Gene	Updated Gene Annotation	Description	Mutations _	No. Observed mutations	
				NL	PL
0099	SYNWH7803_RS00480	hypothetical protein	¹D237N (<u>G</u> AC→ <u>A</u> AC)	0	1
intergenic			² intergenic (+21/-38)	2	0
0102	SYNWH7803_RS00485	rfbA; glucose-1-phosphate thymidylyltransferase	^{3†} F135L (TT <u>C</u> →TT <u>A</u>); ⁴ P166P (CC <u>C</u> →CC <u>T</u>)	3	2
0103	SYNWH7803_RS00490	rfbC; dTDP-4-dehydrorhamnose 3,5-epimerase	⁵ Q12H (CAA→CA <u>I</u>); ⁶ E34K (<u>G</u> AG→AAG); ⁷ A95A (GC <u>C</u> →GC <u>I</u>); ⁸ R104C (<u>C</u> GC→ <u>I</u> GC)	0	4
0104	SYNWH7803_RS00495	rfbD; dTDP-4-keto-L-rhamnose reductase	⁹ Δ1 bp; ¹⁰ G157G (GG <mark>C</mark> →GG I); ¹¹ A158S (<u>G</u> CG→ I CG); ¹² N162I (A <u>A</u> C→A I C)	0	5
0105	SYNWH7803_RS00500	sugar transferase	¹³ W305* (TG <u>G</u> →TG <u>A</u>); ¹⁴ V312I (<u>G</u> TC→ <u>A</u> TC)	2	0
0140	SYNWH7803_RS00670	glycosyl transferase family protein	¹⁵ 2 bp→AC	0	1
0219	SYNWH7803_RS01065	homology to GtrA, YdjC-family	¹⁶ L340P (C <u>I</u> T→C <u>C</u> T); ¹⁷ H275N (<u>C</u> AC→ <u>A</u> AC) ; ¹⁸ E102* (<u>G</u> AA→ <u>I</u> AA); ¹⁹ Y13* (TA <u>C</u> →TA <u>G</u>)	4	0
0232	SYNWH7803_RS01130	hypothetical protein/glcosyltransferase type A	²⁰ P58T (<mark>C</mark> CG→ <u>A</u> CG)	0	1
1386	SYNWH7803_RS06815	two-component system sensor histidine kinase	²¹ Δ1 bp; ²² G344* (\underline{G} GA \rightarrow \underline{T} GA); ²³ S341I (A \underline{G} T \rightarrow ATT); ²⁴ Δ1 bp	5	1