

A1: Summary Report

Patient Details

Patient Name:	GINA REGAL	Barcode:	1002
DOB:	1/02/00	NHI:	NHX1212
Gender:	F	Ethnicity:	NZ EUROPEAN
Location:	HAMILTON	Sample Type:	SPUTUM
Sample Date:	9/11/16	Culture Method:	MGIT CULTURED ISOLATES
Reporting Lab:	LABPLUS	Technician:	DR ROD JACKSON
Sequencing Date:	19/05/20	Sequencing:	WGS
Requested by:	DR. EXAMPLE	Contact:	EMAIL.ADDRESS@DHB.GOV.T.NZ

Resistance Profile

Drug Class	Resistance	Drug	Mutation
First Line	Sensitive	Ethambutol	No resistance predicted
		Pyrazinamide	No resistance predicted
	Resistance	Isoniazid Rifampicin	fabG1 — rpoB Asp435Tyr
Second Line	Sensitive	Amikacin	No resistance predicted
		Capreomycin	No resistance predicted
		Kanamycin	No resistance predicted
		Linezolid	No resistance predicted
		Para-aminosalicylic	No resistance predicted
	Resistance	Ethionamide Fluoroquinolones Streptomycin	fabG1 — gyrB Asn499Asp rpsL Lys43Arg

A2: Whole Genome Sequencing Report

Genotypic Report

Drug	Genomics Position	Allele		Gene (Mutation)	High Confidence	PMID
		Reference	Alternative			
Isoniazid	1673425	C	T	fabG1 —	yes	26104204
Rifampicin	761109	G	T	rpoB Asp435Tyr	yes	11880449
Ethionamide	1673425	C	T	fabG1 —	yes	26104204
Streptomycin	781687	A	G	rpsL Lys43Arg	yes	22646308
Fluoroquinolones	6734	A	G	gyrB Asn499Asp	yes	17412727
Fluoroquinolones	6738	C	A	gyrB Thr500Asn	yes	24055765
Fluoroquinolones	6741	A	T	gyrB Glu501Val	yes	19721073
Fluoroquinolones	6750	C	T	gyrB Ala504Val	yes	24055765
Fluoroquinolones	7563	G	T	gyrA Gly88Cys	yes	24055765
Fluoroquinolones	7564	G	C	gyrA Gly88Ala	yes	24055765
Fluoroquinolones	7572	T	C	gyrA Ser91Pro	yes	21562102
Fluoroquinolones	7582	A	G	gyrA Asp94Gly	yes	21300839
Fluoroquinolones	7582	A	C	gyrA Asp94Ala	yes	21300839
Fluoroquinolones	6749	G	A	gyrB Ala504Thr	no	19470506
Fluoroquinolones	7582	A	T	gyrA Asp94Val	no	17434825

NOTE: This genotypic report is a prediction of drug resistance based on the detection of known resistance conferring mutations, and should only be used as a guide. The actual drug susceptibility may differ from what is stated in this report.

A3: Technical Report

Pipeline

Pipeline	Resistance V0.01a	Genome	H37Rv
Phred Score	30	Masking	NO

Forward Statistics: WRX.R1.fastq.gz

Pre-Trimming		Post-Trimming	
Total Reads	VALUE	Total Reads	VALUE (%)
Average Quality	VALUE	Average Quality	VALUE
GC%	VALUE	GC%	VALUE
Mean read length	VALUE	Mean read length	VALUE
Read range	VALUE	Read range	VALUE

Reverse Statistics

File name	NAME		
Pre-Trimming		Post-Trimming	
Total Reads	VALUE	Total Reads	VALUE (%)
Average Quality	VALUE	Average Quality	VALUE
GC%	VALUE	GC%	VALUE
Mean read length	VALUE	Mean read length	VALUE
Read range	VALUE	Read range	VALUE

Alignment Statistics

Notes