

Integrase Resistance Mutations in Treatment-Naive Persons with Advanced Immunodeficiency Initiating Dual Therapy with Dolutegravir and Lamivudine

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PURPOSE

Integrase and/or Nucleoside Reverse Transcriptase Inhibitor (NRTI) resistance mutations in treatment-naïve individuals may impact the efficacy of dolutegravir/lamivudine (DTG/3TC). This study assessed the prevalence of baseline integrase mutations and their association with virologic outcomes in severely immunocompromised individuals initiating DTG/3TC dual therapy.

METHODS

- The DOLCE study enrolled treatment-naïve, HIV-positive individuals with advanced disease (CD4 count less than 200 cells/mm³) who initiated DTG/3TC (DT) or DTG plus TDF/XTC (TT), randomized 2:1, between 2021 and 2023 in Argentina and Brazil. Virological suppression rates were similar among DT and TT initiators.
- Genotypic resistance testing was performed at screening; results were unavailable at treatment initiation. Mutations were classified per IAS-USA guidelines.
- Protocol-defined exclusionary mutations were G118R, Q148H/K/R, R263K (integrase), and M184V/I (NRTI). Logistic regression evaluated associations between integrase mutations and virologic efficacy.

CONCLUSIONS

Baseline integrase mutations, mostly accessory or polymorphic, were uncommon and did not impact virologic outcomes. We didn't find an association between virological failure and baseline resistance mutations. These findings suggest that routine baseline integrase resistance testing may not be necessary before starting DTG/3TC dual therapy in this population with advanced disease.

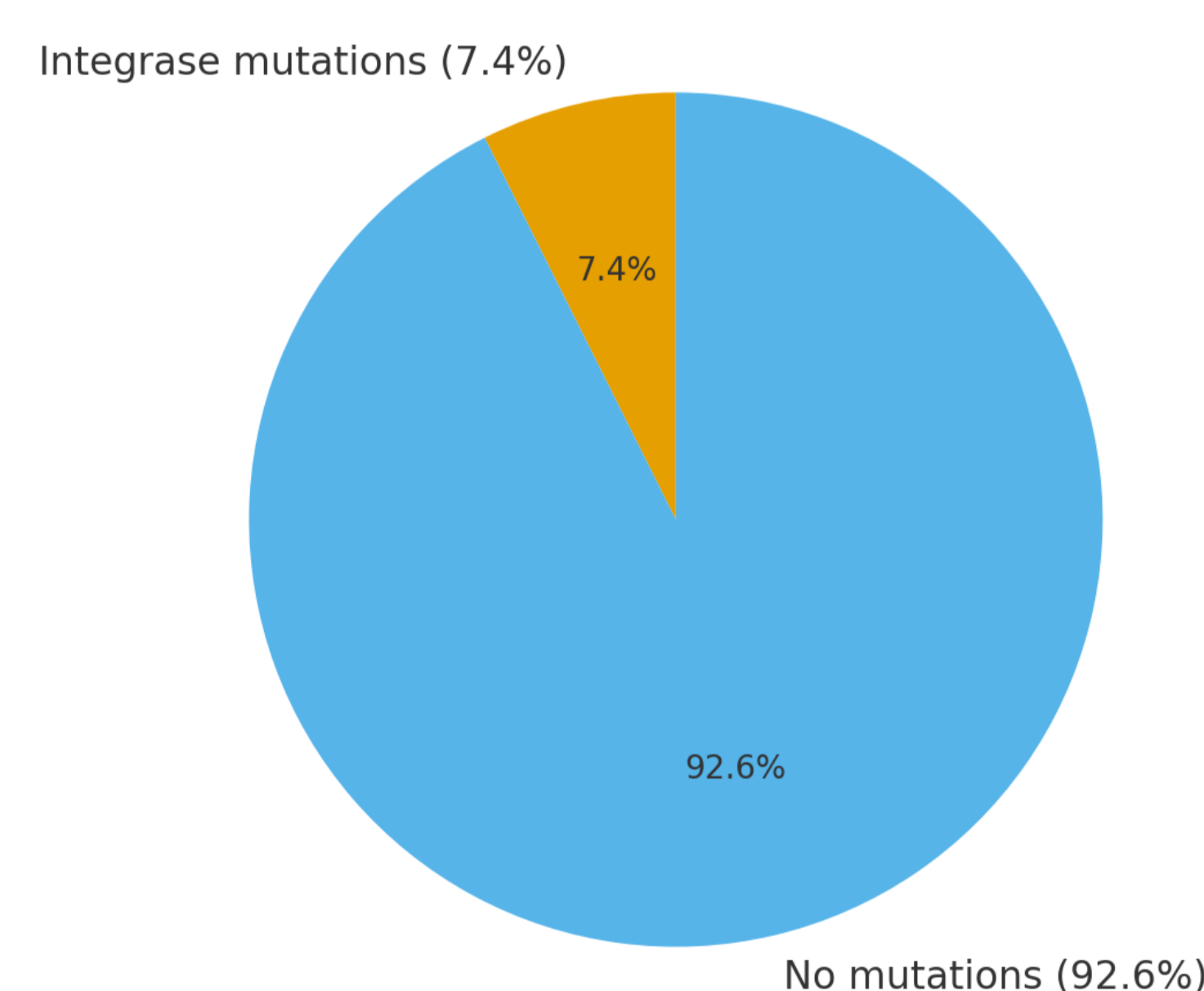
ACKNOWLEDGEMENTS

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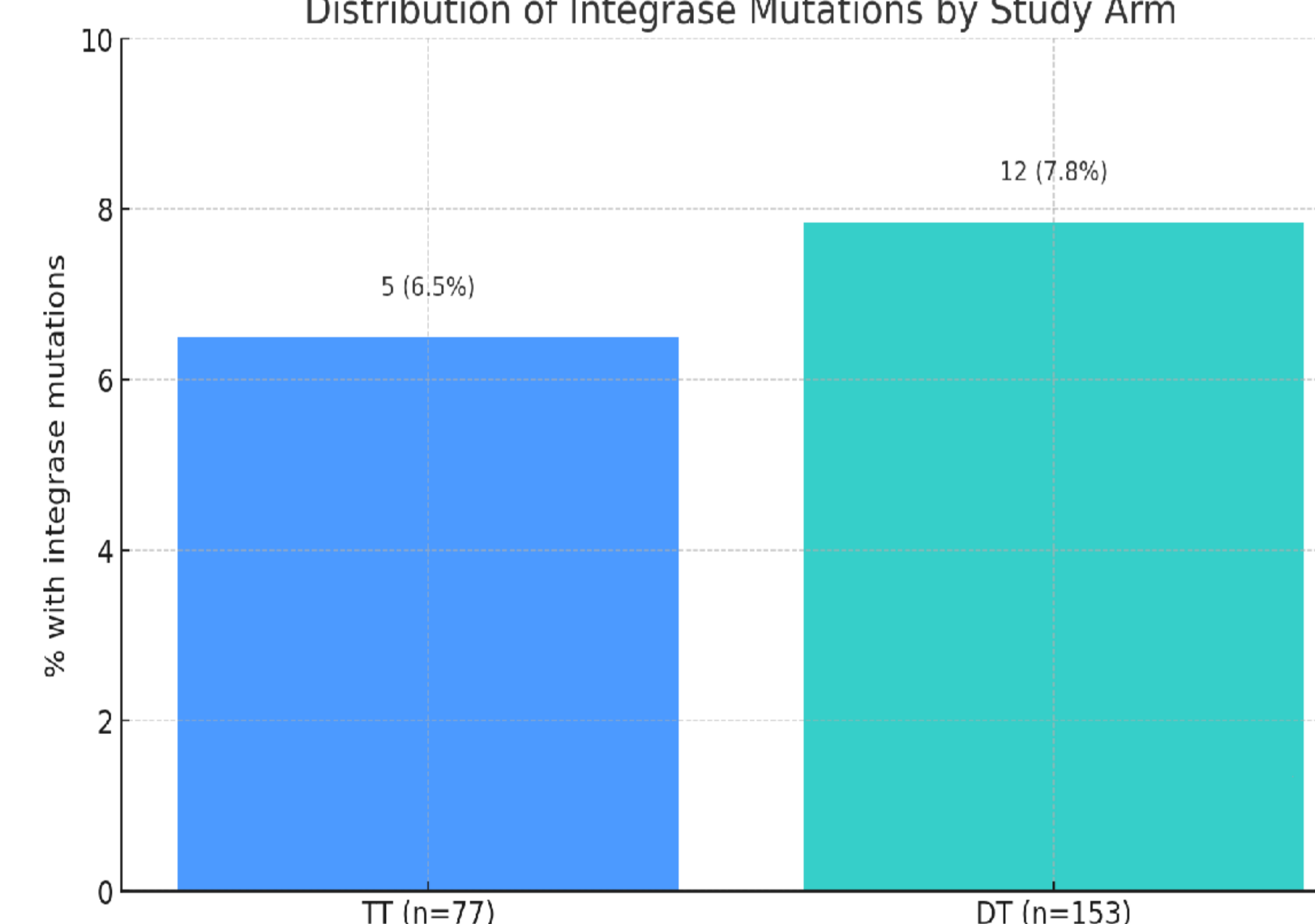
RESULTS

- Among 230 randomized participants, 17 (7.4%) had accessory or polymorphic integrase mutations at baseline: 5 (6.5%) in TT and 12 (7.8%) in DT.
- No major or exclusionary defined by protocol integrase mutations were detected. Variants included G163R, G163K, G163GR, E157EQ, Q146QK, Q95K, D232N, T97A, and E157Q.
- One DT participant harbored the M184V mutation (0.43%).
- At week 48, 14/17 participants with baseline integrase mutations (82.4%; 95% CI: 56–95) achieved viral suppression, similar to 173/213 without integrase mutations (81.2%; 95% CI: 76–96).
- Three participants with integrase mutations experienced protocol-defined virologic failure (63, 114, 94 copies/mL); all were in DT, with high baseline viral loads and good adherence.
- The participant with the M184V mutation achieved sustained virologic suppression. No significant differences in efficacy were observed based on baseline integrase mutations after adjusting for viral load, CD4 count, and HIV-1 subtype ($p = 0.8$).

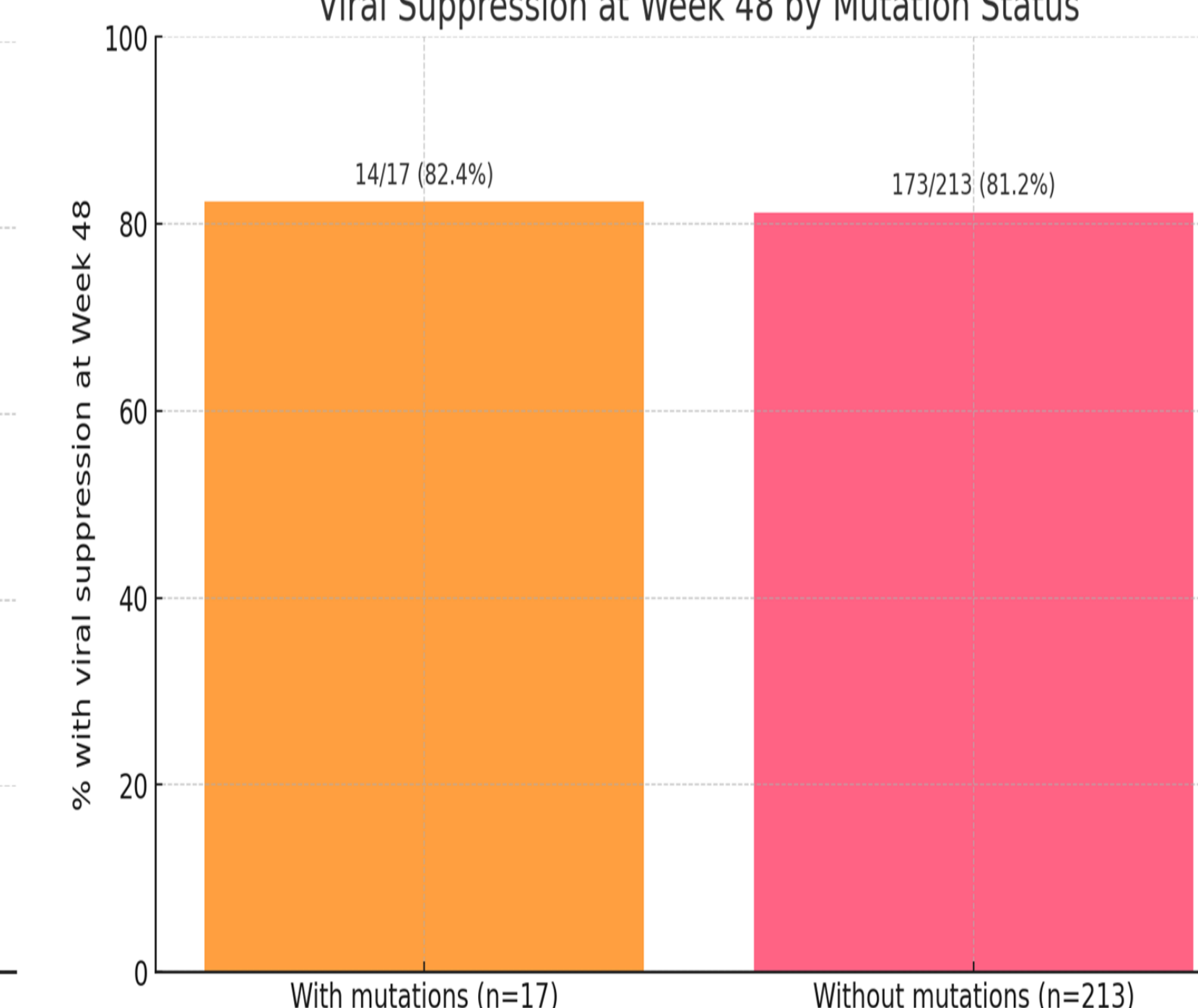
Baseline Integrase Resistance Mutations (n=230)



Distribution of Integrase Mutations by Study Arm



Viral Suppression at Week 48 by Mutation Status



REFERENCES

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