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## IL-8 Protease SpyCEP/ScpC Promotes Group A Streptococcal Resistance to Neutrophil Killing

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## **Supplemental Figures**



**Figure S1:** Mutagenesis and heterologous expression of IL-8 protease SpyCEP. (A) Scheme for precise, in-frame allelic replacement of the *cepA* gene in M1 GAS; (B) Western blot confirms loss of SpyCEP expression upon targeted mutagenesis in M1 GAS, complementation of the mutant *in trans*, and heterologous expression of SpyCEP in *Lactococcus lactis*; (C) Presence of SpyCEP correlates with IL-8 protease activity by SDS-PAGE.

## **Supplementary Figure S2**



**Figure S2:** Elimination of SpyCEP does not affect M1 protein or DNase expression. (A) M1 protein expression is quantified by dot blot for the WT M1 GAS parent strain (with and without the vector pDCerm), isogenic  $\Delta cepA$  mutant and the complemented  $\Delta cepA$  mutant. (**B**) DNase activity

assessed by degradation of calf thymus DNA by supernatants of WT M1 GAS parent vs. isogenic  $\Delta cepA$  mutant as compared by agarose gel electrophoresis.



**Figure S3:** Additional studies on the role of SpyCEP in GAS phagocyte resistance. (A) No difference in human whole blood survival observed between the WT M1 GAS strain and the isogenic  $\Delta cepA$  mutant. Experiments performed in triplicate and repeated three times with similar results; representative experiment, mean  $\pm$  SEM. (B) WT GAS parent strain shows enhanced neutrophil survival compared to the isogenic  $\Delta cepA$  mutant when phagocytosis is inhibited by cytochalasin D (Cyt D) or NET degradation is inhibited by G-actin. Experiments performed in triplicate and repeated three times with similar results; representative experiment shown, mean  $\pm$  SEM . (C) In the presence of excess exogenous IL-8 differences in induction of neutrophil extracellular traps (NETs) in response to bacterial exposure is abolished; quantitative enumeration of NETs, experiments performed in triplicate and repeated two to three times with similar results; representative experiment  $\pm$  SEM, two-sided T test.

## Amino acid alignment of Group A Streptococcal SpyCEP and the homologous Streptococcus iniae protease, Cepl

S. iniae Cepl GAS SpyCEP	MEKKERFSLRKYKSGIV SVLIGTVFFAGASQVSADQLTTVSIAEETPSLLTKEVDGQTENSNQDKLTM
S. iniae Cepl GAS SpyCEP	L SAD S SAEKE E SWP L L DAP I NKDI H DWI QV SGAWENGYKGOGKVI ALI DT GI DVNHQAMRI SDI SQAKFKI AEDMDQQKAKAKI NYGKWI NQKV DAASMANTGPDATOKSASLP PWNTDI H DWVKTKGAWDKGYKGOGKVVAVIDT GI DPAHQSMRI SDVSTAKVKSKEDMLA ROKAAGI NYGSWI NDKV 101
S. iniae Cepl GAS SpyCEP	I FAHNYVENNDKVKEVKEDFDFDFD I EDDS I LDS I ESTLVQSVDKKRYRVEPKSNSDKPKETV I QINPDDFSHI I DWPSHDDE SQHESHGMHVTG I AVGN VFAHNYVENSDNIKENQFE-DFDEDWENFEFDAEAEPKAIKKHKIYRPQSTQAPKETVIKTEETDGSHDIDWTQTDDDTKYESHGMHVTGIVAGN 201
S. iniae Cepl GAS SpyCEP	PLEASPIGERFLGWAPEAQVIFMRVFANDFMGFGEALWIKAIEDAVALGADAINLSLGGPNGSFLGGNASLMAAIEKAKKAGVSVIVAAGNERLFGSDHA SKEAAATGERFLGIAPEAQVMFMRVFANDVMGSAESLFIKAIEDAVALGADVINLSLGTANGAQLSGSKPLMEAIEKAKKAGVSVVVAAGNERVYGSDHD 301
S. iniae Cepl GAS SpyCEP	D P FASNPDYGIVNSPSTGKIPTSVAAIDNKIIIDRLMKVEGLENRADLDHGKALYTESIDYKKIKEVLSFDNDYDFIYINQPTDQAYIGKEVKGKIVLIE DP LATNPDYGLVGSPSTGRTPTSVAAINSKWVIQRLMTVEELEKRADLNHGKAIYSESVDFKNIKDSLGYDKSHQFAYVKESTDAGYNAQDVKGKIALIE 401
S. iniae Cepl GAS SpyCEP	RHLDH PYVEL I ANAKKHE VAGIL I FNH I PGQ SNRKMRLTS E GQVLPSAFI SHE FGKAMSQLNGNGTGRLRHESKLSKASNOR SQQMNHFSSWGLTSDGYL RDPNKTYDEMIALAKKHGALGVLI FNNKPGQ SNRSMRLTANGMGI PSAFI SHE FGKAMSQLNGNGTGSLEFDSVVSKAP SQKGNEMNHFSNWGLTSDGYL 501
S. iniae Cepl GAS SpyCEP	K P D I TAPGGD I YSTYNDNHY G SQ TG T SMAS PY I AGA SLLEKQY IEAQHPDVKT EEM SDLVKYLLMSNASI HKDPKTQLTTSPRRQGAGLLN VQA A V TSGL K P D I TAPGGD I YSTYNDNHY G SQ TG TSMAS PQ I AGA SLLVKQYLEKTQPNLPKEK 1 ADI VKNLLMSNAQI HVNPEIKT TTSPRQGAGLLN I DG A V TSGL 601
S. iniae Cepl GAS SpyCEP	YE TGSDNYGSISLGNLGEKISFDVTVHNLSNHARNLRYVTDEMTDKYL-EDGRFTLSSVALKSYQGHLVDVPAKGQTHIRVSMDVSEFTKTLTKQMPNGY YVTGKDNYGSISLGNITDTMTFDVTVHNLSNKDKTLRYDTELLTDHVDPQKGRFTLTSRSLKTYQGGEVTVPANGKVTVRVTMDVSQFTKELTKQMPNGY 701
S. iniae Cepl GAS SpyCEP	FLEGFVRHENATDISTRDRVN I PFVGFRGEFQNLAVVEES I YNLKARGERGFYFEESK TFDEI YVGRHYTGLVTI GADANVSTRT I SDNGI HTLGTYRNRD YLEGFVRFRDSODDOLNRVN I PFVGFRGOFENLAVAEES I YRLKSOGRTGFYFDES G PRDDI YVGRHFTGLVTLGSET NVSTRT I SDNGLHTLGTFRNAD 801
S. iniae Cepl GAS SpyCEP	GKEILEKDQSGNVVLAISPNGDKNQDHVAFKGVFLRKYKGLKASVYRADDHKRQQLLWTS-QAHNGEKNYHSDIRHPQSTTLLSTEFSGRSLSGEDLPDG GKFILEKNAQGNPVLAISPNGDNNQDFAAFKGVFLRKYQGLKASVYHASDKEHKNPLWYSPESFKGDKNENSDIRFARSTTLLGTAFSGKSLTGAELPDG 901
S. iniae Cepl GAS SpyCEP	KYQYVVSYYPDVIGAKSQEMVFDVIVDREKPLLTSASFNPETREFKALDVHDRGQSGLLRDSVFYLEEKDGKPYTISINQGFKYVSVADNKVEVGKSKDG YYH <mark>YVVSYYPDVVGAK</mark> RQEMTFDMILDRQKPVLSQATFDPETNRFKPEPLKDRGLAGVRKDSVFYLERKDNKPYTVTINDSYKYVSVEDNKTFVERQADG 1001
S. iniae Cepl GAS SpyCEP	GFILPLDKANLADFYYMVEDFAGNIAIAKLGETLPEAFGKDIFTFTKEGNYQNHHALEDKLQMTANDSGLMTNQSDIMITNRNRPLSRUVKANQSQLIL SFILPLDKAKLGDFYYMVEDFAGNVAIAKLGDHLPQTLGKTPIKLKTDGNYQTKETLKDNLEMTQSDTGLVTNQAQLAVVHRNQPQSQLTKMNQ-DFFI 1101
S. iniae Cepl GAS SpyCEP	S PNDNGNRDF HAFRGLP GOVYHDLKWTVFAADDKLMS SALWTSPQKTTVGDLNTHRWDGKTQMGQKVLSGAWRYVILTYRDA SGMVKRQEHDHLVSHEEPH SPNEDGNKDFVAFKGLENNVYNDLTVNVYAKDDHQKQTPIWSSQAGAGASATESTAWYGIHARGSKVMPGDYQYVVTYRDEHGKEHQKQYTTSVNDKKPM 1201
S. iniae Cepl GAS SpyCEP	I TKAS FOXDGDKEF FKPGKVLDLNOVGIAR E E VFYVLEKE GRKYDIATVDDLVTIS DR RULIPRNADGSYTIPKVEGVTPADFYYLVEDMAGNIVYSSLL I TOGREDTINGVDHETPDNTKALGSSGIVREE VFYLAKKNGRKEDVTEGKDGITVSDNKVYIPKNPDGSYTISKRDGVTLSDYYYLVEDRAGNVSFATLR 1301
S. iniae Cepl GAS SpyCEP	PMRSVGDGQGILDIALVYNNS I ERPKNPFTYL VRDENGQALDKLNYYDDRTQMLILPFGHYSLELLTYDHNVAELVSPRUFNLEISEQNAEVHAEVIRK DLKAVGKDKAVVNFGLDLPVPEDKQINNFTYLVRDADGKPTENLEYYNNSGNSLILPYGKYTVELLTYDTNAAKLESDKIVSFTLSADNEQQVTFKMTM 1401
S. iniae Cepl GAS SpyCEP	SEPAMVRVFENQLLPEGSRVMLQAENGQLIDLDKSLYVPNAYGATVRDGRUTIMVTLPEGYHISGQUSIDALQNDLVESYLQLVSNG-PSVSEGQNVAGS LATSQITAHEDHLLPEGSRVSLKTAQGQLIPLEQSLYVPKAYGKTVQEGTYEVVVSLPKGYRIEGNTKVNTLPNEVHELSLRLVKVGDASDSTGDHKVMS 1501
S. iniae Cepl GAS SpyCEP	1000 VAMLKEFLKNAIRQGE HKASQL PVTGEKQALLSLSSG VALLCLAPLSEFQKRYDKGE KNNSQALTASATPTKTTPATAKAL PSTGEKMGLKLRIVGLVLLGLTCL-ESRKKSTK-D 1601

**Figure S4.** Homologue of GAS SpyCEP identified in *Streptococcus iniae*. Amino acid alignment of CepA (M14 GAS strain JS95) and CepI (*S. iniae* strain 9117) with high level of similarity; identical amino acids shaded in black and similar amino acids shaded in gray. Alignment generated with ClustalW and shaded using BOXSHADE.



**Figure S5.** *Streptococcus iniae* CepI contributes to IL-8 degradation and neutrophil resistance. (A) Insertional mutagenesis scheme for the *S. iniae cepA* homologue, *cepI*. (B) Anti-IL-8 Western blot confirms cleavage of IL-8 by *S. iniae* reduced in  $\Delta cepI$  mutant. (C) SpyCEP contributes to *S. iniae* survival upon co-incubation with human neutrophils, experiments performed in triplicate and repeated twice with similar results; representative experiment  $\pm$  SEM. (D) C57Bl6 mice injected subcutaneously with equivalent inocula of WT *S. iniae* (left flank) or the the isogenic  $\Delta cepI$  mutant (right flank); skin lesion progression measured for 2 days.