### **Supplemental Figures**

# Differential Binding of Human and Murine IgGs to Catalytic and Cell Wall Binding Domains of *Staphylococcus aureus* Peptidoglycan Hydrolases

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В

S-CHAP	230	YTWGQCTYHVFNRRAEIGKGISTYWWNANNWDNAAAADGYTIDNRPTVGSIAQTDVG	286
A-CHAP	508	YPHGQCTWYVYNRMKQFGTSISGDLGDAHNWNNRAQYRDYQVSHTPKRHAAVVFEAGQFG	567
S-CHAP	287	YYGHVMFVERVNNDGSILVSEMNYSAAPGILTYRTVPAYQVNNYRYI 333	
A-CHAP	568	ADQHYGHVAFVEKVNSDGSIVISESNVKGL-GIISHRTINAAAAEELSYI 616	
LytM	184	GHAKDASWLTSRKQLQPYGQYHGGGAHYGVDYAMPENSPVYSLTDGTVVQAGWSNYG	240
LytU	47	GHSKWEDFFRGSRITETFGKYQHSPFDGKHYGIDFALPKGTPIKAPTNGKVTRI-FNNEL	105
LytM	241	GGNQVTIKEANSNNYQWYMHNNRLTVSAGDKVKAGDQIAYSGSTGN-STAPHVHFQRMSG	299
LytU	106	GGKVLQIAEDNGEYHQWYLHLDKYNVKVGDRVKAGDIIAYSGNTGKQTTGAHLHFQRMKG	165
LytM	300	GIGNQYAVDPTSYLQ 314	
LytU	166	G+GN YA DP +++ GVGNAYAEDPKPFIE 180	

#### Supplemental Figure 1. Alignment of homologous domains of Sle1 and Aly.

(A) Alignment of the S-CHAP and A-CHAP domains of Sle1 and Aly, respectively. (B) Alignment of the C-terminal catalytic M23 domains of LytM and LytU. Identical residues are indicated between the aligned sequences and conservative replacements are indicated with +.

## Supplemental Figure 2

A



B



С

























F

#### Supplemental Figure 2. Original gels and Western blots

Panel A shows the full Simply Blue-stained gels of Figure 2 (left panels). Panel B shows the full blots used to generate the Western blots presented in Figure 2 (right panels). Panel C shows the zymograms used to generate Figure 3. Panel D shows the full blots used to generate the Western blots presented in Figure 6.

Panel E shows the full blots used to generate the Western blots presented in Figure 7.

Panel F shows the full blots used to generate the Western blots presented in Figure 8.

All image sections presented in the respective Figures in the main manuscript have been marked with interrupted lines.