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### **Additional file 5: Mutations associated with antimicrobial resistance identified in one clinical sample (CT-36|E)**

<b>Gene</b>	<b>Nucleotide Position</b>	<b>Ref</b>	<b>Allele</b>	<b>Count</b>	<b>Coverage</b>	<b>Frequency (%)</b>	<b>Strand balance</b>	<b>Average quality</b>	<b>Coding region change</b>	<b>Amino acid change</b>
L22	154	G	A	11	11	100	0.4	38	154G>A	Gly52Ser
L22	193	C	T	17	18	94.44	0.5	38.35	193C>T	Arg65Cys
L22	230	T	C	13	14	92.86	0.5	38.14	230T>C	Val77Ala