

Laura Viñuela¹, A. Fuentes-López², A. de Salazar¹, E. Serrano-Conde¹, MJ. Pérez-Elias², J. Olalla³, A. Pinto-Martínez⁴, JA. Iribarren⁵, M. Masía⁶, M. Montero⁷, I. Falces-Romero⁸, JR. Blanco⁹, M. Rivero¹⁰, L. García-Fraile¹¹, N. Espinosa¹², B. Baza¹³, A. Aguilera¹⁴, MD. Maciá¹⁵, M. Martínez¹⁶, A. Iborra¹⁷, A. Imaz¹⁸, JL. Gómez-Sirvent¹⁹, J. Peraire²⁰, I. Portilla²¹, M. Sanchiz²², I Suárez²³ B Alejos²⁴ and F García¹ on behalf of CoRIS

¹Hospital Universitario San Cecilio; Granada, Spain; ²Hospital Ramón y Cajal; Madrid, Spain; ³Hospital Costa del Sol; Marbella (Málaga), Spain; ⁴Hospital Universitario 12 de Octubre; Madrid, Spain; ⁵Hospital Universitario Donostia; San Sebastian, Spain; ⁶Hospital General Universitario de Elche, Universidad Miguel Hernández; Elche, Spain; ⁷Hospital Universitario La Fe; Valencia, Spain; ⁸Hospital Universitario La Paz/IdiPAZ; Madrid, Spain; ⁹Hospital San Pedro; Logroño, Spain; ¹⁰Hospital de Navarra; Pamplona, Spain; ¹¹Hospital La Princesa; Madrid, Spain; ¹²Hospital Virgen del Rocío; Sevilla, Spain; ¹³Centro Sanitario Sandoval; Madrid, Spain; ¹⁴Complejo Hospitalario Santiago Compostela; Universidad de Santiago de Compostela; Santiago de Compostela, Spain; ¹⁵Hospital Son Espases; Mallorca, Spain; ¹⁶Hospital Universitari MutuaTerrassa; Terrasa, Spain; ¹⁷Hospital Virgen Arrixaca; Murcia, Spain; ¹⁸Hospital Universitario de Bellvitge; Barcelona, Spain; ¹⁹Hospital Universitario de Canarias; Las Palmas de Gran Canaria, Spain; ²⁰Hospital Universitari de Tarragona Joan XXIII, IISPV, Universitat Rovira i Virgili, Tarragona, Spain; ²¹Hospital Alicante; Alicante, Spain; ²²Hospital Universitari Vall d'Hebron; Barcelona, Spain; ²³Hospital Infanta Sofia, Madrid, Spain; ²⁴Instituto de Salud Carlos III, Madrid, Spain.

Background & Aim

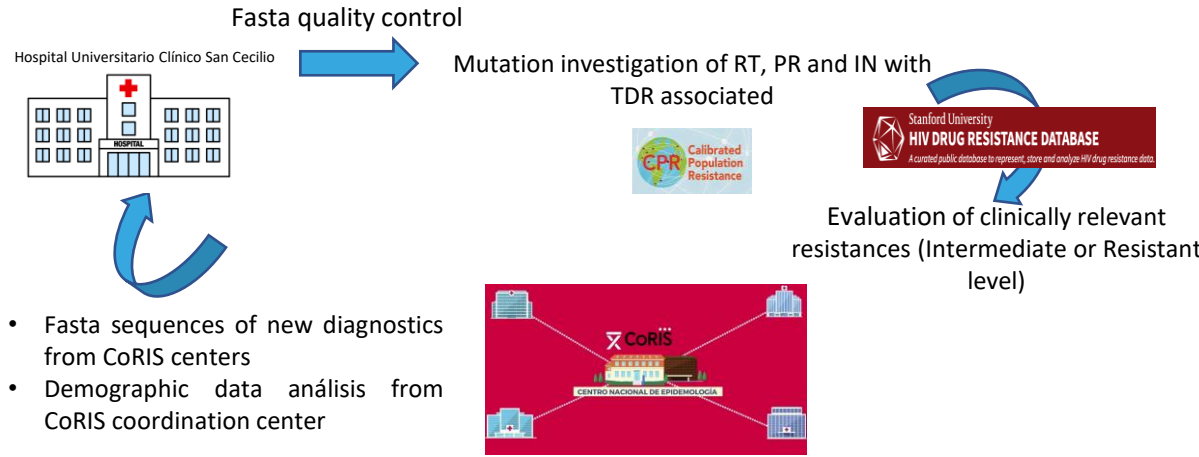
Testing for transmitted drug resistance (TDR) in the reverse transcriptase (RT) and protease (Pro) in newly diagnosed patients with HIV is recommended by clinical guidelines, as a part of the initial clinical assessment. On the other hand, INSTI baseline resistance is not recommended by clinical guidelines, so surveillance programs offering these data in real time are needed.

CoRIS program a platform for HIV research in people living with HIV, was established in 2004 and the transmitted drug resistance (TDR) work package begun in 2007.

- The aims of our study are:
- ✓ To update TDR data in newly diagnosed patients from CoRIS for the period 2019-2021
 - ✓ To present results on transmitted resistance with clinical relevance to drugs included in ART recommended as first line treatment in Spain

Methods

- We analyzed RT and Pro fasta sequences of HIV new diagnostics from 22 CoRIS centers. As integrase is not part of routine testing in NAIVE patients in Spain, we analyzed only a subset of available integrase sequences.
- After fasta quality control, mutations in RT, Pro and integrase associated with TDR were investigated using the CPR-Stanford tool.
- To evaluate clinically relevant resistance (Intermediate or Resistant level) to the drugs currently recommended as first-line treatment in the GESIDA guidelines, as well as to the NNRTIs EFV, DOR and RPV, we used the Stanford v.9.1 HIVDB Algorithm.



Results

A total of **1859** patients were analyzed, 758 (2019), 567 (2020) and 534 (2021); of these, integrase data were available in **651** patients, 264 (2019), 167 (2020) and 220 (2021).

Most of the patients were MSM, aged 26-45, with high school or university education. Half of them presented with less than 10⁵ c/ml HIV-RNA and/or more than 350 CD4 count.

Transmitted drug resistance (TDR) in the period 2019-2021 was: 3.7% for NRTIs, 6.02% for NNRTIs, 0.97% for PIs, and 0.15% for INIs. Of interest, most of the patients with TDR carried singletons

Clinically relevant resistance to first-line drugs in the period 2019-2021 was 1.3% for TDF (of which 1.2% were Intermediates), 2% for ABCs, 0.8% for 3TC/FTCs, 6.5% for EFVs, 2.6% for DOR (1.5% I), 7.1% for RPV, 0.7% for RAL (all Intermediate), 0.05 % for BIC, DTG (n=1, S153F) and DRV

Characteristics	Class	n (%)
Age	<=25	18%
	26-35	43%
	36-45	20%
	>55	6,70%
Education	no studies	0,20%
	primary school	8,80%
	secondary school	43,90%
	university	28,80%
Gender	male	91%
	female	8,90%
	Origin	Spain
Transmission route	Europe (not-Spain)	4,30%
	Africa	4,60%
	Latin/Central America	46,40%
CD4 counts (cells/mm3) (%)	MSM	75,30%
	PWID	0,30%
	MSW	19%
	other/unknown	5,40%
Viral Load (copies/mL) (%)	<200	21,1%
	200-350	25,1%
	350-1000	54,1%
	>1000	1,4%
	Unknown	1,2%
HBV coinfection (%)	<100.000	50,6%
	100.000-500.000	29,2%
	>500.000	17,6%
HCV coinfection (%)	Unknown	2,3%
	Yes	1,8%
	No	70%
Unknown	Unknown	28,1%
	Yes	2,5%
	No	91,4%
Unknown	Unknown	6%

Nº of mutations	NRTIs	NNRTIs*	PIs	INIs**
1 mut	47 (3.4%)	90 (6.6%)	17 (1.2%)	1 (0.15%)
2 mut	13 (1.5%)	22 (1.6%)	1 (0.08%)	-
≥3 mut	3 (0.2%)	-	-	-
Prevalence	3.7%	6.02%	0.97%	0.15%

*E138A not included at WHO list

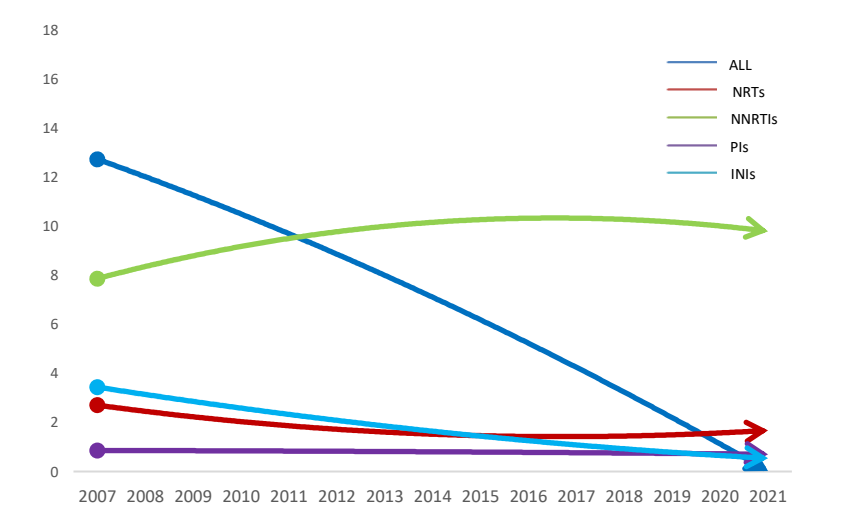
	n	Prevalence (IC 95%)
NRTIs	Tenofovir	25 1.3 (0.9-1.9)
	3TC/FTC	15 0.8 (0.5-1.3)
	Abacavir	39 2.1 (1.5-2.8)
NNRTIs	Efavirenz	120 6.5 (5.3-7.7)
	Rilpivirine	132 7.1 (5.9-8.4)
	Doravirine	48 2.6 (1.9-3.4)

	n	Prevalence (IC 95%)
PIs	Lopinavir	4 0.2 (0.06-0.5)
	Atazanavir	2 0.1 (0.01-0.4)
	Darunavir	1 0.05 (0.0013-0.3)
	Dolutegravir	1 0.05 (0.0013-0.3)
INIs	Raltegravir	13 0.7 (0.4-1.2)
	Bictegravir	1 0.05 (0.0013-0.3)

A detailed description of the most relevant TDR mutations included in current SDRM lists is shown below

	Mutation	n (%)	
NRTIs	M41L	22 (1.6%)	
	M184IV	14 (1.02%)	
	L210W	12 (0.87%)	
	Others*	21 (1.5%)	
	M46IL	9 (0.65%)	
PIs	I47V	2 (0.15%)	
	I85V	2 (0.15%)	
	Others**	6 (0.4%)	
	K103NS	91 (6.6%)	
	K101EP	13 (0.95%)	
NNRTIs	G190AES	13 (0.95%)	
	P225H	5 (0.4%)	
	Others***	8 (0.6%)	
	INIs	E92Q	1 (0.15%)

From 2007, we have assisted to a decrease in clinically relevant resistance (CRR) to all drug classes except for NNRTIs, driven by the introduction of second generation integrase inhibitors as preferred drugs for first line regimens



Conclusions

As in previous years, the highest prevalence of transmitted resistance occurred for NNRTIs, with doravirine being the drug in this family with the lowest levels of TDR. Transmitted resistance to PIs, INIs, and 3TC/FTC continues at very low levels, with no patient presenting full resistance to 2nd generation integrase inhibitors or darunavir.

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*D67NG, K219QENR, K70E, T215SDYQEI, L74IV, T69D; **D30N, N88D, F35Y, G73C, L90M, V82A ***Y188CL, L100I, V179F, Y181C