

SEQUENCE LISTING

<110> Linnane Pharm AB

<120> Novel Therapy

<130> P3095/WO

<150> GB1617548.1

<151> 2016-10-17

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<170> BiSSAP 1.3.6

<210> 1

<211> 330

<212> PRT

<213> Escherichia coli

<220>

<223> Sigma S protein

<400> 1

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Gln Glu Pro Ser Asp Asn Asp Leu Ala Glu Glu Glu Leu Leu Ser Gln
          35          40          45
Gly Ala Thr Gln Arg Val Leu Asp Ala Thr Gln Leu Tyr Leu Gly Glu
          50          55          60
Ile Gly Tyr Ser Pro Leu Leu Thr Ala Glu Glu Glu Val Tyr Phe Ala
65          70          75          80
Arg Arg Ala Leu Arg Gly Asp Val Ala Ser Arg Arg Arg Met Ile Glu
          85          90          95
Ser Asn Leu Arg Leu Val Val Lys Ile Ala Arg Arg Tyr Gly Asn Arg
          100         105         110
Gly Leu Ala Leu Leu Asp Leu Ile Glu Glu Gly Asn Leu Gly Leu Ile
          115         120         125
Arg Ala Val Glu Lys Phe Asp Pro Glu Arg Gly Phe Arg Phe Ser Thr
          130         135         140
Tyr Ala Thr Trp Trp Ile Arg Gln Thr Ile Glu Arg Ala Ile Met Asn
145          150         155         160
Gln Thr Arg Thr Ile Arg Leu Pro Ile His Ile Val Lys Glu Leu Asn
          165         170         175
Val Tyr Leu Arg Thr Ala Arg Glu Leu Ser His Lys Leu Asp His Glu
          180         185         190
Pro Ser Ala Glu Glu Ile Ala Glu Gln Leu Asp Lys Pro Val Asp Asp

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Val	Ser	Arg	Met	Leu	Arg	Leu	Asn	Glu	Arg	Ile	Thr	Ser	Val	Asp	Thr
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Pro	Leu	Gly	Gly	Asp	Ser	Glu	Lys	Ala	Leu	Leu	Asp	Ile	Leu	Ala	Asp
225					230						235				240
Glu	Lys	Glu	Asn	Gly	Pro	Glu	Asp	Thr	Thr	Gln	Asp	Asp	Asp	Met	Lys
			245							250				255	
Gln	Ser	Ile	Val	Lys	Trp	Leu	Phe	Glu	Leu	Asn	Ala	Lys	Gln	Arg	Glu
		260					265						270		
Val	Leu	Ala	Arg	Arg	Phe	Gly	Leu	Leu	Gly	Tyr	Glu	Ala	Ala	Thr	Leu
	275					280					285				
Glu	Asp	Val	Gly	Arg	Glu	Ile	Gly	Leu	Thr	Arg	Glu	Arg	Val	Arg	Gln
	290					295					300				
Ile	Gln	Val	Glu	Gly	Leu	Arg	Arg	Leu	Arg	Glu	Ile	Leu	Gln	Thr	Gln
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<213> Escherichia coli

<220>

<223> Sigma S protein

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Gly	Val	Glu	Val	Phe	Asp	Glu	Lys	Ala	Leu	Val	Glu	Glu	Glu	Pro	Ser
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Asp	Asn	Asp	Leu	Ala	Glu	Glu	Glu	Leu	Leu	Ser	Gln	Gly	Ala	Thr	Gln
	50				55					60					
Arg	Val	Leu	Asp	Ala	Thr	Gln	Leu	Tyr	Leu	Gly	Glu	Ile	Gly	Tyr	Ser
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Pro	Leu	Leu	Thr	Ala	Glu	Glu	Glu	Val	Tyr	Phe	Ala	Arg	Arg	Ala	Leu
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Arg	Gly	Asp	Val	Ala	Ser	Arg	Arg	Arg	Met	Ile	Glu	Ser	Asn	Leu	Arg
	100							105					110		
Leu	Val	Val	Lys	Ile	Ala	Arg	Arg	Tyr	Gly	Asn	Arg	Gly	Leu	Ala	Leu
	115					120						125			
Leu	Asp	Leu	Ile	Glu	Glu	Gly	Asn	Leu	Gly	Leu	Ile	Arg	Ala	Val	Glu
	130				135					140					
Lys	Phe	Asp	Pro	Glu	Arg	Gly	Phe	Arg	Phe	Ser	Thr	Tyr	Ala	Thr	Trp
145				150						155					160
Trp	Ile	Arg	Gln	Thr	Ile	Glu	Arg	Ala	Ile	Met	Asn	Gln	Thr	Arg	Thr
			165					170						175	

Ile Arg Leu Pro Ile His Ile Val Lys Glu Leu Asn Val Tyr Leu Arg
 180 185 190
 Thr Ala Arg Glu Leu Ser His Lys Leu Asp His Glu Pro Ser Ala Glu
 195 200 205
 Glu Ile Ala Glu Gln Leu Asp Lys Pro Val Asp Asp Val Ser Arg Met
 210 215 220
 Leu Arg Leu Asn Glu Arg Ile Thr Ser Val Asp Thr Pro Leu Gly Gly
 225 230 235 240
 Asp Ser Glu Lys Ala Leu Leu Asp Ile Leu Ala Asp Glu Lys Glu Asn
 245 250 255
 Gly Pro Glu Asp Thr Thr Gln Asp Asp Asp Met Lys Gln Ser Ile Val
 260 265 270
 Lys Trp Leu Phe Glu Leu Asn Ala Lys Gln Arg Glu Val Leu Ala Arg
 275 280 285
 Arg Phe Gly Leu Leu Gly Tyr Glu Ala Ala Thr Leu Glu Asp Val Gly
 290 295 300
 Arg Glu Ile Gly Leu Thr Arg Glu Arg Val Arg Gln Ile Gln Val Glu
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 Glu Ala Leu Phe Arg Glu
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 <212> PRT
 <213> Escherichia coli

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 Thr Ala Arg
 35

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 <223> NplD protein

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Pro	Ala	Pro	Val	Ser	Ser	Val	Asn	Gly	Asn	Ala	Pro	Ala	Asn	Thr	Asn
35							40				45				
Ser	Gly	Met	Leu	Ile	Thr	Pro	Pro	Pro	Lys	Met	Gly	Thr	Thr	Ser	Thr
50							55				60				
Ala	Gln	Gln	Pro	Gln	Ile	Gln	Pro	Val	Gln	Gln	Pro	Gln	Ile	Gln	Ala
65							70				75				
Thr	Gln	Gln	Pro	Gln	Ile	Gln	Pro	Met	Gln	Pro	Val	Ala	Gln	Gln	Pro
			85				90				95				
Val	Gln	Met	Glu	Asn	Gly	Arg	Ile	Val	Tyr	Asn	Arg	Gln	Tyr	Gly	Asn
			100				105				110				
Ile	Pro	Lys	Gly	Ser	Tyr	Ser	Gly	Ser	Thr	Tyr	Thr	Val	Lys	Lys	Gly
115							120				125				
Asp	Thr	Leu	Phe	Tyr	Ile	Ala	Trp	Ile	Thr	Gly	Asn	Asp	Phe	Arg	Asp
130							135				140				
Leu	Ala	Gln	Arg	Asn	Asn	Ile	Gln	Ala	Pro	Tyr	Ala	Leu	Asn	Val	Gly
145							150				155				
Gln	Thr	Leu	Gln	Val	Gly	Asn	Ala	Ser	Gly	Thr	Pro	Ile	Thr	Gly	Gly
			165				170				175				
Asn	Ala	Ile	Thr	Gln	Ala	Asp	Ala	Ala	Glu	Gln	Gly	Val	Val	Ile	Lys
180							185				190				
Pro	Ala	Gln	Asn	Ser	Thr	Val	Ala	Val	Ala	Ser	Gln	Pro	Thr	Ile	Thr
195							200				205				
Tyr	Ser	Glu	Ser	Ser	Gly	Glu	Gln	Ser	Ala	Asn	Lys	Met	Leu	Pro	Asn
210							215				220				
Asn	Lys	Pro	Thr	Ala	Thr	Thr	Val	Thr	Ala	Pro	Val	Thr	Val	Pro	Thr
225							230				235				
Ala	Ser	Thr	Thr	Glu	Pro	Ile	Val	Ser	Ser	Thr	Ser	Thr	Ser	Thr	Pro
			245				250				255				
Ile	Ser	Thr	Trp	Arg	Trp	Pro	Thr	Glu	Gly	Lys	Val	Ile	Glu	Thr	Phe
			260				265				270				
Gly	Ala	Ser	Glu	Gly	Gly	Asn	Lys	Gly	Ile	Asp	Ile	Ala	Gly	Ser	Lys
275							280				285				
Gly	Gln	Ala	Ile	Ile	Ala	Thr	Ala	Asp	Gly	Arg	Val	Val	Tyr	Ala	Gly
290							295				300				
Asn	Ala	Leu	Arg	Gly	Tyr	Gly	Asn	Leu	Ile	Ile	Ile	Lys	His	Asn	Asp
305							310				315				
Asp	Tyr	Leu	Ser	Ala	Tyr	Ala	His	Asn	Asp	Thr	Met	Leu	Val	Arg	Glu
			325				330				335				
Gln	Gln	Glu	Val	Lys	Ala	Gly	Gln	Lys	Ile	Ala	Thr	Met	Gly	Ser	Thr
			340				345				350				
Gly	Thr	Ser	Ser	Thr	Arg	Leu	His	Phe	Glu	Ile	Arg	Tyr	Lys	Gly	Lys
355							360				365				
Ser	Val	Asn	Pro	Leu	Arg	Tyr	Leu	Pro	Gln	Arg					
370							375								