

Potential for replacing the boosted-PI with doravirine in ART-experienced subjects successfully treated with darunavir-cobicistat plus dolutegravir dual regimen

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DoDaco Cohort

PURPOSE

Doravirine (DOR) retains full activity against many single resistance-associated mutations (RAMs) selected by older NNRTIs (1,2) and DOR may increase dolutegravir (DTG) exposure (3). Although not formally recommended as dual therapy in people with HIV (PWH), real-life data support such a dual combination as switch option (4-6). We aimed to explore the potential for replacing darunavir-cobicistat DRV-c with DOR in PWH receiving the dual combination of DTG + DRV/c in the DoDaco Cohort, based on the presence of RAMs in their historical genotypes.

METHODS

The DoDaco Cohort includes antiretroviral-experienced PWH on DTG+DRV/c dual regimen from 7 HIV centers. In Italy. Cumulative plasma genotypic resistance testing (GRT) of all participants were evaluated for the presence of NNRTI-associated RAMs (namely: V106A/M, V108I, Y188L, G190S, F227C/L/V, M230I/L, L234I, K103N/P225H, K103N/L100I and K103N/Y181C) and their combinations conferring low, intermediate or high resistance, according to Stanford algorithms.

DRV/c was deemed replaceable by DOR if no major DTG RAMs were concomitantly present, although, in selected cases with low-level DTG resistance, the double DTG dose (i.e. bid) was allowed for the option of DRV/c removal.

CONCLUSIONS

According to historical genotypes, two thirds of PWH included in the DoDaco cohort are eligible for replacing DRV-c with DOR in the combination to DTG, with potential benefits in lipid profile, cardiovascular risk, drug-drug interactions and overall costs.

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RESULTS

A total of **327** individuals (66% males; median age 60 [IQR 55-63] years; median CD4+ count 618 [IQR 386-925] cells/ml, 118 (36%) had AIDS, median (IQR) duration of HIV therapy was 24 (19-28) years, 128 (39%) received more than 8 treatment lines) were included in our cohort. For those with available historical genotypes, RAMs were present in 240 (88%), **191** (70%), 67 (24%), and 18 (6.6%) cases for NRTI, NNRTI, PI and INSTI, respectively. DTG resistance (low to intermediate) was found in 18 (6.6 %) individuals. Figure 1 shows the distribution of NNRTI-associated mutations. Among **219** (67% of the cohort) with documented NNRTI RAMs, 78/219 (35%) had **low, intermediate or high** DOR-associated resistance (Table 2). DTG was not fully active in 2 individuals, so DRV/c may be replaceable by DOR in up to 247/327 (75%) PWH based on the historical genotypes if no specific contraindications. In 31/219 (14%) individuals, DTG bid may be desirable (based on the past INSTI genotype) if DRV/c removal is requested or considered.

Figure 1: Most frequent major NNRTI-resistance mutations (alone or in combination) for DOR

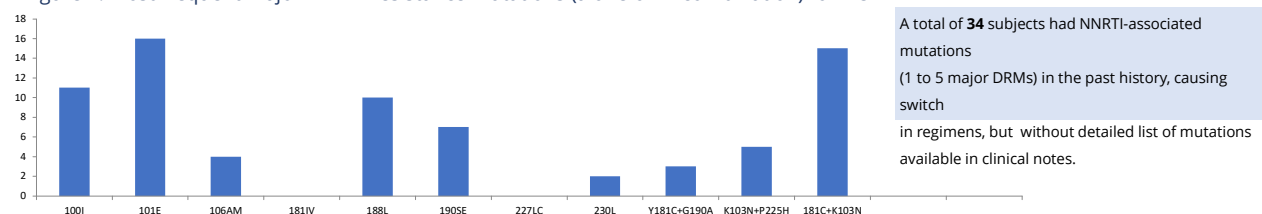


Table 2: distribution of susceptibility to doravirine according to the past genotypic resistance in the DoDaco Cohort

HIV Centre	PWH	PWH with NNRTI RAMs	Grade of susceptibility to doravirine by genotypic resistance (in 113 PWH)			
			Potentially Low	Low	Intermediate	High
Bergamo	124	90	16	15	10	10
Milano (L. Sacco)	59	26	1	1	4	4
Monza	44	28	2	4	1	1
Pavia	50	40	8	3	7	3
Padova	27	19	5	6	3	-
Milano (Santi Paolo e Carlo)	6	4	-	-	-	-
Siena	17	12	3	2	-	4
Total	327	219	35	31	25	22

* according to the Stanford Algorithm interpretation

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