

Supplementary tables

Table S1 Distribution of drug resistance conferring mutations in different genes included in the cluster analysis for INH, RIF, EMB, PZA, SLIDs and FQs among 461 XDR-TB patients diagnosed with XDR-TB in the Western Cape Province of South Africa, 2006 – 2017

Drugs	Gene	Different mutations	Isolates n (%)
INH	<i>inhA</i> promotor	<i>fabG1_c.-15C>T</i>	31 (6.7)
	<i>inhA</i> promotor + <i>ahpC</i>	<i>fabG1_c.-15C>T, ahpC_c.-48G>A</i>	1 (0.2)
	<i>katG + ahpC</i>	<i>katG_p.Ser315Thr, ahpC_c.-48G>A</i>	13 (2.8)
	<i>inhA</i> promotor + <i>inhA</i>	<i>fabG1_c.-15C>T, inhA_p.Ile194Thr</i>	72 (15.6)
	<i>inhA + katG</i>	<i>inhA_p.Ser94Ala, katG_p.Ser315Thr</i>	3 (0.7)
	<i>inhA</i> promotor + <i>katG</i>	<i>fabG1_c.-15C>T, katG_p.Ser315Thr</i>	143 (31.0)
		<i>fabG1_c.-15C>T, katG_p.Thr394Ala</i>	1 (0.2)
		<i>fabG1_c.-17G>T, katG_p.Ser315Thr</i>	153 (33.2)
		<i>fabG1_c.-8T>A, katG_p.Ser315Thr</i>	7 (0.4)
	<i>katG</i>	<i>katG_p.Ser315Thr</i>	37 (8.0)
RIF	<i>rpoB</i>	<i>rpoB_p.Asp435Gly, rpoB_p.Leu452Pro</i>	5 (1.1)
		<i>rpoB_p.Asp435Tyr</i>	4 (0.9)
		<i>rpoB_p.Asp435Val</i>	154 (33.4)
		<i>rpoB_p.Gln432Lys</i>	1 (0.2)
		<i>rpoB_p.His445Asn</i>	1 (0.2)
		<i>rpoB_p.His445Asp</i>	1 (0.2)
		<i>rpoB_p.Ser450Leu</i>	286 (62.0)
		<i>rpoB_p.Ser450Leu, rpoB_p.His445Arg</i>	1 (0.2)
		<i>rpoB_p.Ser450Leu, rpoB_p.Thr400Ala</i>	2 (0.4)
		<i>rpoC_p.Leu527Val, rpoB_p.Ser450Leu</i>	4 (0.9)
		<i>rpoC_p.Phe452Ser, rpoB_p.Ser450Leu</i>	2 (0.4)
EMB*	<i>embA</i>	<i>embA_c.-16C>G</i>	4 (0.9)
	<i>embB</i>	<i>embB_p.Asp354Ala</i>	2 (0.4)
		<i>embB_p.Gln497Arg</i>	5 (1.1)
		<i>embB_p.Gln497Lys, embB_p.Met306Ile</i>	3 (0.7)
		<i>embB_p.Gly406Asp</i>	2 (0.4)
		<i>embB_p.Gly406Ser</i>	2 (0.4)
		<i>embB_p.Met306Ile</i>	304 (65.9)
		<i>embB_p.Met306Ile, embB_p.Asp1024Asn</i>	1 (0.2)
		<i>embB_p.Met306Val</i>	127 (27.5)
	<i>embB + embA</i>	<i>embB_p.Met306Val, embA_c.-16C>T</i>	2 (0.4)
PZA**	<i>pncA</i>	<i>pncA_c.135_136insTG</i>	1 (0.2)
		<i>pncA_c.389_389del</i>	1 (0.2)
		<i>pncA_c.391_392insGG</i>	1 (0.2)
		<i>pncA_c.456_457insC</i>	5 (1.1)
		<i>pncA_c.459_466del</i>	3 (0.7)
		<i>pncA_c.517_518insG</i>	153 (33.2)
		<i>pncA_p.Asp8Asn</i>	33 (7.2)
		<i>pncA_p.Cys14Arg</i>	144 (31.2)
		<i>pncA_p.Cys14Trp</i>	1 (0.2)
		<i>pncA_p.Gln10Pro</i>	3 (0.7)
		<i>pncA_p.Gly132Ala</i>	1 (0.2)

		<i>pncA</i> _p.Met175Val	1 (0.2)
		<i>pncA</i> _p.Phe13Leu	1 (0.2)
		<i>pncA</i> _p.Ser164Pro	6 (1.3)
		<i>pncA</i> _p.Thr135Pro	6 (1.3)
		<i>pncA</i> _p.Thr142Met	1 (0.2)
		<i>pncA</i> _p.Tyr103*	10 (2.2)
		<i>pncA</i> _p.Val130Ala	7 (1.5)
		<i>pncA</i> _p.Val139Gly	1 (0.2)
		<i>pncA</i> _p.Val139Met	13 (2.8)
SLIDs	<i>rrs</i>	<i>rrs</i> _r.1401a>g	448 (97.2)
		<i>rrs</i> _r.1402c>t	1 (0.2)
	<i>eis</i>	<i>eis</i> _c.-10G>A	2 (0.4)
		<i>eis</i> _c.-10G>C	4 (0.9)
		<i>eis</i> _c.-12C>T	2 (0.4)
		<i>eis</i> _c.-14C>T	1 (0.2)
		<i>eis</i> _c.-37G>T	2 (0.4)
		<i>eis</i> _c.-8C>A	1 (0.2)
FQs	<i>gyrA</i>	<i>gyrA</i> _p.Ala90Val	67 (14.5)
		<i>gyrA</i> _p.Asp89Asn	1 (0.2)
		<i>gyrA</i> _p.Asp94Ala	51 (11.1)
		<i>gyrA</i> _p.Asp94Ala, <i>gyrA</i> _p.Ala90Val	2 (0.4)
		<i>gyrA</i> _p.Asp94Asn	55 (11.9)
		<i>gyrA</i> _p.Asp94Gly	206 (44.7)
		<i>gyrA</i> _p.Asp94His	4 (0.9)
		<i>gyrA</i> _p.Asp94Tyr	16 (3.5)
		<i>gyrA</i> _p.Gly88Cys	15 (3.3)
		<i>gyrA</i> _p.Ser91Pro	18 (3.9)
	<i>gyrB</i>	<i>gyrA</i> _p.Ser91Pro, <i>gyrA</i> _p.Ala90Val	1 (0.2)
		<i>gyrA</i> _p.Ser91Pro, <i>gyrA</i> _p.Asp94Gly	1 (0.2)
		<i>gyrB</i> _p.Arg446Cys	6 (1.3)
		<i>gyrB</i> _p.Asn499Asp	1 (0.2)
		<i>gyrB</i> _p.Asn499Lys	1 (0.2)
		<i>gyrB</i> _p.Asp461Asn	4 (0.9)
		<i>gyrB</i> _p.Asp461His	3 (0.7)
	<i>gyrA + gyrB</i>	<i>gyrB</i> _p.Glu501Asp	5 (1.1)
		<i>gyrB</i> _p.Glu501Val	1 (0.2)
		<i>gyrB</i> _p.Arg446Cys, <i>gyrA</i> _p.Ala90Val	1 (0.2)
		<i>gyrB</i> _p.Asp461Asn, <i>gyrA</i> _p.Ala90Val	2 (0.4)

^a NA = not applicable, INH = isoniazid, RIF = rifampicin, EMB = ethambutol, PZA = pyrazinamide, SLIDs = second line injectable drugs, FQs = fluoroquinolones, *isolates susceptible to EMB (n=9), **isolates with no pncA mutations and susceptible to PZA (n=69)

Table S2 All mutations observed in the samples included in the analyses

SAMPLE	RIF	INH	PZA	EMB	FQS	SLIDS
R20784	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R24480	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	pncA_p.Thr135Pro	embB_p.Met306Val	gyrA_p.Ala90Val	rrs_r.1401a>g
R34488	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Asn	rrs_r.1401a>g
R6709	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Asn	rrs_r.1401a>g
R36768	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Asn	rrs_r.1401a>g
R38716	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	-	embB_p.Met306Val	gyrA_p.Asp94Ala, gyrA_p.Ala90Val	rrs_r.1401a>g
R35914	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	-	embB_p.Met306Val	gyrA_p.Asp94Ala	rrs_r.1401a>g
R35861	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R19366	rpoB_p.Asp435Val	katG_p.Ser315Thr	pncA_p.Phe13Leu	embB_p.Met306Ile	gyrA_p.Asp94Asn	rrs_r.1401a>g
R15097	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Tyr	rrs_r.1401a>g
R4274	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Ala90Val	rrs_r.1401a>g
R39526	rpoB_p.Ser450Leu	ahpC_c.-48G>A, katG_p.Ser315Thr	pncA_p.Val139Met	embB_p.Met306Val	gyrA_p.Gly88Cys	rrs_r.1401a>g
R11121	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Tyr	rrs_r.1401a>g
R13403	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	-	embB_p.Met306Val	gyrA_p.Asp94Ala	rrs_r.1401a>g
R13123	rpoB_p.Ser450Leu	fabG1_c.-15C>T	pncA_p.Cys14Trp	embB_p.Met306Val	gyrA_p.Asp94Gly	rrs_r.1401a>g
R23865	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R27545	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R29439	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Ala90Val	rrs_r.1401a>g

R11654	rpoB_p.Ser450Leu	katG_p.Ser315Thr, inhA_p.Ser94Ala	pncA_p.Gln10Pro	embB_p.Gln497Arg	gyrA_p.Asp94Tyr	-
R32587	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R21471	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	-	embB_p.Met306Val	gyrA_p.Asp94Ala	rrs_r.1401a>g
R20376	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R33930	rpoB_p.Ser450Leu	fabG1_c.-15C>T	-	-	gyrA_p.Ala90Val	rrs_r.1401a>g
R37017	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R31116	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Ser91Pro	rrs_r.1401a>g
R10867	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R24220	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Asn	rrs_r.1401a>g
R27133	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Ser91Pro	rrs_r.1401a>g
X116	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Ala90Val	rrs_r.1401a>g
R19367	rpoB_p.Ser450Leu, rpoB_p.Thr400Ala	fabG1_c.-15C>T	-	embB_p.Gly406Ser	gyrA_p.Asp94Ala	rrs_r.1401a>g
R27188	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R4218	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R22010	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R19517	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	pncA_p.Val130Ala	embB_p.Met306Val	gyrA_p.Asp94Gly	rrs_r.1401a>g
R27001	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	-	embB_p.Met306Val	gyrA_p.Asp94Gly	rrs_r.1401a>g
R39203	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R13723	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Ala90Val	rrs_r.1401a>g
R17660	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g

R21670	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R14894	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R18094	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R29343	rpoB_p.Ser450Leu	katG_p.Ser315Thr	pncA_c.391_392insGG	embB_p.Met306Ile	gyrA_p.Ala90Val	rrs_r.1401a>g
R10435	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	pncA_p.Asp8Asn	embB_p.Met306Val	gyrA_p.Asp94Gly	rrs_r.1401a>g
R33907	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Ala90Val	rrs_r.1401a>g
R8695	rpoB_p.Ser450Leu	ahpC_c.-48G>A, katG_p.Ser315Thr	pncA_p.Val139Met	embB_p.Met306Val	gyrA_p.Asp94Gly	-
R16642	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	pncA_p.Thr135Pro	embB_p.Met306Val	gyrA_p.Ala90Val	rrs_r.1401a>g
X167	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	pncA_p.Ser164Pro	embB_p.Met306Val	gyrA_p.Ala90Val	rrs_r.1401a>g
R20666	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrB_p.Asn499Lys	rrs_r.1401a>g
R4250	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R18476	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R20954	rpoB_p.Asp435Tyr	katG_p.Ser315Thr	-	embA_c.-16C>G	gyrA_p.Asp94Gly	rrs_r.1401a>g
R10912	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R9908	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	-	embB_p.Met306Val	gyrA_p.Asp94Ala	rrs_r.1401a>g
R15629	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Asn	rrs_r.1401a>g
R27153	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R24194	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Ala90Val	rrs_r.1401a>g
R19148	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R6265	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	-	embB_p.Met306Val	gyrA_p.Asp94Ala	rrs_r.1401a>g
R34500	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g

R9762	rpoB_p.Ser450Leu, rpoB_p.Thr400Ala	fabG1_c.-15C>T	-	embB_p.Gly406Ser	gyrA_p.Asp94Ala	rrs_r.1401a>g
R10552	rpoB_p.Ser450Leu	katG_p.Ser315Thr, inhA_p.Ser94Ala	pncA_p.Gln10Pro	embB_p.Gln497Arg	gyrA_p.Asp94Tyr	-
R21313	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Thr394Ala	pncA_p.Tyr103*	embB_p.Met306Ile	gyrA_p.Ala90Val	rrs_r.1401a>g
R27420	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R4853	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R13060	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R6297	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrB_p.Asp461Asn	-
R31647	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Ala90Val	rrs_r.1401a>g
R39270	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	pncA_p.Asp8Asn	embB_p.Met306Val	gyrA_p.Ser91Pro	rrs_r.1401a>g
R5235	rpoB_p.Ser450Leu	fabG1_c.-15C>T	-	embB_p.Met306Ile	gyrA_p.Ala90Val	rrs_r.1401a>g
R28124	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R26837	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R35320	rpoB_p.Asp435Tyr	katG_p.Ser315Thr	-	embA_c.-16C>G	gyrA_p.Asp94Gly	rrs_r.1401a>g
R32251	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R30006	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	-	embB_p.Met306Val	gyrA_p.Asp94Ala	rrs_r.1401a>g
R5908	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R26993	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Ala90Val	rrs_r.1401a>g
R23187	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94His	rrs_r.1401a>g
R27711	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R19712	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Asn	rrs_r.1401a>g
R11405	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g

R9964	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R29546	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Ser91Pro	rrs_r.1401a>g
R11581	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R16785	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94His	rrs_r.1401a>g
R4558	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Asn	rrs_r.1401a>g
R14770	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	-	embB_p.Met306Val	gyrA_p.Asp94Ala	rrs_r.1401a>g
R17181	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R5581	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	-	embB_p.Gln497Arg	gyrA_p.Ala90Val	rrs_r.1401a>g
R36647	rpoB_p.Ser450Leu	fabG1_c.-15C>T	pncA_p.Tyr103*	embB_p.Met306Val	gyrA_p.Asp94Gly	-
R10318	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	-	embB_p.Met306Val	gyrA_p.Asp94Ala	rrs_r.1401a>g
R11448	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Ala90Val	rrs_r.1401a>g
R20710	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R34486	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Ala	rrs_r.1401a>g
R33481	rpoB_p.Ser450Leu, rpoC_p.Leu527Val	fabG1_c.-15C>T, inhA_p.Ile194Thr	-	embB_p.Met306Val	gyrA_p.Ala90Val	rrs_r.1401a>g
R27383	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	-	embB_p.Met306Val	gyrA_p.Asp94Ala	rrs_r.1401a>g
R36909	rpoB_p.Ser450Leu	katG_p.Ser315Thr	pncA_p.Asp8Asn	embB_p.Met306Val	gyrA_p.Asp94Gly	rrs_r.1401a>g
R13188	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	-	embB_p.Met306Val	gyrA_p.Asp94Ala	rrs_r.1401a>g
R30371	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Ala90Val	rrs_r.1401a>g
R19868	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R23831	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R22020	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Ala90Val	rrs_r.1401a>g

R15633	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	-	embB_p.Met306Val	gyrA_p.Asp94Ala	rrs_r.1401a>g
R11139	rpoB_p.Ser450Leu	fabG1_c.-15C>T	pncA_c.459_466del	embB_p.Met306Val	gyrA_p.Ser91Pro	rrs_r.1401a>g
X4	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Ser91Pro	rrs_r.1401a>g
R9946	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrB_p.Asp461Asn	rrs_r.1401a>g
R10873	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Asn	rrs_r.1401a>g
R24620	rpoB_p.Ser450Leu	fabG1_c.-15C>T	pncA_c.459_466del	embB_p.Met306Val	gyrA_p.Asp94Gly	rrs_r.1401a>g
R10921	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Asn	rrs_r.1401a>g
R4888	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R35632	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	-	embB_p.Met306Val	gyrA_p.Asp94Ala	rrs_r.1401a>g
R30452	rpoB_p.Ser450Leu	katG_p.Ser315Thr	pncA_p.Asp8Asn	embB_p.Met306Val	gyrA_p.Asp94Gly	rrs_r.1401a>g
R21302	rpoB_p.Ser450Leu	katG_p.Ser315Thr	pncA_p.Asp8Asn	embB_p.Met306Val	gyrA_p.Asp94Gly	rrs_r.1401a>g
R39835	rpoB_p.Ser450Leu	katG_p.Ser315Thr	pncA_p.Asp8Asn	embB_p.Met306Val	gyrA_p.Asp94Gly	rrs_r.1401a>g
R29662	rpoB_p.Ser450Leu	fabG1_c.-15C>T	-	embB_p.Met306Val	gyrB_p.Arg446Cys	rrs_r.1401a>g
R37464	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R27664	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R31345	rpoB_p.Ser450Leu	ahpC_c.-48G>A, katG_p.Ser315Thr	pncA_p.Val139Met	embB_p.Met306Val	gyrA_p.Asp94Gly	rrs_r.1401a>g
R24675	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Ala90Val	rrs_r.1401a>g
R33323	rpoB_p.Ser450Leu	fabG1_c.-15C>T	-	-	gyrA_p.Ala90Val	rrs_r.1401a>g
R3715	rpoB_p.Ser450Leu	katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Asn	rrs_r.1401a>g
R39416	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R18531	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Asn	rrs_r.1401a>g
R3696	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Asn	rrs_r.1401a>g

R12966	rpoB_p.Ser450Leu	fabG1_c.-15C>T	-	embB_p.Met306Val	gyrB_p.Arg446Cys	rrs_r.1401a>g
R17871	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Asn	rrs_r.1401a>g
R27129	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R17207	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	-	embB_p.Met306Val	gyrA_p.Asp94Ala	rrs_r.1401a>g
R24417	rpoB_p.Ser450Leu	katG_p.Ser315Thr	pncA_p.Asp8Asn	embB_p.Met306Val	gyrA_p.Asp94Gly	rrs_r.1401a>g
R26807	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Asn	rrs_r.1401a>g
R39422	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrB_p.Asp461Asn, gyrA_p.Ala90Val	rrs_r.1401a>g
R9127	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R18066	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Tyr	rrs_r.1401a>g
R3896	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
X130	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Asn	rrs_r.1401a>g
R36850	rpoB_p.Asp435Gly, rpoB_p.Leu452Pro	fabG1_c.-8T>A, katG_p.Ser315Thr	pncA_c.456_457insC	embB_p.Met306Val	gyrA_p.Ala90Val	rrs_r.1401a>g
R19234	rpoB_p.Ser450Leu	katG_p.Ser315Thr	-	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R13602	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	pncA_p.Asp8Asn	embB_p.Met306Val	gyrA_p.Ser91Pro	rrs_r.1401a>g
R9882	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Gly88Cys	rrs_r.1401a>g
R8073	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Asn	rrs_r.1401a>g
R17400	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	-	embB_p.Met306Val	gyrA_p.Asp94Ala	rrs_r.1401a>g
R18522	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	-	embB_p.Met306Val	gyrA_p.Asp94Ala	rrs_r.1401a>g
R34305	rpoB_p.Ser450Leu	ahpC_c.-48G>A, katG_p.Ser315Thr	pncA_p.Val139Met	embB_p.Met306Val	gyrB_p.Asp461His	rrs_r.1401a>g
R31322	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Asn	rrs_r.1401a>g
R28413	rpoB_p.Ser450Leu	katG_p.Ser315Thr	pncA_p.Asp8Asn	embB_p.Met306Val	gyrA_p.Asp94Gly	rrs_r.1401a>g

R33922	rpoB_p.Ser450Leu	fabG1_c.-15C>T	pncA_c.459_466del	embB_p.Met306Val	gyrA_p.Ala90Val	rrs_r.1401a>g
R22572	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R9671	rpoB_p.Ser450Leu	ahpC_c.-48G>A, katG_p.Ser315Thr	pncA_p.Val139Met	embB_p.Met306Val, embA_c.-16C>T	gyrA_p.Asp94Gly, gyrA_p.Ser91Pro	rrs_r.1401a>g
R37378	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R6932	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R6702	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Gly88Cys	rrs_r.1401a>g
R21689	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R34001	rpoB_p.Ser450Leu	ahpC_c.-48G>A, katG_p.Ser315Thr	pncA_p.Val139Met	embB_p.Met306Val	gyrA_p.Ala90Val	rrs_r.1401a>g
R18174	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	-	embB_p.Met306Val	gyrA_p.Asp94Ala	rrs_r.1401a>g
R12052	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrB_p.Arg446Cys	rrs_r.1401a>g
R11687	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
X7	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R19048	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R4489	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R31891	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	pncA_p.Asp8Asn	embB_p.Met306Val	gyrA_p.Ser91Pro	rrs_r.1401a>g
R27080	rpoB_p.Ser450Leu, rpoC_p.Leu527Val	fabG1_c.-15C>T, inhA_p.Ile194Thr	-	embB_p.Met306Val	gyrA_p.Ala90Val	rrs_r.1401a>g
R22688	rpoB_p.Ser450Leu	katG_p.Ser315Thr	pncA_p.Asp8Asn	embB_p.Met306Val	gyrA_p.Asp94Gly	rrs_r.1401a>g
R14756	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	pncA_p.Val130Ala	embB_p.Met306Val	gyrA_p.Asp94Gly	rrs_r.1401a>g
R13166	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R15305	rpoB_p.Ser450Leu	fabG1_c.-15C>T	-	-	gyrA_p.Asp94Gly	rrs_r.1401a>g
R30446	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	pncA_p.Asp8Asn	embB_p.Met306Val	gyrA_p.Ser91Pro	rrs_r.1401a>g

R14489	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	pncA_p.Val130Ala	embB_p.Met306Val	gyrA_p.Asp94Gly	rrs_r.1401a>g
R21213	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	pncA_p.Asp8Asn	embB_p.Met306Val	gyrA_p.Asp94Gly	rrs_r.1401a>g
R29897	rpoB_p.Asp435Gly, rpoB_p.Leu452Pro	fabG1_c.-8T>A, katG_p.Ser315Thr	pncA_c.456_457insC	embB_p.Met306Val	gyrA_p.Ala90Val	rrs_r.1401a>g
R37359	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R34514	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Ala90Val	rrs_r.1401a>g
R10398	rpoB_p.Asp435Gly, rpoB_p.Leu452Pro	fabG1_c.-8T>A, katG_p.Ser315Thr	pncA_c.456_457insC	embB_p.Met306Val	gyrA_p.Ala90Val	rrs_r.1401a>g
R17281	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R2969	rpoB_p.Ser450Leu	fabG1_c.-15C>T	pncA_p.Tyr103*	embB_p.Asp354Ala	gyrA_p.Asp94Gly	rrs_r.1401a>g
X91	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94His	rrs_r.1401a>g
R28468	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Asn	rrs_r.1401a>g
R24211	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	pncA_p.Asp8Asn	embB_p.Met306Val	gyrA_p.Ser91Pro	rrs_r.1401a>g
R12425	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Ala90Val	rrs_r.1401a>g
R23866	rpoB_p.Ser450Leu	ahpC_c.-48G>A, katG_p.Ser315Thr	pncA_p.Val139Met	embB_p.Met306Val	gyrA_p.Asp94Asn	rrs_r.1401a>g
R13560	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R18028	rpoB_p.Ser450Leu	fabG1_c.-15C>T	-	embB_p.Met306Val	gyrA_p.Asp94Ala	rrs_r.1401a>g
R36404	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R22911	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R31329	rpoB_p.Ser450Leu	katG_p.Ser315Thr	pncA_p.Asp8Asn	embB_p.Met306Val	gyrA_p.Asp94Gly	rrs_r.1401a>g
X2	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	pncA_p.Asp8Asn	embB_p.Met306Val	gyrB_p.Glu501Asp	rrs_r.1401a>g
R19406	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R15282	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	pncA_p.Ser164Pro	embB_p.Met306Val	gyrA_p.Asp94Ala	rrs_r.1401a>g

R4221	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R27276	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Asn	rrs_r.1401a>g
R34184	rpoB_p.Ser450Leu	katG_p.Ser315Thr	pncA_p.Asp8Asn	embB_p.Met306Val	gyrA_p.Asp94Gly	rrs_r.1401a>g
R18700	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R20367	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	pncA_p.Thr135Pro	embB_p.Met306Val	gyrA_p.Ala90Val	rrs_r.1401a>g
R29551	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Ala90Val	rrs_r.1401a>g
R17581	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Ala90Val	rrs_r.1401a>g
R14876	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R35808	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R9951	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R6705	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R21826	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R26892	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R26725	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	-	embB_p.Met306Val	gyrA_p.Asp94Ala	rrs_r.1401a>g
R33910	rpoB_p.Ser450Leu	fabG1_c.-15C>T	-	-	gyrA_p.Ala90Val	rrs_r.1401a>g
R9941	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Asn	rrs_r.1401a>g
R8666	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	-	embB_p.Gln497Arg	gyrA_p.Ala90Val	rrs_r.1401a>g
R31630	rpoB_p.Ser450Leu	fabG1_c.-8T>A, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile, embB_p.Gln497Lys	gyrA_p.Ser91Pro	rrs_r.1401a>g
R19882	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Asn	rrs_r.1401a>g
R20189	rpoB_p.Ser450Leu	fabG1_c.-15C>T	pncA_p.Tyr103*	-	gyrA_p.Ser91Pro	rrs_r.1401a>g
R29125	rpoB_p.Ser450Leu	katG_p.Ser315Thr	pncA_p.Asp8Asn	embB_p.Met306Val	gyrA_p.Asp94Gly	rrs_r.1401a>g

R20073	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R18369	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Ala90Val	rrs_r.1401a>g
R22740	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Ser91Pro	rrs_r.1401a>g
R29799	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R39462	rpoB_p.Asp435Tyr	katG_p.Ser315Thr	-	embA_c.-16C>G	gyrA_p.Asp94Gly	rrs_r.1401a>g
R38964	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R4933	rpoB_p.Ser450Leu	katG_p.Ser315Thr	pncA_p.Asp8Asn	embB_p.Met306Val	gyrA_p.Asp94Gly	rrs_r.1401a>g
R28700	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Asn	rrs_r.1401a>g
R17064	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Tyr	rrs_r.1401a>g
R4203	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R33225	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Tyr	rrs_r.1401a>g
R23197	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Gly88Cys	rrs_r.1401a>g
R20964	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Asn	rrs_r.1401a>g
R27500	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	pncA_p.Val130Ala	embB_p.Met306Val	gyrA_p.Asp94Gly	rrs_r.1401a>g
R17084	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Tyr	rrs_r.1401a>g
R18920	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Ala	rrs_r.1401a>g
R27303	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	pncA_p.Asp8Asn	embB_p.Met306Val	gyrA_p.Ser91Pro	rrs_r.1401a>g
R17203	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Gly88Cys	rrs_r.1401a>g
R31399	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Ala90Val	rrs_r.1401a>g
R5620	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Ala90Val	rrs_r.1401a>g
X39	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Ala	rrs_r.1401a>g

R19628	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R9673	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Gly88Cys	rrs_r.1401a>g
R19005	rpoB_p.Ser450Leu	katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile, embB_p.Gln497Lys	gyrA_p.Ala90Val	rrs_r.1401a>g
R15310	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Ala	rrs_r.1401a>g
R12550	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R16923	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R28945	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R5317	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R32291	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R19725	rpoB_p.Asp435Gly, rpoB_p.Leu452Pro	fabG1_c.-8T>A, katG_p.Ser315Thr	pncA_c.456_457insC	embB_p.Met306Val	gyrA_p.Ala90Val	rrs_r.1401a>g
R9126	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Asn	rrs_r.1401a>g
R15957	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Ala90Val	rrs_r.1401a>g
R4524	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R9779	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	pncA_p.Thr135Pro	embB_p.Met306Val	gyrA_p.Ala90Val	rrs_r.1401a>g
R33494	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R17926	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Tyr	rrs_r.1401a>g
R33916	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R13673	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Asn	rrs_r.1401a>g
R27501	rpoB_p.Ser450Leu, rpoC_p.Phe452Ser	ahpC_c.-48G>A, katG_p.Ser315Thr	pncA_p.Val139Met	embB_p.Met306Val	gyrB_p.Asp461His	rrs_r.1401a>g
R16826	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g

R32208	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Tyr	rrs_r.1401a>g
X49	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Asn	rrs_r.1401a>g
R9047	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R10854	rpoB_p.Ser450Leu, rpoC_p.Leu527Val	fabG1_c.-15C>T, inhA_p.Ile194Thr	-	embB_p.Met306Val	gyrA_p.Ala90Val	rrs_r.1401a>g
R23384	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R17830	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Tyr	rrs_r.1401a>g
R26851	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R23853	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Ala90Val	rrs_r.1401a>g
R19816	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R3889	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrB_p.Asp461Asn	rrs_r.1401a>g
R16323	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	-	embB_p.Met306Val	gyrA_p.Asp94Ala	rrs_r.1401a>g
R38942	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	-	embB_p.Met306Val	gyrA_p.Asp94Ala	rrs_r.1401a>g
R15100	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	-	embB_p.Met306Val	gyrA_p.Asp94Gly	rrs_r.1401a>g
R33918	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R23604	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	pncA_p.Val130Ala	embB_p.Met306Val	gyrA_p.Asp94Gly	rrs_r.1401a>g
R33917	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R5166	rpoB_p.Ser450Leu	fabG1_c.-15C>T	-	embB_p.Met306Val	gyrB_p.Arg446Cys	rrs_r.1401a>g
R15539	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Asn	rrs_r.1401a>g
X173	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrB_p.Asp461Asn	rrs_r.1401a>g
R27650	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrB_p.Asp461Asn, gyrA_p.Ala90Val	rrs_r.1401a>g

R9345	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	-	embB_p.Met306Val	gyrA_p.Asp94Ala	rrs_r.1401a>g
R13923	rpoB_p.His445Asp	fabG1_c.-15C>T	pncA_p.Thr142Met	-	gyrA_p.Asp94Asn	-
R19726	rpoB_p.Asp435Gly, rpoB_p.Leu452Pro	fabG1_c.-8T>A, katG_p.Ser315Thr	pncA_c.456_457insC	embB_p.Met306Val	gyrA_p.Ala90Val	rrs_r.1401a>g
R23839	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R19512	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Tyr	rrs_r.1401a>g
R3983	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R30937	rpoB_p.Ser450Leu	katG_p.Ser315Thr	pncA_p.Asp8Asn	embB_p.Met306Val	gyrA_p.Asp94Ala	-
X50	rpoB_p.Ser450Leu	fabG1_c.-8T>A, katG_p.Ser315Thr	pncA_c.389_389del	embB_p.Asp1024Asn, embB_p.Met306Ile	gyrA_p.Asp94Asn	rrs_r.1401a>g
R35856	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R22123	rpoB_p.Ser450Leu	katG_p.Ser315Thr	pncA_p.Asp8Asn	embB_p.Met306Val	gyrA_p.Asp94Gly	rrs_r.1401a>g
R11044	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Ala90Val	rrs_r.1401a>g
R22654	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	pncA_p.Val130Ala	embB_p.Met306Val	gyrA_p.Asp94Gly	rrs_r.1401a>g
R15167	rpoB_p.Asp435Tyr	katG_p.Ser315Thr	-	embA_c.-16C>G	gyrB_p.Glu501Asp	rrs_r.1401a>g
R18977	rpoB_p.Ser450Leu	fabG1_c.-15C>T	pncA_p.Tyr103*	embB_p.Met306Val	gyrA_p.Asp94Gly	-
R13121	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Ala	rrs_r.1401a>g
R15531	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Asn	rrs_r.1401a>g
R8194	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R17085	rpoB_p.Ser450Leu	fabG1_c.-15C>T	-	-	gyrB_p.Glu501Val	rrs_r.1401a>g
R32272	rpoB_p.Ser450Leu	fabG1_c.-15C>T	-	embB_p.Met306Val	gyrB_p.Arg446Cys	rrs_r.1401a>g
R39352	rpoB_p.Ser450Leu	katG_p.Ser315Thr	pncA_p.Asp8Asn	embB_p.Met306Val	gyrA_p.Asp94Gly	rrs_r.1401a>g
R38892	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Ala90Val	rrs_r.1401a>g
R28854	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	-	embB_p.Met306Val	gyrA_p.Asp94Ala	rrs_r.1401a>g

R22918	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	pncA_p.Ser164Pro	embB_p.Met306Val	gyrA_p.Ala90Val	rrs_r.1401a>g
R9191	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R4731	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94His	rrs_r.1401a>g
R13071	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R8247	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Ala90Val	rrs_r.1401a>g
R21103	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R3174	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R39195	rpoB_p.Ser450Leu	katG_p.Ser315Thr	pncA_p.Asp8Asn	embB_p.Met306Val	gyrA_p.Asp94Gly	rrs_r.1401a>g
R4164	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	-	embB_p.Met306Val	gyrA_p.Asp94Ala	rrs_r.1401a>g
R28518	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Ala	rrs_r.1402c>t
R20767	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R22116	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R7831	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R26863	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R6596	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Tyr	rrs_r.1401a>g
R30463	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
X154	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Asn	rrs_r.1401a>g
R28887	rpoB_p.Ser450Leu	ahpC_c.-48G>A, katG_p.Ser315Thr	pncA_p.Val139Met	embB_p.Met306Val, embA_c.-16C>T	gyrA_p.Ala90Val	rrs_r.1401a>g
R30207	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R20321	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Asn	rrs_r.1401a>g

R33926	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Gly88Cys	rrs_r.1401a>g
R11138	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	pncA_p.Thr135Pro	embB_p.Met306Val	gyrA_p.Ala90Val	rrs_r.1401a>g
R19351	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Asn	rrs_r.1401a>g
R10319	rpoB_p.Ser450Leu	katG_p.Ser315Thr	-	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R11493	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R33804	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R14766	rpoB_p.Ser450Leu	fabG1_c.-15C>T	-	embB_p.Met306Val	gyrA_p.Ala90Val, gyrB_p.Arg446Cys	rrs_r.1401a>g
R26787	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	pncA_p.Val130Ala	embB_p.Met306Val	gyrA_p.Asp94Gly	rrs_r.1401a>g
R7110	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R38747	rpoB_p.Ser450Leu	ahpC_c.-48G>A, katG_p.Ser315Thr	pncA_p.Val139Met	embB_p.Met306Val	gyrA_p.Ala90Val	-
R18774	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Gly88Cys	rrs_r.1401a>g
R33904	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	pncA_p.Thr135Pro	embB_p.Met306Val	gyrA_p.Ala90Val	rrs_r.1401a>g
R30454	rpoB_p.Gln432Lys	katG_p.Ser315Thr	-	embB_p.Met306Val	gyrA_p.Asp94Asn	rrs_r.1401a>g
R5065	rpoB_p.Ser450Leu	katG_p.Ser315Thr	pncA_p.Gly132Ala	embB_p.Met306Val	gyrA_p.Asp94Gly	rrs_r.1401a>g
R31853	rpoB_p.Ser450Leu	katG_p.Ser315Thr, inhA_p.Ser94Ala	pncA_p.Gln10Pro	embB_p.Gln497Arg	gyrA_p.Asp94Gly	-
R8407	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R6809	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	-	embB_p.Met306Val	gyrA_p.Asp94Ala	rrs_r.1401a>g
R6196	rpoB_p.Ser450Leu	fabG1_c.-15C>T	-	-	gyrA_p.Asp94Asn	rrs_r.1401a>g
R22034	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R5318	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	-	embB_p.Met306Val	gyrA_p.Asp94Ala	rrs_r.1401a>g
R14852	rpoB_p.Ser450Leu	katG_p.Ser315Thr	pncA_p.Val139Gly	embB_p.Met306Val	gyrA_p.Asp94Gly	rrs_r.1401a>g

R21754	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R39730	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Ser91Pro, gyrA_p.Ala90Val	rrs_r.1401a>g
R33486	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Asn	rrs_r.1401a>g
R19404	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R18343	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	-	embB_p.Met306Val	gyrA_p.Asp94Ala	rrs_r.1401a>g
R9157	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Asn	rrs_r.1401a>g
R38779	rpoB_p.Ser450Leu	fabG1_c.-15C>T	pncA_p.Tyr103*	embB_p.Met306Ile	gyrA_p.Asp94Gly	-
R29391	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrB_p.Asn499Asp	rrs_r.1401a>g
R23538	rpoB_p.Ser450Leu	katG_p.Ser315Thr	pncA_p.Asp8Asn	embB_p.Met306Val	gyrA_p.Asp94Gly	rrs_r.1401a>g
R5258	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R27406	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R19077	rpoB_p.Ser450Leu	katG_p.Ser315Thr	-	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R10637	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	pncA_p.Ser164Pro	embB_p.Met306Val	gyrA_p.Ala90Val	rrs_r.1401a>g
R24418	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Ala90Val	rrs_r.1401a>g
R11709	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R36112	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Asn	rrs_r.1401a>g
R32278	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R3731	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Asn	rrs_r.1401a>g
R4704	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Ala90Val	rrs_r.1401a>g
R5252	rpoB_p.Ser450Leu	fabG1_c.-15C>T, ahpC_c.-48G>A	pncA_p.Tyr103*	embB_p.Asp354Ala	gyrA_p.Asp94Gly	rrs_r.1401a>g
R12533	rpoB_p.Ser450Leu, rpoC_p.Phe452Ser	ahpC_c.-48G>A, katG_p.Ser315Thr	pncA_p.Val139Met	embB_p.Met306Val	gyrB_p.Asp461His	rrs_r.1401a>g

R18143	rpoB_p.Ser450Leu	katG_p.Ser315Thr	pncA_p.Met175Val	embB_p.Met306Val	gyrA_p.Ser91Pro	-
R26831	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R24412	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Ala90Val	rrs_r.1401a>g
R22901	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	-	embB_p.Met306Val	gyrA_p.Asp94Ala	rrs_r.1401a>g
R16000	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Ala90Val	rrs_r.1401a>g
R3974	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	-	embB_p.Met306Val	gyrA_p.Asp94Ala	rrs_r.1401a>g
R29794	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
X85	rpoB_p.Ser450Leu	katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile, embB_p.Gln497Lys	gyrA_p.Ala90Val	rrs_r.1401a>g
R21236	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Asn	rrs_r.1401a>g
R28751	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R21413	rpoB_p.Ser450Leu	katG_p.Ser315Thr	pncA_p.Asp8Asn	embB_p.Met306Val	gyrA_p.Asp94Gly	rrs_r.1401a>g
R36725	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R32070	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R19290	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R20863	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	-	embB_p.Met306Val	gyrA_p.Asp94Ala	rrs_r.1401a>g
R17970	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Tyr	rrs_r.1401a>g
R36765	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrB_p.Glu501Asp	rrs_r.1401a>g
R38960	rpoB_p.Ser450Leu	ahpC_c.-48G>A, katG_p.Ser315Thr	pncA_p.Val139Met	embB_p.Met306Val	gyrA_p.Ser91Pro	rrs_r.1401a>g
R2850	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Gly88Cys	rrs_r.1401a>g
R19790	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Ser91Pro	rrs_r.1401a>g
R29690	rpoB_p.Ser450Leu	katG_p.Ser315Thr	pncA_p.Asp8Asn	embB_p.Met306Val	gyrA_p.Asp94Gly	rrs_r.1401a>g

R13449	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R4775	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R4817	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R36351	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrB_p.Glu501Asp	rrs_r.1401a>g
R30040	rpoB_p.Ser450Leu	fabG1_c.-15C>T	pncA_p.Tyr103*	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R19631	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Ala90Val	rrs_r.1401a>g
R11158	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Ala90Val	rrs_r.1401a>g
R17597	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Asn	rrs_r.1401a>g
R17009	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Gly88Cys	rrs_r.1401a>g
R9362	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	pncA_p.Ser164Pro	embB_p.Met306Val	gyrA_p.Asp94Asn	rrs_r.1401a>g
R20890	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Asn	rrs_r.1401a>g
R11689	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Gly88Cys	rrs_r.1401a>g
R15797	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R5954	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R27917	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R23874	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	-	embB_p.Met306Val	gyrA_p.Asp94Ala, gyrA_p.Ala90Val	rrs_r.1401a>g
R3833	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
X66	rpoB_p.Ser450Leu	fabG1_c.-15C>T	-	embB_p.Met306Val	gyrB_p.Arg446Cys	rrs_r.1401a>g
R9950	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	-	embB_p.Met306Val	gyrA_p.Asp94Ala	rrs_r.1401a>g
R2967	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R38180	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Gly88Cys	rrs_r.1401a>g

R21092	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Ala90Val	rrs_r.1401a>g
R5066	rpoB_p.Ser450Leu	fabG1_c.-15C>T	pncA_p.Tyr103*	embB_p.Gly406Asp	gyrA_p.Asp94Gly	rrs_r.1401a>g
X198	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	-	embB_p.Met306Val	gyrA_p.Asp94Ala	rrs_r.1401a>g
R18045	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R33791	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R38864	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Asn	rrs_r.1401a>g
R13332	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Tyr	rrs_r.1401a>g
R11502	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Gly88Cys	rrs_r.1401a>g
R33316	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Ser164Pro	embB_p.Met306Ile	gyrA_p.Asp94Ala	rrs_r.1401a>g
R33506	rpoB_p.Ser450Leu	fabG1_c.-15C>T	pncA_c.135_136insTG	embB_p.Gly406Asp	gyrA_p.Asp94Gly	rrs_r.1401a>g
R5490	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R32275	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R33997	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Ala	rrs_r.1401a>g
R18285	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R16462	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	-	embB_p.Met306Val	gyrA_p.Asp94Ala	rrs_r.1401a>g
R10212	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R29538	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R8081	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Gly88Cys	rrs_r.1401a>g
R4577	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R15871	rpoB_p.Ser450Leu	fabG1_c.-15C>T	-	-	gyrA_p.Asp94Gly	rrs_r.1401a>g
R27659	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Ala90Val	rrs_r.1401a>g

R36851	rpoB_p.Ser450Leu	fabG1_c.-15C>T	pncA_p.Tyr103*	embB_p.Met306Val	gyrA_p.Asp94Asn	rrs_r.1401a>g
R10877	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Ala	rrs_r.1401a>g
R4312	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R17303	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Asn	rrs_r.1401a>g
R3960	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R24216	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	-	embB_p.Met306Val	gyrA_p.Asp94Ala	rrs_r.1401a>g
R4825	rpoB_p.Ser450Leu	katG_p.Ser315Thr	pncA_p.Asp8Asn	embB_p.Met306Val	gyrA_p.Asp94Gly	rrs_r.1401a>g
R27605	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Asn	rrs_r.1401a>g
R33925	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R20825	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Asn	rrs_r.1401a>g
R27920	rpoB_p.His445Arg, rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	pncA_p.Asp8Asn	embB_p.Met306Val	gyrA_p.Asp94Gly	rrs_r.1401a>g
R4488	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R11380	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Ala	rrs_r.1401a>g
R3093	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Gly88Cys	rrs_r.1401a>g
R18647	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R34532	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Ala90Val	rrs_r.1401a>g
R35862	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R15416	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp89Asn	rrs_r.1401a>g
X38	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R35206	rpoB_p.His445Asn	katG_p.Ser315Thr	-	embB_p.Met306Val	gyrA_p.Asp94Gly	rrs_r.1401a>g
R3269	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Asn	rrs_r.1401a>g

R6254	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R38593	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Ala	rrs_r.1401a>g
R3415	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R27725	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R32158	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	-	embB_p.Met306Val	gyrA_p.Asp94Ala	rrs_r.1401a>g
R19442	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R17246	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Ser91Pro	rrs_r.1401a>g
R37936	rpoB_p.Ser450Leu	ahpC_c.-48G>A, katG_p.Ser315Thr	pncA_p.Val139Met	embB_p.Met306Val	gyrA_p.Ala90Val	rrs_r.1401a>g
R5354	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R33511	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R27658	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Ala90Val	rrs_r.1401a>g
R29503	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	pncA_p.Asp8Asn	embB_p.Met306Val	gyrB_p.Glu501Asp	rrs_r.1401a>g
R18095	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	pncA_p.Asp8Asn	embB_p.Met306Val	gyrA_p.Asp94Gly	rrs_r.1401a>g
R37955	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Asn	rrs_r.1401a>g
R22895	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Asn	rrs_r.1401a>g
R31314	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
R30466	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
X3	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Gly	rrs_r.1401a>g
X186	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Ala	rrs_r.1401a>g
R2841	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Ala	rrs_r.1401a>g

R15141	rpoB_p.Ser450Leu	katG_p.Ser315Thr	pncA_p.Asp8Asn	embB_p.Met306Val	gyrA_p.Asp94Gly	rrs_r.1401a>g
R13587	rpoB_p.Ser450Leu, rpoC_p.Leu527Val	fabG1_c.-15C>T, inhA_p.Ile194Thr	-	embB_p.Met306Val	gyrA_p.Ala90Val	rrs_r.1401a>g
R32055	rpoB_p.Ser450Leu	fabG1_c.-15C>T, katG_p.Ser315Thr	pncA_p.Cys14Arg	embB_p.Met306Ile	gyrA_p.Asp94Tyr	rrs_r.1401a>g
R24188	rpoB_p.Ser450Leu	fabG1_c.-15C>T, inhA_p.Ile194Thr	pncA_p.Asp8Asn	embB_p.Met306Val	gyrA_p.Asp94Asn	rrs_r.1401a>g
R19246	rpoB_p.Asp435Val	fabG1_c.-17G>T, katG_p.Ser315Thr	pncA_c.517_518insG	embB_p.Met306Ile	gyrA_p.Asp94Asn	rrs_r.1401a>g

^a INH = isoniazid, RIF = rifampicin, EMB = ethambutol, PZA = pyrazinamide, SLIDs = second line injectable drugs, FQs = fluoroquinolones, - = no mutation

Table S3 The number of clusters after merging on common ancestor using different SNP cutoffs

SNP cutoff	5	8	10	12	15
Clusters after merging on common ancestor (n)	12	11	12	10	10

Supplementary figures

Figure S1 Distribution of cluster sizes with 7 small clusters (2-4 isolates), 1 medium cluster (19 isolates), 3 large clusters (66, 70, 72 isolates) and 1 very large cluster (150 isolates).

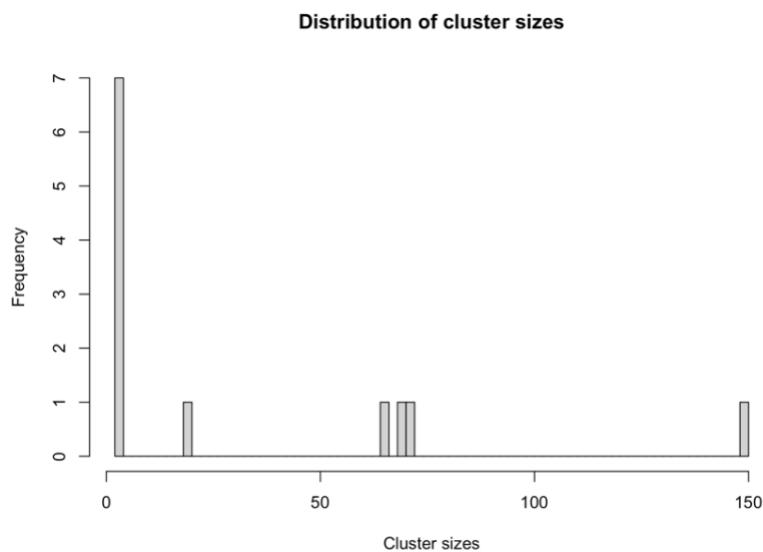


Figure S2 Phylogenetic tree annotated with lineage information (inner circle) and transmission cluster sizes (outer circle). All large clusters are part of lineage 2.2.1 and 2.2.2.

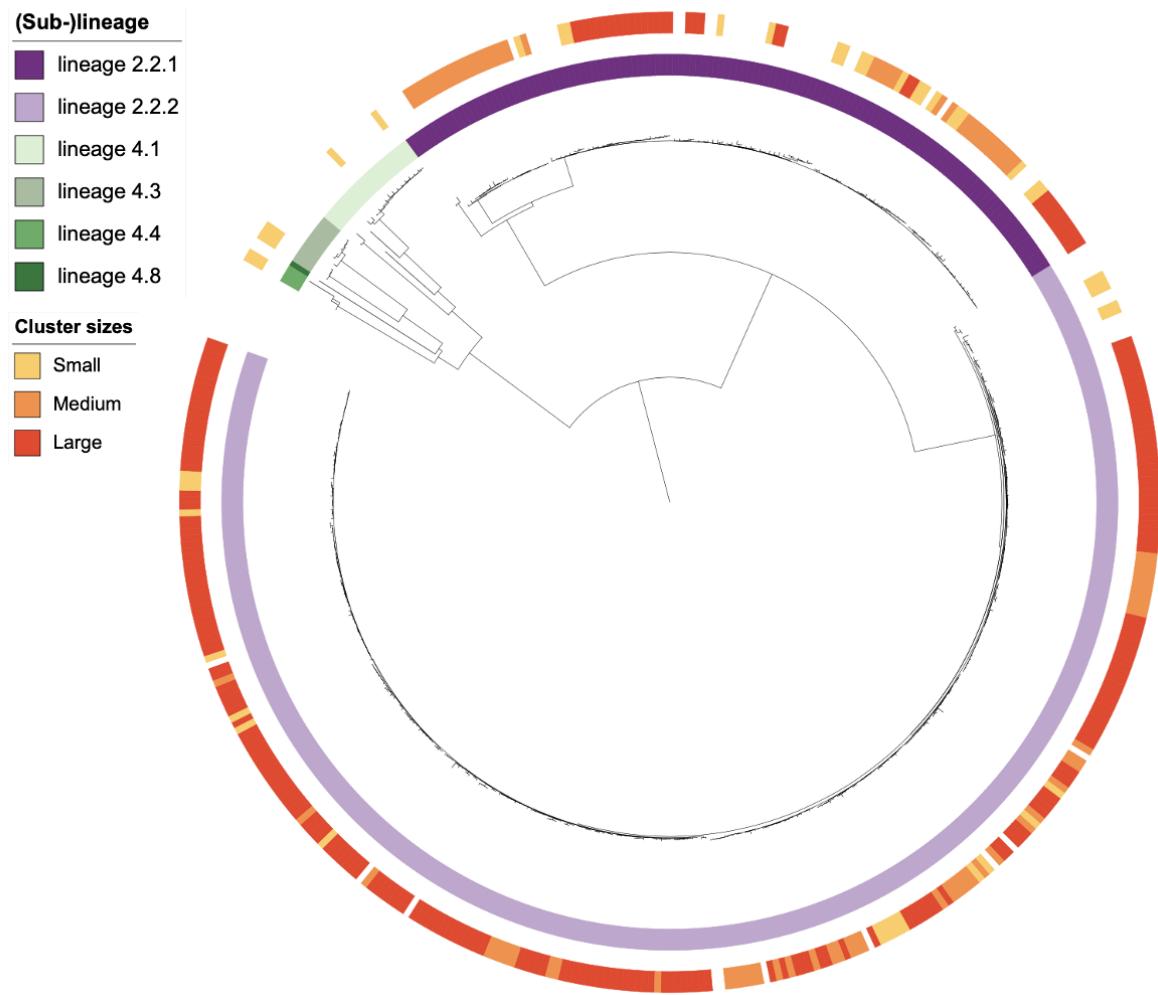


Figure S3a Distribution of gender (inner circle) and cluster sizes after inclusion of INH + RIF + EMB + PZA + FQs + SLIDs resistance conferring mutations (outer circle) across the phylogenetic tree of individual XDR-Mtb isolates collected from 461 patients in the Western Cape Province of South Africa, 2006-2017

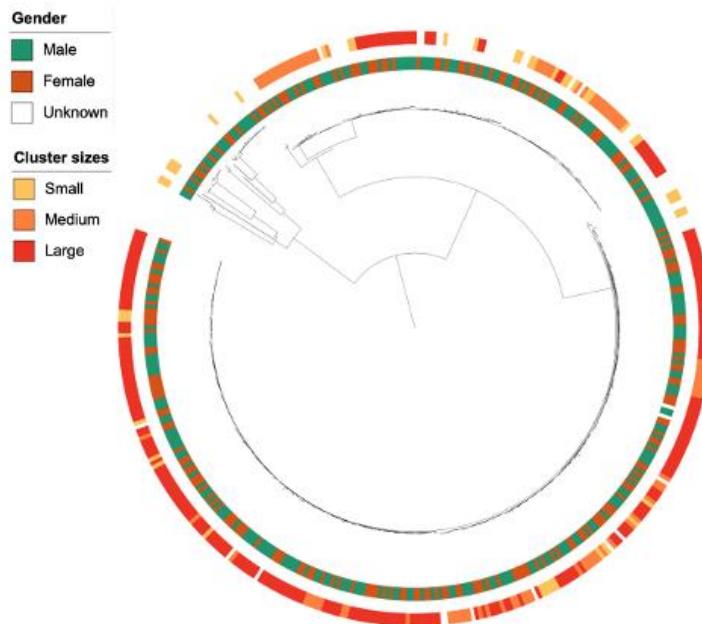


Fig S3bDistribution of age (inner circle) and cluster sizes after inclusion of INH + RIF + EMB + PZA + FQs + SLIDs resistance conferring mutations (outer circle) across the phylogenetic tree of individual XDR-Mtb isolates collected from 461 patients in the Western Cape Province of South Africa, 2006-2017

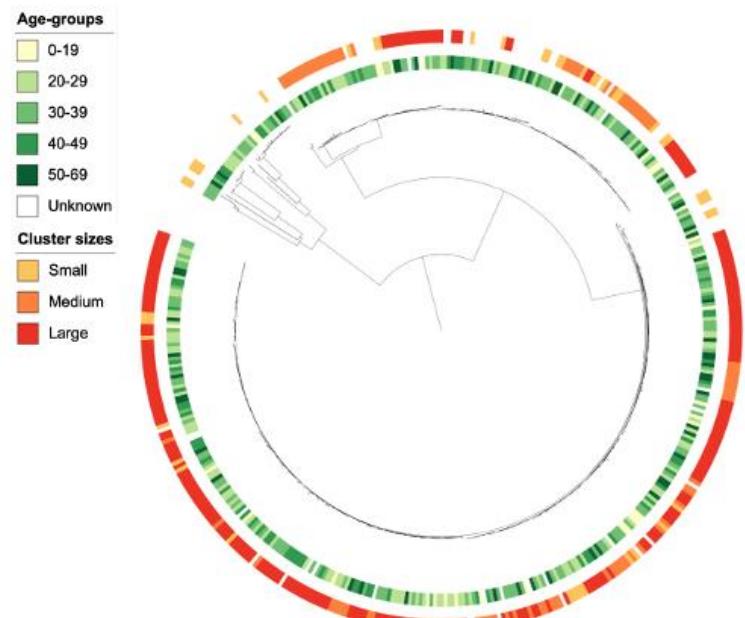


Fig S3c Distribution of district (urban vs. rural) (inner circle) and cluster sizes after inclusion of INH + RIF + EMB + PZA + SLIDs +FQs resistance conferring mutations (outer circle) across the phylogenetic tree of individual XDR-Mtb isolates collected from 461 patients in the Western Cape Province of South Africa, 2006-2017

