

HIV Drug Resistance Report

Patient information

Name: H,T

Sex: \mathbf{F}

DOB/Age: **31/12/1985**

H&C number/SP Number: 9903393725

Lab no: $\mathbf{V33024431}$

Sequence summary

Sequence includes HIV1 IN, codons 1-288 missing: None

HIV subtype: HIV subtype cannot be determined from integrase alone

Drug	Generic Name	Type	Resistance Call
BIC	bictegravir	INSTI	Susceptible
CAB	cabotegravir	INSTI	Susceptible
DTG	dolutegravir	INSTI	Susceptible
EVG	elvitegravir	INSTI	Susceptible
RAL	raltegravir	INSTI	Susceptible

Date of specimen collection: 14/05/2022 Specimen type: EDTA Blood Date received: 05/05/2023

Regional Virus Laboratory/Pathogen Sequencing LG.031 Institute of Pathology Belfast Health and Social Care Trust Grosvenor Road, Belfast, BT12 6BA



Significant mutations

Integrase inhibitor mutations: No mutations of this type were detected

Other mutations

IN other mutations: E11D, S24N, S39M, I72V, L101I, S119P, I135V, Q177L, V201I, L234I, S283G

Significant comments

INT comments:

No comment

Other comments

IN comments:

No comment

Definitions for drug susceptibility

"Susceptible" indicates no evidence of reduced ARV susceptibility compared with a wild-type virus.

"Potential low-level resistance" indicates that the sequence may contain mutations indicating previous ARV exposure or may contain mutation that are associated with drug resistance only when they occur with additional mutations.

"Low-level resistance" indicates that there that the virus encoded by the submitted sequence may have reduced in vitro ARV susceptibility or that patients harboring viruses with the submitted mutations may have a suboptimal virological response to treatment with the ARV. "Intermediate resistance" indicates a high likelihood that a drug's activity will be reduced but that the drug will likely retain significant remaining antiviral activity.

High-level resistance" indicates that the predicted level of resistance is similar to those observed in viruses with the highest levels of in vitro drug resistance or that clinical data exist demonstrating that patients infected with viruses having such mutations usually have little or no virological response to treatment with the ARV. Further information is available here.

Date and time created: 09/10/2023,09:43