

PCT\_sequence\_protocol  
SEQUENCE LISTING

<110> Life Science Inkubator BetriebsGmbH& Co. KG

<120> Mittel zur Endotoxin-Detoxifikation

<130> 54 235 K

<160> 1

<170> BISSAP 1.0

<210> 1

<211> 535

<212> PRT

<213> Homo sapiens

<220>

<221> SOURCE

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<223> /mol\_type="protein"  
/organism="Homo sapiens"

<400> 1

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Leu Gln Leu Ser  Leu Gly Ile Ile  Pro Val Glu Glu Glu Asn Pro Asp
      20      25      30
Phe Trp Asn Arg Glu Ala Ala Glu Ala Leu Gly Ala Ala Lys Lys Leu
      35      40      45
Gln Pro Ala Gln Thr Ala Ala Lys Asn Leu Ile Ile Phe Leu Gly Asp
      50      55      60
Gly Met Gly Val Ser Thr Val Thr Ala Ala Arg Ile Leu Lys Gly Gln
65      70      75      80
Lys Lys Asp Lys Leu Gly Pro Glu Ile Pro Leu Ala Met Asp Arg Phe
      85      90      95
Pro Tyr Val Ala Leu Ser Lys Thr Tyr Asn Val Asp Lys His Val Pro
      100      105      110
Asp Ser Gly Ala Thr Ala Thr Ala Tyr Leu Cys Gly Val Lys Gly Asn
      115      120      125
Phe Gln Thr Ile Gly Leu Ser Ala Ala Ala Arg Phe Asn Gln Cys Asn
      130      135      140
Thr Thr Arg Gly Asn Glu Val Ile Ser Val Met Asn Arg Ala Lys Lys
145      150      155      160
Ala Gly Lys Ser Val Gly Val Val Thr Thr Thr Arg Val Gln His Ala
      165      170      175
Ser Pro Ala Gly Thr Tyr Ala His Thr Val Asn Arg Asn Trp Tyr Ser
      180      185      190
Asp Ala Asp Val Pro Ala Ser Ala Arg Gln Glu Gly Cys Gln Asp Ile
      195      200      205
Ala Thr Gln Leu Ile Ser Asn Met Asp Ile Asp Val Ile Leu Gly Gly
      210      215      220
Gly Arg Lys Tyr Met Phe Arg Met Gly Thr Pro Asp Pro Glu Tyr Pro
225      230      235      240
Asp Asp Tyr Ser Gln Gly Gly Thr Arg Leu Asp Gly Lys Asn Leu Val
      245      250      255
Gln Glu Trp Leu Ala Lys Arg Gln Gly Ala Arg Tyr Val Trp Asn Arg
      260      265      270
Thr Glu Leu Met Gln Ala Ser Leu Asp Pro Ser Val Thr His Leu Met
      275      280      285
Gly Leu Phe Glu Pro Gly Asp Met Lys Tyr Glu Ile His Arg Asp Ser
      290      295      300
Thr Leu Asp Pro Ser Leu Met Glu Met Thr Glu Ala Ala Leu Arg Leu
305      310      315      320
Leu Ser Arg Asn Pro Arg Gly Phe Phe Leu Phe Val Glu Gly Gly Arg
      325      330      335
Ile Asp His Gly His His Glu Ser Arg Ala Tyr Arg Ala Leu Thr Glu
      340      345      350

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Thr	Ile	Met	Phe	Asp	Asp	Ala	Ile	Glu	Arg	Ala	Gly	Gln	Leu	Thr	Ser
		355					360					365			
Glu	Glu	Asp	Thr	Leu	Ser	Leu	Val	Thr	Ala	Asp	His	Ser	His	Val	Phe
	370					375					380				
Ser	Phe	Gly	Gly	Tyr	Pro	Leu	Arg	Gly	Ser	Ser	Ile	Phe	Gly	Leu	Ala
385					390					395					400
Pro	Gly	Lys	Ala	Arg	Asp	Arg	Lys	Ala	Tyr	Thr	Val	Leu	Leu	Tyr	Gly
			405						410					415	
Asn	Gly	Pro	Gly	Tyr	Val	Leu	Lys	Asp	Gly	Ala	Arg	Pro	Asp	Val	Thr
			420					425					430		
Glu	Ser	Glu	Ser	Gly	Ser	Pro	Glu	Tyr	Arg	Gln	Gln	Ser	Ala	Val	Pro
		435					440					445			
Leu	Asp	Glu	Glu	Thr	His	Ala	Gly	Glu	Asp	Val	Ala	Val	Phe	Ala	Arg
	450					455					460				
Gly	Pro	Gln	Ala	His	Leu	Val	His	Gly	Val	Gln	Glu	Gln	Thr	Phe	Ile
465					470					475					480
Ala	His	Val	Met	Ala	Phe	Ala	Ala	Cys	Leu	Glu	Pro	Tyr	Thr	Ala	Cys
			485						490					495	
Asp	Leu	Ala	Pro	Pro	Ala	Gly	Thr	Thr	Asp	Ala	Ala	His	Pro	Gly	Arg
			500					505					510		
Ser	Val	Val	Pro	Ala	Leu	Leu	Pro	Leu	Leu	Ala	Gly	Thr	Leu	Leu	Leu
		515					520						525		
Leu	Glu	Thr	Ala	Thr	Ala	Pro									
	530					535									