

# IgG Endopeptidase SeMac does not Inhibit Opsonophagocytosis of *Streptococcus equi* Subspecies *equi* by Horse Polymorphonuclear Leukocytes

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**Abstract:** The secreted Mac protein made by group A *Streptococcus* (GAS) inhibits opsonophagocytosis of GAS by human polymorphonuclear leukocytes (PMNs). This protein also has the endopeptidase activity against human immunoglobulin G (IgG), and the Cys94, His262 and Asp284 are critical for the enzymatic activity. The horse pathogen *Streptococcus equi* subspecies *equi* produces a homologue of Mac (SeMac). SeMac was characterized to determine whether SeMac has IgG endopeptidase activity and inhibits opsonophagocytosis of *S. equi* by horse PMNs. The gene was cloned and recombinant SeMac was overexpressed in *Escherichia coli* and purified to homogeneity. Mice with experimental *S. equi* infection and horses with strangles caused by *S. equi* seroconverted to SeMac, indicating that SeMac is produced *in vivo* during infection. SeMac has endopeptidase activity against human IgG. However, the protein just cleaves a small fraction, which may be IgG1 only, of horse IgG. Replacement of Cys102 with Ser or His272 with Ala abolishes the enzymatic activity of SeMac, and the Asp294Ala mutation greatly decreases the enzymatic activity. SeMac does not inhibit opsonophagocytosis of *S. equi* by horse PMNs but opsonophagocytosis of GAS by human PMNs. Thus, SeMac is a cysteine endopeptidase with a limited activity against horse IgG and must have other function.

**Keywords:** IgG, Endopeptidase SeMac, opsonophagocytosis, polymorphonuclear leukocytes.

## INTRODUCTION

Gram-positive bacterium *Streptococcus equi* subspecies *equi* (*S. equi*) causes equine strangles, a highly contagious purulent lymphadenitis and one of the most common infectious diseases in horses [1,2]. The infection initially causes nasal discharge and fever and, then, leads to abscess formation in local lymph nodes, causing enormous pain and respiratory difficulty. There is massive infiltration of polymorphonuclear leukocytes (PMNs) to the infection site. However, *S. equi* effectively evades the horse innate immunity by being resistant to phagocytosis by PMNs. Horses recovered from strangles acquire immunity against *S. equi* reinfection [3]. The immunity is primarily mediated by protective antibodies [4], which opsonize and thus enhance phagocytosis of *S. equi* by horse PMNs.

To survive in hosts, bacterial pathogens have evolved multiple mechanisms to evade host defense. For examples, both *S. equi* and group A *Streptococcus* (GAS) produce the hyaluronic acid capsule and surface protein M protein to contribute to resistance to phagocytosis by PMNs. We found that GAS produces a secreted Mac protein (also known as IdeE), which inhibits opsonophagocytosis of GAS by human PMNs [5]. This protein can cleave the heavy chain of human immunoglobulin G (IgG) using Cys94, His262 and Asp284 as its catalytic triad [6-8]. There are two kinds of Mac

produced by GAS isolates [7], which use different mechanisms to block the interaction between IgG and Fc receptor on the surface of PMNs. The type-1 Mac, such as M1 Mac produced by serotype M1 GAS strains, has high enzymatic activity and low affinity to Fc receptor on the surface of PMNs, while the type-2 Mac can bind to the Fc receptor and has lower enzymatic activity [9]. *S. equi* has a homologue of GAS M1 Mac (designated SeMac). In this study, SeMac was prepared and characterized. The results indicate that SeMac is a cysteine endopeptidase but does not inhibit opsonophagocytosis of *S. equi* by horse PMNs, suggesting that SeMac has function other than evading horse acquired immunity against *S. equi* infection.

## MATERIALS AND METHODS

### Materials

Purified horse IgG1 and a mixture of horse IgG1 and IgG4 were kindly provided by Dr. Bettina Wagner at Cornell University. *S. equi*-specific mouse sera were obtained from adult female outbred CD-1 Swiss mice (Charles River Laboratories, Wilmington, Mass.) 21 days after they were inoculated subcutaneously with  $1 \times 10^7$  cfu *S. equi* strain SEM1. Convalescent sera from 3 horses suffering from strangles were obtained 30 days after diagnosis. GAS M1 Mac was prepared as previously described [5].

### Bacterial Strains and Growth

Six of 10 *S. equi* strains used were kindly provided by Dr. James Musser at Methodist Hospital, Houston, Texas,

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and these strains were isolated more than 20 years ago from horses with strangles in the Eastern U.S. (4 strains), Brazil (1 strain), and Finland (1 strain). The other 4 strains were isolated in 2003 from horses with strangles in Livingston (designated strain SEM1) [10], Pony, Great Fall, and Norris in Montana. GAS strain MGAS5005 (serotype M1) has been described [11]. *Escherichia coli* Novablue and BL21(DE3) (Novagen, Madison, Wis.) were used for gene cloning and protein expression, respectively. *S. equi* and GAS strains were routinely grown in Todd-Hewitt broth (Difco Laboratories, Detroit, Mich.) supplemented with 0.2% yeast extract (THY) in 5% CO<sub>2</sub> at 37°C. THY agar and tryptose agar with 5% sheep blood (Becton Dickinson, Cockeysville, Md.) were used as the solid media.

### Gene Cloning and Mutagenesis

The gene fragment encoding mature SeMac was cloned from SEM1 with primers 5'-ACCATGGACGA TTACCAAAGGAATGCTAC-3' and 5'-CGAATTCT TAGCTCAGTTTCTGCCATATG-3'. The protein made from this cloned fragment lacks the presumed secretion signal sequence (amino acids 1-34). The PCR product was digested with *NcoI* and *EcoRI* and ligated into pET-21d (Novagen) to yield recombinant plasmid pSEMAC. The cloned gene was sequenced and had identical DNA sequence with the corresponding open reading frame of *S. equi* genome database [12]. Amino acid replacement of Cys102 and His272 or Asp294 of SeMac with serine and alanine, respectively, was achieved by site-directed mutagenesis using the Quick-Change Mutagenesis kit (Stratagene, La Jolla, Calif.). The entire mutated gene was sequenced to confirm the mutations and rule out spurious mutations.

### Expression and Purification of Recombinant SeMac

Recombinant SeMac was purified from *E. coli* BL21 (DE3) containing plasmid pSEMAC. Bacteria were grown to optical density at 600 nm of 0.5 in 6 liters of Luria-Bertani broth supplemented with 100 mg of ampicillin per liter at 37°C, and SeMac expression was then induced with 0.5 mM IPTG 6 h. Solutions used in purification were buffered with 10 mM Tris-HCl (pH 8.0). Cell paste was sonicated for 15 min at 4°C in 60 ml of the buffer, and centrifuged at 15,000 g for 10 min. The supernatant obtained was loaded onto a DEAE-Sepharose column (2.5 by 10 cm). The column was washed with 50 ml of Tris-HCl, and SeMac was eluted with 200 ml of 50 mM NaCl. SeMac was identified by sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE), and peak fractions were pooled. Ammonium sulfate was added to the pool to a concentration of 1.2 M, and the resulting solution was applied to a phenyl-Sepharose column (1.5 x 10 cm). The column was washed with 50 ml of 1.2 M (NH<sub>4</sub>)<sub>2</sub>SO<sub>4</sub> and eluted with a 100-ml linear gradient of 1.2 to 0.7 M (NH<sub>4</sub>)<sub>2</sub>SO<sub>4</sub>. Fractions containing the protein were pooled. The pooled protein was precipitated with (NH<sub>4</sub>)<sub>2</sub>SO<sub>4</sub> at 70% of saturation, centrifuged, dialyzed against 3 liters of 10 mM Tris-HCl buffer (pH 8.0) overnight, and loaded onto a DEAE-Sepharose column (1.5 by 10 cm). The protein was eluted with a 100-ml linear gradient of 2 to 8 mM NaCl and pooled. The pooled protein was concentrated by (NH<sub>4</sub>)<sub>2</sub>SO<sub>4</sub> precipitation and dialyzed against Tris-HCl as described above.

### DNA Sequencing

Genomic DNA was isolated with a FastDNA SPIN Kit (Qbiogene, Calsbad, Calif.) according to manufacturer's protocol. A DNA fragment containing *S. equi mac* was amplified by PCR using the genomic DNA samples and primers 5'-GCATCTCTACTATCTCATCAC-3' and 5'-ACAGGCACATTAATGTTTAAAC-3'. Sequences of the PCR products were obtained from both DNA strands with an Applied Biosystem 310 automated sequencer (Applied Biosystems, Inc., Foster City, Calif.).

### Anti-SeMac antisera

Four female outbred CD-1 Swiss mice (4-week-old) (Charles River Laboratories) were immunized subcutaneously with 50 µg of recombinant SeMac suspended in 200 µl of saline emulsified in 44 µl of monophosphoryl lipid A-synthetic trehalose dicorynomycolate adjuvant (Corixa, Hamilton, Mont.). The animals were anesthetized by isoflurane inhalation prior to immunization and blood collection. Blood was collected prior to active immunization to prepare control sera. Mice were boosted at weeks 2 and 4 with 50 µg of protein mixed with the adjuvant. Anti-SeMac antiserum was prepared from blood obtained 1 week after the second booster.

### Phagocytosis Assay

Phagocytosis assay was performed as described previously [13, 14]. Briefly, *S. equi* SEM1 and GAS bacteria from exponential growth phase in THY were washed with phosphate-buffered saline (PBS) and labeled with 0.75 µg/ml FITC in PBS at 37°C for 20 min. The labeled bacteria were washed and resuspended at 1 x 10<sup>9</sup> cfu/ml in PBS. Ten µl of the labeled bacteria were mixed with 100 µl of non-immune heparinized horse or human blood with or without 100 µg/ml SeMac and incubated with gentle shaking at 37°C for 5, 10, or 20 min. The samples were immediately processed using an Immunolyse Kit (Beckman Coulter) according to the manufacturer's protocol and analyzed by flow cytometry. The percentage of PMNs with fluorescent bacteria was used as a measure of phagocytosis efficiency.

### Other Assays

To assess IgG endopeptidase activity of SeMac, human, mouse or horse IgG (20 µg) was incubated with *E. coli* lysate containing SeMac proteins or 1 µg purified SeMac or GAS M1 Mac in PBS at 37° for 90 min, and the reaction mixture was analyzed by SDS-PAGE. Western immunoblot analysis, which was performed as described previously [15], was used to assess *in vitro* SeMac production and the presence of SeMac-specific antibody in the horse and mouse convalescent sera. Culture supernatant proteins of *S. equi*, which were used to assess the *in vitro* SeMac production, were prepared by the method of Lei *et al.* [16].

## RESULTS

### *S. equi* Mac Gene

*S. equi* genome encodes a Mac homologue SeMac. The protein shares 62.4% identity in amino acid sequence with Mac made by serotype M1 GAS and a putative secretion

signal sequence (amino acids 1-34), and the catalytic residues of GAS Mac consisting of Cys94, His262 and Asp284 are conserved in SeMac (Cys102, His272 and Asp294) (Fig. 1). To test whether other *S. equi* strains have the *mac* gene, 10 *S. equi* isolates representing various geographic locations were tested with PCR using *mac*-specific primers. All the strains tested had the *mac* gene (Fig. 2). DNA sequencing found that the *mac* gene is 100% conserved in DNA sequence in these strains.

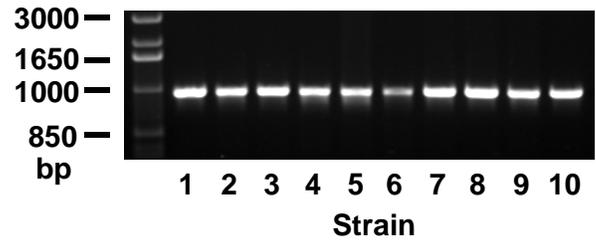
**Recombinant SeMac and *In Vitro* and *In Vivo* Expression of SeMac**

To characterize SeMac, the fragment of *S. equi mac* gene encoding mature SeMac was cloned, and recombinant SeMac was purified to >95% purity as assessed by SDS-PAGE (Fig. 3A). To assess the *in vitro* production of SeMac, culture supernatant of the 10 *S. equi* strains was prepared by the method of Lei *et al.* [16], resolved by SDS-PAGE, and probed by Western immunoblot with anti-SeMac mouse antisera. No SeMac was detected in all the samples (data not shown), suggesting that SeMac is not produced *in vitro*. To test whether SeMac is produced *in vivo* during *S. equi* infection, the presence of SeMac-specific antibody was assessed by Western immunoblot analysis with convalescent sera from 3 horses suffered from strangles and mice with experimental *S. equi* infection. All the convalescent sera tested had SeMac-specific antibody (Fig. 3B), indicating that SeMac is produced *in vivo* during infection.

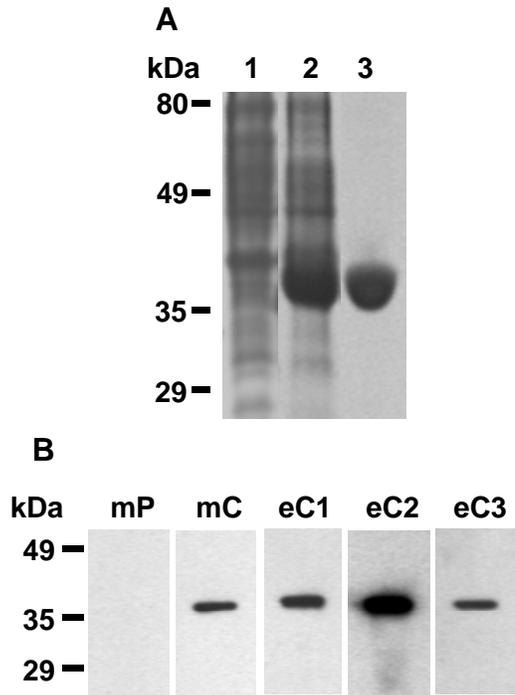
**Endopeptidase Activity of SeMac against Horse IgG1 and Human IgG**

GAS Mac can cleave the heavy chain of human IgG at the lower hinge region between Fab and Fc fragments. A catalytic triad of Cys94, His262, and Asp284 residues is critical for the enzymatic activity of GAS Mac [7, 8]. SeMac possesses putative catalytic residues of Cys102, His272, and Asp294 (Fig. 1). To determine whether SeMac also is a cysteine endopeptidase targeting IgG, Cys102 and His272 or Asp294 of SeMac were replaced with Ser and Ala, respectively, by site-directed mutagenesis, and wild-type and mutant (SeMac<sup>Cys102Ser</sup>, SeMac<sup>His272Ala</sup> and SeMac<sup>Asp294Ala</sup>) SeMac proteins expressed in *E. coli* were tested for IgG endopeptidase activity using human IgG. Wild-type SeMac cleaved the heavy chain of human IgG, while SeMac<sup>Cys102Ser</sup> and SeMac<sup>His272Ala</sup> completely lost the IgG endopeptidase activity, and SeMac<sup>Asp294Ala</sup> had dramatically lower enzymatic activity than the wild-type protein (Fig. 4A). The results indicate that SeMac is a cysteine endopeptidase, which

uses Cys102, His272 and Asp294 as its catalytic triad to cleave human IgG.



**Fig. (2).** Detection of *S. equi mac* gene in clinical isolates by PCR. The picture shows the *mac* PCR product amplified from 10 *S. equi* isolates from horses suffering from strangles.



**Fig. (3).** Recombinant SeMac and evidence for *in vivo* production of SeMac. (A) SDS-PAGE analysis of recombinant SeMac. Lanes: 1, *E. coli* lysate without SeMac (control); 2, *E. coli* lysate containing overexpressed SeMac; 3, purified SeMac. (B) Western immunoblot analysis of SeMac with SeMac antibody present in the convalescent-phase sera from 3 horses with *S. equi* infection (eC1, eC2, eC3) and mice before (mP) and after (mC) experimental *S. equi* infection.

SeMac	MKTTIAYPNKPHSLSGALLTAIAIFSLASSNITYADDDYQRNATEAYAKEVPHQITSVWTKGVTPLTPEQFRY
M1 Mac	.RKRC.ST-----AV.A.VTL.V.SVDRGVI..SFS.A.QEIR.SEVT.YHV.....AN.TQ
SeMac	NNEDFIHAPYL.H...DITKA.DGKDNLLCGAATAGNMLHWFDQNKTEIEAYLSKHPEKQKIIFNNQEL
M1 Mac	G...F...YVANQGWY...TFN...D...DQ.KR..EE.....N..GEQM
SeMac	FDLKAAIDTKDSQTSNQLFNFRDKAFPNLSARQLGVMPDLVLDLDFINGYLLNVFKTQSTDVNRVYQDKDK
M1 Mac	.V.EA...NH.LD.K.E..KE...Y..TKH...F..H.I.....R.SLTNHGP.P.KEG--S...
SeMac	RGGIFDAVFTRGDQTTLLTARHDLKNKGLNDISTIIKQELTEGRALALSHTYANVSIHVINLWGADFNAE
M1 Mac	.....SK...S...F.E.N.KE..DL..K...K..G.....R.N.....DSN
SeMac	GNLEAIYVTDSDANASIGMKKYFVGINAHRHVAISAKKIEGENIGAQVLGLFTLSSGKDIWQKLS
M1 Mac	...KAIYVT.S.S.....V.SAGK.....E.KED.....T.Q.S.NQTN

**Fig. (1).** Amino acid sequence alignment of SeMac and GAS M1 Mac. The conserved Cys (C), His (H), and Asp (D) constituting the catalytic triad are highlighted in red and underlined. Amino acid residues of GAS Mac identical to those in SeMac (·) and gaps introduced to maximize alignment (-) are indicated. The arrow indicates the presumed cleavage site of the secretion signal sequence.



**Table 1. The Amino Acid Sequences of the Lower Hinge Region of Human and Horse IgG Subgroups and Their Cleavability by GAS M1 Mac and SeMac**

IgG Subgroup	AA Sequence in the Lower Hinge Region <sup>a</sup>	Cleaved by <sup>b</sup>	
		M1 Mac	SeMac
Human IgG1	HTCPPCPAPELL <u>GG</u> PSV	+	+
Human IgG2	VE.....PVA- <u>  </u> ....	+	+
Human IgG3	PP..R..... <u>  </u> ....	+	+
Human IgG4	PP..S.....F. <u>  </u> ....	+	+
Horse IgG1	SK..KC.....	+	+
Horse IgG2	PQ...YTHSKF.....	ND	ND
Horse IgG3	CE..K.....	ND	ND
Horse IgG4	GGC.T..PEC.SV....	-	-
Horse IgG5	SP..K.....P.....	ND	ND
Horse IgG6	VIKE..CC.KCP.R...	ND	ND
Horse IgG7	CGGC.TC..C.SV....	ND	ND

<sup>a</sup>Cleavage was prior to the Gly residue underlined and determined by N-terminal amino acid sequencing of the proteolytic products in the reaction catalyzed by GAS M1, Mac and dots indicate identical residues with human IgG1.

<sup>b</sup>+, cleaved; -, not cleaved; ND, not determined.

majority of total horse IgG. The peptide fragment, PELLGG, in the lower hinge region is proposed to bind to the active site of Mac [9], and the cleavage occurs between the two Gly residues (Table 1). This lower hinge region is not well conserved in the seven subgroups of horse IgG [17] (Table 1). Cleavable horse IgG1 has PELLGG, while the non-cleavable horse IgG4 has PECLSVG in the same region, suggesting that the amino acid sequences of the lower hinge region are important for cleavability by SeMac and GAS Mac. If this is true, horse IgG3 may be also cleavable by SeMac, while the other five horse IgG subgroups may not be cleaved. These horse IgG antibodies that cannot be cleaved by SeMac thus still can mediate the opsonophagocytosis of *S. equi*.

SeMac and GAS M1 Mac show similar enzymatic specificity (Fig. 4 and Table 1), confirming the previous finding [18]. SeMac can cleave human IgG and inhibit the opsonophagocytosis of GAS by human PMNs, but it has limited enzymatic activity against horse IgG and is unable to inhibit opsonophagocytosis of *S. equi* by horse PMNs. Thus, there is a correlation between the enzymatic activity of SeMac and its ability to inhibit opsonophagocytosis, suggesting that SeMac functions like GAS M1 Mac in the inhibition of opsonophagocytosis of GAS by human PMNs.

Timoney *et al.* recently reported that IdeE/SeMac reduces the bactericidal activity of isolated equine PMNs for *S. equi* [19]. Our results suggest that the inhibition of the bactericidal activity of PMNs may not be mediated by opsonophagocytosis or may be insignificant in whole blood.

SeMac is not produced *in vitro*, whereas GAS Mac is [16]. The *mac* gene is controlled by the two-component regulatory system CovRS [5], which also controls the expression of many virulence factors including the hyaluronic capsule [20]. The hyaluronic capsule of *S. equi* is highly

produced *in vitro*. These observations suggest that the *semac* and *mac* genes are regulated by different mechanisms. This suggestion is supported by only 29% DNA sequence identity between *S. equi* and M1 GAS in the upstream region of the *semac* and *mac* genes.

*S. equi* is a horse pathogen. The fact that SeMac does not inhibit opsonophagocytosis of *S. equi* by horse PMNs indicates that SeMac is not involved in the evasion of the acquired horse immunity against *S. equi*. This suggests that SeMac has other unknown function.

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