

Sequence analysis of pooled bacterial samples enables identification of strain variation in group A streptococcus

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Supplementary Table S1. List of the GAS strains used in this study and their respective *emm* types

<b><i>emm</i> type</b>	<b>Pool 1</b>	<b>Pool 2</b>	<b>Total</b>
emm1.0	5	5	10
emm1.10	3	3	6
emm1.24	1	0	1
emm1.45	1	1	2
emm1.50	1	0	1
emm11.0	2	2	4
emm119.1	2	2	4
emm12.0	5	4	9
emm2.0	2	3	5
emm22.0	1	1	2
emm22.3	1	1	2
emm28.0	4	5	9
emm6.4	1	0	1
emm73.0	2	3	5
emm75.0	3	4	7
emm76.3	1	1	2
emm77.0	2	3	5
emm78.3	2	2	4
emm84.0	1	1	2
emm212 (st75.0)	1	1	2
emm11.1	1	0	1
emm4.0	1	1	2
emm78.0	2	2	4
emm89.0	5	5	10
<b>Total</b>	<b>50</b>	<b>50</b>	<b>100</b>