



P251: Transmitted drug resistance to antiretroviral therapy: Interim analysis from a cross-sectional study including 11 healthcare centres in Chile



María Elena Ceballos¹, Cinthya Ruiz-Tagle¹, Felipe Castañeda¹, Marcela Ferrés^{1,2}, Carlos Palma², Angélica Domínguez de Landa¹, Manuel Espinoza¹, Alejandro Afani³, María Elvira Balcells¹

¹ Pontificia Universidad Católica de Chile, Santiago, Chile; ² Red de Salud UC-CHRISTUS, Santiago, Chile; ³ Hospital Clínico Universidad de Chile, Santiago, Chile

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Background

- ❖Antiretroviral therapy (ART) has reduced HIV morbi-mortality and its transmission.
- ❖Treatment failure can occur when acquiring a strain with mutations conferring resistance to ART.
- ❖Transmitted drug resistance to ART (TDR) reported nationally was 10.45% in 2018 and is increasing worldwide.
- ❖In Chile, we do not perform baseline genotyping test

2 Objective

Determine the percentage (n=151)/prevalence (n=168) of global TDR and the relevance of incorporating the genotyping study in naïve people living with HIV in Chile.



Methods

- Observational, cross-sectional study
- ❖Inclusion criteria:
- ❖Exclusion criteria:
- ó18 years old
- ✓ HIV viral load <1000 ARN
- √HIV diagnosis <12 months
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- copies/mL
- ✓No prior ART
- ARN genotyping (nested PCR/Sanger sequencing)
- TDR identified according to HIV Drug Resistance Database, Stanford University.





Results

Figure 1. Transmitted resistance to ART families.

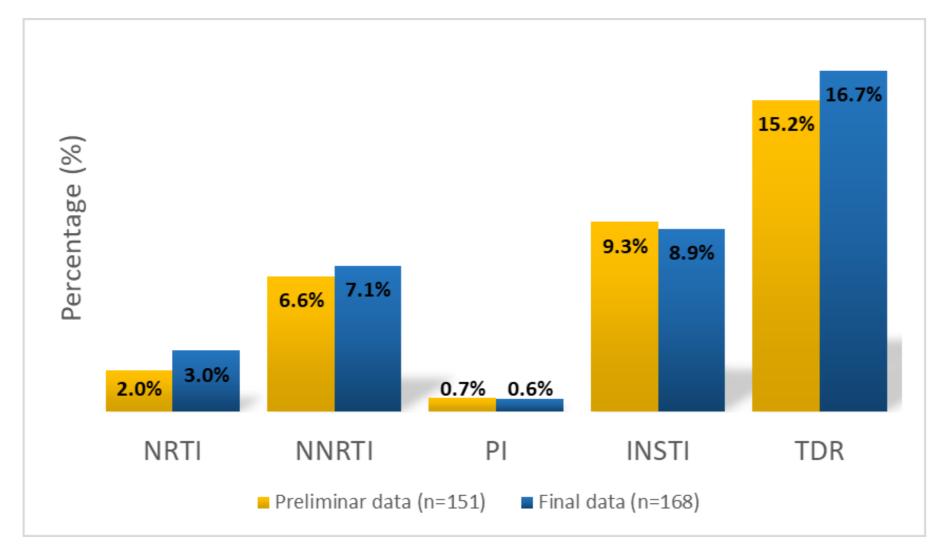
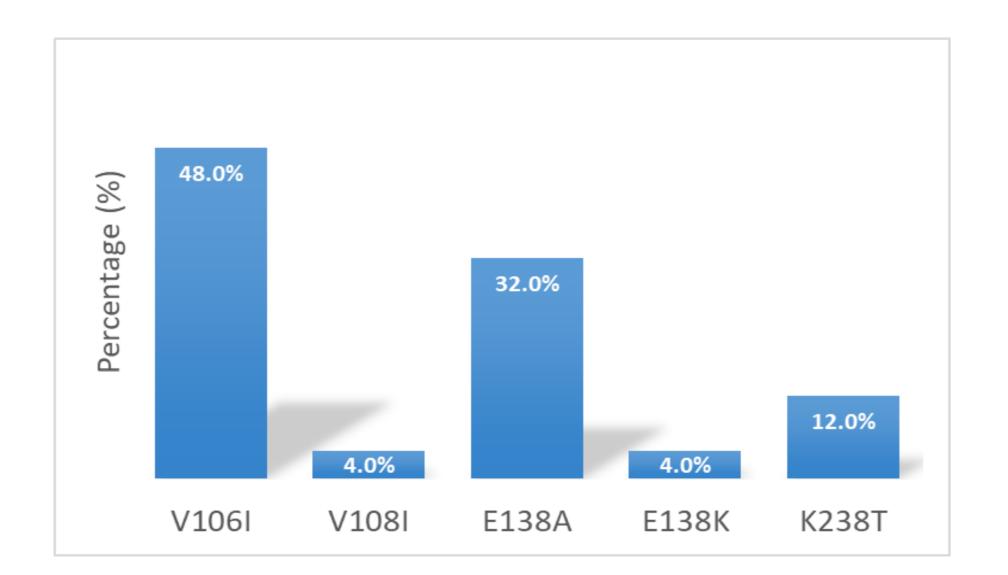


Figure 2. Other NNRTI mutations (not included in the WHO list) conferring transmitted resistance to ART.



Rilpivirine resistance 6,5% (E138A/K, K101E)

Table 1. Transmitted resistance to ART.

	Preliminar	Final data
	data (N=151)	(N=168)
HIV subtype		
В	100 (66.2%)	114 (67.9%)
C	1 (0.7%)	1 (0.6%)
B+F	48 (31.8%)	51 (30.3%)
AG	2 (1.3%)	2 (1.2%)
Overall TDR	23 (15.2%)	28 (16.7%)
TDR to NRTI	3 (2.0%)	5 (3.0%)
NRTI mutations resistance		
M41L	-	1 (14.3%)
D67N	2 (50%)	2 (28.6%)
T69D	-	1 (14.3%)
L210W	1 (25%)	1 (14.3%)
T215E	-	1 (14.3%)
K219Q	1 (25%)	1 (14.3%)
TDR to NNRTI	10 (6.6%)	12 (7.1%)
NNRTI mutations resistance		
K101E	-	1 (9.1%)
K103N	8 (80%)	8 (72.7%)
Combined mutations	2 (20%)	2 (18.2%)
TDR to PI	1 (0.7%)	1 (0.6%)
TDR to INSTI	14 (9.3%)	15 (8.9%)
First generation		20 (11.9%)
Second generation		2 (1.2%)
INSTI mutations resistance		
E92G	1 (5.88%)	1 (5.0%)
T97A	1 (5.88%)	1 (5.0%)
E138A	-	1 (5.0%)
G140S	-	1 (5.0%)
Q148H	-	1 (5.0%)
Q148K	1 (5.88%)	1 (5.0%)
E157Q	2 (11.77%)	2 (10.0%)
G163K	2 (11.77%)	2 (10.0%)
G163R	10 (58.82%)	10 (50.0%)

5 Conclusions

In Chile, an increase is observed in the percentage of

TDR to ART compared to what has been historically reported.

- The main families affected are the NNRTIs and the INSTIs (mostly first generation).
- Considering these preliminary results, it is considered pertinent to incorporate the baseline genotyping study in patients starting ART with both NNRTI efavirenz or rilpivirine and first-generation INSTI.

Funding: FONIS SA22I0035, Agencia Nacional de investigación Desarrollo (ANID).

Acknowledgement: HIV medical care teams from Hospital de Arica, Iquique, Antofagasta, Van Buren, Sótero del Río, Barros Luco Trudeau, Concepción, Talcahuano, Puerto Montt; Red de Salud UC-CHRISTUS and ANID.



Agencia Nacional de Investigación y Desarrollo Ministerio de Ciencia, Tecnología, Conocimiento e Innovación

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Gobierno de Chil

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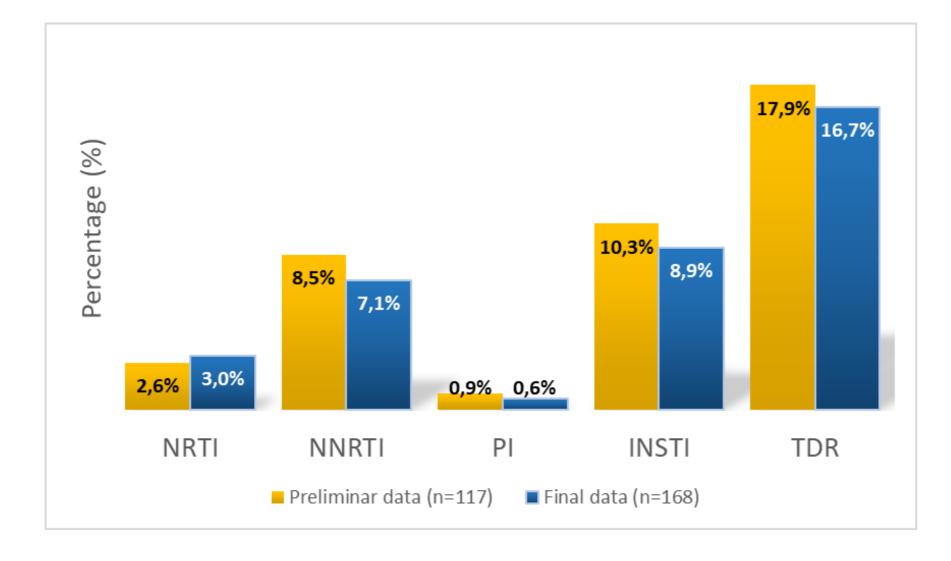
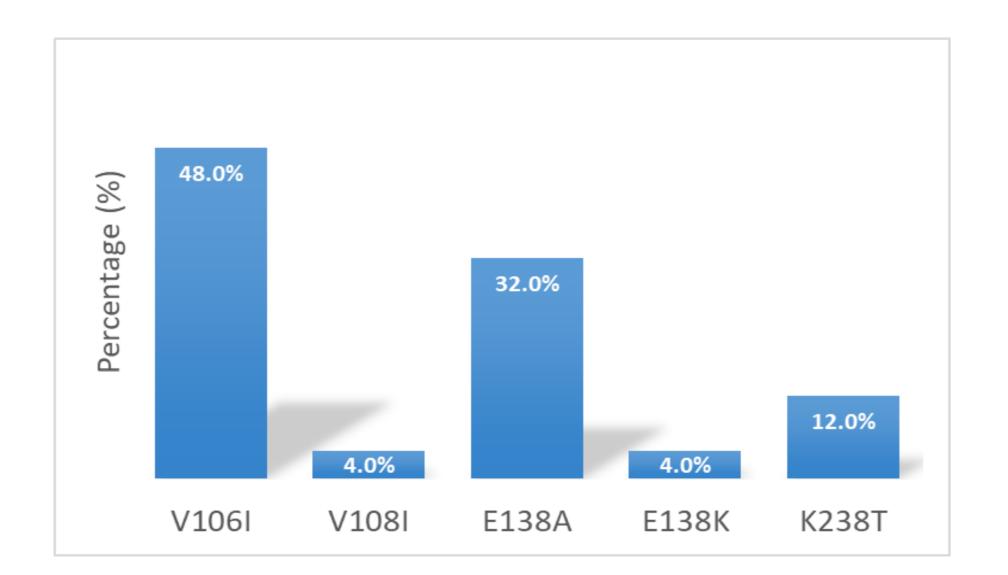


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