

## Supporting information

Identification of a Staphylococcal Complement Inhibitor with broad host specificity in equid *S. aureus* strains

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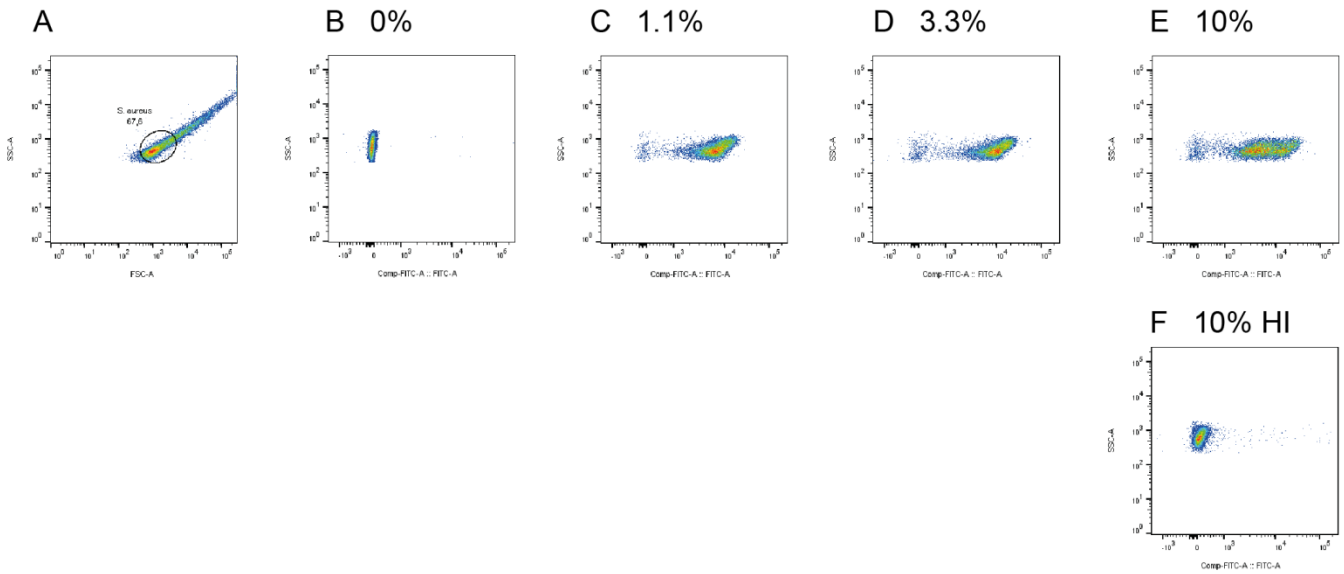
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**Running title:** Identification of staphylococcal equine SCIN

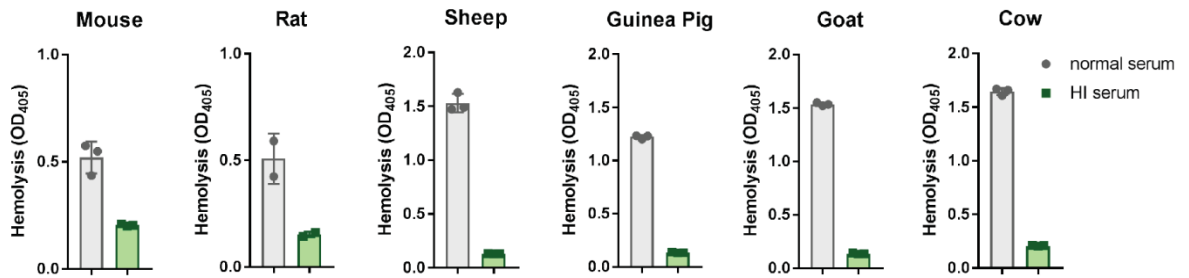
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**Keywords:** Complement system, Equine, Host adaptation, Immune evasion, SCIN

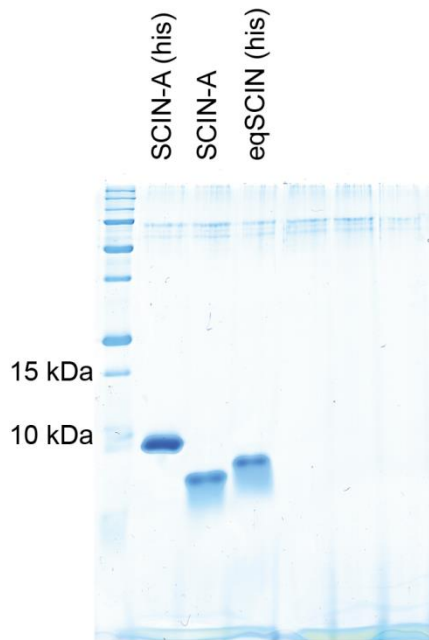
This file contains one Supplemental Table, as well as three Supplemental Figures and their legends



**Figure S1. Representative flow plots for C3b deposition with equine serum.** (A) Representation of gating *S. aureus* Wood in each well. (B-E) Detection of surface-bound C3b with anti-C3 conjugated with FITC and measured by flow cytometry analysis as geometric mean fluorescence of the gated bacteria after addition of various concentrations of equine serum. (F) 10% HI serum does show any C3b deposition, by the lack of FITC signal.



**Figure S2. HI serum lacks ability to lyse erythrocytes compared to normal serum.** The serum of the different animals used in Figure 5D were heat-inactivated and tested in the hemolysis assay. The complement in the sera of all animals lacked activity to lyse erythrocytes, while normal serum had active complement to lyse the erythrocytes.



**Figure S3. Protein gel with purified proteins used in this paper.** Purified SCIN-A without his tag (lane 1) was used in the phagocytosis assay, while the his-tagged protein was used for the C3b deposition and hemolysis assay (lane 2). The protein in lane 3 is eqSCIN with non-cleavable 6xHis tag. The concentration of the proteins is 150  $\mu\text{g/ml}$ .

**Supplementary table 1:** *scn-eq* is present in *Staphylococcus aureus* strains from different clonal complexes (CC) and different countries. For *scn-eq*, the gene location (prophage/chromosome), % identity to the reference (3711), single nucleotide polymorphisms (SNPs) and insertions (INS) are indicated. Furthermore, the presence of *lukPQ* and other *scns* are reported.

Strain ID	CC	Country	Host species	Location	<i>scn-eq</i>										<i>lukPQ</i>		Reference
					Identity (%)	SNPs							INS		<i>lukP</i>	<i>lukQ</i>	
						75 <sup>ns</sup>	103 <sup>ns</sup>	157 <sup>ns</sup>	189 <sup>s</sup>	198 <sup>ns</sup>	254 <sup>ns</sup>	318 <sup>s</sup>	87				
KM595-06	1	Switzerland	Horse	Prophage	98,261	G	A	G	A	C	G	T			+	+	Sieber <i>et al.</i> 2011 (34)
KM777-07	1	Switzerland	Horse	Prophage	98,261	G	A	G	A	C	G	T			+	+	Sieber <i>et al.</i> 2011 (34)
KM489-05	1	Switzerland	Horse	Prophage	98,261	G	A	G	A	C	G	T			+	+	Sieber <i>et al.</i> 2011 (34)
m-25-21 2.3	1	United Kingdom	Cow	Prophage	98,261	G	A	G	A	C	G	T			+	+	Koop <i>et al.</i> 2017 (14)
m-25-21 2.11	1	United Kingdom	Cow	Prophage	98,261	G	A	G	A	C	G	T			+	+	Koop <i>et al.</i> 2017 (14)
8205	1	United Kingdom	Horse	Prophage	98,261	G	A	G	A	C	G	T			+	+	Koop <i>et al.</i> 2017 (14)
1928	1	United Kingdom	Horse	Prophage	98,261	G	A	G	A	C	G	T			+	+	Koop <i>et al.</i> 2017 (14)
8231	1	United Kingdom	Horse	Prophage	98,261	G	A	G	A	C	G	T			+	+	Koop <i>et al.</i> 2017 (14)
8182	1	United Kingdom	Horse	Prophage	98,261	G	A	G	A	C	G	T			+	+	Koop <i>et al.</i> 2017 (14)
VetBz55B	133	Brazil	Buffalo	Chromosome	100	T	G	A	T	G	A	T			+	+	Aires-de-Sousa <i>et al.</i> 2007 (37)
VetBz63	133	Brazil	Buffalo	Chromosome	100	T	G	A	T	G	A	T			+	+	Aires-de-Sousa <i>et al.</i> 2007 (37)
c3388	133	Tunisia	Donkey	Prophage	100	T	G	A	T	G	A	T			+	+	Gharsa <i>et al.</i> 2012 (35)
c3401	133	Tunisia	Donkey	Prophage	100	T	G	A	T	G	A	T			+	+	Gharsa <i>et al.</i> 2012 (35)
c3403	133	Tunisia	Donkey	Prophage	100	T	G	A	T	G	A	T			+	+	Gharsa <i>et al.</i> 2012 (35)
c4439	133	Tunisia	Donkey	Prophage	100	T	G	A	T	G	A	T			+	+	Gharsa <i>et al.</i> 2012 (35)
c4444	133	Tunisia	Donkey	Prophage	100	T	G	A	T	G	A	T			+	+	Gharsa <i>et al.</i> 2012 (35)
c4445	133	Tunisia	Donkey	Prophage	100	T	G	A	T	G	A	T			+	+	Gharsa <i>et al.</i> 2012 (35)
c4451	133	Tunisia	Donkey	Prophage	100	T	G	A	T	G	A	T			+	+	Gharsa <i>et al.</i> 2012 (35)
c4452	133	Tunisia	Donkey	Prophage	100	T	G	A	T	G	A	T			+	+	Gharsa <i>et al.</i> 2012 (35)
c3815	133	Tunisia	Goat	Prophage	100	T	G	A	T	G	A	T			+	+	Gharsa <i>et al.</i> 2012 (35)
7107	133	United Kingdom	Horse	Prophage	100	T	G	A	T	G	A	T			+	+	Koop <i>et al.</i> 2017 (14)
7540	133	United Kingdom	Horse	Prophage	100	T	G	A	T	G	A	T			+	+	Koop <i>et al.</i> 2017 (14)
5431	133	United Kingdom	Horse	Prophage	100	T	G	A	T	G	A	T			+	+	Koop <i>et al.</i> 2017 (14)
8401	133	United Kingdom	Horse	Prophage	100	T	G	A	T	G	A	T			+	+	Koop <i>et al.</i> 2017 (14)
3711	133	United Kingdom	Horse	Prophage	100	T	G	A	T	G	A	T			+	+	Koop <i>et al.</i> 2017 (14)
8572	350	United Kingdom	Horse	Prophage	98,261	G	A	G	A	C	G	T			+	+	Koop <i>et al.</i> 2017 (14)
3507	522	United Kingdom	Horse	Prophage	99,42	T	G	A	A	G	A	C			+	+	Koop <i>et al.</i> 2017 (14)
8571	522	United Kingdom	Horse	Prophage	98,261	G	A	G	A	C	G	T			+	+	Koop <i>et al.</i> 2017 (14)
KM1549-2-06	1660	Switzerland	Horse	Prophage	95,763	G	A	G	A	C	G	T	AAAGC		+	+	Sieber <i>et al.</i> 2011 (34)

<sup>ns</sup> = non-synonymous

<sup>s</sup> = synonymous