

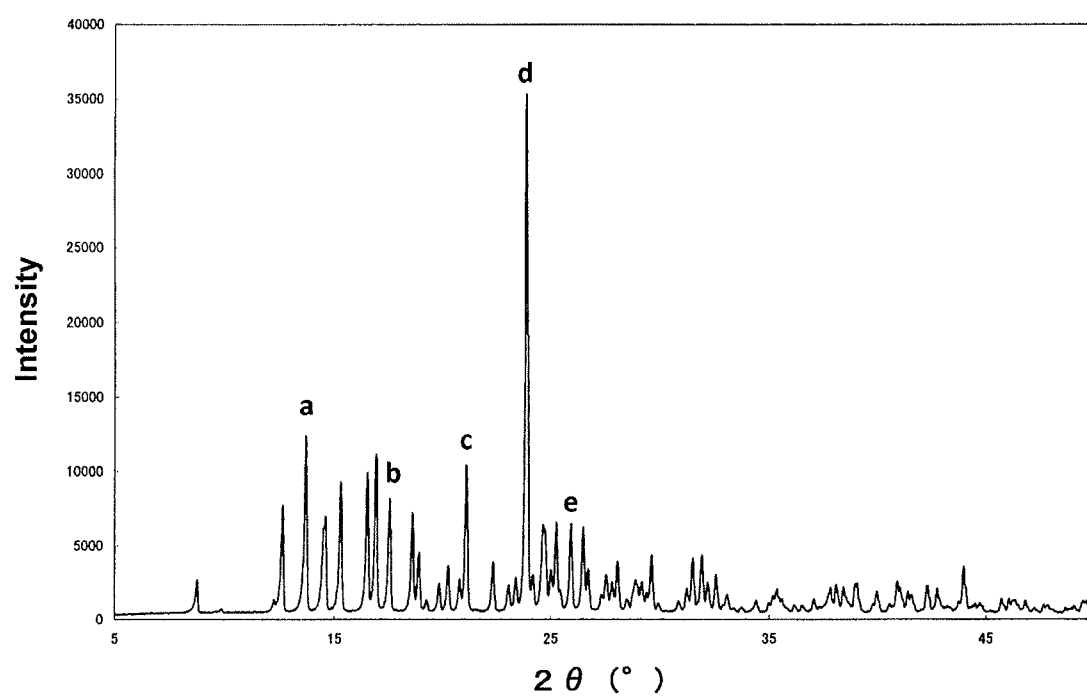


FIG. 1

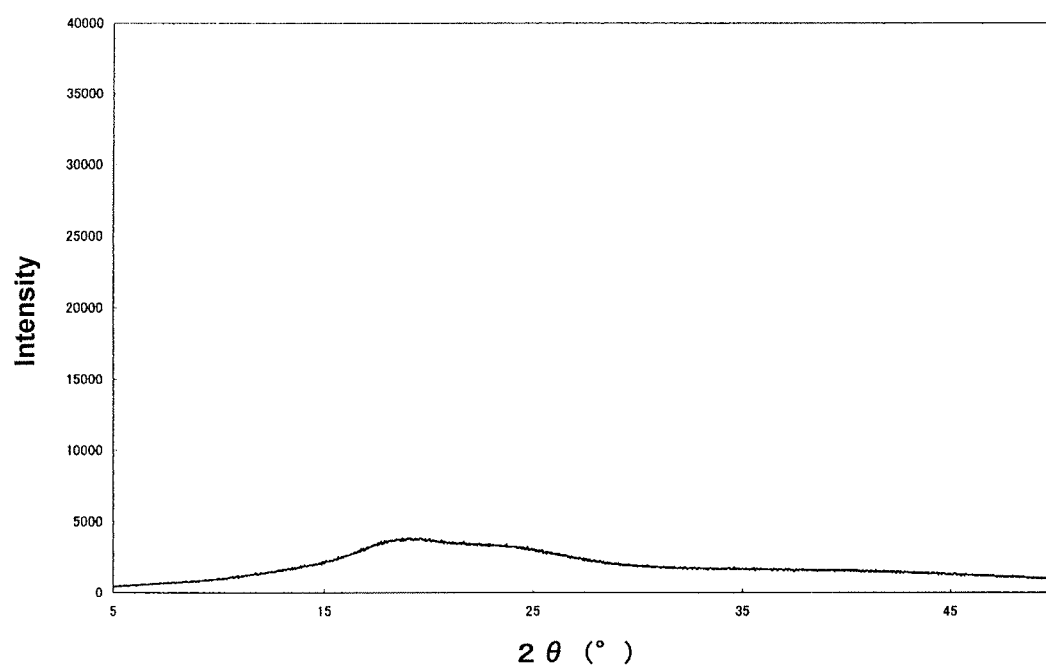


FIG. 2

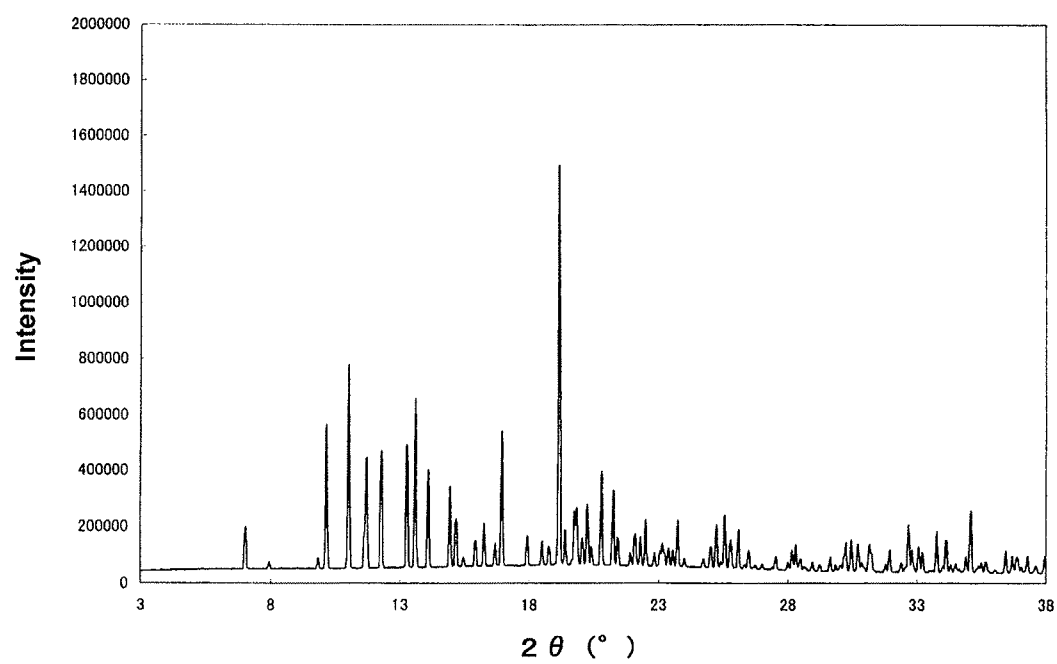


FIG. 3

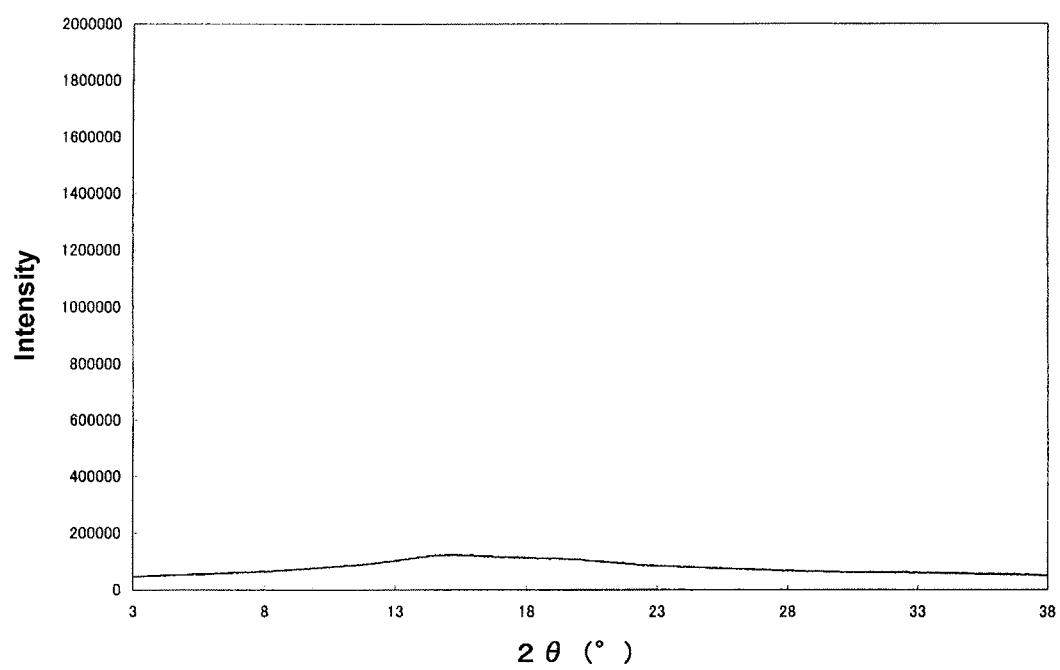


FIG. 4

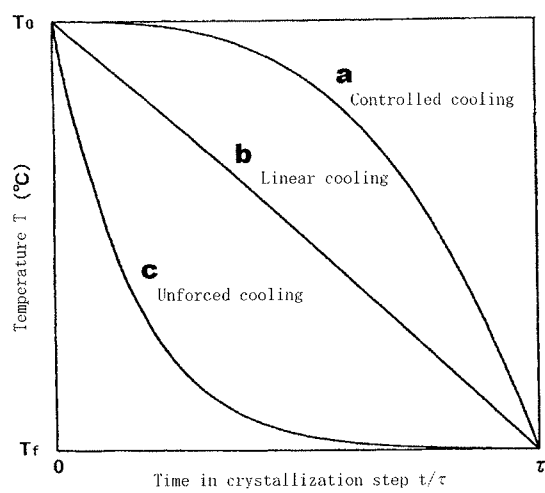


FIG. 5

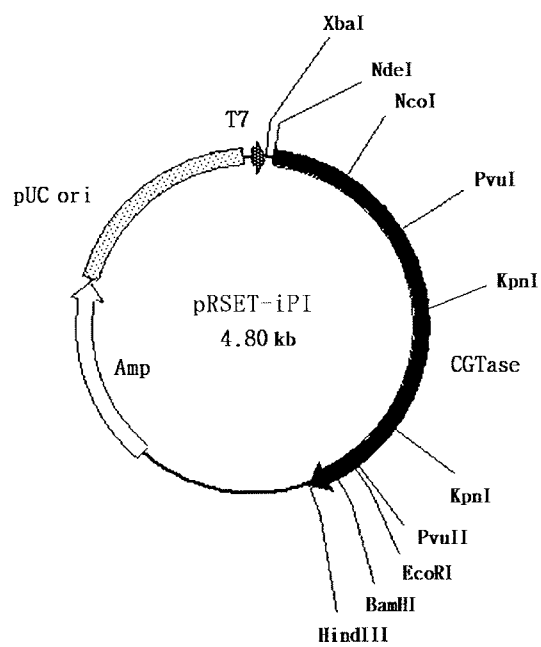


FIG. 6

Sequence-Listing

@@@@@@@@@@@@@@@@SEQUENCE LISTING

<110> Hayashibara Co., Ltd.

<120> Process for producing a particulate composition comprising crystalline alpha, alpha-trehalose di-hydrate

<130> WO1243

<160> 13

<210> 1

<211> 682

<212> PRT

<213> Paenibacillus illinoisensis

<400> 1

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Asp Thr Ala Val Thr Asn Lys Gln Asn Phe Ser Thr Asp Val Ile Tyr
 1      5      10      15
Gln Val Phe Thr Asp Arg Phe Leu Asp Gly Asn Pro Ser Asn Asn Pro
      20      25      30
Thr Gly Ala Ala Phe Asp Gly Thr Cys Ser Asn Leu Lys Leu Tyr Cys
      35      40      45
Gly Gly Asp Trp Gln Gly Leu Ile Asn Lys Ile Asn Asp Asn Tyr Phe
      50      55      60
Ser Asp Leu Gly Val Thr Ala Leu Trp Ile Ser Gln Pro Val Glu Asn
      65      70      75      80
Ile Phe Ala Thr Ile Asn Tyr Ser Gly Val Thr Asn Thr Ala Tyr His
      85      90      95
Gly Tyr Trp Ala Arg Asp Phe Lys Lys Thr Asn Pro Tyr Phe Gly Thr
      100      105      110
Met Thr Asp Phe Gln Asn Leu Val Thr Ser Ala His Ala Lys Gly Ile
      115      120      125
Lys Ile Ile Ile Asp Phe Ala Pro Asn His Thr Phe Pro Ala Met Glu
      130      135      140
Thr Asp Thr Ser Phe Ala Glu Asn Gly Lys Leu Tyr Asp Asn Gly Ser
      145      150      155      160
Leu Val Gly Gly Tyr Thr Asn Asp Thr Asn Gly Tyr Phe His His Asn
      165      170      175
Gly Gly Ser Asp Phe Ser Thr Leu Glu Asn Gly Ile Tyr Lys Asn Leu
      180      185      190
Tyr Asp Leu Ala Asp Leu Asn His Asn Asn Ser Thr Ile Asp Thr Tyr
      195      200      205
Phe Lys Asp Ala Ile Lys Leu Trp Leu Asp Met Gly Val Asp Gly Ile
      210      215      220
Arg Val Asp Ala Val Lys His Met Pro Gln Gly Trp Gln Lys Asn Trp
      225      230      235      240
Met Ser Ser Ile Tyr Ala His Lys Pro Val Phe Thr Phe Gly Glu Trp
      245      250      255
Phe Leu Gly Ser Ala Ala Ser Asp Ala Asp Asn Thr Asp Phe Ala Asn
      260      265      270
Glu Ser Gly Met Ser Leu Leu Asp Phe Arg Phe Asn Ser Ala Val Arg
      275      280      285
Asn Val Phe Arg Asp Asn Thr Ser Asn Met Tyr Ala Leu Asp Ser Met
      290      295      300
Leu Thr Ala Thr Ala Ala Asp Tyr Asn Gln Val Asn Asp Gln Val Thr
      305      310      315      320
Phe Ile Asp Asn His Asp Met Asp Arg Phe Lys Thr Ser Ala Val Asn
      325      330      335
Asn Arg Arg Leu Glu Gln Ala Leu Ala Phe Thr Leu Thr Ser Arg Gly
      340      345      350
Val Pro Ala Ile Tyr Tyr Gly Thr Glu Gln Tyr Leu Thr Gly Asn Gly
      355      360      365
Asp Pro Asp Asn Arg Gly Lys Met Pro Ser Phe Ser Lys Ser Thr Thr
      370      375      380
Ala Phe Ser Val Ile Ser Lys Leu Ala Pro Leu Arg Lys Ser Asn Pro
      385      390      395      400
Ala Ile Ala Tyr Gly Ser Thr Gln Gln Arg Trp Ile Asn Asn Asp Val

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Sequence-Listing

				405					410					415			
Tyr	Ile	Tyr	Glu	Arg	Lys	Phe	Gly	Lys	Ser	Val	Ala	Val	Val	Ala	Val		
			420					425					430				
Asn	Arg	Asn	Leu	Thr	Thr	Pro	Thr	Ser	Ile	Thr	Asn	Leu	Asn	Thr	Ser		
		435					440					445					
Leu	Pro	Ser	Gly	Thr	Tyr	Thr	Asp	Val	Leu	Gly	Gly	Val	Leu	Asn	Gly		
		450				455					460						
Asn	Asn	Ile	Thr	Ser	Ser	Gly	Gly	Asn	Ile	Ser	Ser	Phe	Thr	Leu	Ala		
465				470					475						480		
Ala	Gly	Ala	Thr	Ala	Val	Trp	Gln	Tyr	Thr	Ala	Ser	Glu	Thr	Thr	Pro		
			485					490						495			
Thr	Ile	Gly	His	Val	Gly	Pro	Val	Met	Gly	Lys	Pro	Gly	Asn	Val	Val		
			500					505					510				
Thr	Ile	Asp	Gly	Arg	Gly	Phe	Gly	Ser	Thr	Lys	Gly	Thr	Val	Tyr	Phe		
		515					520					525					
Gly	Thr	Thr	Ala	Val	Thr	Gly	Ser	Ala	Ile	Thr	Ser	Trp	Glu	Asp	Thr		
		530				535					540						
Gln	Ile	Lys	Val	Thr	Ile	Pro	Pro	Val	Ala	Gly	Gly	Asp	Tyr	Ala	Val		
545				550					555						560		
Lys	Val	Ala	Ala	Asn	Gly	Val	Asn	Ser	Asn	Ala	Tyr	Asn	Asp	Phe	Thr		
			565					570						575			
Ile	Leu	Ser	Gly	Asp	Gln	Val	Ser	Val	Arg	Phe	Val	Ile	Asn	Asn	Ala		
			580				585						590				
Thr	Thr	Ala	Leu	Gly	Glu	Asn	Ile	Tyr	Leu	Thr	Gly	Asn	Val	Ser	Glu		
		595				600						605					
Leu	Gly	Asn	Trp	Thr	Thr	Gly	Ala	Ala	Ser	Ile	Gly	Pro	Ala	Phe	Asn		
		610				615					620						
Gln	Val	Ile	His	Ala	Tyr	Pro	Thr	Trp	Tyr	Tyr	Asp	Val	Ser	Val	Pro		
625				630					635						640		
Ala	Gly	Lys	Gln	Leu	Glu	Phe	Lys	Phe	Phe	Lys	Lys	Asn	Gly	Ala	Thr		
			645					650						655			
Ile	Thr	Trp	Glu	Gly	Gly	Ser	Asn	His	Thr	Phe	Thr	Thr	Pro	Thr	Ser		
			660				665						670				
Gly	Thr	Ala	Thr	Val	Thr	Val	Asn	Trp	Gln								
		675					680										

<210> 2

<211> 683

<212> PRT

<213> Paenibacillus pabuli

<400> 2

Asp	Thr	Ala	Val	Thr	Asn	Lys	Gln	Asn	Phe	Ser	Thr	Asp	Val	Ile	Tyr
1				5					10					15	
Gln	Val	Phe	Thr	Asp	Arg	Phe	Leu	Asp	Gly	Asn	Pro	Ser	Asn	Asn	Pro
			20					25					30		
Thr	Gly	Ala	Ala	Tyr	Asp	Ser	Ser	Cys	Thr	Asn	Leu	Lys	Leu	Tyr	Cys
		35					40					45			
Gly	Gly	Asp	Trp	Gln	Gly	Leu	Val	Asn	Lys	Ile	Asn	Asp	Asn	Tyr	Phe
		50				55					60				
Thr	Asp	Leu	Gly	Ile	Thr	Ala	Leu	Trp	Ile	Ser	Gln	Pro	Val	Glu	Asn
		65			70					75				80	
Ile	Tyr	Ser	Leu	Ile	Asn	Tyr	Ser	Gly	Val	Asn	Asn	Ser	Ala	Tyr	His
			85						90					95	
Gly	Tyr	Trp	Ala	Arg	Asp	Phe	Lys	Lys	Thr	Asn	Pro	Ala	Phe	Gly	Thr
		100						105					110		
Met	Thr	Asp	Phe	Gln	Asn	Leu	Ile	Asn	Thr	Ala	His	Ala	Lys	Gly	Ile
		115					120					125			
Lys	Val	Ile	Ile	Asp	Phe	Ala	Pro	Asn	His	Thr	Ser	Pro	Ala	Met	Glu
		130				135					140				
Thr	Asp	Thr	Ser	Phe	Ala	Glu	Asn	Gly	Lys	Leu	Tyr	Asn	Asn	Gly	Thr
				150						155					160
Leu	Leu	Gly	Gly	Tyr	Thr	Asn	Asp	Thr	Asn	Lys	Phe	Phe	His	His	Asn
				165				170						175	
Gly	Gly	Ser	Asp	Phe	Ser	Ser	Leu	Glu	Asn	Gly	Ile	Tyr	Lys	Asn	Leu
			180					185					190		
Tyr	Asp	Leu	Ala	Asp	Leu	Asn	His	Asn	Asn	Ser	Thr	Ile	Asp	Thr	Tyr
		195					200					205			

Sequence-Listing

Phe	Lys	Asp	Ala	Ile	Lys	Leu	Trp	Leu	Asp	Met	Gly	Ile	Asp	Gly	Ile
210	210					215					220				
Arg	Val	Asp	Ala	Val	Lys	His	Met	Pro	Met	Gly	Trp	Gln	Lys	Asn	Trp
225					230					235					240
Met	Ser	Ser	Ile	Tyr	Gly	Tyr	Lys	Pro	Val	Phe	Thr	Phe	Gly	Glu	Trp
				245					250					255	
Phe	Leu	Gly	Ser	Ser	Ala	Ser	Asp	Ala	Asp	Asn	Thr	Asn	Phe	Ala	Asn
			260					265					270		
Gln	Ser	Gly	Met	Ser	Leu	Leu	Asp	Phe	Arg	Phe	Asn	Asn	Glu	Val	Arg
		275					280					285			
Asn	Val	Phe	Arg	Asp	Asn	Thr	Ser	Thr	Met	Val	Ala	Leu	Asp	Ser	Met
	290					295					300				
Ile	Thr	Ser	Thr	Ala	Ala	Asp	Tyr	Ala	Gln	Val	Asn	Asp	Gln	Val	Thr
305					310					315					320
Phe	Ile	Asp	Asn	His	Asp	Met	Asp	Arg	Phe	Lys	Thr	Ser	Ala	Val	Asn
				325					330					335	
Asn	Arg	Arg	Leu	Glu	Gln	Ala	Leu	Ala	Phe	Thr	Leu	Thr	Ser	Arg	Gly
			340					345					350		
Val	Pro	Ala	Ile	Tyr	Tyr	Gly	Thr	Glu	Gln	Tyr	Leu	Thr	Gly	Asn	Gly
		355					360					365			
Asp	Pro	Asp	Asn	Arg	Ala	Lys	Met	Pro	Ser	Phe	Ser	Lys	Thr	Thr	Thr
	370					375					380				
Ala	Phe	Asn	Val	Ile	Ser	Lys	Leu	Ala	Pro	Leu	Arg	Lys	Thr	Asn	Pro
385					390					395					400
Ala	Ile	Ala	Tyr	Gly	Thr	Thr	Gln	Gln	Arg	Trp	Ile	Asn	Asn	Asp	Val
				405					410					415	
Tyr	Val	Tyr	Glu	Arg	Lys	Phe	Gly	Asn	Asn	Val	Ala	Val	Val	Ala	Val
			420					425					430		
Asn	Arg	Asn	Leu	Ser	Thr	Pro	Thr	Ser	Ile	Ser	Gly	Leu	Thr	Thr	Ser
		435					440					445			
Leu	Pro	Ser	Gly	Thr	Tyr	Asn	Asp	Val	Leu	Ala	Gly	Ala	Leu	Ser	Gly
	450					455					460				
Asn	Asn	Ile	Thr	Ser	Thr	Gly	Gly	Asn	Val	Ala	Asn	Phe	Thr	Leu	Ala
465					470					475					480
Ala	Gly	Ala	Thr	Ala	Val	Trp	Gln	Tyr	Thr	Ala	Asn	Thr	Thr	Thr	Pro
				485					490					495	
Thr	Ile	Gly	His	Val	Gly	Pro	Thr	Met	Gly	Lys	Ala	Gly	Asn	Thr	Val
			500					505					510		
Thr	Ile	Asp	Gly	Arg	Gly	Phe	Gly	Thr	Thr	Lys	Gly	Thr	Val	Tyr	Phe
		515					520					525			
Gly	Thr	Thr	Ala	Val	Thr	Gly	Ser	Ala	Ile	Thr	Ser	Trp	Glu	Asp	Thr
	530					535					540				
Gln	Ile	Lys	Val	Thr	Ile	Pro	Ala	Val	Ala	Ala	Gly	Asn	Tyr	Ala	Val
545					550				555						560
Lys	Val	Ala	Ala	Gly	Gly	Val	Asn	Ser	Asn	Thr	Tyr	Asn	Asn	Phe	Thr
				565					570					575	
Ile	Leu	Ser	Gly	Asn	Gln	Val	Ser	Val	Arg	Phe	Val	Ile	Asn	Asn	Ala
			580					585					590		
Ser	Thr	Thr	Leu	Gly	Gln	Asn	Leu	Tyr	Leu	Thr	Gly	Asn	Val	Ala	Glu
		595					600					605			
Leu	Gly	Asn	Trp	Ser	Thr	Gly	Pro	Leu	Ala	Ile	Gly	Pro	Ala	Phe	Asn
	610					615					620				
Gln	Val	Ile	Tyr	Ser	Tyr	Pro	Thr	Trp	Tyr	Tyr	Asp	Val	Ser	Val	Pro
625					630					635					640
Ala	Gly	Thr	Asn	Leu	Glu	Phe	Lys	Phe	Phe	Lys	Lys	Asn	Gly	Ser	Thr
				645					650					655	
Ile	Thr	Trp	Glu	Asn	Gly	Asn	Asn	His	Thr	Phe	Thr	Thr	Pro	Thr	Ser
			660					665					670		
Gly	Thr	Ala	Thr	Val	Thr	Val	Asp	Trp	Gln	Pro					
		675					680								

<210> 3
 <211> 682
 <212> PRT
 <213> Paenibacillus amylolyticus

<400> 3
 Asp Thr Ala Val Thr Asn Lys Gln Asn Phe Ser Thr Asp Val Ile Tyr

Sequence-Listing

1				5				10					15			
Gln	Ile	Phe	Thr	Asp	Arg	Phe	Leu	Asp	Gly	Asn	Pro	Ser	Asn	Asn	Pro	
			20					25					30			
Thr	Gly	Ala	Ala	Tyr	Asp	Ala	Thr	Cys	Ser	Asn	Leu	Lys	Leu	Tyr	Cys	
		35					40					45				
Gly	Gly	Asp	Trp	Gln	Gly	Leu	Ile	Asn	Lys	Ile	Asn	Asp	Asn	Tyr	Phe	
	50					55					60					
Ser	Asp	Leu	Gly	Val	Thr	Ala	Leu	Trp	Ile	Ser	Gln	Pro	Val	Glu	Asn	
65					70					75					80	
Ile	Phe	Ala	Thr	Ile	Asn	Tyr	Gly	Gly	Val	Ile	Asn	Thr	Ala	Tyr	His	
			85						90					95		
Gly	Tyr	Trp	Ala	Arg	Asp	Phe	Lys	Lys	Thr	Asn	Pro	Tyr	Phe	Gly	Thr	
			100					105					110			
Met	Ala	Asp	Phe	Gln	Asn	Leu	Ile	Thr	Thr	Ala	His	Ala	Lys	Gly	Ile	
		115					120					125				
Lys	Ile	Val	Ile	Asp	Phe	Ala	Pro	Asn	His	Thr	Ser	Pro	Ala	Met	Glu	
	130					135					140					
Thr	Asp	Thr	Ser	Phe	Ala	Glu	Asn	Gly	Lys	Leu	Tyr	Asp	Asn	Gly	Asn	
145					150					155					160	
Leu	Val	Gly	Gly	Tyr	Thr	Asn	Asp	Thr	Asn	Gly	Tyr	Phe	His	His	Asn	
				165					170					175		
Gly	Gly	Ser	Asp	Phe	Ser	Ser	Leu	Glu	Asn	Gly	Ile	Tyr	Lys	Asn	Leu	
			180					185					190			
Tyr	Asp	Leu	Ala	Asp	Leu	Asn	His	Asn	Asn	Ser	Thr	Ile	Asp	Gln	Tyr	
		195					200					205				
Phe	Lys	Asp	Ala	Ile	Lys	Leu	Trp	Leu	Asp	Met	Gly	Val	Asp	Gly	Ile	
	210					215					220					
Arg	Val	Asp	Ala	Val	Lys	His	Met	Pro	Leu	Gly	Trp	Gln	Lys	Ser	Trp	
225					230					235					240	
Met	Ser	Ser	Ile	Tyr	Ala	His	Lys	Pro	Val	Phe	Thr	Phe	Gly	Glu	Trp	
				245					250					255		
Phe	Leu	Gly	Ser	Ala	Ala	Ser	Asp	Ala	Asp	Asn	Thr	Glu	Phe	Ala	Asn	
			260					265					270			
Glu	Ser	Gly	Met	Ser	Leu	Leu	Asp	Phe	Arg	Phe	Asn	Ser	Ala	Val	Arg	
		275					280					285				
Asp	Val	Phe	Arg	Asp	Asn	Thr	Ser	Asn	Met	Tyr	Ala	Leu	Asp	Ser	Met	
	290					295					300					
Ile	Thr	Gly	Thr	Ala	Ala	Asp	Tyr	Asn	Gln	Val	Asn	Asp	Gln	Val	Thr	
305					310					315					320	
Phe	Ile	Asp	Asn	His	Asp	Met	Asp	Arg	Phe	Lys	Thr	Ser	Ala	Val	Asn	
				325					330					335		
Asn	Arg	Arg	Leu	Glu	Gln	Ala	Leu	Ala	Phe	Thr	Leu	Thr	Ser	Arg	Gly	
			340					345					350			
Val	Pro	Ala	Ile	Tyr	Tyr	Gly	Thr	Glu	Gln	Tyr	Leu	Thr	Gly	Asn	Gly	
		355					360					365				
Asp	Pro	Asp	Asn	Arg	Ala	Lys	Met	Pro	Ser	Phe	Ser	Lys	Thr	Thr	Thr	
	370					375					380					
Ala	Phe	Asn	Val	Ile	Ser	Lys	Leu	Ala	Pro	Leu	Arg	Lys	Ser	Asn	Pro	
385					390					395					400	
Ala	Ile	Ala	Tyr	Gly	Ser	Thr	Gln	Gln	Arg	Trp	Ile	Asn	Asn	Asp	Val	
				405				410						415		
Tyr	Val	Tyr	Glu	Arg	Lys	Phe	Gly	Lys	Ser	Val	Ala	Val	Val	Ala	Val	
			420					425					430			
Asn	Arg	Asn	Leu	Ser	Thr	Pro	Ala	Ser	Ile	Ala	Asn	Leu	Ser	Thr	Ser	
		435					440					445				
Leu	Pro	Thr	Gly	Asn	Tyr	Thr	Asp	Val	Leu	Gly	Gly	Ala	Leu	Asn	Gly	
	450					455					460					
Ser	Asn	Ile	Thr	Ser	Thr	Asn	Gly	Asn	Val	Ser	Ser	Phe	Thr	Leu	Ala	
465					470					475					480	
Ala	Gly	Ala	Thr	Ala	Val	Trp	Gln	Tyr	Thr	Thr	Ser	Glu	Thr	Thr	Pro	
				485					490					495		
Thr	Ile	Gly	His	Val	Gly	Pro	Val	Met	Gly	Lys	Pro	Gly	Asn	Val	Val	
			500					505					510			
Thr	Ile	Ser	Gly	Arg	Gly	Phe	Gly	Ser	Thr	Lys	Gly	Thr	Val	Tyr	Phe	
	515						520					525				
Gly	Thr	Thr	Ala	Val	Thr	Gly	Ala	Ala	Ile	Thr	Ser	Trp	Glu	Asp	Thr	
	530					535				540						
Gln	Ile	Lys	Val	Thr	Ile	Pro	Ala	Val	Ala	Ala	Gly	Asn	Tyr	Ala	Val	

Sequence-Listing

545	Lys	Val	Ala	Ala	Asn	Gly	Val	Asn	Ser	Asn	Ala	Tyr	Asn	Asn	Phe	Thr	560
					565					570						575	
	Ile	Leu	Thr	Gly	Asp	Gln	Val	Thr	Val	Arg	Phe	Val	Ile	Asn	Asn	Ala	
				580					585					590			
	Ser	Thr	Thr	Leu	Gly	Gln	Asn	Ile	Tyr	Leu	Thr	Gly	Asn	Val	Ala	Glu	
				595				600					605				
	Leu	Gly	Asn	Trp	Ser	Thr	Gly	Thr	Thr	Ala	Ile	Gly	Pro	Ala	Phe	Asn	
		610					615					620					
	Gln	Val	Ile	His	Ala	Tyr	Pro	Thr	Trp	Tyr	Tyr	Asp	Val	Ser	Val	Pro	
					630						635					640	
	Ala	Gly	Lys	Gln	Leu	Glu	Phe	Lys	Phe	Phe	Lys	Lys	Asn	Gly	Ala	Thr	
					645					650					655		
	Ile	Thr	Trp	Glu	Gly	Gly	Ser	Asn	His	Thr	Phe	Thr	Thr	Pro	Thr	Ser	
			660					665						670			
	Gly	Thr	Ala	Thr	Val	Asn	Val	Asn	Trp	Gln							
			675					680									

<210> 4

<211> 680

<212> PRT

<213> Geobacillus stearothermophilus

<400> 4

Ala	Gly	Asn	Leu	Asn	Lys	Val	Asn	Phe	Thr	Ser	Asp	Val	Val	Tyr	Gln	
				5					10					15		
Ile	Val	Val	Asp	Arg	Phe	Val	Asp	Gly	Asn	Thr	Ser	Asn	Asn	Pro	Ser	
			20					25					30			
Gly	Ala	Leu	Phe	Ser	Ser	Gly	Cys	Thr	Asn	Leu	Arg	Lys	Tyr	Cys	Gly	
		35					40					45				
Gly	Asp	Trp	Gln	Gly	Ile	Ile	Asn	Lys	Ile	Asn	Asp	Gly	Tyr	Leu	Thr	
		50				55					60					
Asp	Met	Gly	Val	Thr	Ala	Ile	Trp	Ile	Ser	Gln	Pro	Val	Glu	Asn	Val	
		65			70					75				80		
Phe	Ser	Val	Met	Asn	Asp	Ala	Ser	Gly	Ser	Ala	Ser	Tyr	His	Gly	Tyr	
				85					90					95		
Trp	Ala	Arg	Asp	Phe	Lys	Lys	Pro	Asn	Pro	Phe	Phe	Gly	Thr	Leu	Ser	
			100					105					110			
Asp	Phe	Gln	Arg	Leu	Val	Asp	Ala	Ala	His	Ala	Lys	Gly	Ile	Lys	Val	
		115					120					125				
Ile	Ile	Asp	Phe	Ala	Pro	Asn	His	Thr	Ser	Pro	Ala	Ser	Glu	Thr	Asn	
		130				135					140					
Pro	Ser	Tyr	Met	Glu	Asn	Gly	Arg	Leu	Tyr	Asp	Asn	Gly	Thr	Leu	Leu	
		145			150					155				160		
Gly	Gly	Tyr	Thr	Asn	Asp	Ala	Asn	Met	Tyr	Phe	His	His	Asn	Gly	Gly	
				165					170					175		
Thr	Thr	Phe	Ser	Ser	Leu	Glu	Asp	Gly	Ile	Tyr	Arg	Asn	Leu	Phe	Asp	
		180						185					190			
Leu	Ala	Asp	Leu	Asn	His	Gln	Asn	Pro	Val	Ile	Asp	Arg	Tyr	Leu	Lys	
		195					200					205				
Asp	Ala	Val	Lys	Met	Trp	Ile	Asp	Met	Gly	Ile	Asp	Gly	Ile	Arg	Met	
		210				215					220					
Asp	Ala	Val	Lys	His	Met	Pro	Phe	Gly	Trp	Gln	Lys	Ser	Leu	Met	Asp	
		225			230					235				240		
Glu	Ile	Asp	Asn	Tyr	Arg	Pro	Val	Phe	Thr	Phe	Gly	Glu	Trp	Phe	Leu	
				245					250					255		
Ser	Glu	Asn	Glu	Val	Asp	Ala	Asn	Asn	His	Tyr	Phe	Ala	Asn	Glu	Ser	
		260						265					270			
Gly	Met	Ser	Leu	Leu	Asp	Phe	Arg	Phe	Gly	Gln	Lys	Leu	Arg	Gln	Val	
		275					280					285				
Leu	Arg	Asn	Asn	Ser	Asp	Asn	Trp	Tyr	Gly	Phe	Asn	Gln	Met	Ile	Gln	
		290				295					300					
Asp	Thr	Ala	Ser	Ala	Tyr	Asp	Glu	Val	Leu	Asp	Gln	Val	Thr	Phe	Ile	
		305			310					315				320		
Asp	Asn	His	Asp	Met	Asp	Arg	Phe	Met	Ile	Asp	Gly	Gly	Asp	Pro	Arg	
				325					330					335		
Lys	Val	Asp	Met	Ala	Leu	Ala	Val	Leu	Leu	Thr	Ser	Arg	Gly	Val	Pro	
			340					345					350			

Sequence-Listing

Asn	Ile	Tyr	Tyr	Gly	Thr	Glu	Gln	Tyr	Met	Thr	Gly	Asn	Gly	Asp	Pro
	355						360					365			
Asn	Asn	Arg	Lys	Met	Met	Ser	Ser	Phe	Asn	Lys	Asn	Thr	Arg	Ala	Tyr
	370					375					380				
Gln	Val	Ile	Gln	Lys	Leu	Ser	Ser	Leu	Arg	Arg	Asn	Asn	Pro	Ala	Leu
	385				390					395					400
Ala	Tyr	Gly	Asp	Thr	Glu	Gln	Arg	Trp	Ile	Asn	Gly	Asp	Val	Tyr	Val
			405						410					415	
Tyr	Glu	Arg	Gln	Phe	Gly	Lys	Asp	Val	Val	Leu	Val	Ala	Val	Asn	Arg
			420					425					430		
Ser	Ser	Ser	Ser	Asn	Tyr	Ser	Ile	Thr	Gly	Leu	Phe	Thr	Ala	Leu	Pro
		435					440					445			
Ala	Gly	Thr	Tyr	Thr	Asp	Gln	Leu	Gly	Gly	Leu	Leu	Asp	Gly	Asn	Thr
	450					455					460				
Ile	Gln	Val	Gly	Ser	Asn	Gly	Ser	Val	Asn	Ala	Phe	Asp	Leu	Gly	Pro
	465				470					475					480
Gly	Glu	Val	Gly	Val	Trp	Ala	Tyr	Ser	Ala	Thr	Glu	Ser	Thr	Pro	Ile
				485					490					495	
Ile	Gly	His	Val	Gly	Pro	Met	Met	Gly	Gln	Val	Gly	His	Gln	Val	Thr
			500					505					510		
Ile	Asp	Gly	Glu	Gly	Phe	Gly	Thr	Asn	Thr	Gly	Thr	Val	Lys	Phe	Gly
		515					520					525			
Thr	Thr	Ala	Ala	Asn	Val	Val	Ser	Trp	Ser	Asn	Asn	Gln	Ile	Val	Val
	530					535					540				
Ala	Val	Pro	Asn	Val	Ser	Pro	Gly	Lys	Tyr	Asn	Ile	Thr	Val	Gln	Ser
	545				550					555					560
Ser	Ser	Gly	Gln	Thr	Ser	Ala	Ala	Tyr	Asp	Asn	Phe	Glu	Val	Leu	Thr
				565					570					575	
Asn	Asp	Gln	Val	Ser	Val	Arg	Phe	Val	Val	Asn	Asn	Ala	Thr	Thr	Asn
			580					585					590		
Leu	Gly	Gln	Asn	Ile	Tyr	Ile	Val	Gly	Asn	Val	Tyr	Glu	Leu	Gly	Asn
		595					600					605			
Trp	Asp	Thr	Ser	Lys	Ala	Ile	Gly	Pro	Met	Phe	Asn	Gln	Val	Val	Tyr
	610					615					620				
Ser	Tyr	Pro	Thr	Trp	Tyr	Ile	Asp	Val	Ser	Val	Pro	Glu	Gly	Lys	Thr
	625				630					635					640
Ile	Glu	Phe	Lys	Phe	Ile	Lys	Lys	Asp	Ser	Gln	Gly	Asn	Val	Thr	Trp
				645					650					655	
Glu	Ser	Gly	Ser	Asn	His	Val	Tyr	Thr	Thr	Pro	Thr	Asn	Thr	Thr	Gly
			660					665					670		
Lys	Ile	Ile	Val	Asp	Trp	Gln	Asn								
		675					680								

<210> 5
 <211> 686
 <212> PRT
 <213> Bacillus macerans

<400> 5

Ser	Pro	Asp	Thr	Ser	Val	Asn	Asn	Lys	Leu	Asn	Phe	Ser	Thr	Asp	Thr
1				5					10					15	
Val	Tyr	Gln	Ile	Val	Thr	Asp	Arg	Phe	Val	Asp	Gly	Asn	Ser	Ala	Asn
			20					25					30		
His	Pro	Thr	Gly	Ala	Ala	Phe	Ser	Asp	His	Ser	Asn	Leu	Lys	Leu	
		35					40				45				
Tyr	Phe	Gly	Gly	Asp	Trp	Gln	Gly	Ile	Thr	Asn	Lys	Ile	Asn	Asp	Gly
	50					55					60				
Tyr	Leu	Thr	Gly	Met	Gly	Ile	Thr	Ala	Leu	Trp	Ile	Ser	Gln	Pro	Val
	65				70					75				80	
Glu	Asn	Ile	Thr	Ala	Val	Ile	Lys	Tyr	Ser	Gly	Val	Asn	Asn	Thr	Ala
				85					90					95	
Tyr	His	Gly	Tyr	Trp	Pro	Arg	Asp	Phe	Lys	Lys	Thr	Asn	Ala	Ala	Phe
			100					105					110		
Gly	Ser	Phe	Thr	Asp	Phe	Ser	Asn	Leu	Ile	Ala	Ala	Ala	His	Ser	His
		115					120					125			
Asn	Ile	Lys	Val	Val	Met	Asp	Phe	Ala	Pro	Asn	His	Thr	Asn	Pro	Ala
	130					135					140				
Ser	Ser	Thr	Asp	Pro	Ser	Phe	Ala	Glu	Asn	Gly	Ala	Leu	Tyr	Asn	Asn

145	Sequence-Listing													160	
Gly	Thr	Leu	Leu	Gly	Lys	Tyr	Ser	Asn	Asp	Thr	Ala	Gly	Leu	Phe	His
His	Asn	Gly	Gly	Thr	Asp	Phe	Ser	Thr	Thr	Glu	Ser	Gly	Ile	Tyr	Lys
Asn	Leu	Tyr	Asp	Leu	Ala	Asp	Ile	Asn	Gln	Asn	Asn	Asn	Thr	Ile	Asp
Ser	Tyr	Leu	Lys	Glu	Ser	Ile	Gln	Leu	Trp	Leu	Asn	Leu	Gly	Val	Asp
Gly	Ile	Arg	Phe	Asp	Ala	Val	Lys	His	Met	Pro	Gln	Gly	Trp	Gln	Lys
Ser	Tyr	Val	Ser	Ser	Ile	Tyr	Ser	Ser	Ala	Asn	Pro	Val	Phe	Thr	Phe
Gly	Glu	Trp	Phe	Leu	Gly	Pro	Asp	Glu	Met	Thr	Gln	Asp	Asn	Ile	Asn
Phe	Ala	Asn	Gln	Ser	Gly	Met	His	Leu	Leu	Asp	Phe	Ala	Phe	Ala	Gln
Glu	Ile	Arg	Glu	Val	Phe	Arg	Asp	Lys	Ser	Glu	Thr	Met	Thr	Asp	Leu
Asn	Ser	Val	Ile	Ser	Ser	Thr	Gly	Ser	Ser	Tyr	Asn	Tyr	Ile	Asn	Asn
Met	Val	Thr	Phe	Ile	Asp	Asn	His	Asp	Met	Asp	Arg	Phe	Gln	Gln	Ala
Gly	Ala	Ser	Thr	Arg	Pro	Thr	Glu	Gln	Ala	Leu	Ala	Val	Thr	Leu	Thr
Ser	Arg	Gly	Val	Pro	Ala	Ile	Tyr	Tyr	Gly	Thr	Glu	Gln	Tyr	Met	Thr
Gly	Asn	Gly	Asp	Pro	Asn	Asn	Arg	Gly	Met	Met	Thr	Gly	Phe	Asp	Thr
Asn	Lys	Thr	Ala	Tyr	Lys	Val	Ile	Lys	Ala	Leu	Ala	Pro	Leu	Arg	Lys
Ser	Asn	Pro	Ala	Leu	Ala	Tyr	Gly	Ser	Thr	Thr	Gln	Arg	Trp	Val	Asn
Ser	Asp	Val	Tyr	Val	Tyr	Glu	Arg	Lys	Phe	Gly	Ser	Asn	Val	Ala	Leu
Val	Ala	Val	Asn	Arg	Ser	Ser	Thr	Thr	Ala	Tyr	Pro	Ile	Ser	Gly	Ala
Leu	Thr	Ala	Leu	Pro	Asn	Gly	Thr	Tyr	Thr	Asp	Val	Leu	Gly	Gly	Leu
Leu	Asn	Gly	Asn	Ser	Ile	Thr	Val	Asn	Gly	Gly	Thr	Val	Ser	Asn	Phe
Thr	Leu	Ala	Ala	Gly	Gly	Thr	Ala	Val	Trp	Gln	Tyr	Thr	Thr	Thr	Glu
Ser	Ser	Pro	Ile	Ile	Gly	Asn	Val	Gly	Pro	Thr	Met	Gly	Lys	Pro	Gly
Asn	Thr	Ile	Thr	Ile	Asp	Gly	Arg	Gly	Phe	Gly	Thr	Thr	Lys	Asn	Lys
Val	Thr	Phe	Gly	Thr	Thr	Ala	Val	Thr	Gly	Ala	Asn	Ile	Val	Ser	Trp
Glu	Asp	Thr	Glu	Ile	Lys	Val	Lys	Val	Pro	Asn	Val	Ala	Ala	Gly	Asn
Thr	Ala	Val	Thr	Val	Thr	Asn	Ala	Ala	Gly	Thr	Ser	Ala	Ala	Ala	Phe
Asn	Asn	Phe	Asn	Val	Leu	Thr	Ala	Asp	Gln	Val	Thr	Val	Arg	Phe	Lys
Val	Asn	Asn	Ala	Thr	Thr	Ala	Leu	Gly	Gln	Asn	Val	Tyr	Leu	Thr	Gly
Asn	Val	Ala	Glu	Leu	Gly	Asn	Trp	Thr	Ala	Ala	Asn	Ala	Ile	Gly	Pro
Met	Tyr	Asn	Gln	Val	Glu	Ala	Ser	Tyr	Pro	Thr	Trp	Tyr	Phe	Asp	Val
Ser	Val	Pro	Ala	Asn	Thr	Ala	Leu	Gln	Phe	Lys	Phe	Ile	Lys	Val	Asn
Gly	Ser	Thr	Val	Thr	Trp	Glu	Gly	Gly	Asn	Asn	His	Thr	Phe	Thr	Ser
Pro	Ser	Ser	Gly	Val	Ala	Thr	Val	Thr	Val	Asp	Trp	Gln	Asn		

Sequence-Listing

<210> 6
 <211> 681
 <212> PRT
 <213> Thermoanaerobacter thermosulfurigenes

<400> 6
 Ala Pro Asp Thr Ser Val Ser Asn Val Val Asn Tyr Ser Thr Asp Val
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 Ile Tyr Gln Ile Val Thr Asp Arg Phe Leu Asp Gly Asn Pro Ser Asn
 20 25 30
 Asn Pro Thr Gly Asp Leu Tyr Asp Pro Thr His Thr Ser Leu Lys Lys
 35 40 45
 Tyr Phe Gly Gly Asp Trp Gln Gly Ile Ile Asn Lys Ile Asn Asp Gly
 50 55 60
 Tyr Leu Thr Gly Met Gly Ile Thr Ala Ile Trp Ile Ser Gln Pro Val
 65 70 75 80
 Glu Asn Ile Tyr Ala Val Leu Pro Asp Ser Thr Phe Gly Gly Ser Thr
 85 90 95
 Ser Tyr His Gly Tyr Trp Ala Arg Asp Phe Lys Lys Pro Asn Pro Phe
 100 105 110
 Phe Gly Ser Phe Thr Asp Phe Gln Asn Leu Ile Ala Thr Ala His Ala
 115 120 125
 His Asn Ile Lys Val Ile Ile Asp Phe Ala Pro Asn His Thr Ser Pro
 130 135 140
 Ala Ser Glu Thr Asp Pro Thr Tyr Gly Glu Asn Gly Arg Leu Tyr Asp
 145 150 155 160
 Asn Gly Val Leu Leu Gly Gly Tyr Thr Asn Asp Thr Asn Gly Tyr Phe
 165 170 175
 His His Tyr Gly Gly Thr Asn Phe Ser Ser Tyr Glu Asp Gly Ile Tyr
 180 185 190
 Arg Asn Leu Phe Asp Leu Ala Asp Leu Asp Gln Gln Asn Ser Thr Ile
 195 200 205
 Asp Ser Tyr Leu Lys Ala Ala Ile Lys Leu Trp Leu Asp Met Gly Ile
 210 215 220
 Asp Gly Ile Arg Met Asp Ala Val Lys His Met Ala Phe Gly Trp Gln
 225 230 235 240
 Lys Asn Phe Met Asp Ser Ile Leu Ser Tyr Arg Pro Val Phe Thr Phe
 245 250 255
 Gly Glu Trp Tyr Leu Gly Thr Asn Glu Val Asp Pro Asn Asn Thr Tyr
 260 265 270
 Phe Ala Asn Glu Ser Gly Met Ser Leu Leu Asp Phe Arg Phe Ala Gln
 275 280 285
 Lys Val Arg Gln Val Phe Arg Asp Asn Thr Asp Thr Met Tyr Gly Leu
 290 295 300
 Asp Ser Met Ile Gln Ser Thr Ala Ala Asp Tyr Asn Phe Ile Asn Asp
 305 310 315 320
 Met Val Thr Phe Ile Asp Asn His Asp Met Asp Arg Phe Tyr Thr Gly
 325 330 335
 Gly Ser Thr Arg Pro Val Glu Gln Ala Leu Ala Phe Thr Leu Thr Ser
 340 345 350
 Arg Gly Val Pro Ala Ile Tyr Tyr Gly Thr Glu Gln Tyr Met Thr Gly
 355 360 365
 Asn Gly Asp Pro Tyr Asn Arg Ala Met Met Thr Ser Phe Asp Thr Thr
 370 375 380
 Thr Thr Ala Tyr Asn Val Ile Lys Lys Leu Ala Pro Leu Arg Lys Ser
 385 390 395 400
 Asn Pro Ala Ile Ala Tyr Gly Thr Gln Lys Gln Arg Trp Ile Asn Asn
 405 410 415
 Asp Val Tyr Ile Tyr Glu Arg Gln Phe Gly Asn Asn Val Ala Leu Val
 420 425 430
 Ala Ile Asn Arg Asn Leu Ser Thr Ser Tyr Tyr Ile Thr Gly Leu Tyr
 435 440 445
 Thr Ala Leu Pro Ala Gly Thr Tyr Ser Asp Met Leu Gly Gly Leu Leu
 450 455 460
 Asn Gly Ser Ser Ile Thr Val Ser Ser Asn Gly Ser Val Thr Pro Phe
 465 470 475 480
 Thr Leu Ala Pro Gly Glu Val Ala Val Trp Gln Tyr Val Ser Thr Thr
 485 490 495

Sequence-Listing

Asn	Pro	Pro	Leu	Ile	Gly	His	Val	Gly	Pro	Thr	Met	Thr	Lys	Ala	Gly
			500					505					510		
Gln	Thr	Ile	Thr	Ile	Asp	Gly	Arg	Gly	Phe	Gly	Thr	Thr	Ala	Gly	Gln
		515					520					525			
Val	Leu	Phe	Gly	Thr	Thr	Pro	Ala	Thr	Ile	Val	Ser	Trp	Glu	Asp	Thr
	530					535					540				
Glu	Val	Lys	Val	Lys	Val	Pro	Ala	Leu	Thr	Pro	Gly	Lys	Tyr	Asn	Ile
	545				550					555					560
Thr	Leu	Lys	Thr	Ala	Ser	Gly	Val	Thr	Ser	Asn	Ser	Tyr	Asn	Asn	Ile
			565						570					575	
Asn	Val	Leu	Thr	Gly	Asn	Gln	Val	Cys	Val	Arg	Phe	Val	Val	Asn	Asn
			580					585					590		
Ala	Thr	Thr	Val	Trp	Gly	Glu	Asn	Val	Tyr	Leu	Thr	Gly	Asn	Val	Ala
		595					600					605			
Glu	Leu	Gly	Asn	Trp	Asp	Thr	Ser	Lys	Ala	Ile	Gly	Pro	Met	Phe	Asn
	610					615					620				
Gln	Val	Val	Tyr	Gln	Tyr	Pro	Thr	Trp	Tyr	Tyr	Asp	Val	Ser	Val	Pro
	625				630					635					640
Ala	Gly	Thr	Thr	Ile	Glu	Phe	Ile	Lys	Lys	Asn	Gly	Ser	Thr	Val	Thr
				645					650					655	
Trp	Glu	Gly	Gly	Tyr	Asn	His	Val	Tyr	Thr	Thr	Pro	Thr	Ser	Gly	Thr
			660					665					670		
Ala	Thr	Val	Ile	Val	Asp	Trp	Gln	Pro							
		675					680								

<210> 7
 <211> 2154
 <212> DNA
 <213> Paenibacillus illinoisensis

<400> 7	
atg ttt caa atg gcc aaa cgc gtt ctc ctc agt acc acg cta acg ttc	48
Met Phe Gln Met Ala Lys Arg Val Leu Leu Ser Thr Thr Leu Thr Phe	
1 5 10 15	
agc ctg ctt gcc ggc agt gca ttg cct ttc ctg cct gcc tcc gcg att	96
Ser Leu Leu Ala Gly Ser Ala Leu Pro Phe Leu Pro Ala Ser Ala Ile	
20 25 30	
tat gcc gat gcg gat acg gct gtc acc aac aag caa aat ttc agt acc	144
Tyr Ala Asp Ala Asp Thr Ala Val Thr Asn Lys Gln Asn Phe Ser Thr	
35 40 45	
gat gtc atc tat caa gtt ttt acg gac cgg ttt ctg gac ggt aac cca	192
Asp Val Ile Tyr Gln Val Phe Thr Asp Arg Phe Leu Asp Gly Asn Pro	
50 55 60	
tcc aac aac ccc act gga gct gct ttt gat ggc aca tgc agc aac ctg	240
Ser Asn Asn Pro Thr Gly Ala Ala Phe Asp Gly Thr Cys Ser Asn Leu	
65 70 75 80	
aaa ctg tac tgc ggc ggc gac tgg cag ggg ctg att aac aaa atc aat	288
Lys Leu Tyr Cys Gly Gly Asp Trp Gln Gly Leu Ile Asn Lys Ile Asn	
85 90 95	
gac aac tat ttc agt gac ctg ggt gtc aca gcc ctc tgg atc tcc cag	336
Asp Asn Tyr Phe Ser Asp Leu Gly Val Thr Ala Leu Trp Ile Ser Gln	
100 105 110	
cct gtc gaa aat att ttc gct acc atc aac tac agc ggt gta acc aac	384
Pro Val Glu Asn Ile Phe Ala Thr Ile Asn Tyr Ser Gly Val Thr Asn	
115 120 125	
act gct tat cac ggc tat tgg gca cgg gat ttc aag aag acc aat cca	432
Thr Ala Tyr His Gly Tyr Trp Ala Arg Asp Phe Lys Lys Thr Asn Pro	
130 135 140	
tat ttc gga acc atg acc gat ttt cag aat ctg gtg acc tcc gcc cat	480

Sequence-Listing															
Tyr 145	Phe	Gly	Thr	Met	Thr 150	Asp	Phe	Gln	Asn	Leu 155	Val	Thr	Ser	Ala	His 160
gct Ala	aaa Lys	ggc Gly	atc Ile	aaa Lys 165	atc Ile	att Ile	att Ile	gat Asp	ttc Phe 170	gcg Ala	cca Pro	aac Asn	cat His	acg Thr 175	ttc Phe
cct Pro	gcc Ala	atg Met	gaa Glu 180	acc Thr	gat Asp	acc Thr	tcc Ser	ttc Phe 185	gct Ala	gaa Glu	aac Asn	ggc Gly	aaa Lys 190	ctg Leu	tac Tyr
gat Asp	aac Asn	ggc Gly 195	agc Ser	ctg Leu	gtg Val	ggc Gly	ggg Gly 200	tac Tyr	acc Thr	aat Asn	gat Asp	acg Thr 205	aac Asn	gga Gly	tat Tyr
ttt Phe	cac His 210	cac His	aat Asn	ggc Gly	ggc Gly	tcc Ser 215	gat Asp	ttc Phe	tcc Ser	act Thr	ctt Leu 220	gag Glu	aat Asn	ggc Gly	att Ile
tac Tyr 225	aaa Lys	aac Asn	ctc Leu	tac Tyr	gat Asp 230	ctg Leu	gcc Ala	gat Asp	ctg Leu	aat Asn 235	cac His	aat Asn	aac Asn	agc Ser	acg Thr 240
atc Ile	gat Asp	aca Thr	tat Tyr	ttc Phe 245	aaa Lys	gac Asp	gcc Ala	atc Ile	aag Lys 250	ctg Leu	tgg Trp	ctg Leu	gat Asp	atg Met 255	ggc Gly
gtg Val	gac Asp	ggc Gly	atc Ile	cgt Arg 260	gtc Val	gat Asp	gcg Ala	gtc Val	aag Lys 265	cac His	atg Met	cca Pro	cag Gln	gga Gly 270	tgg Trp
cag Gln	aag Lys	aac Asn	tgg Trp 275	atg Met	tca Ser	tcc Ser	atc Ile	tat Tyr 280	gca Ala	cac His	aag Lys	ccg Pro	gta Val 285	ttt Phe	acc Thr
ttc Phe	ggc Gly 290	gaa Glu	tgg Trp	ttc Phe	ctg Leu	gga Gly 295	tct Ser	gct Ala	gca Ala	tcc Ser	gat Asp 300	gcg Ala	gat Asp	aac Asn	aca Thr
gat Asp 305	ttt Phe	gcc Ala	aat Asn	gaa Glu	tcc Ser 310	ggt Gly	atg Met	agt Ser	ttg Leu	ctt Leu 315	gat Asp	ttt Phe	cgt Arg	ttc Phe	aat Asn 320
tcg Ser	gct Ala	gtc Val	cgc Arg	aac Asn 325	gtg Val	ttc Phe	cgg Arg	gat Asp	aac Asn 330	aca Thr	tcc Ser	aac Asn	atg Met	tac Tyr 335	gcg Ala
ctg Leu	gat Asp	tcc Ser	atg Met 340	ctt Leu	acg Thr	gct Ala	acg Thr	gca Ala 345	gca Ala	gat Asp	tac Tyr	aat Asn	caa Gln 350	gtg Val	aat Asn
gac Asp	caa Gln 355	gtc Val	act Thr	ttc Phe	att Ile	gac Asp 360	aac Asn	cat His	gat Asp	atg Met	gac Asp	cgt Arg 365	ttc Phe	aaa Lys	aca Thr
agt Ser	gcg Ala 370	gtg Val	aac Asn	aac Asn	cgc Arg	cgt Arg 375	ctg Leu	gaa Glu	cag Gln	gct Ala	ctg Leu 380	gcc Ala	ttc Phe	acg Thr	ctg Leu
acc Thr 385	tca Ser	cgc Arg	ggc Gly	gta Val	cct Pro 390	gcc Ala	atc Ile	tat Tyr	tat Tyr	ggt Gly 395	acc Thr	gag Glu	cag Gln	tat Tyr	ctg Leu 400
acc Thr	ggg Gly	aat Asn	ggt Gly	gac Asp 405	ccg Pro	gat Asp	aac Asn	cgg Arg	ggc Gly 410	aaa Lys	atg Met	cct Pro	tcc Ser	ttc Phe 415	tcc Ser
aaa	tcg	acc	aca	gcg	ttc	agc	gtg	atc	agc	aag	ctg	gct	cct	ctg	cgc

Sequence-Listing															
Lys	Ser	Thr	Thr	Ala	Phe	Ser	Val	Ile	Ser	Lys	Leu	Ala	Pro	Leu	Arg
			420					425					430		
aaa	tcc	aac	ccg	gcg	att	gcc	tac	ggt	tcc	aca	cag	cag	cgc	tgg	atc
Lys	Ser	Asn	Pro	Ala	Ile	Ala	Tyr	Gly	Ser	Thr	Gln	Gln	Arg	Trp	Ile
		435					440					445			1344
aac	aat	gat	gta	tat	atc	tat	gag	cgc	aag	ttt	ggc	aaa	agc	gtt	gcc
Asn	Asn	Asp	Val	Tyr	Ile	Tyr	Glu	Arg	Lys	Phe	Gly	Lys	Ser	Val	Ala
	450					455					460				1392
gtt	gtt	gcc	gtt	aac	cgc	aat	ctc	acg	acg	cca	acc	agt	atc	acg	aac
Val	Val	Ala	Val	Asn	Arg	Asn	Leu	Thr	Thr	Pro	Thr	Ser	Ile	Thr	Asn
465					470					475					1440
ctg	aat	acg	tcc	ctt	cca	tca	gga	aca	tac	acc	gat	gtg	ctg	ggc	ggc
Leu	Asn	Thr	Ser	Leu	Pro	Ser	Gly	Thr	Tyr	Thr	Asp	Val	Leu	Gly	Gly
				485					490					495	1488
gtg	ctg	aac	gga	aac	aac	att	act	tca	agt	gga	ggc	aac	att	tct	tcc
Val	Leu	Asn	Gly	Asn	Asn	Ile	Thr	Ser	Ser	Gly	Gly	Asn	Ile	Ser	Ser
			500					505					510		1536
ttc	acg	ctc	gca	gca	gga	gct	acc	gct	gtg	tgg	cag	tat	acg	gca	agt
Phe	Thr	Leu	Ala	Ala	Gly	Ala	Thr	Ala	Val	Trp	Gln	Tyr	Thr	Ala	Ser
		515					520					525			1584
gaa	acg	acg	cca	acc	atc	ggt	cac	gtt	ggc	cct	gta	atg	ggt	aaa	cct
Glu	Thr	Thr	Pro	Thr	Ile	Gly	His	Val	Gly	Pro	Val	Met	Gly	Lys	Pro
	530					535					540				1632
ggt	aac	gtc	gtt	acc	atc	gac	gga	cgg	ggg	ttc	ggc	tcc	acc	aaa	ggt
Gly	Asn	Val	Val	Thr	Ile	Asp	Gly	Arg	Gly	Phe	Gly	Ser	Thr	Lys	Gly
545					550					555					560
acc	gtc	tac	ttc	ggt	aca	aca	gcc	ggt	acg	ggc	tct	gcc	atc	act	tca
Thr	Val	Tyr	Phe	Gly	Thr	Thr	Ala	Val	Thr	Gly	Ser	Ala	Ile	Thr	Ser
					565					570					575
tgg	gaa	gac	act	cag	atc	aaa	gtc	acc	att	cca	cca	gta	gca	ggc	ggt
Trp	Glu	Asp	Thr	Gln	Ile	Lys	Val	Thr	Ile	Pro	Pro	Val	Ala	Gly	Gly
			580					585					590		1776
gac	tat	gca	gtg	aaa	gtc	gct	gcc	aat	ggt	gtg	aac	agc	aat	gcc	tat
Asp	Tyr	Ala	Val	Lys	Val	Ala	Ala	Asn	Gly	Val	Asn	Ser	Asn	Ala	Tyr
		595					600					605			1824
aac	gat	ttc	acg	atc	cta	agc	ggc	gat	cag	gtc	tcg	gtt	cgg	ttc	gtc
Asn	Asp	Phe	Thr	Ile	Leu	Ser	Gly	Asp	Gln	Val	Ser	Val	Arg	Phe	Val
	610					615					620				1872
atc	aat	aat	gcc	aca	act	gca	ctg	ggt	gaa	aac	atc	tac	ctg	acg	ggc
Ile	Asn	Asn	Ala	Thr	Thr	Ala	Leu	Gly	Glu	Asn	Ile	Tyr	Leu	Thr	Gly
625					630					635					640
aac	gtg	tcc	gaa	ctg	ggg	aac	tgg	acc	aca	ggt	gcg	gct	tcc	att	gga
Asn	Val	Ser	Glu	Leu	Gly	Asn	Trp	Thr	Thr	Gly	Ala	Ala	Ser	Ile	Gly
				645				650						655	1968
ccg	gct	ttc	aat	cag	gtc	atc	cac	gcc	tac	ccg	act	tgg	tat	tat	gac
Pro	Ala	Phe	Asn	Gln	Val	Ile	His	Ala	Tyr	Pro	Thr	Trp	Tyr	Tyr	Asp
			660					665					670		2016
gta	agt	gtt	cca	gcc	ggg	aaa	cag	ctg	gaa	ttc	aag	ttc	ttc	aag	aaa
Val	Ser	Val	Pro	Ala	Gly	Lys	Gln	Leu	Glu	Phe	Lys	Phe	Phe	Lys	Lys
		675					680					685			2064
aac	ggg	gct	acc	att	acg	tgg	gag	ggt	gga	tcc	aat	cac	acc	ttt	aca
															2112

Sequence-Listing

Asn	Gly	Ala	Thr	Ile	Thr	Trp	Glu	Gly	Gly	Ser	Asn	His	Thr	Phe	Thr	
690						695					700					
aca	ccg	acc	agc	ggg	act	gcc	act	gta	aca	gta	aac	tgg	caa			2154
Thr	Pro	Thr	Ser	Gly	Thr	Ala	Thr	Val	Thr	Val	Asn	Trp	Gln			
705					710					715						

<210> 8
 <211> 30
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> sense primer for PCR

<400> 8
 ttcaccacaa tggccgctcc gatttctcca 30

<210> 9
 <211> 30
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> antisense primer for PCR

<400> 9
 tggagaaatc ggagcggcca ttgtggtgaa 30

<210> 10
 <211> 30
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> sense primer for PCR

<400> 10
 cttccatcag gaacacacac cgatgtgctg 30

<210> 11
 <211> 30
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> antisense primer for PCR

<400> 11
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<210> 12
 <211> 682
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> mutated CGTase(G178R)

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 Gln Val Phe Thr Asp Arg Phe Leu Asp Gly Asn Pro Ser Asn Asn Pro
 20 25 30
 Thr Gly Ala Ala Phe Asp Gly Thr Cys Ser Asn Leu Lys Leu Tyr Cys
 35 40 45
 Gly Gly Asp Trp Gln Gly Leu Ile Asn Lys Ile Asn Asp Asn Tyr Phe

Sequence-Listing

50	55	60													
Ser	Asp	Leu	Gly	Val	Thr	Ala	Leu	Trp	Ile	Ser	Gln	Pro	Val	Glu	Asn
65					70					75					80
Ile	Phe	Ala	Thr	Ile	Asn	Tyr	Ser	Gly	Val	Thr	Asn	Thr	Ala	Tyr	His
				85					90					95	
Gly	Tyr	Trp	Ala	Arg	Asp	Phe	Lys	Lys	Thr	Asn	Pro	Tyr	Phe	Gly	Thr
			100					105					110		
Met	Thr	Asp	Phe	Gln	Asn	Leu	Val	Thr	Ser	Ala	His	Ala	Lys	Gly	Ile
		115				120						125			
Lys	Ile	Ile	Ile	Asp	Phe	Ala	Pro	Asn	His	Thr	Phe	Pro	Ala	Met	Glu
	130					135					140				
Thr	Asp	Thr	Ser	Phe	Ala	Glu	Asn	Gly	Lys	Leu	Tyr	Asp	Asn	Gly	Ser
145					150					155					160
Leu	Val	Gly	Gly	Tyr	Thr	Asn	Asp	Thr	Asn	Gly	Tyr	Phe	His	His	Asn
				165					170					175	
Gly	Arg	Ser	Asp	Phe	Ser	Thr	Leu	Glu	Asn	Gly	Ile	Tyr	Lys	Asn	Leu
			180					185					190		
Tyr	Asp	Leu	Ala	Asp	Leu	Asn	His	Asn	Asn	Ser	Thr	Ile	Asp	Thr	Tyr
		195					200					205			
Phe	Lys	Asp	Ala	Ile	Lys	Leu	Trp	Leu	Asp	Met	Gly	Val	Asp	Gly	Ile
	210					215					220				
Arg	Val	Asp	Ala	Val	Lys	His	Met	Pro	Gln	Gly	Trp	Gln	Lys	Asn	Trp
225					230					235					240
Met	Ser	Ser	Ile	Tyr	Ala	His	Lys	Pro	Val	Phe	Thr	Phe	Gly	Glu	Trp
				245					250					255	
Phe	Leu	Gly	Ser	Ala	Ala	Ser	Asp	Ala	Asp	Asn	Thr	Asp	Phe	Ala	Asn
			260					265					270		
Glu	Ser	Gly	Met	Ser	Leu	Leu	Asp	Phe	Arg	Phe	Asn	Ser	Ala	Val	Arg
		275					280					285			
Asn	Val	Phe	Arg	Asp	Asn	Thr	Ser	Asn	Met	Tyr	Ala	Leu	Asp	Ser	Met
	290					295					300				
Leu	Thr	Ala	Thr	Ala	Ala	Asp	Tyr	Asn	Gln	Val	Asn	Asp	Gln	Val	Thr
305					310					315					320
Phe	Ile	Asp	Asn	His	Asp	Met	Asp	Arg	Phe	Lys	Thr	Ser	Ala	Val	Asn
				325					330					335	
Asn	Arg	Arg	Leu	Glu	Gln	Ala	Leu	Ala	Phe	Thr	Leu	Thr	Ser	Arg	Gly
			340					345					350		
Val	Pro	Ala	Ile	Tyr	Tyr	Gly	Thr	Glu	Gln	Tyr	Leu	Thr	Gly	Asn	Gly
		355					360					365			
Asp	Pro	Asp	Asn	Arg	Gly	Lys	Met	Pro	Ser	Phe	Ser	Lys	Ser	Thr	Thr
	370					375					380				
Ala	Phe	Ser	Val	Ile	Ser	Lys	Leu	Ala	Pro	Leu	Arg	Lys	Ser	Asn	Pro
385					390					395					400
Ala	Ile	Ala	Tyr	Gly	Ser	Thr	Gln	Gln	Arg	Trp	Ile	Asn	Asn	Asp	Val
				405					410					415	
Tyr	Ile	Tyr	Glu	Arg	Lys	Phe	Gly	Lys	Ser	Val	Ala	Val	Val	Ala	Val
			420					425					430		
Asn	Arg	Asn	Leu	Thr	Thr	Pro	Thr	Ser	Ile	Thr	Asn	Leu	Asn	Thr	Ser
		435					440					445			
Leu	Pro	Ser	Gly	Thr	Tyr	Thr	Asp	Val	Leu	Gly	Gly	Val	Leu	Asn	Gly
	450					455					460				
Asn	Asn	Ile	Thr	Ser	Ser	Gly	Gly	Asn	Ile	Ser	Ser	Phe	Thr	Leu	Ala
465					470					475					480
Ala	Gly	Ala	Thr	Ala	Val	Trp	Gln	Tyr	Thr	Ala	Ser	Glu	Thr	Thr	Pro
				485					490					495	
Thr	Ile	Gly	His	Val	Gly	Pro	Val	Met	Gly	Lys	Pro	Gly	Asn	Val	Val
			500					505					510		
Thr	Ile	Asp	Gly	Arg	Gly	Phe	Gly	Ser	Thr	Lys	Gly	Thr	Val	Tyr	Phe
		515					520					525			
Gly	Thr	Thr	Ala	Val	Thr	Gly	Ser	Ala	Ile	Thr	Ser	Trp	Glu	Asp	Thr
	530					535					540				
Gln	Ile	Lys	Val	Thr	Ile	Pro	Pro	Val	Ala	Gly	Gly	Asp	Tyr	Ala	Val
545					550					555					560
Lys	Val	Ala	Ala	Asn	Gly	Val	Asn	Ser	Asn	Ala	Tyr	Asn	Asp	Phe	Thr
				565					570					575	
Ile	Leu	Ser	Gly	Asp	Gln	Val	Ser	Val	Arg	Phe	Val	Ile	Asn	Asn	Ala
			580					585					590		
Thr	Thr	Ala	Leu	Gly	Glu	Asn	Ile	Tyr	Leu	Thr	Gly	Asn	Val	Ser	Glu

Sequence-Listing

595	600	605													
Leu	Gly	Asn	Trp	Thr	Thr	Gly	Ala	Ala	Ser	Ile	Gly	Pro	Ala	Phe	Asn
610	615	620													
Gln	Val	Ile	His	Ala	Tyr	Pro	Thr	Trp	Tyr	Tyr	Asp	Val	Ser	Val	Pro
625					630					635					640
Ala	Gly	Lys	Gln	Leu	Glu	Phe	Lys	Phe	Phe	Lys	Lys	Asn	Gly	Ala	Thr
				645					650						655
Ile	Thr	Trp	Glu	Gly	Gly	Ser	Asn	His	Thr	Phe	Thr	Thr	Pro	Thr	Ser
			660					665					670		
Gly	Thr	Ala	Thr	Val	Thr	Val	Asn	Trp	Gln						
675							680								

<210> 13
 <211> 682
 <212> PRT
 <213> Artificial sequence

<220>
 <223> mutated CGTase(Y454H)

<400> 13

Asp	Thr	Ala	Val	Thr	Asn	Lys	Gln	Asn	Phe	Ser	Thr	Asp	Val	Ile	Tyr
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Gln	Val	Phe	Thr	Asp	Arg	Phe	Leu	Asp	Gly	Asn	Pro	Ser	Asn	Asn	Pro
			20					25					30		
Thr	Gly	Ala	Ala	Phe	Asp	Gly	Thr	Cys	Ser	Asn	Leu	Lys	Leu	Tyr	Cys
		35					40					45			
Gly	Gly	Asp	Trp	Gln	Gly	Leu	Ile	Asn	Lys	Ile	Asn	Asp	Asn	Tyr	Phe
	50					55					60				
Ser	Asp	Leu	Gly	Val	Thr	Ala	Leu	Trp	Ile	Ser	Gln	Pro	Val	Glu	Asn
65					70					75					80
Ile	Phe	Ala	Thr	Ile	Asn	Tyr	Ser	Gly	Val	Thr	Asn	Thr	Ala	Tyr	His
				85					90					95	
Gly	Tyr	Trp	Ala	Arg	Asp	Phe	Lys	Lys	Thr	Asn	Pro	Tyr	Phe	Gly	Thr
			100					105					110		
Met	Thr	Asp	Phe	Gln	Asn	Leu	Val	Thr	Ser	Ala	His	Ala	Lys	Gly	Ile
		115					120					125			
Lys	Ile	Ile	Ile	Asp	Phe	Ala	Pro	Asn	His	Thr	Phe	Pro	Ala	Met	Glu
	130					135					140				
Thr	Asp	Thr	Ser	Phe	Ala	Glu	Asn	Gly	Lys	Leu	Tyr	Asp	Asn	Gly	Ser
145					150					155					160
Leu	Val	Gly	Gly	Tyr	Thr	Asn	Asp	Thr	Asn	Gly	Tyr	Phe	His	His	Asn
				165					170					175	
Gly	Gly	Ser	Asp	Phe	Ser	Thr	Leu	Glu	Asn	Gly	Ile	Tyr	Lys	Asn	Leu
			180					185					190		
Tyr	Asp	Leu	Ala	Asp	Leu	Asn	His	Asn	Asn	Ser	Thr	Ile	Asp	Thr	Tyr
		195					200					205			
Phe	Lys	Asp	Ala	Ile	Lys	Leu	Trp	Leu	Asp	Met	Gly	Val	Asp	Gly	Ile
	210					215					220				
Arg	Val	Asp	Ala	Val	Lys	His	Met	Pro	Gln	Gly	Trp	Gln	Lys	Asn	Trp
225					230					235					240
Met	Ser	Ser	Ile	Tyr	Ala	His	Lys	Pro	Val	Phe	Thr	Phe	Gly	Glu	Trp
				245					250					255	
Phe	Leu	Gly	Ser	Ala	Ala	Ser	Asp	Ala	Asp	Asn	Thr	Asp	Phe	Ala	Asn
			260					265					270		
Glu	Ser	Gly	Met	Ser	Leu	Leu	Asp	Phe	Arg	Phe	Asn	Ser	Ala	Val	Arg
		275					280					285			
Asn	Val	Phe	Arg	Asp	Asn	Thr	Ser	Asn	Met	Tyr	Ala	Leu	Asp	Ser	Met
	290					295					300				
Leu	Thr	Ala	Thr	Ala	Ala	Asp	Tyr	Asn	Gln	Val	Asn	Asp	Gln	Val	Thr
305					310					315					320
Phe	Ile	Asp	Asn	His	Asp	Met	Asp	Arg	Phe	Lys	Thr	Ser	Ala	Val	Asn
				325					330					335	
Asn	Arg	Arg	Leu	Glu	Gln	Ala	Leu	Ala	Phe	Thr	Leu	Thr	Ser	Arg	Gly
			340					345					350		
Val	Pro	Ala	Ile	Tyr	Tyr	Gly	Thr	Glu	Gln	Tyr	Leu	Thr	Gly	Asn	Gly
		355					360					365			

Sequence-Listing

Asp	Pro	Asp	Asn	Arg	Gly	Lys	Met	Pro	Ser	Phe	Ser	Lys	Ser	Thr	Thr
	370					375					380				
Ala	Phe	Ser	Val	Ile	Ser	Lys	Leu	Ala	Pro	Leu	Arg	Lys	Ser	Asn	Pro
385					390					395					400
Ala	Ile	Ala	Tyr	Gly	Ser	Thr	Gln	Gln	Arg	Trp	Ile	Asn	Asn	Asp	Val
				405					410					415	
Tyr	Ile	Tyr	Glu	Arg	Lys	Phe	Gly	Lys	Ser	Val	Ala	Val	Val	Ala	Val
			420					425					430		
Asn	Arg	Asn	Leu	Thr	Thr	Pro	Thr	Ser	Ile	Thr	Asn	Leu	Asn	Thr	Ser
		435					440					445			
Leu	Pro	Ser	Gly	Thr	His	Thr	Asp	Val	Leu	Gly	Gly	Val	Leu	Asn	Gly
	450					455					460				
Asn	Asn	Ile	Thr	Ser	Ser	Gly	Gly	Asn	Ile	Ser	Ser	Phe	Thr	Leu	Ala
465					470					475					480
Ala	Gly	Ala	Thr	Ala	Val	Trp	Gln	Tyr	Thr	Ala	Ser	Glu	Thr	Thr	Pro
				485					490						495
Thr	Ile	Gly	His	Val	Gly	Pro	Val	Met	Gly	Lys	Pro	Gly	Asn	Val	Val
			500					505					510		
Thr	Ile	Asp	Gly	Arg	Gly	Phe	Gly	Ser	Thr	Lys	Gly	Thr	Val	Tyr	Phe
		515					520					525			
Gly	Thr	Thr	Ala	Val	Thr	Gly	Ser	Ala	Ile	Thr	Ser	Trp	Glu	Asp	Thr
						535					540				
Gln	Ile	Lys	Val	Thr	Ile	Pro	Pro	Val	Ala	Gly	Gly	Asp	Tyr	Ala	Val
545					550					555					560
Lys	Val	Ala	Ala	Asn	Gly	Val	Asn	Ser	Asn	Ala	Tyr	Asn	Asp	Phe	Thr
				565					570					575	
Ile	Leu	Ser	Gly	Asp	Gln	Val	Ser	Val	Arg	Phe	Val	Ile	Asn	Asn	Ala
			580					585					590		
Thr	Thr	Ala	Leu	Gly	Glu	Asn	Ile	Tyr	Leu	Thr	Gly	Asn	Val	Ser	Glu
		595					600					605			
Leu	Gly	Asn	Trp	Thr	Thr	Gly	Ala	Ala	Ser	Ile	Gly	Pro	Ala	Phe	Asn
	610					615					620				
Gln	Val	Ile	His	Ala	Tyr	Pro	Thr	Trp	Tyr	Tyr	Asp	Val	Ser	Val	Pro
625					630					635					640
Ala	Gly	Lys	Gln	Leu	Glu	Phe	Lys	Phe	Phe	Lys	Lys	Asn	Gly	Ala	Thr
				645					650					655	
Ile	Thr	Trp	Glu	Gly	Gly	Ser	Asn	His	Thr	Phe	Thr	Thr	Pro	Thr	Ser
			660					665					670		
Gly	Thr	Ala	Thr	Val	Thr	Val	Asn	Trp	Gln						
		675					680								