

**Table S1**

<b>Desirable strain characteristics</b>	<b>GAS M75 611024</b>																																																		
<i>A definite but uncommon contemporary cause of symptomatic pharyngitis</i>		<ul style="list-style-type: none"> <li>• Isolated in 2011 from the throat of a child with pharyngitis in Melbourne(1)</li> <li>• Pre-existing immunity in adults is unknown (no correlate of protection)</li> <li>• <i>emm75</i> &lt;5% of global GAS isolates (all sites and settings)(2)</li> <li>• <i>emm75</i> strains in selected GAS pharyngitis series since 2000</li> </ul>																																																	
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<i>Should cause skin infection</i>	<ul style="list-style-type: none"> <li>• E pattern ‘generalist’ (throat &amp; skin infections)(17)</li> <li>• Cluster E6 - linked phylogenetically to D pattern skin isolates(17, 18)</li> </ul>																																																		
<i>An uncommon cause of invasive GAS disease and immunological sequelae</i>	<ul style="list-style-type: none"> <li>• <i>emm75</i> strains very rarely associated with ARF/RHD or APSGN(19-26)</li> <li>• <i>emm75</i> strains in selected invasive GAS series since 2000</li> </ul> <table border="1"> <thead> <tr> <th>Location</th><th>n</th><th><i>emm75</i></th></tr> </thead> <tbody> <tr><td>Portugal (27)</td><td>160</td><td>0</td></tr> <tr><td>Northern Territory, Australia (28)</td><td>82</td><td>0</td></tr> <tr><td>France (11)</td><td>125</td><td>0.8%</td></tr> <tr><td>Alberta, Canada (29)</td><td>3105</td><td>1%</td></tr> <tr><td>Finland (30)</td><td>1122</td><td>1.2%</td></tr> <tr><td>Kilifi, Kenya (31)</td><td>357</td><td>1.4%</td></tr> <tr><td>Denmark (5)</td><td>201</td><td>1.5%</td></tr> <tr><td>Ireland (32)</td><td>473</td><td>1.5%</td></tr> <tr><td>Fiji (33)</td><td>55</td><td>1.8%</td></tr> <tr><td>Sydney, Australia (34)</td><td>55</td><td>1.8%</td></tr> <tr><td>Scotland (35)</td><td>357</td><td>2%</td></tr> <tr><td>USA (36, 37)*</td><td>17,002</td><td>2.4%</td></tr> <tr><td>Norway (38)</td><td>756</td><td>2.1%</td></tr> <tr><td>Japan (10)</td><td>74</td><td>4.1%</td></tr> <tr><td>Thailand (39)</td><td>234</td><td>5.1%</td></tr> </tbody> </table>		Location	n	<i>emm75</i>	Portugal (27)	160	0	Northern Territory, Australia (28)	82	0	France (11)	125	0.8%	Alberta, Canada (29)	3105	1%	Finland (30)	1122	1.2%	Kilifi, Kenya (31)	357	1.4%	Denmark (5)	201	1.5%	Ireland (32)	473	1.5%	Fiji (33)	55	1.8%	Sydney, Australia (34)	55	1.8%	Scotland (35)	357	2%	USA (36, 37)*	17,002	2.4%	Norway (38)	756	2.1%	Japan (10)	74	4.1%	Thailand (39)	234	5.1%	
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<p style="text-align: center;">*CDC Active Bacterial Core surveillance 2000-2016 (Chris A. Van Beneden, Personal Communication, September 11<sup>th</sup>, 2018)</p>																																																			
<i>Should have predictable and limited virulence, and be suitable for use in animal models</i>	<ul style="list-style-type: none"> <li>• covR/S virulence regulator wildtype (non-mutant)</li> <li>• Does not bind plasminogen and fibrinogen(18)</li> <li>• <i>emm75</i> strains have been used in murine models of nasopharyngitis and invasive disease, and have been successfully transformed for bioluminescence(40)</li> </ul>																																																		
<i>Should have limited antibiotic resistance</i>	<ul style="list-style-type: none"> <li>• See main text</li> </ul>																																																		
<i>The challenge strain should possess a wide array of candidate vaccine antigens</i>	<ul style="list-style-type: none"> <li>• See main text</li> </ul>																																																		

ARF: acute rheumatic fever, APSGN: acute post-streptococcal glomerulonephritis, GAS: group A *Streptococcus*; RHD: rheumatic heart disease

## References for Supplemental Table S1

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