

**S13 Table. Number of peptide matches for S2 Figure, showing the species assignments using the Initial vs the Curated Databases**

PEPTIDE MATCHES DATABASE CURATED												
	<i>S. pneumoniae</i>			<i>S. pseudopneumoniae</i>			<i>S. mitis</i>			<i>S. pyogenes</i>		
	CCUG 28588T	CCUG 7206	CCUG 35180	CCUG 49455T	CCUG 63747	CCUG 62647	CCUG 31611T	CCUG 63687	CCUG 69183	CCUG 4207T	CCUG 25570	CCUG 47803
<i>S. pneumoniae</i>	220	175	214									
<i>S. pseudopneumoniae</i>				433	245	250						
<i>S. mitis</i>						7	272	287	500			
<i>S. pyogenes</i>										314	351	418
<i>S. suis</i>	7								6			

PEPTIDE MATCHES DATABASE INITIAL												
	<i>S. pneumoniae</i>			<i>S. pseudopneumoniae</i>			<i>S. mitis</i>			<i>S. pyogenes</i>		
	CCUG 28588T	CCUG 7206	CCUG 35180	CCUG 49455T	CCUG 63747	CCUG 62647	CCUG 31611T	CCUG 63687	CCUG 69183	CCUG 4207T	CCUG 25570	CCUG 47803
<i>S. pneumoniae</i>	347	286	380	74	26	30	34	21	35			
<i>S. pseudopneumoniae</i>				230	183	189	21	17	37			
<i>S. mitis</i>				18		11	272	144	134			
<i>S. pyogenes</i>										319	360	427
<i>S. suis</i>	7								6			
<i>S. oralis</i>				7	5		16	12				
<i>S. parasanguinis</i>					18							
<i>B. proteoclasticus</i>							34	7	15			