30 NM_026232 17301722 8,124253 8,683105 8,155062 8,36409 0,558852 0,209028 Bin3 NM_026232 17340332 7,602308 8,161225 7,995773 8,164858 0,558917 0,169085 Klraq1 NM_028658 17228860 8,883883 9,444352 9,288135 9,294095 0,560469 0,00596 Aph1a NM_146104 17350869 8,287262 8,852432 8,620236 8,517699 0,56517 -0,102537 Isoc1 NM_025478 <td< td=""><td></td><td></td><td>_</td><td></td><td>,</td><td>,</td><td>,</td><td>Eif1</td><td>_</td></td<>			_		,	,	,	Eif1	_
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17230256 8,095563 8,652717 8,205182 8,29938 0,557154 0,094198 Adss ENSMUST00000016105 17308714 7,069029 7,627498 7,17178 7,386947 0,558469 0,215167 Slc25a30 NM_026232 17301722 8,124253 8,683105 8,155062 8,36409 0,558852 0,209028 Bin3 NM_021328 17340332 7,602308 8,161225 7,995773 8,164858 0,558917 0,169085 Klraq1 NM_028658 17228860 8,883883 9,444352 9,288135 9,294095 0,560469 0,00596 Aph1a NM_146104 17350869 8,287262 8,852432 8,620236 8,517699 0,56517 -0,102537 Isoc1 NM_025478 17282723 9,014027 9,57972 9,130428 9,300289 0,565693 0,169861 Tmed10 NM_026775 17256414 6,971656 7,537413 6,230683 6,543866 0,565757 0,313183 Dnajc7 NM_0119795 173320									
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1732/1204 6/2/127 7/002201 7/212005 7/227040 0/27024 0/02/154 Dbd1 NA 042620									_
רייט איז אראינטן איז אראינטן איז אראינטן דאיז אראינטן איז אראינטן איז אראינטן דאיז אראינטן איז אראינטן דאיז אראינטן איז אראינטן איז אראינטן דאיז דאיז אראינטן דאיז אראינטן דאיז אראינטן דאיז אראינטן דאיז דאיז אראינטן דאיז אראינטן דאיז אראינטן דאיז אראינטן דאיז אראינען דיין דאיז אראינען דיין דאיז אראינען דיין דאיז אראינען דיין דאיין דאיז אראינען דיין דאיז אראינען דיין דאיז אראינען דיין דאיז אראינען דאיז אראינען דיין דאיז אראינען דאין דיין דאיז אראינען דיין דאיין דיין דאיז אראינען דיין דיין דיין דיין דיין דיין דיין די	17334304	6,424127	7,003391	7,312895	7,337049	0,579264	0,024154	Pkd1	NM_013630

17322559	7,589106	8,170588	8,151337	8,246286	0,581482	0,094949	Hmox2	ENSMUST00000004172
17229607	8,324808	8,906755	8,643161	9,13483	0,581947	0,491669	Fcgr2b	NM_001077189
17331564	6,93132	7,513807	7,777517	7,399642	0,582487	-0,377875	Nrip1	ENSMUST00000121927
17215968	8,010756	8,597423	8,612072	8,691059	0,586667	0,078987	Ppp1r7	ENSMUST00000027494
17233226	9,697678	10,29039	7,937216	9,196314	0,592712	1,259098	Lilrb4	NM 013532
17320307	4,24427	4,838446	6,409072	6,939296	0,594176	0,530224	Mapk12	NM 013871
17236836	9,096325	9,690732	9,711331	9,98899	0,594407	0,277659	Atp2b1	NM 026482
17354995	6,832865	7,431355	7,303247	7,224578	0,59849	-0,078669	Fech	NM 007998
17272913	6,895566	7,496176	6,706542	7,210859	0,60061	0,504317	Sgsh	NM 018822
17240053	6,671791	7,430170	6,979804	7,210033	0,60155	0,304317	Ncoa7	NM 172495
	6,949783	7,563349	8,69096					
17255026	<u> </u>			8,800943 5,139194	0,613566	0,109983	Trim25	NM_009546
17221756	6,687934	7,302514	4,588058		0,61458	0,551136	Ogfrl1	NM_001081079
17334803	7,194066	7,810008	8,133793	8,183661	0,615942	0,049868	Jmjd8	ENSMUST00000133595
17500275	6,681298	7,298223	6,380334	6,72434	0,616925	0,344006	Erlin2	NM_153592
17269911	7,105235	7,722378	6,891325	7,549478	0,617143	0,658153	Vat1	NM_012037
17238594	6,633016	7,261329	4,334667	4,435435	0,628313	0,100768	Cd63	NM_001042580
17211774	6,76707	7,395988	7,526197	7,704848	0,628918	0,178651	Plekhb2	NM_145516
17254166	7,885752	8,515647	7,827438	7,978152	0,629895	0,150714	Slfn2	NM_011408
17282563	9,275988	9,909496	9,244076	9,457718	0,633508	0,213642	Npc2	NM_023409
17353337	6,619605	7,255728	5,833216	5,505816	0,636123	-0,3274	Gypc	NM_001048207
17238482	7,058269	7,695981	7,205977	7,405742	0,637712	0,199765	Rnf41	NM_001164237
17305685	7,825674	8,46529	8,024742	8,034334	0,639616	0,009592	Ddhd1	NM_001042719
17272437	6,803088	7,444184	6,783096	7,200078	0,641096	0,416982	Rhbdf2	NM 172572
17304906	7,347307	7,990122	9,332798	9,596867	0,642815	0,264069	Wdfy4	NM 001146022
17350740	6,228748	6,871938	7,455481	7,048126	0,64319	-0,407355	Lmnb1	NM 010721
17243022	6,709941	7,353891	6,884367	6,797133	0,64395	-0,087234	Btbd2	NM 145361
17324553	8,52689	9,178108	8,052605	8,686935	0,651218	0,63433	Ccdc50	ENSMUST00000100026
17342101	6,474099	7,125337	7,112027	7,171763	0,651238	0,059736	Hn1l	ENSMUST00000024981
17372101								
17286254								
17286254	6,041173	6,695262	6,192604	6,299879	0,654089	0,107275	Mboat1	NM_153546
17383467	6,041173 6,406147	6,695262 7,060637	6,192604 5,634828	6,299879 6,380713	0,654089 0,65449	0,107275 0,745885	Mboat1 Ntng2	NM_153546 NM_133501
17383467 17281971	6,041173 6,406147 7,931169	6,695262 7,060637 8,585807	6,192604 5,634828 8,325535	6,299879 6,380713 8,447086	0,654089 0,65449 0,654638	0,107275 0,745885 0,121551	Mboat1 Ntng2 Sgpp1	NM_153546 NM_133501 NM_030750
17383467 17281971 17213428	6,041173 6,406147 7,931169 7,147408	6,695262 7,060637 8,585807 7,802353	6,192604 5,634828 8,325535 6,492986	6,299879 6,380713 8,447086 7,086036	0,654089 0,65449 0,654638 0,654945	0,107275 0,745885 0,121551 0,59305	Mboat1 Ntng2 Sgpp1 Cyp20a1	NM_153546 NM_133501 NM_030750 ENSMUST00000060608
17383467 17281971 17213428 17227483	6,041173 6,406147 7,931169 7,147408 6,560924	6,695262 7,060637 8,585807 7,802353 7,21599	6,192604 5,634828 8,325535 6,492986 7,155493	6,299879 6,380713 8,447086 7,086036 7,298692	0,654089 0,65449 0,654638 0,654945 0,655066	0,107275 0,745885 0,121551 0,59305 0,143199	Mboat1 Ntng2 Sgpp1 Cyp20a1 Camsap2	NM_153546 NM_133501 NM_030750 ENSMUST00000060608 NM_001081360
17383467 17281971 17213428 17227483 17343035	6,041173 6,406147 7,931169 7,147408 6,560924 7,200163	6,695262 7,060637 8,585807 7,802353 7,21599 7,857397	6,192604 5,634828 8,325535 6,492986 7,155493 7,815231	6,299879 6,380713 8,447086 7,086036 7,298692 7,948244	0,654089 0,65449 0,654638 0,654945 0,655066 0,657234	0,107275 0,745885 0,121551 0,59305 0,143199 0,133013	Mboat1 Ntng2 Sgpp1 Cyp20a1 Camsap2 Mtch1	NM_153546 NM_133501 NM_030750 ENSMUST00000060608 NM_001081360 NM_019880
17383467 17281971 17213428 17227483 17343035 17283930	6,041173 6,406147 7,931169 7,147408 6,560924 7,200163 7,566288	6,695262 7,060637 8,585807 7,802353 7,21599 7,857397 8,236075	6,192604 5,634828 8,325535 6,492986 7,155493 7,815231 7,678843	6,299879 6,380713 8,447086 7,086036 7,298692 7,948244 7,862485	0,654089 0,65449 0,654638 0,654945 0,655066 0,657234 0,669787	0,107275 0,745885 0,121551 0,59305 0,143199 0,133013 0,183642	Mboat1 Ntng2 Sgpp1 Cyp20a1 Camsap2 Mtch1 Wars	NM_153546 NM_133501 NM_030750 ENSMUST00000060608 NM_001081360 NM_019880 NM_001164314
17383467 17281971 17213428 17227483 17343035 17283930 17346349	6,041173 6,406147 7,931169 7,147408 6,560924 7,200163 7,566288 6,036273	6,695262 7,060637 8,585807 7,802353 7,21599 7,857397 8,236075 6,710959	6,192604 5,634828 8,325535 6,492986 7,155493 7,815231 7,678843 7,346448	6,299879 6,380713 8,447086 7,086036 7,298692 7,948244 7,862485 7,848524	0,654089 0,65449 0,654638 0,654945 0,655066 0,657234 0,669787 0,674686	0,107275 0,745885 0,121551 0,59305 0,143199 0,133013 0,183642 0,502076	Mboat1 Ntng2 Sgpp1 Cyp20a1 Camsap2 Mtch1 Wars Rfx2	NM_153546 NM_133501 NM_030750 ENSMUST00000060608 NM_001081360 NM_019880 NM_001164314 NM_027787
17383467 17281971 17213428 17227483 17343035 17283930	6,041173 6,406147 7,931169 7,147408 6,560924 7,200163 7,566288 6,036273 7,694138	6,695262 7,060637 8,585807 7,802353 7,21599 7,857397 8,236075 6,710959 8,369534	6,192604 5,634828 8,325535 6,492986 7,155493 7,815231 7,678843 7,346448 5,215715	6,299879 6,380713 8,447086 7,086036 7,298692 7,948244 7,862485 7,848524 6,375921	0,654089 0,65449 0,654638 0,654945 0,655066 0,657234 0,669787	0,107275 0,745885 0,121551 0,59305 0,143199 0,133013 0,183642 0,502076 1,160206	Mboat1 Ntng2 Sgpp1 Cyp20a1 Camsap2 Mtch1 Wars	NM_153546 NM_133501 NM_030750 ENSMUST00000060608 NM_001081360 NM_0119880 NM_001164314 NM_027787 NM_010188
17383467 17281971 17213428 17227483 17343035 17283930 17346349	6,041173 6,406147 7,931169 7,147408 6,560924 7,200163 7,566288 6,036273	6,695262 7,060637 8,585807 7,802353 7,21599 7,857397 8,236075 6,710959	6,192604 5,634828 8,325535 6,492986 7,155493 7,815231 7,678843 7,346448	6,299879 6,380713 8,447086 7,086036 7,298692 7,948244 7,862485 7,848524	0,654089 0,65449 0,654638 0,654945 0,655066 0,657234 0,669787 0,674686	0,107275 0,745885 0,121551 0,59305 0,143199 0,133013 0,183642 0,502076	Mboat1 Ntng2 Sgpp1 Cyp20a1 Camsap2 Mtch1 Wars Rfx2	NM_153546 NM_133501 NM_030750 ENSMUST00000060608 NM_001081360 NM_019880 NM_001164314 NM_027787
17383467 17281971 17213428 17227483 17343035 17283930 17346349 17229620	6,041173 6,406147 7,931169 7,147408 6,560924 7,200163 7,566288 6,036273 7,694138	6,695262 7,060637 8,585807 7,802353 7,21599 7,857397 8,236075 6,710959 8,369534	6,192604 5,634828 8,325535 6,492986 7,155493 7,815231 7,678843 7,346448 5,215715	6,299879 6,380713 8,447086 7,086036 7,298692 7,948244 7,862485 7,848524 6,375921	0,654089 0,65449 0,654638 0,654945 0,655066 0,657234 0,669787 0,674686 0,675396	0,107275 0,745885 0,121551 0,59305 0,143199 0,133013 0,183642 0,502076 1,160206	Mboat1 Ntng2 Sgpp1 Cyp20a1 Camsap2 Mtch1 Wars Rfx2 Fcgr3	NM_153546 NM_133501 NM_030750 ENSMUST00000060608 NM_001081360 NM_0119880 NM_001164314 NM_027787 NM_010188
17383467 17281971 17213428 17227483 17343035 17283930 17346349 17229620 17277387	6,041173 6,406147 7,931169 7,147408 6,560924 7,200163 7,566288 6,036273 7,694138 7,395317	6,695262 7,060637 8,585807 7,802353 7,21599 7,857397 8,236075 6,710959 8,369534 8,075942	6,192604 5,634828 8,325535 6,492986 7,155493 7,815231 7,678843 7,346448 5,215715 7,802212	6,299879 6,380713 8,447086 7,086036 7,298692 7,948244 7,862485 7,848524 6,375921 8,297579	0,654089 0,65449 0,654638 0,654945 0,655066 0,657234 0,669787 0,674686 0,675396	0,107275 0,745885 0,121551 0,59305 0,143199 0,133013 0,183642 0,502076 1,160206 0,495367	Mboat1 Ntng2 Sgpp1 Cyp20a1 Camsap2 Mtch1 Wars Rfx2 Fcgr3 Fos	NM_153546 NM_133501 NM_030750 ENSMUST00000060608 NM_001081360 NM_0119880 NM_001164314 NM_027787 NM_010188 NM_010234
17383467 17281971 17213428 17227483 17343035 17283930 17346349 17229620 17277387 17268729	6,041173 6,406147 7,931169 7,147408 6,560924 7,200163 7,566288 6,036273 7,694138 7,395317 7,084195	6,695262 7,060637 8,585807 7,802353 7,21599 7,857397 8,236075 6,710959 8,369534 8,075942 7,767599	6,192604 5,634828 8,325535 6,492986 7,155493 7,815231 7,678843 7,346448 5,215715 7,802212 7,934349	6,299879 6,380713 8,447086 7,086036 7,298692 7,948244 7,862485 7,848524 6,375921 8,297579 7,90494	0,654089 0,65449 0,654638 0,654945 0,655066 0,657234 0,669787 0,674686 0,675396 0,680625 0,683404	0,107275 0,745885 0,121551 0,59305 0,143199 0,133013 0,183642 0,502076 1,160206 0,495367 -0,029409	Mboat1 Ntng2 Sgpp1 Cyp20a1 Camsap2 Mtch1 Wars Rfx2 Fcgr3 Fos Fbxl20	NM_153546 NM_133501 NM_030750 ENSMUST00000060608 NM_001081360 NM_0119880 NM_001164314 NM_027787 NM_010188 NM_010234 NM_028149
17383467 17281971 17213428 17227483 17343035 17283930 17346349 17229620 17277387 17268729 17335297	6,041173 6,406147 7,931169 7,147408 6,560924 7,200163 7,566288 6,036273 7,694138 7,395317 7,084195 6,714341	6,695262 7,060637 8,585807 7,802353 7,21599 7,857397 8,236075 6,710959 8,369534 8,075942 7,767599 7,400104	6,192604 5,634828 8,325535 6,492986 7,155493 7,815231 7,678843 7,346448 5,215715 7,802212 7,934349 6,843205	6,299879 6,380713 8,447086 7,086036 7,298692 7,948244 7,862485 7,848524 6,375921 8,297579 7,90494 7,077991	0,654089 0,65449 0,654638 0,654945 0,655066 0,657234 0,669787 0,674686 0,675396 0,680625 0,683404 0,685763	0,107275 0,745885 0,121551 0,59305 0,143199 0,133013 0,183642 0,502076 1,160206 0,495367 -0,029409 0,234786	Mboat1 Ntng2 Sgpp1 Cyp20a1 Camsap2 Mtch1 Wars Rfx2 Fcgr3 Fos Fbxi20 Ppard	NM_153546 NM_133501 NM_030750 ENSMUST00000060608 NM_001081360 NM_019880 NM_001164314 NM_027787 NM_010188 NM_010234 NM_028149 NM_011145
17383467 17281971 17213428 17227483 17343035 17283930 17346349 17229620 17277387 17268729 17335297	6,041173 6,406147 7,931169 7,147408 6,560924 7,200163 7,566288 6,036273 7,694138 7,395317 7,084195 6,714341 6,393589	6,695262 7,060637 8,585807 7,802353 7,21599 7,857397 8,236075 6,710959 8,369534 8,075942 7,767599 7,400104 7,080763	6,192604 5,634828 8,325535 6,492986 7,155493 7,815231 7,678843 7,346448 5,215715 7,802212 7,934349 6,843205 6,808092	6,299879 6,380713 8,447086 7,086036 7,298692 7,948244 7,862485 7,848524 6,375921 8,297579 7,90494 7,077991 6,772245	0,654089 0,65449 0,654638 0,654945 0,655066 0,657234 0,669787 0,674686 0,675396 0,680625 0,683404 0,685763 0,687174	0,107275 0,745885 0,121551 0,59305 0,143199 0,133013 0,183642 0,502076 1,160206 0,495367 -0,029409 0,234786 -0,035847	Mboat1 Ntng2 Sgpp1 Cyp20a1 Camsap2 Mtch1 Wars Rfx2 Fcgr3 Fos Fbxl20 Ppard Parl	NM_153546 NM_133501 NM_030750 ENSMUST00000060608 NM_001081360 NM_019880 NM_001164314 NM_027787 NM_010188 NM_010234 NM_028149 NM_011145 NM_001005767
17383467 17281971 17213428 17227483 17343035 17283930 17346349 17229620 17277387 17268729 17335297 17329079 17321597	6,041173 6,406147 7,931169 7,147408 6,560924 7,200163 7,566288 6,036273 7,694138 7,395317 7,084195 6,714341 6,393589 7,093781	6,695262 7,060637 8,585807 7,802353 7,21599 7,857397 8,236075 6,710959 8,369534 8,075942 7,767599 7,400104 7,080763 7,781806	6,192604 5,634828 8,325535 6,492986 7,155493 7,815231 7,678843 7,346448 5,215715 7,802212 7,934349 6,843205 6,808092 7,982627	6,299879 6,380713 8,447086 7,086036 7,298692 7,948244 7,862485 7,848524 6,375921 8,297579 7,90494 7,077991 6,772245 8,048676	0,654089 0,65449 0,654638 0,654945 0,655066 0,657234 0,669787 0,674686 0,675396 0,680625 0,683404 0,685763 0,687174 0,688025	0,107275 0,745885 0,121551 0,59305 0,143199 0,133013 0,183642 0,502076 1,160206 0,495367 -0,029409 0,234786 -0,035847 0,066049	Mboat1 Ntng2 Sgpp1 Cyp20a1 Camsap2 Mtch1 Wars Rfx2 Fcgr3 Fos Fbx120 Ppard Racgap1	NM_153546 NM_133501 NM_030750 ENSMUST00000060608 NM_001081360 NM_019880 NM_001164314 NM_027787 NM_010188 NM_010234 NM_028149 NM_011145 NM_01105767 NM_012025
17383467 17281971 17213428 17227483 17343035 17283930 17346349 17229620 17277387 17268729 17335297 17329079 17321597 17274249 17223799	6,041173 6,406147 7,931169 7,147408 6,560924 7,200163 7,566288 6,036273 7,694138 7,395317 7,084195 6,714341 6,393589 7,093781 6,888887 6,701841	6,695262 7,060637 8,585807 7,802353 7,21599 7,857397 8,236075 6,710959 8,369534 8,075942 7,767599 7,400104 7,080763 7,781806 7,586748 7,402831	6,192604 5,634828 8,325535 6,492986 7,155493 7,815231 7,678843 7,346448 5,215715 7,802212 7,934349 6,843205 6,808092 7,982627 6,835749 7,965491	6,299879 6,380713 8,447086 7,086036 7,298692 7,948244 7,862485 7,848524 6,375921 8,297579 7,90494 7,077991 6,772245 8,048676 7,072966 8,125345	0,654089 0,65449 0,654638 0,654945 0,655066 0,657234 0,669787 0,674686 0,675396 0,680625 0,683404 0,685763 0,687174 0,688025 0,69861	0,107275 0,745885 0,121551 0,59305 0,143199 0,133013 0,183642 0,502076 1,160206 0,495367 -0,029409 0,234786 -0,035847 0,066049 0,237217 0,159854	Mboat1 Ntng2 Sgpp1 Cyp20a1 Camsap2 Mtch1 Wars Rfx2 Fcgr3 Fos Fbx120 Ppard Parl Racgap1 Pdia6	NM_153546 NM_133501 NM_030750 ENSMUST00000060608 NM_001081360 NM_019880 NM_001164314 NM_027787 NM_010188 NM_010234 NM_028149 NM_011145 NM_001005767 NM_012025 NM_027959 NM_001039493
17383467 17281971 17213428 17227483 17343035 17346349 17229620 17277387 17268729 17335297 17321597 1721597 17223799 17347163	6,041173 6,406147 7,931169 7,147408 6,560924 7,200163 7,566288 6,036273 7,694138 7,395317 7,084195 6,714341 6,393589 7,093781 6,888887 6,701841 8,48396	6,695262 7,060637 8,585807 7,802353 7,21599 7,857397 8,236075 6,710959 8,369534 8,075942 7,767599 7,400104 7,080763 7,781806 7,586748 7,402831 9,185109	6,192604 5,634828 8,325535 6,492986 7,155493 7,815231 7,678843 7,346448 5,215715 7,802212 7,934349 6,843205 6,808092 7,982627 6,835749 7,965491 5,920551	6,299879 6,380713 8,447086 7,086036 7,298692 7,948244 7,862485 7,848524 6,375921 8,297579 7,90494 7,077991 6,772245 8,048676 7,072966 8,125345 6,959862	0,654089 0,65449 0,654638 0,654945 0,655066 0,657234 0,669787 0,674686 0,675396 0,680625 0,683404 0,685763 0,687174 0,688025 0,697861 0,70099 0,701149	0,107275 0,745885 0,121551 0,59305 0,143199 0,133013 0,183642 0,502076 1,160206 0,495367 -0,029409 0,234786 -0,035847 0,066049 0,237217 0,159854 1,039311	Mboat1 Ntng2 Sgpp1 Cyp20a1 Camsap2 Mtch1 Wars Rfx2 Fcgr3 Fos Fbx120 Ppard Parl Racgap1 Pdia6 Plekhm3 Xdh	NM_153546 NM_133501 NM_030750 ENSMUST00000060608 NM_001081360 NM_01164314 NM_027787 NM_010188 NM_010188 NM_011145 NM_028149 NM_011145 NM_001005767 NM_012025 NM_027959 NM_001039493 NM_011723
17383467 17281971 17213428 17227483 17343035 17346349 17229620 17277387 17268729 17335297 17321597 17274249 17223799 17347163 17244140	6,041173 6,406147 7,931169 7,147408 6,560924 7,200163 7,566288 6,036273 7,694138 7,395317 7,084195 6,714341 6,393589 7,093781 6,888887 6,701841 8,48396 7,058547	6,695262 7,060637 8,585807 7,802353 7,21599 7,857397 8,236075 6,710959 8,369534 8,075942 7,767599 7,400104 7,080763 7,781806 7,586748 7,402831 9,185109 7,760675	6,192604 5,634828 8,325535 6,492986 7,155493 7,815231 7,678843 7,346448 5,215715 7,802212 7,934349 6,843205 6,808092 7,982627 6,835749 7,965491 5,920551 6,987411	6,299879 6,380713 8,447086 7,086036 7,298692 7,948244 7,862485 7,848524 6,375921 8,297579 7,90494 7,077991 6,772245 8,048676 7,072966 8,125345 6,959862 6,805999	0,654089 0,65449 0,654638 0,654945 0,655066 0,657234 0,669787 0,674686 0,675396 0,680625 0,683404 0,685763 0,687174 0,688025 0,697861 0,70099 0,701149 0,702128	0,107275 0,745885 0,121551 0,59305 0,143199 0,133013 0,183642 0,502076 1,160206 0,495367 -0,029409 0,234786 -0,035847 0,066049 0,237217 0,159854 1,039311 -0,181412	Mboat1 Ntng2 Sgpp1 Cyp20a1 Camsap2 Mtch1 Wars Rfx2 Fcgr3 Fos Fbxl20 Ppard Parl Racgap1 Pdia6 Plekhm3 Xdh Apaf1	NM_153546 NM_133501 NM_030750 ENSMUST00000060608 NM_001081360 NM_019880 NM_001164314 NM_027787 NM_010188 NM_010234 NM_028149 NM_011145 NM_001005767 NM_012025 NM_027959 NM_001039493 NM_011723 NM_011723 NM_0101042558
17383467 17281971 17213428 17227483 17343035 17283930 17346349 17229620 17277387 17268729 17335297 17321597 17274249 17223799 17347163 17244140 17262357	6,041173 6,406147 7,931169 7,147408 6,560924 7,200163 7,566288 6,036273 7,694138 7,395317 7,084195 6,714341 6,393589 7,093781 6,888887 6,701841 8,48396 7,058547 6,425956	6,695262 7,060637 8,585807 7,802353 7,21599 7,857397 8,236075 6,710959 8,369534 8,075942 7,767599 7,400104 7,080763 7,781806 7,586748 7,402831 9,185109 7,760675 7,133457	6,192604 5,634828 8,325535 6,492986 7,155493 7,815231 7,678843 7,346448 5,215715 7,802212 7,934349 6,808092 7,982627 6,835749 7,965491 5,920551 6,987411 6,850734	6,299879 6,380713 8,447086 7,086036 7,298692 7,948244 7,862485 7,848524 6,375921 8,297579 7,90494 7,077991 6,772245 8,048676 7,072966 8,125345 6,959862 6,805999 6,953052	0,654089 0,65449 0,654638 0,654945 0,655066 0,657234 0,669787 0,674686 0,675396 0,680625 0,683404 0,685763 0,687174 0,688025 0,697861 0,70099 0,701149 0,702128 0,707501	0,107275 0,745885 0,121551 0,59305 0,143199 0,133013 0,183642 0,502076 1,160206 0,495367 -0,029409 0,234786 -0,035847 0,066049 0,237217 0,159854 1,039311 -0,181412 0,102318	Mboat1 Ntng2 Sgpp1 Cyp20a1 Camsap2 Mtch1 Wars Rfx2 Fcgr3 Fos Fbx120 Ppard Parl Racgap1 Pdia6 Plekhm3 Xdh Apaf1 Rufy1	NM_153546 NM_133501 NM_030750 ENSMUST00000060608 NM_001081360 NM_019880 NM_001164314 NM_027787 NM_010188 NM_0101234 NM_028149 NM_011145 NM_001005767 NM_012025 NM_027959 NM_011723 NM_011723 NM_011723 NM_011723 NM_011723
17383467 17281971 17213428 17227483 17343035 17283930 17346349 17229620 17277387 17268729 17335297 17321597 17214249 17223799 17347163 17244140 17262357 17281843	6,041173 6,406147 7,931169 7,147408 6,560924 7,200163 7,566288 6,036273 7,694138 7,995317 7,084195 6,714341 6,393589 7,093781 6,888887 6,701841 8,48396 7,055547 6,425956 7,454978	6,695262 7,060637 8,585807 7,802353 7,21599 7,857397 8,236075 6,710959 8,369534 8,075942 7,767599 7,400104 7,78806 7,586748 7,402831 9,185109 7,760675 7,133457 8,168287	6,192604 5,634828 8,325535 6,492986 7,155493 7,815231 7,678843 7,346448 5,215715 7,802212 7,934349 6,843205 6,808092 7,982627 6,835749 7,965491 5,920551 6,987411 6,850734 6,600838	6,299879 6,380713 8,447086 7,086036 7,298692 7,948244 7,862485 7,848524 6,375921 8,297579 7,90494 7,077991 6,772245 8,048676 7,072966 8,125345 6,959862 6,805999 6,953052 7,186118	0,654089 0,65449 0,654638 0,654945 0,655066 0,657234 0,669787 0,674686 0,675396 0,680625 0,683404 0,685763 0,687174 0,688025 0,697861 0,70099 0,701149 0,702128 0,707501 0,713309	0,107275 0,745885 0,121551 0,59305 0,143199 0,133013 0,183642 0,502076 1,160206 0,495367 -0,029409 0,234786 -0,035847 0,066049 0,237217 0,159854 1,039311 -0,181412 0,102318	Mboat1 Ntng2 Sgpp1 Cyp20a1 Camsap2 Mtch1 Wars Rfx2 Fcgr3 Fos Fbxl20 Ppard Parl Racgap1 Pdia6 Plekhm3 Xdh Apaf1 Rufy1 Dhrs7	NM_153546 NM_133501 NM_030750 ENSMUST0000006008 NM_001081360 NM_019880 NM_001164314 NM_027787 NM_010188 NM_010234 NM_028149 NM_011145 NM_001005767 NM_012025 NM_027959 NM_001039493 NM_011723 NM_011723 NM_011723 NM_0117257 NM_025522
17383467 17281971 17213428 17227483 17343035 17283930 17346349 17229620 17277387 17268729 17335297 17329079 17321597 17223799 17347163 17244140 17262357 17281843 17350611	6,041173 6,406147 7,931169 7,147408 6,560924 7,200163 7,566288 6,036273 7,694138 7,395317 7,084195 6,714341 6,393589 7,093781 6,888887 6,701841 8,48396 7,058547 6,425956 7,454978 8,248373	6,695262 7,060637 8,585807 7,802353 7,21599 7,857397 8,236075 6,710959 8,369534 8,075942 7,767599 7,400104 7,080763 7,781806 7,781806 7,781809 7,760575 7,133457 8,168287 8,962262	6,192604 5,634828 8,325535 6,492986 7,155493 7,815231 7,678843 7,346448 5,215715 7,802212 7,934349 6,843205 6,808092 7,986247 6,9835749 7,965491 5,920551 6,87411 6,850734 6,600838 9,42992	6,299879 6,380713 8,447086 7,086036 7,298692 7,948244 7,862485 7,848524 6,375921 8,297579 7,90494 7,077991 6,772245 8,048676 7,072966 8,125345 6,959862 6,805999 6,953052 7,186118 9,756411	0,654089 0,65449 0,654638 0,654945 0,655066 0,657234 0,669787 0,674686 0,675396 0,680625 0,683404 0,685763 0,697861 0,70099 0,701149 0,702128 0,707501 0,713309 0,713889	0,107275 0,745885 0,121551 0,59305 0,143199 0,133013 0,183642 0,502076 1,160206 0,495367 -0,029409 0,234786 -0,035847 0,066049 0,237217 0,159854 1,039311 -0,181412 0,102318 0,58528	Mboat1 Ntng2 Sgpp1 Cyp20a1 Camsap2 Mtch1 Wars Rfx2 Fcgr3 Fos Fbxl20 Ppard Parl Racgap1 Pdia6 Plekhm3 Xdh Apaf1 Rufy1 Dhrs7 Snx2	NM_153546 NM_133501 NM_030750 ENSMUST0000006008 NM_001081360 NM_019880 NM_001164314 NM_027787 NM_010188 NM_010234 NM_028149 NM_011145 NM_001005767 NM_012025 NM_027959 NM_01039493 NM_011723 NM_011723 NM_011723 NM_011723 NM_0117257 NM_025522 NM_026386
17383467 17281971 17213428 17227483 17343035 17283930 17346349 17229620 17277387 17268729 17335297 17329079 17321597 17274249 1723799 17347163 17244140 17262357 17281843 17350611 17272619	6,041173 6,406147 7,931169 7,147408 6,560924 7,200163 7,566288 6,036273 7,694138 7,395317 7,084195 6,714341 6,393589 7,093781 6,701841 8,48396 7,058547 6,425956 7,454978 8,248373 5,99809	6,695262 7,060637 8,585807 7,802353 7,21599 7,857397 8,236075 6,710959 8,369534 8,075942 7,767599 7,400104 7,080763 7,781806 7,781806 7,781807 7,781809 7,760675 7,133457 8,168287 8,962262 6,712692	6,192604 5,634828 8,325535 6,492986 7,155493 7,815231 7,678843 7,346448 5,215715 7,802212 7,934349 6,843205 6,808092 7,982627 6,835749 7,965491 5,920551 6,87411 6,850734 6,600838 9,42992 7,665372	6,299879 6,380713 8,447086 7,086036 7,298692 7,948244 7,862485 7,848524 6,375921 8,297579 7,90494 7,077991 6,772245 8,048676 7,072966 8,125345 6,959862 6,805999 6,953052 7,186118 9,756411 7,954583	0,654089 0,65449 0,65449 0,654638 0,654945 0,655066 0,657234 0,669787 0,674686 0,675396 0,680625 0,683404 0,685763 0,687174 0,70099 0,701149 0,702128 0,707501 0,713309 0,713889 0,714602	0,107275 0,745885 0,121551 0,59305 0,143199 0,133013 0,183642 0,502076 1,160206 0,495367 -0,029409 0,234786 -0,035847 0,066049 0,237217 0,159854 1,039311 -0,181412 0,102318 0,58528 0,326491 0,289211	Mboat1 Ntng2 Sgpp1 Cyp20a1 Camsap2 Mtch1 Wars Rfx2 Fcgr3 Fos Fbxl20 Ppard Parl Racgap1 Pdia6 Plekhm3 Xdh Apaf1 Rufy1 Dhrs7 Snx2 Socs3	NM_153546 NM_133501 NM_030750 ENSMUST0000006008 NM_001081360 NM_019880 NM_01164314 NM_027787 NM_010188 NM_010234 NM_028149 NM_011145 NM_001005767 NM_012025 NM_027959 NM_001039493 NM_011723 NM_011723 NM_011723 NM_011723 NM_0117257 NM_025522 NM_026386 NM_007707
17383467 17281971 17213428 17227483 17343035 17346349 17229620 17277387 17268729 17335297 17329079 17321597 17274249 17223799 17347163 17244140 17262357 17281843 17350611 17272619 17214142	6,041173 6,406147 7,931169 7,147408 6,560924 7,200163 7,566288 6,036273 7,694138 7,395317 7,084195 6,714341 6,393589 7,093781 6,828887 6,701841 8,48396 7,058547 6,425956 7,454978 8,248373 5,99809 7,242959	6,695262 7,060637 8,585807 7,802353 7,21599 7,857397 8,236075 6,710959 8,369534 8,075942 7,767599 7,400104 7,080763 7,781806 7,781806 7,760675 7,133457 8,168287 8,962262 6,712692 7,962944	6,192604 5,634828 8,325535 6,492986 7,155493 7,815231 7,678843 7,346448 5,215715 7,802212 7,934349 6,843205 6,808092 7,982627 6,825749 7,965491 5,920551 6,80838 9,42992 7,665372 4,939416	6,299879 6,380713 8,447086 7,086036 7,298692 7,948244 7,862485 7,848524 6,375921 8,297579 7,90494 7,077991 6,772245 8,048676 7,072966 8,125345 6,959862 6,805999 6,953052 7,186118 9,756411 7,954583 5,328135	0,654089 0,65449 0,654498 0,654638 0,654945 0,655066 0,657234 0,669787 0,674686 0,675396 0,680625 0,683404 0,685763 0,687174 0,688025 0,70099 0,701149 0,702128 0,707501 0,713309 0,713889 0,714602 0,719985	0,107275 0,745885 0,121551 0,59305 0,143199 0,133013 0,183642 0,502076 1,160206 0,495367 -0,029409 0,234786 -0,035847 0,066049 0,237217 0,159854 1,039311 -0,181412 0,102318 0,58528 0,326491 0,289211 0,388719	Mboat1 Ntng2 Sgpp1 Cyp20a1 Camsap2 Mtch1 Wars Rfx2 Fcgr3 Fos Fbxl20 Ppard Parl Racgap1 Pdia6 Plekhm3 Xdh Apaf1 Rufy1 Dhrs7 Snx2 Socs3 Cxcr2	NM_153546 NM_133501 NM_030750 ENSMUST0000006008 NM_001081360 NM_019880 NM_01164314 NM_027787 NM_010188 NM_010234 NM_028149 NM_011145 NM_01005767 NM_012025 NM_027959 NM_001039493 NM_011723 NM_011723 NM_011723 NM_0117257 NM_025522 NM_026386 NM_007707 NM_009909
17383467 17281971 17213428 17227483 17343035 17346349 17229620 17277387 17268729 17335297 17321597 17321597 17321597 17274249 1732799 17347163 17244140 17262357 17281843 17350611 17272619 17214142 17326895	6,041173 6,406147 7,931169 7,147408 6,560924 7,200163 7,566288 6,036273 7,694138 7,395317 7,084195 6,714341 6,393589 7,093781 6,701841 8,48396 7,058547 6,425956 7,454978 8,248373 5,99809 7,242959 7,664179	6,695262 7,060637 8,585807 7,802353 7,21599 7,857397 8,236075 6,710959 8,369534 8,075942 7,767599 7,400104 7,080763 7,781806 7,586748 7,402831 9,185109 7,760675 7,133457 8,168287 8,962262 6,712692 7,962944 8,386475	6,192604 5,634828 8,325535 6,492986 7,155493 7,815231 7,678843 7,346448 5,215715 7,802212 7,934349 6,843205 6,808092 7,982627 6,835749 7,965491 5,920551 6,987411 6,850734 6,600838 9,42992 7,665372 4,939416 7,982733	6,299879 6,380713 8,447086 7,086036 7,298692 7,948244 7,862485 7,848524 6,375921 8,297579 7,90494 7,077991 6,772245 8,048676 7,072966 8,125345 6,959862 6,805999 6,953052 7,186118 9,756411 7,954583 5,328135 7,482109	0,654089 0,65449 0,65449 0,654638 0,654945 0,655066 0,657234 0,669787 0,674686 0,675396 0,680625 0,683404 0,685763 0,687174 0,688025 0,697861 0,70099 0,701149 0,702128 0,707501 0,713309 0,713889 0,714602 0,719985 0,722296	0,107275 0,745885 0,121551 0,59305 0,143199 0,133013 0,183642 0,502076 1,160206 0,495367 -0,029409 0,234786 -0,035847 0,066049 0,237217 0,159854 1,039311 -0,181412 0,102318 0,58528 0,326491 0,289211 0,388719 -0,500624	Mboat1 Ntng2 Sgpp1 Cyp20a1 Camsap2 Mtch1 Wars Rfx2 Fcgr3 Fos Fbxl20 Ppard Parl Racgap1 Pdia6 Plekhm3 Xdh Apaf1 Rufy1 Dhrs7 Snx2 Socs3 Cxcr2 Bach1	NM_153546 NM_133501 NM_030750 ENSMUST0000006008 NM_001081360 NM_019880 NM_01164314 NM_027787 NM_010188 NM_010234 NM_028149 NM_011145 NM_01005767 NM_012025 NM_027959 NM_001093493 NM_011723 NM_011723 NM_011723 NM_011723 NM_0117257 NM_025522 NM_026386 NM_007707 NM_009909 NM_007520
17383467 17281971 17213428 17227483 17343035 17346349 17229620 17277387 17268729 17335297 17321597 17321597 17321597 17274249 1732799 17347163 17244140 17262357 17281843 17350611 17272619 17214142 17326895 17317858	6,041173 6,406147 7,931169 7,147408 6,560924 7,200163 7,566288 6,036273 7,694138 7,395317 7,084195 6,714341 6,393589 7,093781 6,88887 6,701841 8,48396 7,058547 6,425956 7,454978 8,248373 5,99809 7,242959 7,664179 6,285686	6,695262 7,060637 8,585807 7,802353 7,21599 7,857397 8,236075 6,710959 8,369534 8,075942 7,767599 7,400104 7,080763 7,781806 7,586748 7,402831 9,185109 7,760675 7,133457 8,168287 8,962262 6,712692 7,962944 8,386475 7,018915	6,192604 5,634828 8,325535 6,492986 7,155493 7,815231 7,678843 7,346448 5,215715 7,802212 7,934349 6,843205 6,808092 7,982627 6,835749 7,965491 5,920551 6,987411 6,850734 6,600838 9,42992 7,665372 4,939416 7,982733 5,862745	6,299879 6,380713 8,447086 7,086036 7,298692 7,948244 7,862485 7,848524 6,375921 8,297579 7,90494 7,077991 6,772245 8,048676 7,072966 8,125345 6,959862 6,805999 6,953052 7,186118 9,756411 7,954583 5,328135 7,482109 6,228966	0,654089 0,65449 0,65449 0,654638 0,654945 0,655066 0,657234 0,669787 0,674686 0,675396 0,680625 0,683404 0,685763 0,687174 0,688025 0,697861 0,70099 0,701149 0,702128 0,707501 0,713309 0,713889 0,714602 0,719985 0,722296	0,107275 0,745885 0,121551 0,59305 0,143199 0,133013 0,183642 0,502076 1,160206 0,495367 -0,029409 0,234786 -0,035847 0,066049 0,237217 0,159854 1,039311 -0,181412 0,102318 0,58528 0,326491 0,289211 0,388719 -0,500624 0,366221	Mboat1 Ntng2 Sgpp1 Cyp20a1 Camsap2 Mtch1 Wars Rfx2 Fcgr3 Fos Fbxl20 Ppard Parl Racgap1 Pdia6 Plekhm3 Xdh Apaf1 Rufy1 Dhrs7 Snx2 Socs3 Cxcr2 Bach1 Ptk2	NM_153546 NM_133501 NM_030750 ENSMUST0000006008 NM_001081360 NM_019880 NM_01164314 NM_027787 NM_010188 NM_010234 NM_028149 NM_011145 NM_001005767 NM_012025 NM_027959 NM_001039493 NM_011723 NM_011723 NM_011723 NM_011723 NM_0117257 NM_025522 NM_026386 NM_007707 NM_009909 NM_007520 NM_007520 NM_00750
17383467 17281971 17213428 17227483 17343035 17346349 17229620 17277387 17268729 17335297 17321597 17321597 17274249 17223799 17347163 17244140 17262357 17281843 17350611 1727619 17214142 17326895 17317858 17268120	6,041173 6,406147 7,931169 7,147408 6,560924 7,200163 7,566288 6,036273 7,694138 7,395317 7,084195 6,714341 6,393589 7,093781 6,701841 8,48396 7,05847 6,425956 7,454978 8,248373 5,99809 7,242959 7,664179 6,285686 6,054599	6,695262 7,060637 8,585807 7,802353 7,21599 7,857397 8,236075 6,710959 8,369534 8,075942 7,767599 7,400104 7,080763 7,781806 7,586748 7,402831 9,185109 7,6055 7,133457 8,168287 8,962262 6,712692 7,962944 8,386475 7,018915 6,788635	6,192604 5,634828 8,325535 6,492986 7,155493 7,815231 7,678843 7,346448 5,215715 7,802212 7,934349 6,843205 6,808092 7,982627 6,835749 7,965491 5,920551 6,80838 9,42992 7,665372 4,939416 7,982733 5,862745 5,679688	6,299879 6,380713 8,447086 7,086036 7,298692 7,948244 7,862485 7,848524 6,375921 8,297579 7,90494 7,077991 6,772245 8,048676 7,072966 8,125345 6,959862 6,85999 6,953052 7,186118 9,756411 7,954583 5,328135 7,482109 6,228966 5,405785	0,654089 0,65449 0,65449 0,654638 0,654945 0,655066 0,657234 0,669787 0,674686 0,675396 0,680625 0,683404 0,685763 0,687174 0,688025 0,697861 0,70099 0,701149 0,702128 0,707501 0,713309 0,713889 0,714602 0,719985 0,792296 0,733229 0,734036	0,107275 0,745885 0,121551 0,59305 0,143199 0,133013 0,183642 0,502076 1,160206 0,495367 -0,029409 0,234786 -0,035847 0,066049 0,237217 0,159854 1,039311 -0,181412 0,102318 0,58528 0,326491 0,289211 0,388719 -0,500624 0,366221 -0,273903	Mboat1 Ntng2 Sgpp1 Cyp20a1 Camsap2 Mtch1 Wars Rfx2 Fcgr3 Fos Fbxl20 Ppard Parl Racgap1 Pdia6 Plekhm3 Xdh Apaf1 Rufy1 Dhrs7 Snx2 Socs3 Cxcr2 Bach1 Ptk2 Pdk2	NM_153546 NM_133501 NM_030750 ENSMUST0000006008 NM_001081360 NM_019880 NM_01164314 NM_027787 NM_010188 NM_010234 NM_028149 NM_011145 NM_01105767 NM_012025 NM_0210255 NM_027959 NM_001039493 NM_011723 NM_001042558 NM_172557 NM_025522 NM_026386 NM_007707 NM_009909 NM_007520 NM_007982 NM_133667
17383467 17281971 17213428 17227483 17343035 17346349 17229620 17277387 17268729 17335297 17321597 17321597 17321597 17274249 1732799 17347163 17244140 17262357 17281843 17350611 17272619 17214142 17326895 17317858	6,041173 6,406147 7,931169 7,147408 6,560924 7,200163 7,566288 6,036273 7,694138 7,395317 7,084195 6,714341 6,393589 7,093781 6,88887 6,701841 8,48396 7,058547 6,425956 7,454978 8,248373 5,99809 7,242959 7,664179 6,285686	6,695262 7,060637 8,585807 7,802353 7,21599 7,857397 8,236075 6,710959 8,369534 8,075942 7,767599 7,400104 7,080763 7,781806 7,586748 7,402831 9,185109 7,760675 7,133457 8,168287 8,962262 6,712692 7,962944 8,386475 7,018915	6,192604 5,634828 8,325535 6,492986 7,155493 7,815231 7,678843 7,346448 5,215715 7,802212 7,934349 6,843205 6,808092 7,982627 6,835749 7,965491 5,920551 6,987411 6,850734 6,600838 9,42992 7,665372 4,939416 7,982733 5,862745	6,299879 6,380713 8,447086 7,086036 7,298692 7,948244 7,862485 7,848524 6,375921 8,297579 7,90494 7,077991 6,772245 8,048676 7,072966 8,125345 6,959862 6,805999 6,953052 7,186118 9,756411 7,954583 5,328135 7,482109 6,228966	0,654089 0,65449 0,65449 0,654638 0,654945 0,655066 0,657234 0,669787 0,674686 0,675396 0,680625 0,683404 0,685763 0,687174 0,688025 0,697861 0,70099 0,701149 0,702128 0,707501 0,713309 0,713889 0,714602 0,719985 0,722296	0,107275 0,745885 0,121551 0,59305 0,143199 0,133013 0,183642 0,502076 1,160206 0,495367 -0,029409 0,234786 -0,035847 0,066049 0,237217 0,159854 1,039311 -0,181412 0,102318 0,58528 0,326491 0,289211 0,388719 -0,500624 0,366221	Mboat1 Ntng2 Sgpp1 Cyp20a1 Camsap2 Mtch1 Wars Rfx2 Fcgr3 Fos Fbxl20 Ppard Parl Racgap1 Pdia6 Plekhm3 Xdh Apaf1 Rufy1 Dhrs7 Snx2 Socs3 Cxcr2 Bach1 Ptk2	NM_153546 NM_133501 NM_030750 ENSMUST0000006008 NM_001081360 NM_019880 NM_01164314 NM_027787 NM_010188 NM_010234 NM_028149 NM_011145 NM_001005767 NM_012025 NM_027959 NM_001039493 NM_011723 NM_011723 NM_011723 NM_011723 NM_0117257 NM_025522 NM_026386 NM_007707 NM_009909 NM_007520 NM_007520 NM_00750

47056740	7.550500		= .=000.	= 000==0	0.740000	0.000455	51.14	
17356743	7,550539	8,299877	7,470304	7,830759	0,749338	0,360455	Ehd1	ENSMUST00000025684
17330523	7,282652	8,031994	7,324233	7,411808	0,749342	0,087575	Atp6v1a	ENSMUST00000114666
17239856	6,563095	7,317572	5,219164	5,768072	0,754477	0,548908	Akap7	NM_018747
17284065	6,995052	7,752645	6,307638	6,753062	0,757593	0,445424	Cdc42bpb	NM_183016
17233022	8,649394	9,408406	10,18995	10,30242	0,759012	0,11247	Cd24a	NM_009846
17258140	7,088554	7,850896	7,233706	7,325498	0,762342	0,091792	Tmem104	NM_001033393
17353948	7,367426	8,132895	7,098341	7,399754	0,765469	0,301413	Gnpda1	NM_011937
17295588	7,378003	8,144991	6,284079	7,173254	0,766988	0,889175	Naip5	NM_010870
17226771	6,500519	7,274589	7,985357	8,278639	0,77407	0,293282	Ikbke	NM_019777
17224270	6,357535	7,132519	5,406294	5,685338	0,774984	0,279044	Tmbim1	ENSMUST00000113796
17329316	7,046887	7,823175	6,171291	5,734249	0,776288	-0,437042	Dgkg	NM_138650
17356002	8,611594	9,397997	9,788301	10,01997	0,786403	0,231669	Unc93b1	NM_019449
17327995	5,837494	6,632035	5,799664	5,709239	0,794541	-0,090425	Carhsp1	NM_025821
17217182	6,123728	6,919407	6,311366	6,757595	0,795679	0,446229	Nuak2	NM_001195025
17231003	6,039546	6,836799	6,965693	6,812791	0,797253	-0,152902	Mfsd7b	NM 001081259
17222527	6,725369	7,527747	5,307161	5,970162	0,802378	0,663001	Tbc1d8	NM 018775
17314190	5,157845	5,963087	3,793981	4,35394	0,805242	0,559959	Shank3	ENSMUST00000109309
17306856	8,234545	9,04537	8,680914	8,794653	0,810825	0,113739	Dhrs1	NM 026819
17234175	6,619415	7,435453	7,658133	7,635149	0,816038	-0,022984	Ipmk	NM 027184
17329723	8,176059	8,995285	8,857574	8,928021	0,819226	0,070447	Acap2	NM 030138
17223932	6,304934	7,131555	7,383703	7,247778	0,826621	-0,135925	Lancl1	NM 001190985
17329105	6,106488	6,943847	7,068325	6,671293	0,837359	-0,397032	Abcc5	NM 013790
17231891	7,221858	8,069007	7,877102	8,209234	0,847149	0,332132	Map3k5	NM 008580
17295569	6,106658		5,05092	5,99629	0,849509	0,94537	Naip2	NM 010872
17310882	6,954261	6,956167 7,810472	4,857493	5,534661	0,856211	0,677168	'	NM 018755
		7,502509	7,536669				Pgcp	
17241130	6,644825	<i>'</i>		7,574985	0,857684	0,038316	Slc29a3	NM_023596
17307518	5,183063	6,041934	5,804796	5,325121	0,858871	-0,479675	Mir15a	NR_029733
17316956	6,317077	7,179224	6,322385	6,346126	0,862147	0,023741	Trps1	NM_032000
17268227	8,052107	8,916366	8,515485	8,32166	0,864259	-0,193825	Abi3	NM_025659
17238424	7,118496	7,98763	7,802932	7,833811	0,869134	0,030879	Cnpy2	NM_019953
17289889	7,116374	7,995862	9,424937	9,405313	0,879488	-0,019624	Il6st	NM_010560
17293675	6,341447	7,221475	6,929873	7,138278	0,880028	0,208405	Cdc14b	NM_172587
17354434	5,892123	6,772179	7,430119	6,851239	0,880056	-0,57888	Aldh7a1	ENSMUST00000174518
17345038	7,846705	8,730124	9,022942	9,170952	0,883419	0,14801	Cd2ap	NM_009847
17259679	6,965156	7,860522	4,195534	5,027074	0,895366	0,83154	Metrnl	NM_144797
17314564	5,600322	6,506842	6,385886	6,640608	0,90652	0,254722	Tmem106c	NM_201359
17254188	5,412025	6,322153	5,455376	5,577021	0,910128	0,121645	Slfn3	NM_011409
17240330	6,607723	7,51919	5,977628	6,075196	0,911467	0,097568	Slc16a10	NM_001114332
17319192	7,224529	8,138163	8,049911	8,133665	0,913634	0,083754	Tmem184b	NM_172608
17322359	6,138439	7,054343	6,01297	6,382364	0,915904	0,369394	Zfp385a	NM_013866
17236882	7,854383	8,775178	7,749078	7,589499	0,920795	-0,159579	Dusp6	NM_026268
17342581	6,098496	7,021397	6,800559	7,020883	0,922901	0,220324	Decr2	ENSMUST00000040907
17328625	7,766894	8,697867	7,211559	7,803964	0,930973	0,592405	Sdf2l1	NM_022324
17304740	5,949855	6,893625	7,758348	8,148488	0,94377	0,39014	Sh3bp5	ENSMUST00000091903
17241692	6,390726	7,340114	7,735371	7,111695	0,949388	-0,623676	Cdk1	ENSMUST00000020099
17325979	6,764509	7,716749	3,795039	3,63881	0,95224	-0,156229	Gcet2	NM 001159297
17281025	7,576966	8,534466	6,64538	7,030034	0,9575	0,384654	Heatr5a	NM_177171
17301176	6,63165	7,589678	7,300785	7,309306	0,958028	0,008521	Wdfy2	NM_175546
17214368	6,435288	7,393861	5,127243	6,001647	0,958573	0,874404	Cyp27a1	NM 024264
17279404	8,648836	9,608633	9,588039	9,983295	0,959797	0,395256	Pld4	NM_178911
17315349	7,686694	8,657921	7,844229	8,241407	0,971227	0,397178	Mfsd5	NM_134100
17313343	6,641037	7,61781	6,244538	6,365293	0,976773	0,120755	Baiap2	ENSMUST00000026436
17334205	6,402025	7,389976	7,43251	7,628057	0,987951	0,120733	Abca3	NM_013855
17334545	6,172929	7,16865	5,96417	6,071802	0,987931	0,107632	Clcn7	NM_011930
17334343	6,216537	7,10803	6,482624	6,624117	0,997542	0,107032	Mocs1	NM 020042
		9,394493			1,000152			NM 001199296
17269521	8,394341	J,JJ4493	8,395644	8,536424	1,000152	0,14078	Acly	IAIAI_001133530

				,				
17245481	6,554946	7,564241	6,578812	7,174853	1,009295	0,596041	Rassf3	NM_138956
17226194	6,070665	7,089951	5,571286	5,8742	1,019286	0,302914	Ralb	NM_022327
17256473	6,311055	7,330361	6,429249	6,996138	1,019306	0,566889	Atp6v0a1	NM_016920
17326069	7,671528	8,691006	8,573864	9,122976	1,019478	0,549112	Retnla	NM_020509
17286587	9,106037	10,13084	9,893567	10,20862	1,024803	0,315053	Ly86	NM 010745
17355092	7,401188	8,427783	7,078792	7,462524	1,026595	0,383732	Lman1	ENSMUST00000120461
17228544	7,521658	8,549862	8,855238	8,532072	1,028204	-0,323166	Soat1	NM 009230
17327137	6,727843	7,756936	5,314826	6,305424	1,029093	0,990598	ltsn1	NM 010587
17268909	6,529805	7,561562	8,153514	7,592195	1,031757	-0,561319	Top2a	NM 011623
17222549	6,339431	7,371259	6,03305	6,253771	1,031828	0,220721	Rnf149	NM 001033135
17265247	6,164906	7,204969	6,547497	6,518496	1,040063	-0,029001	Pelp1	NM 029231
17357810	8,090467	9,148258	6,651259	7,503748	1,057791	0,852489	Mpeg1	NM 010821
17343263	7,407168	8,468616	10,74929	10,60341	1,061448	-0,14588	Sik1	NM 010831
17326567	7,278197	8,340281	5,883628	6,524311	1,062084	0,640683	Gbe1	ENSMUST00000163832
17339772	5,710759	6,774759	6,600175	7,158906	1,064	0,558731	Rasgrp3	NM 207246
17266952	8,274776	9,343382	7,103242	8,3866	1,068606	1,283358	Ccl9	NM 011338
17289249	6,82862	7,900139	7,103242	7,641573	1,071519	0,197003	Wdr41	NM 172590
17243617	6,006196	7,081518	6,506867	7,016105	1,075322	0,509238	Ckap4	NM 175451
							·	_
17227278	6,988655	8,064938	6,762311	6,969554	1,076283	0,207243	Rnpep	NM_145417
17301558	6,743875	7,822746	6,363808	6,524606	1,078871	0,160798	Kctd9	NM_001111028
17266960	7,722705	8,802958	7,50527	8,615627	1,080253	1,110357	Ccl6	NM_009139
17274540	5,590309	6,672502	7,932548	7,402667	1,082193	-0,529881	Rrm2	NM_009104
17483577	9,616143	10,70433	7,455223	8,969418	1,088187	1,514195	Itgam	NM_001082960
17337269	6,68471	7,778625	7,515759	7,904148	1,093915	0,388389	Nrm	NM_134122
17338378	6,255622	7,349814	4,208319	4,381313	1,094192	0,172994	Treml4	NM_001033922
17324420	6,366271	7,462177	9,181595	9,36886	1,095906	0,187265	St6gal1	NM_145933
17332783	6,353034	7,453197	7,25614	7,651301	1,100163	0,395161	Snx9	ENSMUST00000002436
17337741	7,85463	8,959998	5,732572	6,887778	1,105368	1,155206	Tnfrsf21	NM_178589
17295607	7,041655	8,148333	6,025925	6,868568	1,106678	0,842643	Naip6	NM_010871
17346528	8,400613	9,507429	6,433522	7,582873	1,106816	1,149351	C3	NM_009778
17298364	5,621681	6,730614	8,257797	7,871689	1,108933	-0,386108	Nt5dc2	NM_027289
17352439	6,639665	7,75432	7,213013	7,18357	1,114655	-0,029443	Arhgap12	NM_001039692
17320538	5,56133	6,706966	5,428593	5,737524	1,145636	0,308931	Arsa	NM_009713
17243382	6,169161	7,323791	4,692461	5,629131	1,15463	0,93667	Gna15	NM_010304
17211713	5,217495	6,376225	5,350207	5,945888	1,15873	0,595681	Ptpn18	NM_011206
17345570	6,301345	7,461648	7,271828	7,444485	1,160303	0,172657	Cnpy3	NM_028065
17292915	7,216951	8,382987	8,474016	8,961411	1,166036	0,487395	Dok3	NM_013739
17326075	8,644369	9,815186	2,864657	3,653805	1,170817	0,789148	Retnlg	NM_181596
17252995	6,22764	7,399069	7,652436	7,768938	1,171429	0,116502	Slc43a2	NM_001199284
17217440	5,392055	6,574775	5,77127	6,165281	1,18272	0,394011	Cyb5r1	ENSMUST00000027726
17289527	7,148633	8,346066	8,875358	9,398293	1,197433	0,522935	Cd180	ENSMUST00000022124
17233536	6,521334	7,724533	6,937801	7,14242	1,203199	0,204619	P4ha1	NM_011030
17318713	6,414796	7,618502	6,516696	6,902083	1,203706	0,385387	Arhgap39	NM 001168288
17342642	8,622986	9,830132	9,820033	10,36827	1,207146	0,548237	Dusp1	ENSMUST00000025025
17218261	8,163346	9,377582	7,343568	8,148682	1,214236	0,805114	Ncf2	NM 010877
17493869	5,886518	7,102915	4,188938	4,768302	1,216397	0,579364	P2ry6	NM 183168
17347042	5,738036	6,97031	7,259377	7,023698	1,232274	-0,235679	Ndc80	NM_023294
17284114	6,446747	7,691731	6,741036	7,076285	1,244984	0,335249	Ckb	NM_021273
17251485	5,840549	7,089443	7,596467	7,207796	1,248894	-0,388671	Aurkb	NM 011496
17327084	7,157288	8,430135	8,590483	8,410227	1,272847	-0,180256	Ifngr2	ENSMUST00000023687
17314260	5,927593	7,203182	7,438414	7,803335	1,275589	0,364921	Lrrk2	NM 025730
17292215	5,337922	6,61362	5,065978	5,438004	1,275698	0,372026	Kif13a	NM 010617
17289794	6,033621	7,311998	6,687064	7,047769	1,278377	0,360705	Plk2	NM 152804
17327465	5,596242	6,878515	9,28458	8,627478	1,282273	-0,657102	Ets2	NM 011809
17507288	3,029332	4,312334	4,89523	4,306369	1,283002	-0,588861	Shcbp1	NM 011369
17211043	6,718893	8,007373	7,757994	7,954595	1,28848	0,196601	Sgk3	NM_133220
1,511043	0,7 10033	0,001313	1,131337	1,557555	1,20070	0,130001	201/2	11111_133220

		_						
17307774	4,501289	5,799735	6,180698	5,702818	1,298446	-0,47788	Esco2	ENSMUST00000022613
17330918	6,676832	7,979583	6,008067	6,857502	1,302751	0,849435	Alcam	ENSMUST00000023312
17257444	7,074046	8,37866	4,131145	4,410233	1,304614	0,279088	Ace	NM_207624
17281084	5,923933	7,233448	7,229249	6,513737	1,309515	-0,715512	Egln3	NM_028133
17321761	5,846561	7,156965	5,773309	6,243011	1,310404	0,469702	Smagp	NM_174992
17338371	6,432153	7,771784	4,300385	4,798179	1,339631	0,497794	Trem3	NM_021407
17337228	7,08053	8,422618	5,938332	7,754211	1,342088	1,815879	ler3	NM 133662
17231033	5,928178	7,270733	8,94608	9,293113	1,342555	0,347033	Atf3	NM 007498
17324446	5,672898	7,022024	5,948678	6,04128	1,349126	0,092602	Rtp4	NM 023386
17383216	6,999491	8,35165	6,253881	6,973873	1,352159	0,719992	Slc2a6	NM 172659
17319738	6,51567	7,870815	6,718919	7,367139	1,355145	0,64822	Nfam1	ENSMUST00000109503
17218349	8,267183	9,688818	8,028639	8,442446	1,421635	0,413807	Glul	NM 008131
17211286	7,80631	9,230669	7,81525	8,192135	1,424359	0,376885	Ly96	NM 016923
17497525	5,802816	7,24529	4,69452	5,212566	1,442474	0,518046	Adam8	ENSMUST00000106069
17310673	7,272774	8,723026	5.864444	7,024254	1,450252	1,15981	Ank	NM 020332
17316677	6,577716	8,028574	4,366122	5,948697	1,450858	1,582575	Slc7a8	NM 016972
17300477	6,038124	7,504622	6,597444	6,818183	1,466498	0,220739	Uap1	NM 133806
	5,379539							_
17217399		6,853388	3,245401	3,312111	1,473849	0,06671	Chi3l1	NM_007695
17331918	5,833219	7,310307	5,329571	5,875249	1,477088	0,545678	Tiam1	ENSMUST00000114124
17357640	5,071126	6,553259	3,785595	3,955008	1,482133	0,169413	Ms4a4a	XM_986941
17279349	6,666887	8,149317	5,968636	6,547713	1,48243	0,579077	Adssl1	NM_007421
17315352	4,780274	6,275684	5,709484	5,763056	1,49541	0,053572	Espl1	NM_001014976
17495839	9,259654	10,76009	6,737537	8,84012	1,500436	2,102583	Igsf6	NM_030691
17242376	6,038552	7,549084	6,534376	6,793246	1,510532	0,25887	Pfkl	NM_008826
17291694	6,182758	7,694995	5,813475	6,515891	1,512237	0,702416	Serpinb1a	NM_025429
17254153	5,52689	7,062624	5,935768	6,12475	1,535734	0,188982	Slfn5	NM_183201
17241623	7,299825	8,846231	7,749156	8,083022	1,546406	0,333866	Reep3	NM_001204915
17493949	4,177262	5,726651	4,754477	5,378238	1,549389	0.633761	Folk?	NM 008035
	1)177202	3,720031	7,737777	3,370230	1,343303	0,623761	Folr2	14141_006033
17338416	5,163425	6,725471	3,648878	4,07131	1,562046	0,623761	Trem2	NM_031254
17338416 17256784								NM_031254 ENSMUST00000100403
	5,163425	6,725471	3,648878	4,07131	1,562046	0,422432	Trem2	NM_031254
17256784	5,163425 6,155197	6,725471 7,719026	3,648878 6,158118	4,07131 6,359836	1,562046 1,563829	0,422432 0,201718	Trem2 Tmem106a	NM_031254 ENSMUST00000100403
17256784 17333731	5,163425 6,155197 5,183623	6,725471 7,719026 6,759561	3,648878 6,158118 3,251913	4,07131 6,359836 3,167332	1,562046 1,563829 1,575938	0,422432 0,201718 -0,084581	Trem2 Tmem106a Fpr2	NM_031254 ENSMUST00000100403 NM_008039
17256784 17333731 17310044	5,163425 6,155197 5,183623 5,666428	6,725471 7,719026 6,759561 7,247518	3,648878 6,158118 3,251913 7,503334	4,07131 6,359836 3,167332 8,121386	1,562046 1,563829 1,575938 1,58109	0,422432 0,201718 -0,084581 0,618052	Trem2 Tmem106a Fpr2 Lifr	NM_031254 ENSMUST00000100403 NM_008039 NM_013584
17256784 17333731 17310044 17256502	5,163425 6,155197 5,183623 5,666428 5,577532	6,725471 7,719026 6,759561 7,247518 7,166973	3,648878 6,158118 3,251913 7,503334 5,237371	4,07131 6,359836 3,167332 8,121386 5,780448	1,562046 1,563829 1,575938 1,58109 1,589441	0,422432 0,201718 -0,084581 0,618052 0,543077	Trem2 Tmem106a Fpr2 Lifr Naglu	NM_031254 ENSMUST00000100403 NM_008039 NM_013584 NM_013792
17256784 17333731 17310044 17256502 17270877	5,163425 6,155197 5,183623 5,666428 5,577532 6,13765	6,725471 7,719026 6,759561 7,247518 7,166973 7,762251	3,648878 6,158118 3,251913 7,503334 5,237371 6,442562	4,07131 6,359836 3,167332 8,121386 5,780448 6,914905	1,562046 1,563829 1,575938 1,58109 1,589441 1,624601	0,422432 0,201718 -0,084581 0,618052 0,543077 0,472343	Trem2 Tmem106a Fpr2 Lifr Naglu Icam2	NM_031254 ENSMUST00000100403 NM_008039 NM_013584 NM_013792 ENSMUST00000001055
17256784 17333731 17310044 17256502 17270877 17335467	5,163425 6,155197 5,183623 5,666428 5,577532 6,13765 6,474934	6,725471 7,719026 6,759561 7,247518 7,166973 7,762251 8,102732	3,648878 6,158118 3,251913 7,503334 5,237371 6,442562 6,59338	4,07131 6,359836 3,167332 8,121386 5,780448 6,914905 7,968711	1,562046 1,563829 1,575938 1,58109 1,589441 1,624601 1,627798	0,422432 0,201718 -0,084581 0,618052 0,543077 0,472343 1,375331	Trem2 Tmem106a Fpr2 Lifr Naglu Icam2 Cdkn1a	NM_031254 ENSMUST00000100403 NM_008039 NM_013584 NM_013792 ENSMUST00000001055 NM_007669
17256784 17333731 17310044 17256502 17270877 17335467 17339108	5,163425 6,155197 5,183623 5,666428 5,577532 6,13765 6,474934 7,99635	6,725471 7,719026 6,759561 7,247518 7,166973 7,762251 8,102732 9,625868	3,648878 6,158118 3,251913 7,503334 5,237371 6,442562 6,59338 8,59907	4,07131 6,359836 3,167332 8,121386 5,780448 6,914905 7,968711 8,968794	1,562046 1,563829 1,575938 1,58109 1,589441 1,624601 1,627798 1,629518	0,422432 0,201718 -0,084581 0,618052 0,543077 0,472343 1,375331 0,369724	Trem2 Tmem106a Fpr2 Lifr Naglu Icam2 Cdkn1a Man2a1	NM_031254 ENSMUST00000100403 NM_008039 NM_013584 NM_013792 ENSMUST00000001055 NM_007669 ENSMUST00000086723
17256784 17333731 17310044 17256502 17270877 17335467 17339108 17211790	5,163425 6,155197 5,183623 5,666428 5,577532 6,13765 6,474934 7,99635 6,050535	6,725471 7,719026 6,759561 7,247518 7,166973 7,762251 8,102732 9,625868 7,68284	3,648878 6,158118 3,251913 7,503334 5,237371 6,442562 6,59338 8,59907 8,140153	4,07131 6,359836 3,167332 8,121386 5,780448 6,914905 7,968711 8,968794 7,975585	1,562046 1,563829 1,575938 1,58109 1,589441 1,624601 1,627798 1,629518 1,632305	0,422432 0,201718 -0,084581 0,618052 0,543077 0,472343 1,375331 0,369724 -0,164568	Trem2 Tmem106a Fpr2 Lifr Naglu Icam2 Cdkn1a Man2a1 Hs6st1	NM_031254 ENSMUST00000100403 NM_008039 NM_013584 NM_013792 ENSMUST00000001055 NM_007669 ENSMUST00000086723 NM_015818
17256784 17333731 17310044 17256502 17270877 17335467 17339108 17211790 17258107	5,163425 6,155197 5,183623 5,666428 5,577532 6,13765 6,474934 7,99635 6,050535 6,020436	6,725471 7,719026 6,759561 7,247518 7,166973 7,762251 8,102732 9,625868 7,68284 7,67225	3,648878 6,158118 3,251913 7,503334 5,237371 6,442562 6,59338 8,59907 8,140153 4,786313	4,07131 6,359836 3,167332 8,121386 5,780448 6,914905 7,968711 8,968794 7,975585 5,079708	1,562046 1,563829 1,575938 1,58109 1,589441 1,624601 1,627798 1,629518 1,632305 1,651814	0,422432 0,201718 -0,084581 0,618052 0,543077 0,472343 1,375331 0,369724 -0,164568 0,293395	Trem2 Tmem106a Fpr2 Lifr Naglu Icam2 Cdkn1a Man2a1 Hs6st1 Cd300a	NM_031254 ENSMUST00000100403 NM_008039 NM_013584 NM_013792 ENSMUST00000001055 NM_007669 ENSMUST00000086723 NM_015818 NM_170758
17256784 17333731 17310044 17256502 17270877 17335467 17339108 17211790 17258107 17354764	5,163425 6,155197 5,183623 5,666428 5,577532 6,13765 6,474934 7,99635 6,050535 6,020436 5,438623	6,725471 7,719026 6,759561 7,247518 7,166973 7,762251 8,102732 9,625868 7,68284 7,67225 7,092581	3,648878 6,158118 3,251913 7,503334 5,237371 6,442562 6,59338 8,59907 8,140153 4,786313 4,133324	4,07131 6,359836 3,167332 8,121386 5,780448 6,914905 7,968711 8,968794 7,975585 5,079708 4,284307	1,562046 1,563829 1,575938 1,58109 1,589441 1,624601 1,627798 1,629518 1,632305 1,651814 1,653958	0,422432 0,201718 -0,084581 0,618052 0,543077 0,472343 1,375331 0,369724 -0,164568 0,293395 0,150983	Trem2 Tmem106a Fpr2 Lifr Naglu Icam2 Cdkn1a Man2a1 Hs6st1 Cd300a Arhgef37	NM_031254 ENSMUST00000100403 NM_008039 NM_013584 NM_013792 ENSMUST00000001055 NM_007669 ENSMUST00000086723 NM_015818 NM_170758 NM_177828
17256784 17333731 17310044 17256502 17270877 17335467 17339108 17211790 17258107 17354764 17256959	5,163425 6,155197 5,183623 5,666428 5,577532 6,13765 6,474934 7,99635 6,050535 6,020436 5,438623 8,210927	6,725471 7,719026 6,759561 7,247518 7,166973 7,762251 8,102732 9,625868 7,68284 7,67225 7,092581 9,874049	3,648878 6,158118 3,251913 7,503334 5,237371 6,442562 6,59338 8,59907 8,140153 4,786313 4,133324 7,967088	4,07131 6,359836 3,167332 8,121386 5,780448 6,914905 7,968711 8,968794 7,975585 5,079708 4,284307 8,891502	1,562046 1,563829 1,575938 1,58109 1,589441 1,624601 1,627798 1,629518 1,632305 1,651814 1,653958 1,663122	0,422432 0,201718 -0,084581 0,618052 0,543077 0,472343 1,375331 0,369724 -0,164568 0,293395 0,150983 0,924414	Trem2 Tmem106a Fpr2 Lifr Naglu Icam2 Cdkn1a Man2a1 Hs6st1 Cd300a Arhgef37 Grn	NM_031254 ENSMUST00000100403 NM_008039 NM_013584 NM_013792 ENSMUST00000001055 NM_007669 ENSMUST00000086723 NM_015818 NM_170758 NM_177828 NM_008175
17256784 17333731 17310044 17256502 17270877 17335467 17339108 17211790 17258107 17354764 17256959 17245700	5,163425 6,155197 5,183623 5,666428 5,577532 6,13765 6,474934 7,99635 6,050535 6,020436 5,438623 8,210927 6,863377 7,993747	6,725471 7,719026 6,759561 7,247518 7,166973 7,762251 8,102732 9,625868 7,68284 7,67225 7,092581 9,874049 8,531673 9,678206	3,648878 6,158118 3,251913 7,503334 5,237371 6,442562 6,59338 8,59907 8,140153 4,786313 4,133324 7,967088 8,376373	4,07131 6,359836 3,167332 8,121386 5,780448 6,914905 7,968711 8,968794 7,975585 5,079708 4,284307 8,891502 8,715645	1,562046 1,563829 1,575938 1,58109 1,589441 1,624601 1,627798 1,629518 1,632305 1,651814 1,653958 1,663122 1,668296	0,422432 0,201718 -0,084581 0,618052 0,543077 0,472343 1,375331 0,369724 -0,164568 0,293395 0,150983 0,924414 0,339272	Trem2 Tmem106a Fpr2 Lifr Naglu Icam2 Cdkn1a Man2a1 Hs6st1 Cd300a Arhgef37 Grn Tspan31	NM_031254 ENSMUST00000100403 NM_008039 NM_013584 NM_013792 ENSMUST00000001055 NM_007669 ENSMUST00000086723 NM_015818 NM_170758 NM_177828 NM_008175 NM_008175 NM_025982
17256784 17333731 17310044 17256502 17270877 17335467 17339108 17211790 17258107 17354764 17256959 17245700 17259907	5,163425 6,155197 5,183623 5,666428 5,577532 6,13765 6,474934 7,99635 6,050535 6,020436 5,438623 8,210927 6,863377 7,993747 8,243145	6,725471 7,719026 6,759561 7,247518 7,166973 7,762251 8,102732 9,625868 7,68284 7,67225 7,092581 9,874049 8,531673	3,648878 6,158118 3,251913 7,503334 5,237371 6,442562 6,59338 8,59907 8,140153 4,786313 4,133324 7,967088 8,376373 7,242939	4,07131 6,359836 3,167332 8,121386 5,780448 6,914905 7,968711 8,968794 7,975585 5,079708 4,284307 8,891502 8,715645 8,690989	1,562046 1,563829 1,575938 1,58109 1,589441 1,624601 1,627798 1,629518 1,632305 1,651814 1,653958 1,663122 1,668296 1,684459 1,688139	0,422432 0,201718 -0,084581 0,618052 0,543077 0,472343 1,375331 0,369724 -0,164568 0,293395 0,150983 0,924414 0,339272 1,44805 2,839857	Trem2 Tmem106a Fpr2 Lifr Naglu Icam2 Cdkn1a Man2a1 Hs6st1 Cd300a Arhgef37 Grn Tspan31 Tcn2	NM_031254 ENSMUST00000100403 NM_008039 NM_013584 NM_013792 ENSMUST00000001055 NM_007669 ENSMUST00000086723 NM_015818 NM_170758 NM_177828 NM_008175 NM_008175 NM_025982 ENSMUST00000109993
17256784 1733731 17310044 17256502 17270877 17335467 17339108 17211790 17258107 17354764 17256959 17245700 17259907 17216458 17233394	5,163425 6,155197 5,183623 5,666428 5,577532 6,13765 6,474934 7,99635 6,050535 6,020436 5,438623 8,210927 6,863377 7,993747 8,243145 7,953105	6,725471 7,719026 6,759561 7,247518 7,166973 7,762251 8,102732 9,625868 7,68284 7,68284 9,874049 8,531673 9,678206 9,931284 9,650748	3,648878 6,158118 3,251913 7,503334 5,237371 6,442562 6,59338 8,59907 8,140153 4,786313 4,786313 4,133324 7,967088 8,376373 7,242939 5,985657 8,129412	4,07131 6,359836 3,167332 8,121386 5,780448 6,914905 7,968711 8,968794 7,975585 5,079708 4,284307 8,891502 8,715645 8,690989 8,825514 9,027002	1,562046 1,563829 1,575938 1,58109 1,589441 1,624601 1,627798 1,629518 1,632305 1,651814 1,633958 1,663122 1,668296 1,684459 1,688139 1,697643	0,422432 0,201718 -0,084581 0,618052 0,543077 0,472343 1,375331 0,369724 -0,164568 0,293395 0,150983 0,924414 0,339272 1,44805 2,839857 0,89759	Trem2 Tmem106a Fpr2 Lifr Naglu Icam2 Cdkn1a Man2a1 Hs6st1 Cd300a Adhgef37 Grn Tspan31 Tcn2 Serpinb2 Smpdl3a	NM_031254 ENSMUST00000100403 NM_008039 NM_013584 NM_013792 ENSMUST00000001055 NM_007669 ENSMUST00000086723 NM_015818 NM_170758 NM_177828 NM_008175 NM_008175 NM_025982 ENSMUST00000109993 NM_011111 ENSMUST00000020022
17256784 17333731 17310044 17256502 17270877 17335467 17339108 17211790 17258107 17354764 17256959 17245700 17259907 17216458 17233394 17506991	5,163425 6,155197 5,183623 5,666428 5,577532 6,13765 6,474934 7,99635 6,050535 6,020436 5,438623 8,210927 6,863377 7,993747 8,243145 7,953105 5,190357	6,725471 7,719026 6,759561 7,247518 7,166973 7,762251 8,102732 9,625868 7,68284 7,67225 9,874049 8,531673 9,678206 9,931284 9,650748 6,888694	3,648878 6,158118 3,251913 7,503334 5,237371 6,442562 6,59338 8,59907 8,140153 4,786313 4,133324 7,967088 8,376373 7,242939 5,985657 8,129412 6,126982	4,07131 6,359836 3,167332 8,121386 5,780448 6,914905 7,968711 8,968794 7,975585 5,079708 4,284307 8,891502 8,715645 8,690989 8,825514 9,027002 5,972383	1,562046 1,563829 1,575938 1,58109 1,589441 1,624601 1,627798 1,629518 1,632305 1,651814 1,653958 1,663122 1,668296 1,684459 1,688139 1,697643 1,698337	0,422432 0,201718 -0,084581 0,618052 0,543077 0,472343 1,375331 0,369724 -0,164568 0,293395 0,150983 0,924414 0,339272 1,44805 2,839857 0,89759 -0,154599	Trem2 Trem106a Fpr2 Lifr Naglu lcam2 Cdkn1a Man2a1 Hs6st1 Cd300a Afgef37 Grn Tspan31 Tcn2 Serpinb2 Smpdl3a Nrp1	NM_031254 ENSMUST00000100403 NM_008039 NM_013584 NM_013792 ENSMUST00000001055 NM_007669 ENSMUST00000008723 NM_01588 NM_170758 NM_177828 NM_008175 NM_025982 ENSMUST00000109993 NM_011111 ENSMUST00000020022 NM_008737
17256784 17333731 17310044 17256502 17270877 17335467 17339108 17211790 17258107 17354764 17256959 17245700 17259907 17216458 17233394 17506991 17254395	5,163425 6,155197 5,183623 5,666428 5,577532 6,13765 6,474934 7,99635 6,050335 6,020436 5,4310927 6,863377 7,993747 8,243145 7,953105 5,190357 5,580338	6,725471 7,719026 6,759561 7,247518 7,166973 7,762251 8,102732 9,625868 7,68284 7,69225 9,874049 8,531673 9,678206 9,931284 9,650748 6,888694 7,292683	3,648878 6,158118 3,251913 7,503334 5,237371 6,442562 6,59338 8,59907 8,140153 4,786313 4,133324 7,967088 8,376373 7,242939 5,985657 8,129412 6,126982 5,96351	4,07131 6,359836 3,167332 8,121386 5,780448 6,914905 7,968711 8,968794 7,975585 5,079708 4,284307 8,891502 8,715645 8,690989 8,825514 9,027002 5,972383 6,275317	1,562046 1,563829 1,575938 1,58109 1,589441 1,624601 1,627798 1,629518 1,632305 1,651814 1,653958 1,663122 1,668296 1,684459 1,688139 1,697643 1,698337 1,712345	0,422432 0,201718 -0,084581 0,618052 0,543077 0,472343 1,375331 0,369724 -0,164568 0,293395 0,150983 0,924414 0,339272 1,44805 2,839857 0,89759 -0,154599 0,311807	Trem2 Tmem106a Fpr2 Lifr Naglu Icam2 Cdkn1a Man2a1 Hs6st1 Cd300a Arhgef37 Grn Tspan31 Tcn2 Serpinb2 Smpdl3a Nrp1 Acaca	NM_031254 ENSMUST00000100403 NM_008039 NM_013584 NM_013792 ENSMUST00000001055 NM_007669 ENSMUST0000086723 NM_170758 NM_170758 NM_177828 NM_008175 NM_025982 ENSMUST00000109993 NM_011111 ENSMUST00000020022 NM_008737 NM_133360
17256784 17333731 17310044 17256502 17270877 17335467 17339108 17211790 17258107 17354764 17256959 17245700 17259907 17216458 17233394 17506991 17254395 17264835	5,163425 6,155197 5,183623 5,666428 5,577532 6,13765 6,474934 7,99635 6,050535 6,020436 5,438623 8,210927 6,863377 7,993747 8,243145 7,953105 5,190357 5,580338 8,017208	6,725471 7,719026 6,759561 7,247518 7,166973 7,762251 8,102732 9,625868 7,68284 7,69225 7,092581 9,678206 9,931284 9,650748 6,888694 7,292683 9,732675	3,648878 6,158118 3,251913 7,503334 5,237371 6,442562 6,59338 8,59907 8,140153 4,786313 4,133324 7,967088 8,376373 7,242939 5,985657 8,129412 6,126982 5,96351 7,214764	4,07131 6,359836 3,167332 8,121386 5,780448 6,914905 7,968711 8,968794 7,975585 5,079708 4,284307 8,891502 8,715645 8,690989 8,825514 9,027002 5,972383 6,275317 8,459773	1,562046 1,563829 1,575938 1,58109 1,589441 1,624601 1,627798 1,629518 1,632305 1,651814 1,653958 1,663122 1,6684459 1,688139 1,697643 1,698337 1,712345	0,422432 0,201718 -0,084581 0,618052 0,543077 0,472343 1,375331 0,369724 -0,164568 0,293395 0,150983 0,924414 0,339272 1,44805 2,839857 0,89759 -0,154599 0,311807 1,245009	Trem2 Trem106a Fpr2 Lifr Naglu Icam2 Cdkn1a Man2a1 Hs6st1 Cd300a Arhg6r37 Grn Tspan31 Tcn2 Serpinb2 Smpdl3a Nrp1 Acaca Cd68	NM_031254 ENSMUST00000100403 NM_008039 NM_013584 NM_013792 ENSMUST00000001055 NM_007669 ENSMUST000000086723 NM_015818 NM_170788 NM_177828 NM_008175 NM_025982 ENSMUST00000109993 NM_011111 ENSMUST0000010002022 NM_008737 NM_133360 NM_009853
17256784 1733731 17310044 17256502 17270877 17335467 17339108 17211790 17258107 17354764 17256959 17245700 17259907 17216458 17233394 17506991 17254395 17264835 17358214	5,163425 6,155197 5,183623 5,666428 5,577532 6,13765 6,474934 7,99635 6,050535 6,020436 5,438623 8,210927 6,863377 7,993747 8,243145 7,953105 5,190357 5,580338 8,017208 6,490279	6,725471 7,719026 6,759561 7,247518 7,166973 7,762251 8,102732 9,625868 7,68284 7,67225 7,092581 9,874049 8,531673 9,678206 9,931284 9,650748 6,888694 7,292683 9,732675 8,208713	3,648878 6,158118 3,251913 7,503334 5,237371 6,442562 6,59338 8,59907 8,140153 4,786313 4,133324 7,967088 8,376373 7,242939 5,985657 8,129412 6,126982 5,96351 7,214764 6,423411	4,07131 6,359836 3,167332 8,121386 5,780448 6,914905 7,968711 8,968794 7,975585 5,079708 4,284307 8,891502 8,715645 8,690989 8,825514 9,027002 5,972383 6,275317 8,459773 7,027411	1,562046 1,563829 1,575938 1,58109 1,589441 1,624601 1,627798 1,629518 1,632305 1,651814 1,653958 1,663122 1,668296 1,684459 1,688139 1,697643 1,698337 1,712345 1,715467	0,422432 0,201718 -0,084581 0,618052 0,543077 0,472343 1,375331 0,369724 -0,164568 0,293395 0,150983 0,924414 0,339272 1,44805 2,839857 0,89759 -0,154599 0,311807 1,245009 0,604	Trem2 Tmem106a Fpr2 Lifr Naglu Icam2 Cdkn1a Man2a1 Hs6st1 Cd300a Arhgef37 Grn Tspan31 Tcn2 Serpinb2 Smpdl3a Nrp1 Acaca Cd68 Klf9	NM_031254 ENSMUST00000100403 NM_008039 NM_013584 NM_013792 ENSMUST00000001055 NM_007669 ENSMUST00000086723 NM_015818 NM_1770758 NM_1770758 NM_008175 NM_025982 ENSMUST00000109993 NM_011111 ENSMUST0000001000100010993 NM_011111 ENSMUST000000100010001001000100100100100100100
17256784 1733731 17310044 17256502 17270877 17335467 17339108 17211790 17258107 17354764 17256959 17245700 17259907 17216458 17233394 17506991 17254395 17264835 17358214 17301213	5,163425 6,155197 5,183623 5,666428 5,577532 6,13765 6,474934 7,99635 6,050535 6,020436 5,438623 8,210927 6,863377 7,993747 8,243145 7,953105 5,190357 5,580338 8,017208 6,490279 8,71772	6,725471 7,719026 6,759561 7,247518 7,166973 7,762251 8,102732 9,625868 7,68284 7,67225 7,092581 9,874049 8,531673 9,678206 9,931284 9,650748 6,888694 7,292683 9,732675 8,208713 10,44006	3,648878 6,158118 3,251913 7,503334 5,237371 6,442562 6,59338 8,59907 8,140153 4,786313 4,133324 7,967088 8,376373 7,242939 5,985657 8,129412 6,126982 5,96351 7,214764 6,423411 9,27864	4,07131 6,359836 3,167332 8,121386 5,780448 6,914905 7,968711 8,968794 7,975585 5,079708 4,284307 8,891502 8,715645 8,690989 8,825514 9,027002 5,972383 6,275317 8,459773 7,027411 9,959899	1,562046 1,563829 1,575938 1,58109 1,589441 1,624601 1,627798 1,629518 1,632305 1,651814 1,653958 1,663122 1,668296 1,684139 1,697643 1,698337 1,712345 1,715467 1,718434 1,72234	0,422432 0,201718 -0,084581 0,618052 0,543077 0,472343 1,375331 0,369724 -0,164568 0,293395 0,150983 0,924414 0,339272 1,44805 2,839857 0,89759 -0,154599 0,311807 1,245009 0,604 0,681259	Trem2 Trem106a Fpr2 Lifr Naglu Icam2 Cdkn1a Man2a1 Hs6st1 Cd300a Arhgef37 Grn Tspan31 Tcn2 Serpinb2 Smpd13a Nrp1 Acaca Cd68 Klf9 Ctsb	NM_031254 ENSMUST00000100403 NM_008039 NM_013584 NM_013792 ENSMUST00000001055 NM_007669 ENSMUST00000086723 NM_015818 NM_170758 NM_170758 NM_015818 NM_170758 NM_008175 NM_025982 ENSMUST00000109993 NM_011111 ENSMUST000001002022 NM_008737 NM_133360 NM_009853 NM_010638 NM_07798
17256784 1733731 17310044 17256502 17270877 17335467 17339108 17211790 17258107 17354764 17256959 17245700 17259907 17216488 1723394 17506991 17254395 17264835 17358214 17301213 17254171	5,163425 6,155197 5,183623 5,666428 5,577532 6,13765 6,474934 7,99635 6,050535 6,020436 5,438623 8,210927 6,863377 7,993747 8,243145 7,953105 5,190357 5,580338 8,017208 6,490279 8,71772 6,384534	6,725471 7,719026 6,759561 7,247518 7,166973 7,762251 8,102732 9,625868 7,68284 7,67225 7,0927649 8,531673 9,678206 9,678206 6,888694 7,292683 9,732675 8,208713 10,44006 8,106915	3,648878 6,158118 3,251913 7,503334 5,237371 6,442562 6,59338 8,59907 8,140153 4,786313 4,133324 7,967088 8,376373 7,242939 5,985657 8,129412 6,126982 5,96351 7,214764 6,423411 9,27864 7,432618	4,07131 6,359836 3,167332 8,121386 5,780448 6,914905 7,968711 8,968794 7,975585 5,079708 4,284307 8,891502 8,715645 8,690989 8,825514 9,027002 5,972383 6,275317 8,459773 7,027411 9,959899 7,901112	1,562046 1,563829 1,575938 1,58109 1,589441 1,624601 1,627798 1,629518 1,632305 1,651814 1,653958 1,663122 1,668296 1,684459 1,697643 1,698337 1,712345 1,715467 1,718434 1,72234 1,722381	0,422432 0,201718 -0,084581 0,618052 0,543077 0,472343 1,375331 0,369724 -0,164568 0,293395 0,150983 0,924414 0,339272 1,44805 2,839857 0,89759 -0,154599 0,311807 1,245009 0,604 0,681259 0,468494	Trem2 Trem106a Fpr2 Lifr Naglu Icam2 Cdkn1a Man2a1 Hs6st1 Cd300a Arhgef37 Grn Tspan31 Tcn2 Serpinb2 Smpdl3a Nrp1 Acaca Cd68 Klf9 Ctsb	NM_031254 ENSMUST00000100403 NM_008039 NM_013584 NM_013792 ENSMUST00000001055 NM_007669 ENSMUST00000086723 NM_015818 NM_170758 NM_170758 NM_170788 NM_208175 NM_025982 ENSMUST00000109993 NM_011111 ENSMUST0000010020022 NM_008737 NM_133360 NM_009853 NM_010638 NM_007798 NM_007798 NM_011407
17256784 1733731 17310044 17256502 17270877 17335467 17339108 17211790 17258107 17354764 17256959 17245700 17259907 17216458 17233394 17506991 17254395 17264835 17358214 17301213 17254171 17351465	5,163425 6,155197 5,183623 5,666428 5,577532 6,13765 6,474934 7,99635 6,050535 6,020436 5,438623 8,210927 6,863377 7,993747 8,243145 7,953105 5,190357 5,580338 8,017208 6,490279 8,71772 6,384534 6,21597	6,725471 7,719026 6,759561 7,247518 7,166973 7,762251 8,102732 9,625868 7,68284 7,67225 7,09276049 8,531673 9,678206 9,87931284 9,650748 6,888694 7,292683 9,732675 8,208713 10,44006 8,106915 7,972548	3,648878 6,158118 3,251913 7,503334 5,237371 6,442562 6,59338 8,59907 8,140153 4,786313 4,133324 7,967088 8,376373 7,242939 5,985657 8,129412 6,126982 5,96351 7,214764 6,423411 9,27864 7,432618 7,30476	4,07131 6,359836 3,167332 8,121386 5,780448 6,914905 7,968711 8,968794 7,975585 5,079708 4,284307 8,891502 8,715645 8,69089 9,027002 5,972383 6,275317 8,459773 7,027411 9,959899 7,901112 7,812508	1,562046 1,563829 1,575938 1,58109 1,589441 1,624601 1,627798 1,629518 1,632305 1,651814 1,653958 1,663122 1,668296 1,684459 1,697643 1,698337 1,712345 1,715467 1,718434 1,72234 1,722381 1,756578	0,422432 0,201718 -0,084581 0,618052 0,543077 0,472343 1,375331 0,369724 -0,164568 0,293395 0,150983 0,924414 0,339272 1,44805 2,4839857 0,89759 -0,154599 0,311807 1,245009 0,604 0,681259 0,468494 0,507748	Trem2 Trem106a Fpr2 Lifr Naglu Icam2 Cdkn1a Man2a1 Hs6st1 Cd300a Arhgef37 Grn Tspan31 Tcn2 Serpinb2 Smpdl3a Nrp1 Acaca Cd68 Kif9 Ctsb Slfn1 Tubb6	NM_031254 ENSMUST00000100403 NM_008039 NM_013584 NM_013792 ENSMUST00000001055 NM_007669 ENSMUST00000086723 NM_015818 NM_170758 NM_170782 NM_025982 ENSMUST00000109993 NM_011111 ENSMUST00000020022 NM_008737 NM_133360 NM_008737 NM_133360 NM_009853 NM_0110638 NM_007798 NM_007798 NM_011407 NM_026473
17256784 1733731 17310044 17256502 17270877 17335467 17339108 17211790 17258107 17354764 17256959 17245700 17259907 17216458 172543394 17506991 17254395 17264835 17358214 17301213 17254171 17351465 17328937	5,163425 6,155197 5,183623 5,666428 5,577532 6,13765 6,474934 7,99635 6,050535 6,020436 5,438623 8,210927 6,863377 7,993747 8,243145 7,953105 5,190357 5,580338 8,017208 6,490279 8,71772 6,384534 6,21597 6,019662	6,725471 7,719026 6,759561 7,247518 7,166973 7,762251 8,102732 9,625868 7,68284 7,67225 7,092581 9,874049 8,531673 9,678208 9,931284 9,550748 6,888694 7,292683 9,732675 8,208713 10,44006 8,106915 7,972548 7,808821	3,648878 6,158118 3,251913 7,503334 5,237371 6,442562 6,59338 8,59907 8,140153 4,786313 4,133324 7,967088 8,376373 7,242939 5,985657 8,129412 6,126982 5,96351 7,214764 6,423411 9,27864 7,432618 7,30476 6,604642	4,07131 6,359836 3,167332 8,121386 5,780448 6,914905 7,968711 8,968794 7,975585 5,079708 4,284307 8,891502 8,715645 8,69089 9,027002 5,972383 6,275317 8,459773 7,027411 9,959899 7,901112 7,812508 7,289473	1,562046 1,563829 1,575938 1,58109 1,589441 1,624601 1,627798 1,629518 1,632305 1,651814 1,653958 1,663122 1,668296 1,684459 1,688139 1,697643 1,698337 1,712345 1,715467 1,718434 1,72234 1,722381 1,756578 1,789159	0,422432 0,201718 -0,084581 0,618052 0,543077 0,472343 1,375331 0,369724 -0,164568 0,293395 0,150983 0,924414 0,339272 1,44805 2,839857 0,89759 -0,154599 0,311807 1,245009 0,604 0,681259 0,468494 0,507748	Trem2 Trem106a Fpr2 Lifr Naglu Icam2 Cdkn1a Man2a1 Hs6st1 Cd300a Arhgef37 Grn Tspan31 Tcrpinb2 Smpdl3a Nrp1 Acaca Cd68 Kif9 Ctsb Slfn1 Tubb6 Comt	NM_031254 ENSMUST00000100403 NM_008039 NM_013584 NM_013792 ENSMUST00000001055 NM_007669 ENSMUST00000086723 NM_015818 NM_170758 NM_170758 NM_177828 NM_025982 ENSMUST00000109993 NM_011111 ENSMUST000000100020022 NM_008737 NM_133360 NM_008737 NM_133360 NM_009853 NM_010638 NM_070798 NM_011407 NM_026473 ENSMUST00000115609
17256784 1733731 17310044 17256502 17270877 17335467 17339108 17211790 17258107 17354764 17256959 17245700 17259907 17216488 1723394 17506991 17254375 17264835 17358214 17301213 17254171 17351465 17328937 17306344	5,163425 6,155197 5,183623 5,666428 5,577532 6,13765 6,474934 7,99635 6,050535 6,020436 5,438623 8,210927 6,863377 7,993747 8,243145 7,953105 5,190357 5,580338 8,017208 6,490279 8,71772 6,384534 6,21597 6,019662 5,515233	6,725471 7,719026 6,759561 7,247518 7,166973 7,762251 8,102732 9,625868 7,68284 7,67225 7,092581 9,874049 8,531673 9,678208 4,767226 7,92583 9,732675 8,208713 10,44006 8,106915 7,972548 7,808821 7,305635	3,648878 6,158118 3,251913 7,503334 5,237371 6,442562 6,59338 8,59907 8,140153 4,786313 4,133324 7,967088 8,376373 7,242939 5,985657 8,129412 6,126982 5,96351 7,214764 6,423411 9,27864 7,432618 7,30476 6,604642 6,787159	4,07131 6,359836 3,167332 8,121386 5,780448 6,914905 7,968711 8,968794 7,975585 5,079708 4,284307 8,891502 8,715645 8,69089 8,825514 9,027002 5,972383 6,275317 8,459773 7,027411 9,959899 7,901112 7,812508 7,289473 7,292461	1,562046 1,563829 1,575938 1,58109 1,589441 1,624601 1,627798 1,629518 1,632305 1,651814 1,653958 1,663122 1,668296 1,684459 1,688139 1,697643 1,698337 1,712345 1,715467 1,718434 1,722381 1,756578 1,789159 1,790402	0,422432 0,201718 -0,084581 0,618052 0,543077 0,472343 1,375331 0,369724 -0,164568 0,293395 0,150983 0,924414 0,339272 1,44805 2,839857 -0,154599 0,311807 1,245009 0,604 0,681259 0,468494 0,507748 0,684831 0,505302	Trem2 Trem106a Fpr2 Lifr Naglu Icam2 Cdkn1a Man2a1 Hs6st1 Cd300a Arhgef37 Grn Tspan31 Tcn2 Serpid3a Nrp1 Acaca Cd68 Kif9 Ctsb Slfn1 Tubb6 Comt	NM_031254 ENSMUST00000100403 NM_008039 NM_013584 NM_013792 ENSMUST00000001055 NM_007669 ENSMUST00000086723 NM_015818 NM_170758 NM_177758 NM_0158175 NM_028175 NM_028175 NM_029982 ENSMUST00000109993 NM_011111 ENSMUST000000109993 NM_0111111 ENSMUST00000020022 NM_008737 NM_133360 NM_008737 NM_133360 NM_009853 NM_010638 NM_010638 NM_011407 NM_026473 ENSMUST00000115609 NM_011405
17256784 1733731 17310044 17256502 17270877 17335467 17339108 17211790 17258107 17354764 17256959 17245700 17259907 17216458 1723394 17256959 17254395 17264835 17358214 17301213 17254171 17351465 17328937 17306344 17279131	5,163425 6,155197 5,183623 5,666428 5,577532 6,13765 6,474934 7,99635 6,050535 6,020436 5,438623 8,210927 6,863377 7,993747 8,243145 7,953105 5,190357 5,580338 8,017208 6,490279 8,71772 6,384534 6,21597 6,019662 5,515233 5,345794	6,725471 7,719026 6,759561 7,247518 7,166973 7,762251 8,102732 9,625868 7,68284 7,67225 7,092581 9,874049 8,531673 9,678206 9,931284 9,650748 6,888694 7,829683 9,732675 8,208713 10,44006 8,106915 7,972548 7,808821 7,305635 7,144465	3,648878 6,158118 3,251913 7,503334 5,237371 6,442562 6,59338 8,59907 8,140153 4,786313 4,133324 7,967088 8,376373 7,242939 5,985657 8,129412 6,126982 5,96351 7,214764 6,423411 9,27864 7,432618 7,30476 6,604642 6,787159 4,219324	4,07131 6,359836 3,167332 8,121386 5,780448 6,914905 7,968711 8,968794 7,975585 5,079708 4,284307 8,891502 8,715645 8,69089 8,825514 9,027002 5,972383 6,275317 8,459773 7,027411 9,959899 7,901112 7,812508 7,289473 7,292461 4,739029	1,562046 1,563829 1,575938 1,58109 1,589441 1,624601 1,627798 1,629518 1,632305 1,651814 1,653958 1,663122 1,668296 1,684459 1,688139 1,697643 1,712345 1,715467 1,718434 1,72234 1,722381 1,756578 1,789159 1,790402 1,798671	0,422432 0,201718 -0,084581 0,618052 0,543077 0,472343 1,375331 0,369724 -0,164568 0,293395 0,150983 0,924414 0,339272 1,44805 2,839857 -0,154599 0,311807 1,245009 0,604 0,681259 0,468494 0,507748 0,684831 0,505302 0,519705	Trem2 Trem106a Fpr2 Lifr Naglu Icam2 Cdkn1a Man2a1 Hs6st1 Cd300a Arhgef37 Grn Tspan31 Tcn2 Serpinb2 Smpd13a Araca Cd68 Klf9 Ctsb Slfn1 Tubb6 Comt Slc7a7	NM_031254 ENSMUST00000100403 NM_008039 NM_013584 NM_013792 ENSMUST0000001055 NM_007669 ENSMUST00000086723 NM_015818 NM_170758 NM_017828 NM_08175 NM_025982 ENSMUST000000109993 NM_0111111 ENSMUST00000020022 NM_008737 NM_033360 NM_009853 NM_010638 NM_010638 NM_010798 NM_011407 NM_026473 ENSMUST00000115609 NM_011405 ENSMUST00000102745
17256784 1733731 17310044 17256502 17270877 17335467 17339108 17211790 17258107 17354764 17256959 17245700 17259907 17216488 1723394 17506991 17254375 17264835 17358214 17301213 17254171 17351465 17328937 17306344	5,163425 6,155197 5,183623 5,666428 5,577532 6,13765 6,474934 7,99635 6,050535 6,020436 5,438623 8,210927 6,863377 7,993747 8,243145 7,953105 5,190357 5,580338 8,017208 6,490279 8,71772 6,384534 6,21597 6,019662 5,515233	6,725471 7,719026 6,759561 7,247518 7,166973 7,762251 8,102732 9,625868 7,68284 7,67225 7,092581 9,874049 8,531673 9,678208 4,767226 7,92583 9,732675 8,208713 10,44006 8,106915 7,972548 7,808821 7,305635	3,648878 6,158118 3,251913 7,503334 5,237371 6,442562 6,59338 8,59907 8,140153 4,786313 4,133324 7,967088 8,376373 7,242939 5,985657 8,129412 6,126982 5,96351 7,214764 6,423411 9,27864 7,432618 7,30476 6,604642 6,787159	4,07131 6,359836 3,167332 8,121386 5,780448 6,914905 7,968711 8,968794 7,975585 5,079708 4,284307 8,891502 8,715645 8,69089 8,825514 9,027002 5,972383 6,275317 8,459773 7,027411 9,959899 7,901112 7,812508 7,289473 7,292461	1,562046 1,563829 1,575938 1,58109 1,589441 1,624601 1,627798 1,629518 1,632305 1,651814 1,653958 1,663122 1,668296 1,684459 1,688139 1,697643 1,698337 1,712345 1,715467 1,718434 1,722381 1,756578 1,789159 1,790402	0,422432 0,201718 -0,084581 0,618052 0,543077 0,472343 1,375331 0,369724 -0,164568 0,293395 0,150983 0,924414 0,339272 1,44805 2,839857 -0,154599 0,311807 1,245009 0,604 0,681259 0,468494 0,507748 0,684831 0,505302	Trem2 Trem106a Fpr2 Lifr Naglu Icam2 Cdkn1a Man2a1 Hs6st1 Cd300a Arhgef37 Grn Tspan31 Tcn2 Serpid3a Nrp1 Acaca Cd68 Kif9 Ctsb Slfn1 Tubb6 Comt	NM_031254 ENSMUST00000100403 NM_008039 NM_013584 NM_013792 ENSMUST00000001055 NM_007669 ENSMUST00000086723 NM_015818 NM_170758 NM_177758 NM_0158175 NM_028175 NM_028175 NM_029982 ENSMUST00000109993 NM_011111 ENSMUST000000109993 NM_0111111 ENSMUST00000020022 NM_008737 NM_133360 NM_008737 NM_133360 NM_009853 NM_010638 NM_010638 NM_011407 NM_026473 ENSMUST00000115609 NM_011405

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17230484	5,482994	7,297328	7,097932	7,433383	1,814334	0,335451	Ephx1	ENSMUST00000036928
17346749	7,109239	8,936438	7,586003	8,136797	1,827199	0,550794	Rab31	NM_133685
17218845	8,684654	10,54799	5,951889	8,218878	1,863336	2,266989	Selp	NM 011347
17328508	4,895505	6,779959	3,39824	4,194564	1,884454	0,796324	Fgd4	ENSMUST00000161861
17268884	5,382138	7,294327	5,402496	5,179055	1,912189	-0,223441	Nr1d1	ENSMUST00000064941
17272770	7,820322	9,740899	7,653145	8,406399	1,920577	0,753254	Timp2	ENSMUST00000017610
17337796	6,515199	8,448761	3,791042	5,289989	1,933562	1,498947	Pla2g7	ENSMUST00000024706
17254176	6,803511	8,744821	4,172538	5,549662	1,94131	1,377124	Slfn4	NM 011410
17308257	5,514946	7,508637	4,630305	5,77283	1,993691	1,142525	Sorbs3	NM 011366
		<u> </u>	- 1			-		-
17321823	6,024487	8,031663	5,396936	6,444937	2,007176	1,048001	Krt80	NM_028770
17335506	4,984896	6,998934	3,606941	4,711351	2,014038	1,10441	Pi16	ENSMUST00000114701
17484167	4,92322	6,94167	3,622232	4,609194	2,01845	0,986962	Dock1	NM_001033420
17504444	7,363705	9,384002	7,331205	8,30427	2,020297	0,973065	Cmtm3	NM_024217
17342625	6,138542	8,164219	6,193325	7,181438	2,025677	0,988113	Itfg3	ENSMUST00000114988
17223863	7,332762	9,363936	5,971759	7,213199	2,031174	1,24144	Idh1	NM_001111320
17341963	5,082928	7,122298	4,866551	5,847231	2,03937	0,98068	Slc9a3r2	NM_023055
17320337	6,498953	8,556587	5,765083	6,91497	2,057634	1,149887	Plxnb2	NM_138749
17360977	5,13719	7,201246	5,988608	5,752959	2,064056	-0,235649	Lrp5	NM_008513
17382920	5,425421	7,492276	5,788852	5,950367	2,066855	0,161515	Card9	ENSMUST00000100303
17532569	5,433601	7,501606	4,842346	5,687298	2,068005	0,844952	Ccr1	NM 009912
17265268	4,82003	6,891633	5,573634	6,423313	2,071603	0,849679	Cxcl16	ENSMUST00000019064
17331284	5,035634	7,118655	5,050541	5,839437	2,083021	0,788896	Crybg3	NM 174848
17245648	4,778471	6,872993	5,455342	6,217944	2,094522	0,762602	Slc16a7	ENSMUST00000063318
17247984	6,669252	8,801469	7,202742	7,396691	2,132217	0,193949	Dapk1	NM 029653
17325608	5,745573	7,883711	8,067148	8,45969	2,132217	0,392542	Cd80	NM 009855
17219199			3,82971			0,683401		NM 144559
-	4,591934	6,73474	· ·	4,513111	2,142806	-	Fcgr4	-
17267601	6,844967	9,005307	8,231976	8,786457	2,16034	0,554481	Scpep1	NM_029023
17239227	6,203192	8,364386	5,573036	6,849443	2,161194	1,276407	Rab32	NM_026405
17287827	8,064057	10,23659	6,699467	8,574282	2,172533	1,874815	Tgfbi	NM_009369
17281721	6,115036	8,306716	4,734724	5,437407	2,19168	0,702683	Pygl	NM_133198
17293706	6,27047	8,481782	7,921008	7,387086	2,211312	-0,533922	Ctsl	NM_009984
17243868	5,512076	7,731533	4,430312	5,457442	2,219457	1,02713	Dram1	NM_027878
17256870	6,160548	8,420387	5,475978	6,459056	2,259839	0,983078	Cd300lg	NM_001160711
17530243	8,057837	10,36986	6,63758	8,406939	2,312023	1,769359	Trf	ENSMUST00000035158
17267846	4,160737	6,493881	4,103273	4,544879	2,333144	0,441606	Abcc3	NM_029600
17351053	7,33147	9,683061	5,816728	7,594163	2,351591	1,777435	Csf1r	NM_001037859
17245231	7,095935	9,523712	7,83452	9,248249	2,427777	1,413729	Lyz1	NM_013590
17352036	5,825021	8,292607	6,430522	7,063242	2,467586	0,63272	Pstpip2	NM 013831
17216469	7,171445	9,643112	4,307654	7,325596	2,471667	3,017942	Serpinb10	NM 198028
17418519	4,526365	7,008114	4,384923	4,817927	2,481749	0,433004	Oscp1	NM 172701
17352985	5,241015	7,729548	6,069003	6,77262	2,488533	0,703617	B4galt6	NM 019737
17363470	7,012671	9,504123	6,018223	7,360303	2,491452	1,34208	Gda	NM 010266
17271443	3,522923	6,043066	3,723533	4,006015	2,520143	0,282482	Abca6	NM 147218
17344064	4,624222		4,037507	4,788486	2,544339		Cfb	NM 008198
		7,168561	- 1			0,750979		-
17214197	6,67437	9,223215	5,558557	7,310296	2,548845	1,751739	Slc11a1	NM_013612
17271724	6,024632	8,583055	4,934464	6,156918	2,558423	1,222454	Cd300ld	ENSMUST00000045075
17394153	7,226619	9,802204	6,05878	8,637745	2,575585	2,578965	Slpi	NM_011414
17474974	5,801637	8,388187	8,296436	8,64188	2,58655	0,345444	Plaur	ENSMUST00000002284
17375997	4,300755	6,926266	3,954049	5,023597	2,625511	1,069548	Mertk	ENSMUST00000014505
17224071	9,361686	11,98952	9,819113	11,4637	2,627834	1,644587	Fn1	NM_010233
17245923	6,965324	9,654119	5,746463	7,690616	2,688795	1,944153	Lrp1	NM_008512
17338617	6,84933	9,546226	4,613992	6,609534	2,696896	1,995542	Emr4	NM_139138
17354101	4,581501	7,300237	4,509981	5,189752	2,718736	0,679771	Dpysl3	ENSMUST00000025379
17462796	5,921942	8,655614	4,523855	6,376474	2,733672	1,852619	Clec4d	NM_010819
17326801	3,329509	6,063628	3,6471	4,286726	2,734119	0,639626	Jam2	NM_023844
17258912	5,31407	8,090231	5,673363	6,435722	2,776161	0,762359	Engase	ENSMUST00000135383
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17312219	4,749608	7,54222	3,160961	4,289943	2,792612	1,128982	Ly6g	XM_001475753
17437198	5,976456	8,78579	5,464659	6,862984	2,809334	1,398325	Bst1	NM_009763
17283445	6,1225	8,958448	6,456817	7,529007	2,835948	1,07219	Lgmn	ENSMUST00000021607
17470627	6,484612	9.331956	4.695291	6,978383	2,847344	2,283092	Clec4e	NM 019948
17346975	7,814077	10,67872	6,754671	8,967196	2,864643	2,212525	Emilin2	NM 145158
17394297	6,799634	9,664432	7,768081	9,093259	2,864798	1,325178	Pltp	NM 011125
		-		,			·	
17452705	4,974228	7,871373	3,449855	3,655536	2,897145	0,205681	Niacr1	NM_030701
17421033	3,180879	6,086344	3,582338	4,79048	2,905465	1,208142	Epha2	NM_010139
17309935	4,500522	7,419732	5,610706	6,881321	2,91921	1,270615	Dab2	ENSMUST00000080880
17537081	5,258751	8,185442	3,977983	5,680908	2,926691	1,702925	Tlr13	NM_205820
17320652	5,600131	8,528525	3,209014	5,221596	2,928394	2,012582	Abcd2	NM 011994
17489464	4,886589	7,835878	3,756119	5,557484	2,949289	1.801365	Scn1b	NM 011322
17528692	3,771353	6,72083	4,298397	5,512817	2,949477	1,21442	Cgnl1	NM_026599
17501633	5,008878	7,981215	4,4476	6,149157	2,972337	1,701557	Lpl	ENSMUST00000015712
	· ·	7,491356			2,990629	0,430515		
17495610	4,500727	-	5,18361	5,614125			Gprc5b	NM_022420
17432014	4,822825	7,81644	4,236468	5,888321	2,993615	1,651853	Padi4	ENSMUST00000026381
17433265	4,030719	7,082666	4,115662	4,548475	3,051947	0,432813	Car6	ENSMUST00000030817
17419277	7,101611	10,15654	6,737793	8,714136	3,054929	1,976343	Sdc3	NM_011520
17329093	3,908152	7,009271	3,974698	4,711331	3,101119	0,736633	Cyp2ab1	ENSMUST00000023513
17393868	5,326731	8,458218	5,588224	6,421997	3,131487	0,833773	Mafb	NM 010658
17346150	5,298428	8,488145	5,146091	6,499998	3,189717	1,353907	Lrg1	NM 029796
17540521	7,243188	10,48279	7,798534	9,109869	3,239602	1,311335	Cfp	ENSMUST00000001156
17453611	3,670662	7,043983	5,203271		3,373321	-	Ccl24	
	· ·		· ·	6,483829	-	1,280558		NM_019577
17348282	4,081113	7,489085	4,375387	5,452421	3,407972	1,077034	Colec12	NM_130449
17331669	6,665879	10,08151	6,294654	8,15694	3,415631	1,862286	App	NM_001198823
17384100	4,364265	7,820585	4,102789	5,519053	3,45632	1,416264	Garnl3	ENSMUST00000102810
17297010	6,441812	9,958832	7,053974	8,353849	3,51702	1,299875	Flnb	NM_134080
17307134	5,541005	9,061911	7,208986	7,926934	3,520906	0,717948	Cryl1	NM 030004
17512611	5,553152	9,090222	5,014908	7,112585	3,53707	2,097677	Dpep2	ENSMUST00000081998
17289547	2,738994	6,299932	3,008013	4,123499	3,560938	1,115486	Etv1	NM 007960
17378827	4,49807	8.070959	4,286355	6,114587	3,572889	1,828232	Lbp	NM 008489
17378827	4,345096	-,-	5,378677	7,35069		-	Cd14	ENSMUST00000061829
		7,920608	-		3,575512	1,972013		
17500108	5,774091	9,350451	6,08082	8,100088	3,57636	2,019268	Fgfr1	NM_010206
17271399	5,123266	8,70355	4,708789	6,508837	3,580284	1,800048	Abca9	ENSMUST00000044850
17382737	3,750983	7,336894	4,89595	5,993488	3,585911	1,097538	Fcna	ENSMUST00000028307
17262316	3,776561	7,455116	3,999305	5,740701	3,678555	1,741396	Ltc4s	NM_008521
17493875	4,209774	7,933028	3,677578	4,965712	3,723254	1,288134	P2ry2	NM 008773
17227696	5,992113	9,793705	5,389149	7,728874	3,801592	2,339725	Cfh	NM 009888
17339013	6,258738	10,14557	6,355885	8,713228	3,886832	2,357343	Emr1	NM 010130
17295627	3,240003	7,136444	4,639411	5,331629	3,896441	0,692218	Naip1	ENSMUST00000022142
	-				-		·	
17403255	4,512371	8,417868	4,408272	6,102687	3,905497	1,694415	Gbp1	NM_010259
17508850	5,667226	9,581429	5,931888	8,213323	3,914203	2,281435	Msr1	NM_001113326
17439464	4,76975	8,707665	3,835568	5,545413	3,937915	1,709845	Anxa3	NM_013470
17227892	6,655327	10,61322	8,47919	10,64473	3,957893	2,16554	Prg4	NM_021400
17507321	3,4493	7,459631	4,167992	6,074322	4,010331	1,90633	Efnb2	NM_010111
17439367	6,685216	10,72537	8,271397	9,868114	4,040154	1,596717	Cxcl13	NM_018866
17265229	7,235255	11,32977	8,009497	10,42574	4,094515	2,416243	Alox15	NM 009660
17431612	3,94887	8,179264	4,72221	6,658095	4,230394	1,935885	C1qc	ENSMUST00000046332
17451012	3,747929	8,005649	3,390403	4,830787	4,25772	1,440384	Vmn2r26	NM 019917
17431607	6,319914	10,66183	7,430313	9,350456	4,341916	1,920143	C1qb	NM_009777
17394679	3,58815	8,251455	4,372238	6,3088	4,663305	1,936562	Ptgis	ENSMUST00000018113
17348400	4,695361	9,367277	3,952312	6,803017	4,671916	2,850705	Gata6	NM_010258
17230830	4,154748	9,029202	5,322719	7,337743	4,874454	2,015024	Tgfb2	NM_009367
17309268	5,688822	10,60203	5,931916	8,575165	4,913208	2,643249	Ednrb	NM_007904
17343918	5,139817	10,07237	6,479048	8,46674	4,932553	1,987692	C4b	NM_009780
17233384	4,944606	9,913021	6,418943	8,491428	4,968415	2,072485	Fabp7	ENSMUST00000020024
17233384	4,944606	9,913021	6,418943	8,491428	4,968415	2,072485	Fabp7	ENSMUST00000020024
17438995	4,485271	9,603101	6,947411	8,889437			Cxcl2	ENSMUST00000075433
					5,11783	1,942026		
17404091	4,209189	9,333618	4,371784	6,457975	5,124429	2,086191	Fabp4	NM_024406
17248860	4,040284	9,207533	5,307802	7,554366	5,167249	2,246564	Timd4	NM_178759
17491193	5,357244	10,82792	6,483482	9,568161	5,470676	3,084679	Saa3	NM_011315
17543365	3,184662	8,985921	5,298122	7,64684	5,801259	2,348718	Vsig4	NM_177789
17398734	3,011252	8,858755	3,914339	6,889477	5,847503	2,975138	Cd5l	NM_009690
	•			•	•	•		

Table S2.

Gene	Forward	Reverse
Abca6	ATGATGACTGGTTTCCTGGTCA	GCAACAGAACAGCCATCAGG
Abcc3	GGACCATGAAGCCTTGCAAAAT	GGCTGGCTCATTGTCTGTCA
Galectin-1	CTCAAAGTTCGGGGAGAGGT	CATTGAAGCGAGGATTGAAGT
Galectin-3	GCCTACCCCAGTGCTCCT	GGTCATAGGGCACCGTCA
Galectin-9	GCATTGGTTCCCCTGAGAATA	TCCAGTAAAGGGGATGATCG
Gapdh	ATGGTGAAGGTCGGTGTGA	AATCTCCACTTTGCCACTGC
Gzm B	GCTGCTCACTGTGAAGGAAGT	TGGGGAATGCATTTTACCAT
Gzm C	TCCTGATTCTCCTGACCCTACT	GACGGGAATGTGGACTGATCT
Gzm D	CTGCAAAAACAGCTCTGTCCAA	TGATCTGCTGTGTCTCCTCCT
Gzm E	GGTAATAGGAGATACTGTGGAGG	TGTGACTGTCATTGTCCTGTTC
Gzm G	AGAAAGCATGGCACCAATGAC	ATTGGGTCTGGGCAACTTGA
H2-q1	CTACGGCTGGGAAATGAGAC	GCACCTCAGGGTGACTTCAT
Ifn-γ	ATCTGGAGGAACTGGCAAAA	TTCAAGACTTCAAAGAGTCTGAGGTA
Prf 1	TGAGAAGACCTATCAGGACCAG	GTCAAGGTGGAGGTT
Rae-1β	CAGCAAATGCCACTGAAGTGAA	GGTCTTGTGAGTGTCCACTTTG
Rae-18	CTCCTACCCCAGCAGATGAAG	CCCTGGGTCACCTGAAGTC
Rae-1ɛ	GACCCACAGACCAAATGGCA	CTCTGTCCTTTGAGCTTCTTGC
β2М	GAATGGGAGCCGAACATAC	CCGTTCTTCAGCATTTGGAT
β-actin	GATGTGGATCAGCAAGCAGGA	AGCTCAGTAACAGTCCGCCTA

Chapter 3

Summary

The idea that some chemotherapeutic agents, including classical chemotherapeutic compounds, may exert immunostimulatory effects has generate considerable interest in their combination with immunotherapy. We are testing this strategy in a clinical trial (DENDR1 - EUDRACT N° 2008-005035-15) currently active at our Institute. Patients affected by newly diagnosis of glioblastoma (GBM), enrolled in DENDR1 clinical trial and treated with dendritic cells (DCs) in combination with temozolomide (TMZ), showed a significant increase of peripheral cytotoxic NK cell percentage but a failure of CD8⁺ T lymphocyte response. From these clinical observations, we moved to evaluate the potential direct effect of TMZ on NK cell response by using our preclinical malignant glioma model.

We treated GL261 glioma-bearing mice with five intraperitoneal injections of a low-dose of TMZ 9 days after intracranial tumor implantation. Peripheral blood lymphocytes (PBLs) and tumor-infiltrating lymphocytes (TILs) were analyzed by flow cytometry to characterize a potential direct effect of TMZ on immune effectors. TMZ induced a rapid and reversible lymphopenia of CD8⁺ T cells with a concomitant limited homing to the brain. Conversely, the frequency of peripheral blood NK cells from TMZ-treated mice increased significantly at early time point and remained higher than controls throughout the entire treatment.

Microarray analysis performed on blood-purified NK cells revealed that Abcc3 was one of the most up-regulated genes in TMZ-treated mice compared to controls during chemotherapy administration. Abcc3, a

member of the ATP-binding cassette transporter superfamily, is implicated in acquired multidrug-resistance in different types of tumors. In the present study, we provided the evidence that Abcc3 plays an important role in protecting blood-derived NK cells from the cytotoxic effects of the alkylating agent TMZ. Indeed, we found that Abcc3 was over-expressed and functionally active in response to TMZ in NK cells, and not in CD8⁺ T lymphocytes, from naïve and gliomabearing mice. Moreover, our results indicated that NK cells displayed a high expression of this transporter in physiological conditions suggesting a potential basal drug-resistant phenotype.

The major finding of this study was the demonstration of the critical role of Abcc3 in protecting NK cells from apoptosis. NK cells negative for Abcc3 were more prone to undergo apoptosis during chemotherapy treatment and the percentage of apoptotic cells was higher compared to NK cells Abcc3 positive. The inhibition of Abcc3 activity with the specific molecule MK571 significantly induced apoptosis in NK cells. Because the pathways involving Akt activation promotes lymphocyte survival, we investigated Akt phosphorylation in response to chemotherapy. Only NK cells expressing Abcc3 presented Akt activation during TMZ treatment.

We also found that TMZ is able to influence positively the maturation status and functions of NK cells. During TMZ treatment, there was a significant enrichment of the NK cell subsets with migratory potential resulting in a greater tumor-infiltration ability. NK cells infiltrate the tumor mass early leading to an increase and persistent expression of cytotoxic signals including Ifn- γ , granzymes and perforin. Additionally,

TMZ modified the glioma microenvironment reducing the strong immunosuppressive components and providing the optimal condition for NK cell recruitment and effector function.

Finally, we found that cytotoxicity, evaluated as IFN-γ production and specific lytic activity against GL261 cells by peripheral NK cells, significantly increased compared to control mice.

These findings demonstrated for the first time the direct involvement of Abcc3 in conferring NK cell resistance to chemotherapy and the influence of TMZ on immune effector functions and glioma microenvironment, observations to consider for trials where glioblastoma patients are treated with chemo-immunotherapy.

Conclusions

Glioblastoma (GBM) remains a particularly devastating solid tumor for which standard treatment options offer only modest efficacy. Cancer immunotherapy, the scientific breakthrough of the year 2013 by Science ¹, represents the most promising cancer treatment approach with a potential to achieve a complete, long-lasting remission with few or no side effects. The advances in cancer immunotherapy are mostly related to the treatment of melanoma and other solid tumors ^{2,3}. However, a growing amount of studies indicates that immunotherapy is a well-suited opportunity to target glioma cells for destruction with high specificity and efficiency improving the outcome of patients ^{4,5}. Unfortunately, one of the major limitation of immune-related strategies in GBM is the highly immunosuppressive tumor microenvironment.

Despite conventional dogma that indicates chemotherapy as immunosuppressive, now it is proposed its ability to overcome or realistically to reduce the immune suppressive factors, responsible for T cell anergy and amplification of the regulatory T (Treg) cells and myeloid-derived suppressor cells (MDSCs) ⁶, produced by cancer cells. In fact, chemotherapy could be used to influence the immune system and tumor to create an environment where cancer vaccines have a better chance of success ^{7,8}.

TMZ is proposed as a possible immune adjuvant and already administered with immune vaccines ^{9,10}. Although TMZ reaches its anti-tumor effects through direct killing of tumor cells, it demonstrated several direct and indirect effects also on immune cells by inducing lymphopenia and modulating tumor microenvironment ^{11–13}.

In this scenario of co-administration with immune approaches, an important goal was represented by characterization of the potential direct effects of chemotherapy on immune system. Gaining better insights of the molecular mechanisms induced by TMZ on immune cells is essential to design efficient therapeutic strategies of chemo-immunotherapy.

Our experience with DC-based immunotherapy started in 2008 in close collaboration with the Unit of Cell Therapy Production located within our institute ¹⁴. Currently, we have two clinical studies for DC immunotherapy (DENDR1, EUDRACT 2008-005035-15 and DENDR2, EUDRACT 2008-005038-62, respectively for the treatment of first diagnosis and recurrent GBM) coordinated by Dr. Finocchiaro and approved by the Italian Ministry of Health. In both studies, we are testing the effects of immunotherapy with DCs loaded with tumor lysate and simultaneous treatment with TMZ ¹⁵.

Analyses on blood lymphocytes before and after DC vaccines indicated an increased NK, but not CD8⁺ T, cell cytotoxic response significantly associated with prolonged survival.

Consistent with these clinical studies, we found a great increase in peripheral trafficking and local homing of NK cells in GL261 gliomabearing mice treated with TMZ. In fact, murine NK cells can efficiently overcome the drug-mediated toxicity of chemotherapy and infiltrate tumors producing an efficient anti-glioma immune response.

First, we found that the expression and function of Abcc3 transporter is a protective mechanism activated by NK cells to escape death during TMZ administration. Abcc3 expression is already associated in acquired MDR in different tumors ^{16,17}. Its relative expression in

lymphocytes is still controversial ^{18–20}. However, it is commonly accepted that in leukocytes its efflux function could be involved in protection against toxins and chemotherapy compounds. Indeed, Abcc3 expression is associated to the strong refractory to chemotherapy in patients with leukemia suggesting its active role in efflux of anticancer agents ^{21,22}. The expression and activity of MDR transporters could be one important point to take into account when the outcome of combination therapy regimens is considered.

Another important goal was to highlight the positive influence of TMZ on NK cell anti-tumor activity. The effects of different chemotherapeutic drugs on NK cell activity have been already observed suggesting negative or no modulation of NK cell cytotoxic functions ^{11,23}. While the direct effect of TMZ on NK cell effector activity has not been well investigated, another alkylating agent as cyclophosphamide has been proposed as a potent inductor of NK-dependent antitumor immunity ^{24–26}. Our results demonstrate a direct effect of TMZ on influencing NK cell activation, migration and specific anti-glioma cytotoxicity.

These important findings suggest a translational potential for the definition of novel chemo-immunotherapy protocols for GBM. Several groups, including ours, proposed the relevance of NK cells in inhibiting glioma growth ^{27–30}. Since NK cells have been shown to positively influence multiple aspects of immune response ^{31–34}, their activation by chemotherapy could be exploited for enhancing the magnitude of both innate and adaptive immunity against tumors resulting in a better clinical outcome.

These findings highlighted two important aspects poorly examine in depth: the molecular mechanisms mediating chemo-resistance in immune cells during chemotherapy treatment and the role of NK cells in immune approaches to treat brain tumors.

Overall our observations may help to set up the best conditions for a clinical trial taking into account the strong different influence of TMZ on NK cells and CD8⁺ T lymphocytes which are, normally, protagonists of effector response during a DC-based immunotherapy.

Future perspectives

During the last four years, we have observed one of the most important revolutions in all the history of the medicine. First of all, in 2011 Ralph Steinman received the Nobel Prize for his discovery of the dendritic cells. Only two years later, immunotherapy, considered for many years to be more a theory than therapeutic strategy, was defined the breakthrough of the year 2013 ¹. The main advances were preferentially related to the successful obtained from the use of checkpoints inhibitors in some solid tumors, especially metastatic melanoma ³⁵.

Our immune system contains all the potential solutions we need to ward off the cancer, however it is not perfect. Immunotherapy can potentiate specific anti-tumor immune responses and the efficacy of this strategy may be increased by secret and unexpected allies ³⁶. Recent studies suggest that some chemotherapeutic agents possess immune stimulatory effects ³⁷ and can effectively favor immune response against tumors. However, a detailed understanding of the cellular and molecular basis of relationship between chemotherapy and immune system is essential for planning the optimal strategy for combining immune approaches with standard therapies for cancer.

Our preclinical data showed that temozolomide (TMZ), the chemotherapeutic drug used as standard therapy for patients affected by GBM, influences the activity and the phenotype of NK cells. Most important, Abcc3 transporter with its efflux activity confers to NK cells chemo-resistance to TMZ protecting them from apoptosis.

These observations encourage us to investigate the potential involvement of ABCC3 in influencing the immune response, and

specifically the NK cell response of GBM patients enrolled in our clinical trial DENDR1. We assessed ABCC3 expression by real-time PCR on PBLs of 17 GBM patients and found a significant higher expression in responders (PFS \geq 12 and high NK cell response) compared to non-responders (p = 0.0006). ABCC3 expression also increases significantly at the time of first vaccine after radiochemotherapy (p = 0.01), and after concomitant vaccine and TMZ administration (p = 0.01) (Figure 1). In order to measure the ABCC3 expression by flow cytometry specifically on NK and CD8⁺ T cells, we used an anti-human ABCC3 antibody. Preliminary results indicated that ABCC3 is differentially expressed in NK cells compared to CD8⁺ T cells also in human PBLs before TMZ administration (68.6 \pm 16.8 vs 1.2 ± 0.8 , NK vs CD8⁺ T cells respectively, p = 0.0002; Figure 2). Now we are going to investigate ABCC3 expression in PBLs before, after and during DC immunotherapy. Specific attention will be given to last four vaccines co-administrated with adjuvant TMZ in order to understand whether ABCC3 plays the same protection role against chemotherapy potentiating NK cell anti-glioma response.

Next, we plan to characterize widely the peripheral effector response in our DENDR1 patients. Flow cytometry analyses showed an increased cytotoxic NK cell frequency correlated with a prolonged PFS. Unexpectedly, we have a lack of an efficient antitumor CD8⁺ T cell response. Although several studies suggest as NK cell activation promote CTL antitumor response ^{32,38}, other data affirm that a strong NK cell response may be inhibitory ^{39,40}. Based on these data we aim to study and characterize some potential effect of the robust NK cell cytotoxicity on CD8⁺ T activation and expansion.

Our pre-clinical observations indicated a more sensitivity of CD8⁺ T cells to cytotoxic effects of TMZ in contrast with a potent NK cell activation. To further improve the survival of our patients and delay the unavoidable cancer immunoediting ^{41,42} and consequently the recurrence, we need to trigger an efficient CTL response. In addition, since a critical requirement of an efficient cancer immunotherapy is the generation of long-lasting and specific memory, we plan to investigate the CD8⁺ T (effector) memory formation during DC vaccinations and concomitant TMZ administration.

Therefore, our main challenge is to figure out any issues related to the current treatment schedule evaluating the possibility to limit repeated exposure to TMZ during DC vaccines.

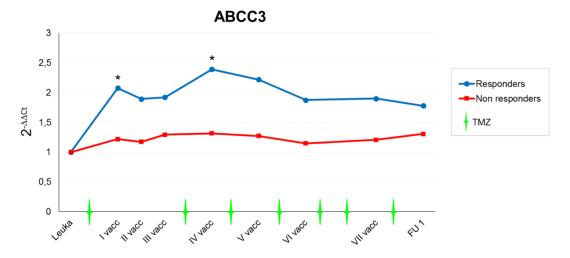


Figure 1. ABCC3 expression in PBLs of DENDR1 patients. ABCC3 transporter is differentially expressed in responders (patients with a PFS \geq 12 and high NK cell response) and non-responders. ABCC3 is over-expressed at the time of first vaccine after radio-TMZ, and after the third vaccine in concomitance with adjuvant chemotherapy. * p = 0.01.

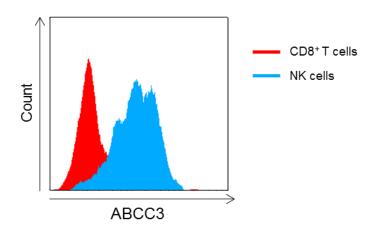


Figure 2. Differential expression of ABCC3 in PBLs of a GBM patient enrolled in DENDR1 study. Representative dot plot of basal ABCC3 positivity expression in NK cells versus CD8⁺ T lymphocytes.

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