

X20729 Sequence Listing_ST25
SEQUENCE LISTING

<110> Eli Lilly and Company
Audion Therapeutics

<120> NOTCH PATHWAY SIGNALING INHIBITOR COMPOUNDS

<130> X20729

<150> 62/189,393

<151> 2015-07-07

<160> 2

<170> PatentIn version 3.5

<210> 1

<211> 2531

<212> PRT

<213> Mus musculus

<400> 1

Met Pro Arg Leu Leu Thr Pro Leu Leu Cys Leu Thr Leu Leu Pro Ala
1 5 10 15

Leu Ala Ala Arg Gly Leu Arg Cys Ser Gln Pro Ser Gly Thr Cys Leu
20 25 30

Asn Gly Gly Arg Cys Glu Val Ala Asn Gly Thr Glu Ala Cys Val Cys
35 40 45

Ser Gly Ala Phe Val Gly Gln Arg Cys Gln Asp Ser Asn Pro Cys Leu
50 55 60

Ser Thr Pro Cys Lys Asn Ala Gly Thr Cys His Val Val Asp His Gly
65 70 75 80

Gly Thr Val Asp Tyr Ala Cys Ser Cys Pro Leu Gly Phe Ser Gly Pro
85 90 95

Leu Cys Leu Thr Pro Leu Asp Asn Ala Cys Leu Ala Asn Pro Cys Arg
100 105 110

Asn Gly Gly Thr Cys Asp Leu Leu Thr Leu Thr Glu Tyr Lys Cys Arg
115 120 125

Cys Pro Pro Gly Trp Ser Gly Lys Ser Cys Gln Gln Ala Asp Pro Cys
130 135 140

Ala Ser Asn Pro Cys Ala Asn Gly Gly Gln Cys Leu Pro Phe Glu Ser
145 150 155 160

x20729 Sequence Listing_ST25

Ser Tyr Ile Cys Arg Cys Pro Pro Gly Phe His Gly Pro Thr Cys Arg
165 170 175

Gln Asp Val Asn Glu Cys Ser Gln Asn Pro Gly Leu Cys Arg His Gly
180 185 190

Gly Thr Cys His Asn Glu Ile Gly Ser Tyr Arg Cys Ala Cys Arg Ala
195 200 205

Thr His Thr Gly Pro His Cys Glu Leu Pro Tyr Val Pro Cys Ser Pro
210 215 220

Ser Pro Cys Gln Asn Gly Gly Thr Cys Arg Pro Thr Gly Asp Thr Thr
225 230 235 240

His Glu Cys Ala Cys Leu Pro Gly Phe Ala Gly Gln Asn Cys Glu Glu
245 250 255

Asn Val Asp Asp Cys Pro Gly Asn Asn Cys Lys Asn Gly Gly Ala Cys
260 265 270

Val Asp Gly Val Asn Thr Tyr Asn Cys Arg Cys Pro Pro Glu Trp Thr
275 280 285

Gly Gln Tyr Cys Thr Glu Asp Val Asp Glu Cys Gln Leu Met Pro Asn
290 295 300

Ala Cys Gln Asn Gly Gly Thr Cys His Asn Thr His Gly Gly Tyr Asn
305 310 315 320

Cys Val Cys Val Asn Gly Trp Thr Gly Glu Asp Cys Ser Glu Asn Ile
325 330 335

Asp Asp Cys Ala Ser Ala Ala Cys Phe Gln Gly Ala Thr Cys His Asp
340 345 350

Arg Val Ala Ser Phe Tyr Cys Glu Cys Pro His Gly Arg Thr Gly Leu
355 360 365

Leu Cys His Leu Asn Asp Ala Cys Ile Ser Asn Pro Cys Asn Glu Gly
370 375 380

Ser Asn Cys Asp Thr Asn Pro Val Asn Gly Lys Ala Ile Cys Thr Cys
385 390 395 400

Pro Ser Gly Tyr Thr Gly Pro Ala Cys Ser Gln Asp Val Asp Glu Cys
405 410 415

x20729 Sequence Listing_ST25

Ala Leu Gly Ala Asn Pro Cys Glu His Ala Gly Lys Cys Leu Asn Thr
420 425 430

Leu Gly Ser Phe Glu Cys Gln Cys Leu Gln Gly Tyr Thr Gly Pro Arg
435 440 445

Cys Glu Ile Asp Val Asn Glu Cys Ile Ser Asn Pro Cys Gln Asn Asp
450 455 460

Ala Thr Cys Leu Asp Gln Ile Gly Glu Phe Gln Cys Ile Cys Met Pro
465 470 475 480

Gly Tyr Glu Gly Val Tyr Cys Glu Ile Asn Thr Asp Glu Cys Ala Ser
485 490 495

Ser Pro Cys Leu His Asn Gly His Cys Met Asp Lys Ile Asn Glu Phe
500 505 510

Gln Cys Gln Cys Pro Lys Gly Phe Asn Gly His Leu Cys Gln Tyr Asp
515 520 525

Val Asp Glu Cys Ala Ser Thr Pro Cys Lys Asn Gly Ala Lys Cys Leu
530 535 540

Asp Gly Pro Asn Thr Tyr Thr Cys Val Cys Thr Glu Gly Tyr Thr Gly
545 550 555 560

Thr His Cys Glu Val Asp Ile Asp Glu Cys Asp Pro Asp Pro Cys His
565 570 575

Tyr Gly Ser Cys Lys Asp Gly Val Ala Thr Phe Thr Cys Leu Cys Gln
580 585 590

Pro Gly Tyr Thr Gly His His Cys Glu Thr Asn Ile Asn Glu Cys His
595 600 605

Ser Gln Pro Cys Arg His Gly Gly Thr Cys Gln Asp Arg Asp Asn Ser
610 615 620

Tyr Leu Cys Leu Cys Leu Lys Gly Thr Thr Gly Pro Asn Cys Glu Ile
625 630 635 640

Asn Leu Asp Asp Cys Ala Ser Asn Pro Cys Asp Ser Gly Thr Cys Leu
645 650 655

Asp Lys Ile Asp Gly Tyr Glu Cys Ala Cys Glu Pro Gly Tyr Thr Gly
660 665 670

x20729 Sequence Listing_ST25

Ser Met Cys Asn Val Asn Ile Asp Glu Cys Ala Gly Ser Pro Cys His
675 680 685

Asn Gly Gly Thr Cys Glu Asp Gly Ile Ala Gly Phe Thr Cys Arg Cys
690 695 700

Pro Glu Gly Tyr His Asp Pro Thr Cys Leu Ser Glu Val Asn Glu Cys
705 710 715 720

Asn Ser Asn Pro Cys Ile His Gly Ala Cys Arg Asp Gly Leu Asn Gly
725 730 735

Tyr Lys Cys Asp Cys Ala Pro Gly Trp Ser Gly Thr Asn Cys Asp Ile
740 745 750

Asn Asn Asn Glu Cys Glu Ser Asn Pro Cys Val Asn Gly Gly Thr Cys
755 760 765

Lys Asp Met Thr Ser Gly Tyr Val Cys Thr Cys Arg Glu Gly Phe Ser
770 775 780

Gly Pro Asn Cys Gln Thr Asn Ile Asn Glu Cys Ala Ser Asn Pro Cys
785 790 795 800

Leu Asn Gln Gly Thr Cys Ile Asp Asp Val Ala Gly Tyr Lys Cys Asn
805 810 815

Cys Pro Leu Pro Tyr Thr Gly Ala Thr Cys Glu Val Val Leu Ala Pro
820 825 830

Cys Ala Thr Ser Pro Cys Lys Asn Ser Gly Val Cys Lys Glu Ser Glu
835 840 845

Asp Tyr Glu Ser Phe Ser Cys Val Cys Pro Thr Gly Trp Gln Gly Gln
850 855 860

Thr Cys Glu Val Asp Ile Asn Glu Cys Val Lys Ser Pro Cys Arg His
865 870 875 880

Gly Ala Ser Cys Gln Asn Thr Asn Gly Ser Tyr Arg Cys Leu Cys Gln
885 890 895

Ala Gly Tyr Thr Gly Arg Asn Cys Glu Ser Asp Ile Asp Asp Cys Arg
900 905 910

Pro Asn Pro Cys His Asn Gly Gly Ser Cys Thr Asp Gly Ile Asn Thr

x20729 Sequence Listing_ST25
920 925

915

Ala Phe Cys Asp Cys Leu Pro Gly Phe Gln Gly Ala Phe Cys Glu Glu
930 935 940

Asp Ile Asn Glu Cys Ala Ser Asn Pro Cys Gln Asn Gly Ala Asn Cys
945 950 955 960

Thr Asp Cys Val Asp Ser Tyr Thr Cys Thr Cys Pro Val Gly Phe Asn
965 970 975

Gly Ile His Cys Glu Asn Asn Thr Pro Asp Cys Thr Glu Ser Ser Cys
980 985 990

Phe Asn Gly Gly Thr Cys Val Asp Gly Ile Asn Ser Phe Thr Cys Leu
995 1000 1005

Cys Pro Pro Gly Phe Thr Gly Ser Tyr Cys Gln Tyr Asp Val Asn
1010 1015 1020

Glu Cys Asp Ser Arg Pro Cys Leu His Gly Gly Thr Cys Gln Asp
1025 1030 1035

Ser Tyr Gly Thr Tyr Lys Cys Thr Cys Pro Gln Gly Tyr Thr Gly
1040 1045 1050

Leu Asn Cys Gln Asn Leu Val Arg Trp Cys Asp Ser Ala Pro Cys
1055 1060 1065

Lys Asn Gly Gly Arg Cys Trp Gln Thr Asn Thr Gln Tyr His Cys
1070 1075 1080

Glu Cys Arg Ser Gly Trp Thr Gly Val Asn Cys Asp Val Leu Ser
1085 1090 1095

Val Ser Cys Glu Val Ala Ala Gln Lys Arg Gly Ile Asp Val Thr
1100 1105 1110

Leu Leu Cys Gln His Gly Gly Leu Cys Val Asp Glu Gly Asp Lys
1115 1120 1125

His Tyr Cys His Cys Gln Ala Gly Tyr Thr Gly Ser Tyr Cys Glu
1130 1135 1140

Asp Glu Val Asp Glu Cys Ser Pro Asn Pro Cys Gln Asn Gly Ala
1145 1150 1155

x20729 Sequence Listing_ST25

Thr	Cys	Thr	Asp	Tyr	Leu	Gly	Gly	Phe	Ser	Cys	Lys	Cys	Val	Ala
	1160					1165					1170			
Gly	Tyr	His	Gly	Ser	Asn	Cys	Ser	Glu	Glu	Ile	Asn	Glu	Cys	Leu
	1175					1180					1185			
Ser	Gln	Pro	Cys	Gln	Asn	Gly	Gly	Thr	Cys	Ile	Asp	Leu	Thr	Asn
	1190					1195					1200			
Ser	Tyr	Lys	Cys	Ser	Cys	Pro	Arg	Gly	Thr	Gln	Gly	Val	His	Cys
	1205					1210					1215			
Glu	Ile	Asn	Val	Asp	Asp	Cys	His	Pro	Pro	Leu	Asp	Pro	Ala	Ser
	1220					1225					1230			
Arg	Ser	Pro	Lys	Cys	Phe	Asn	Asn	Gly	Thr	Cys	Val	Asp	Gln	Val
	1235					1240					1245			
Gly	Gly	Tyr	Thr	Cys	Thr	Cys	Pro	Pro	Gly	Phe	Val	Gly	Glu	Arg
	1250					1255					1260			
Cys	Glu	Gly	Asp	Val	Asn	Glu	Cys	Leu	Ser	Asn	Pro	Cys	Asp	Pro
	1265					1270					1275			
Arg	Gly	Thr	Gln	Asn	Cys	Val	Gln	Arg	Val	Asn	Asp	Phe	His	Cys
	1280					1285					1290			
Glu	Cys	Arg	Ala	Gly	His	Thr	Gly	Arg	Arg	Cys	Glu	Ser	Val	Ile
	1295					1300					1305			
Asn	Gly	Cys	Arg	Gly	Lys	Pro	Cys	Lys	Asn	Gly	Gly	Val	Cys	Ala
	1310					1315					1320			
Val	Ala	Ser	Asn	Thr	Ala	Arg	Gly	Phe	Ile	Cys	Arg	Cys	Pro	Ala
	1325					1330					1335			
Gly	Phe	Glu	Gly	Ala	Thr	Cys	Glu	Asn	Asp	Ala	Arg	Thr	Cys	Gly
	1340					1345					1350			
Ser	Leu	Arg	Cys	Leu	Asn	Gly	Gly	Thr	Cys	Ile	Ser	Gly	Pro	Arg
	1355					1360					1365			
Ser	Pro	Thr	Cys	Leu	Cys	Leu	Gly	Ser	Phe	Thr	Gly	Pro	Glu	Cys
	1370					1375					1380			
Gln	Phe	Pro	Ala	Ser	Ser	Pro	Cys	Val	Gly	Ser	Asn	Pro	Cys	Tyr
	1385					1390					1395			

x20729 Sequence Listing_ST25

Asn	Gln	Gly	Thr	Cys	Glu	Pro	Thr	Ser	Glu	Asn	Pro	Phe	Tyr	Arg
1400						1405					1410			
Cys	Leu	Cys	Pro	Ala	Lys	Phe	Asn	Gly	Leu	Leu	Cys	His	Ile	Leu
1415						1420					1425			
Asp	Tyr	Ser	Phe	Thr	Gly	Gly	Ala	Gly	Arg	Asp	Ile	Pro	Pro	Pro
1430						1435					1440			
Gln	Ile	Glu	Glu	Ala	Cys	Glu	Leu	Pro	Glu	Cys	Gln	Val	Asp	Ala
1445						1450					1455			
Gly	Asn	Lys	Val	Cys	Asn	Leu	Gln	Cys	Asn	Asn	His	Ala	Cys	Gly
1460						1465					1470			
Trp	Asp	Gly	Gly	Asp	Cys	Ser	Leu	Asn	Phe	Asn	Asp	Pro	Trp	Lys
1475						1480					1485			
Asn	Cys	Thr	Gln	Ser	Leu	Gln	Cys	Trp	Lys	Tyr	Phe	Ser	Asp	Gly
1490						1495					1500			
His	Cys	Asp	Ser	Gln	Cys	Asn	Ser	Ala	Gly	Cys	Leu	Phe	Asp	Gly
1505						1510					1515			
Phe	Asp	Cys	Gln	Leu	Thr	Glu	Gly	Gln	Cys	Asn	Pro	Leu	Tyr	Asp
1520						1525					1530			
Gln	Tyr	Cys	Lys	Asp	His	Phe	Ser	Asp	Gly	His	Cys	Asp	Gln	Gly
1535						1540					1545			
Cys	Asn	Ser	Ala	Glu	Cys	Glu	Trp	Asp	Gly	Leu	Asp	Cys	Ala	Glu
1550						1555					1560			
His	Val	Pro	Glu	Arg	Leu	Ala	Ala	Gly	Thr	Leu	Val	Leu	Val	Val
1565						1570					1575			
Leu	Leu	Pro	Pro	Asp	Gln	Leu	Arg	Asn	Asn	Ser	Phe	His	Phe	Leu
1580						1585					1590			
Arg	Glu	Leu	Ser	His	Val	Leu	His	Thr	Asn	Val	Val	Phe	Lys	Arg
1595						1600					1605			
Asp	Ala	Gln	Gly	Gln	Gln	Met	Ile	Phe	Pro	Tyr	Tyr	Gly	His	Glu
1610						1615					1620			
Glu	Glu	Leu	Arg	Lys	His	Pro	Ile	Lys	Arg	Ser	Thr	Val	Gly	Trp
1625						1630					1635			

x20729 Sequence Listing_ST25

Ala	Thr	Ser	Ser	Leu	Leu	Pro	Gly	Thr	Ser	Gly	Gly	Arg	Gln	Arg
1640						1645					1650			
Arg	Glu	Leu	Asp	Pro	Met	Asp	Ile	Arg	Gly	Ser	Ile	Val	Tyr	Leu
1655						1660					1665			
Glu	Ile	Asp	Asn	Arg	Gln	Cys	Val	Gln	Ser	Ser	Ser	Gln	Cys	Phe
1670						1675					1680			
Gln	Ser	Ala	Thr	Asp	Val	Ala	Ala	Phe	Leu	Gly	Ala	Leu	Ala	Ser
1685						1690					1695			
Leu	Gly	Ser	Leu	Asn	Ile	Pro	Tyr	Lys	Ile	Glu	Ala	Val	Lys	Ser
1700						1705					1710			
Glu	Pro	Val	Glu	Pro	Pro	Leu	Pro	Ser	Gln	Leu	His	Leu	Met	Tyr
1715						1720					1725			
Val	Ala	Ala	Ala	Ala	Phe	Val	Leu	Leu	Phe	Phe	Val	Gly	Cys	Gly
1730						1735					1740			
Val	Leu	Leu	Ser	Arg	Lys	Arg	Arg	Arg	Gln	His	Gly	Gln	Leu	Trp
1745						1750					1755			
Phe	Pro	Glu	Gly	Phe	Lys	Val	Ser	Glu	Ala	Ser	Lys	Lys	Lys	Arg
1760						1765					1770			
Arg	Glu	Pro	Leu	Gly	Glu	Asp	Ser	Val	Gly	Leu	Lys	Pro	Leu	Lys
1775						1780					1785			
Asn	Ala	Ser	Asp	Gly	Ala	Leu	Met	Asp	Asp	Asn	Gln	Asn	Glu	Trp
1790						1795					1800			
Gly	Asp	Glu	Asp	Leu	Glu	Thr	Lys	Lys	Phe	Arg	Phe	Glu	Glu	Pro
1805						1810					1815			
Val	Val	Leu	Pro	Asp	Leu	Ser	Asp	Gln	Thr	Asp	His	Arg	Gln	Trp
1820						1825					1830			
Thr	Gln	Gln	His	Leu	Asp	Ala	Ala	Asp	Leu	Arg	Met	Ser	Ala	Met
1835						1840					1845			
Ala	Pro	Thr	Pro	Pro	Gln	Gly	Glu	Val	Asp	Ala	Asp	Cys	Met	Asp
1850						1855					1860			
Val	Asn	Val	Arg	Gly	Pro	Asp	Gly	Phe	Thr	Pro	Leu	Met	Ile	Ala

x20729 Sequence Listing_ST25

1865						1870						1875
Ser	Cys	Ser	Gly	Gly	Gly	Leu	Glu	Thr	Gly	Asn	Ser	Glu
1880						1885					1890	
Glu	Asp	Ala	Pro	Ala	Val	Ile	Ser	Asp	Phe	Ile	Tyr	Gln
1895						1900					1905	
Ser	Leu	His	Asn	Gln	Thr	Asp	Arg	Thr	Gly	Glu	Thr	Ala
1910						1915					1920	
Leu	Ala	Ala	Arg	Tyr	Ser	Arg	Ser	Asp	Ala	Ala	Lys	Arg
1925						1930					1935	
Glu	Ala	Ser	Ala	Asp	Ala	Asn	Ile	Gln	Asp	Asn	Met	Gly
1940						1945					1950	
Pro	Leu	His	Ala	Ala	Val	Ser	Ala	Asp	Ala	Gln	Gly	Val
1955						1960					1965	
Ile	Leu	Leu	Arg	Asn	Arg	Ala	Thr	Asp	Leu	Asp	Ala	Arg
1970						1975					1980	
Asp	Gly	Thr	Thr	Pro	Leu	Ile	Leu	Ala	Ala	Arg	Leu	Ala
1985						1990					1995	
Gly	Met	Leu	Glu	Asp	Leu	Ile	Asn	Ser	His	Ala	Asp	Val
2000						2005					2010	
Val	Asp	Asp	Leu	Gly	Lys	Ser	Ala	Leu	His	Trp	Ala	Ala
2015						2020					2025	
Asn	Asn	Val	Asp	Ala	Ala	Val	Val	Leu	Leu	Lys	Asn	Gly
2030						2035					2040	
Lys	Asp	Met	Gln	Asn	Asn	Lys	Glu	Glu	Thr	Pro	Leu	Phe
2045						2050					2055	
Ala	Arg	Glu	Gly	Ser	Tyr	Glu	Thr	Ala	Lys	Val	Leu	Leu
2060						2065					2070	
Phe	Ala	Asn	Arg	Asp	Ile	Thr	Asp	His	Met	Asp	Arg	Leu
2075						2080					2085	
Asp	Ile	Ala	Gln	Glu	Arg	Met	His	His	Asp	Ile	Val	Arg
2090						2095					2100	

x20729 Sequence Listing_ST25

Asp	Glu	Tyr	Asn	Leu	Val	Arg	Ser	Pro	Gln	Leu	His	Gly	Thr	Ala
	2105					2110					2115			
Leu	Gly	Gly	Thr	Pro	Thr	Leu	Ser	Pro	Thr	Leu	Cys	Ser	Pro	Asn
	2120					2125					2130			
Gly	Tyr	Leu	Gly	Asn	Leu	Lys	Ser	Ala	Thr	Gln	Gly	Lys	Lys	Ala
	2135					2140					2145			
Arg	Lys	Pro	Ser	Thr	Lys	Gly	Leu	Ala	Cys	Gly	Ser	Lys	Glu	Ala
	2150					2155					2160			
Lys	Asp	Leu	Lys	Ala	Arg	Arg	Lys	Lys	Ser	Gln	Asp	Gly	Lys	Gly
	2165					2170					2175			
Cys	Leu	Leu	Asp	Ser	Ser	Ser	Met	Leu	Ser	Pro	Val	Asp	Ser	Leu
	2180					2185					2190			
Glu	Ser	Pro	His	Gly	Tyr	Leu	Ser	Asp	Val	Ala	Ser	Pro	Pro	Leu
	2195					2200					2205			
Leu	Pro	Ser	Pro	Phe	Gln	Gln	Ser	Pro	Ser	Met	Pro	Leu	Ser	His
	2210					2215					2220			
Leu	Pro	Gly	Met	Pro	Asp	Thr	His	Leu	Gly	Ile	Ser	His	Leu	Asn
	2225					2230					2235			
Val	Ala	Ala	Lys	Pro	Glu	Met	Ala	Ala	Leu	Ala	Gly	Gly	Ser	Arg
	2240					2245					2250			
Leu	Ala	Phe	Glu	Pro	Pro	Pro	Pro	Arg	Leu	Ser	His	Leu	Pro	Val
	2255					2260					2265			
Ala	Ser	Ser	Ala	Ser	Thr	Val	Leu	Ser	Thr	Asn	Gly	Thr	Gly	Ala
	2270					2275					2280			
Met	Asn	Phe	Thr	Val	Gly	Ala	Pro	Ala	Ser	Leu	Asn	Gly	Gln	Cys
	2285					2290					2295			
Glu	Trp	Leu	Pro	Arg	Leu	Gln	Asn	Gly	Met	Val	Pro	Ser	Gln	Tyr
	2300					2305					2310			
Asn	Pro	Leu	Arg	Pro	Gly	Val	Thr	Pro	Gly	Thr	Leu	Ser	Thr	Gln
	2315					2320					2325			
Ala	Ala	Gly	Leu	Gln	His	Ser	Met	Met	Gly	Pro	Leu	His	Ser	Ser
	2330					2335					2340			

x20729 Sequence Listing_ST25

Leu	Ser	Thr	Asn	Thr	Leu	Ser	Pro	Ile	Ile	Tyr	Gln	Gly	Leu	Pro
	2345					2350					2355			
Asn	Thr	Arg	Leu	Ala	Thr	Gln	Pro	His	Leu	Val	Gln	Thr	Gln	Gln
	2360					2365					2370			
Val	Gln	Pro	Gln	Asn	Leu	Gln	Leu	Gln	Pro	Gln	Asn	Leu	Gln	Pro
	2375					2380					2385			
Pro	Ser	Gln	Pro	His	Leu	Ser	Val	Ser	Ser	Ala	Ala	Asn	Gly	His
	2390					2395					2400			
Leu	Gly	Arg	Ser	Phe	Leu	Ser	Gly	Glu	Pro	Ser	Gln	Ala	Asp	Val
	2405					2410					2415			
Gln	Pro	Leu	Gly	Pro	Ser	Ser	Leu	Pro	Val	His	Thr	Ile	Leu	Pro
	2420					2425					2430			
Gln	Glu	Ser	Gln	Ala	Leu	Pro	Thr	Ser	Leu	Pro	Ser	Ser	Met	Val
	2435					2440					2445			
Pro	Pro	Met	Thr	Thr	Thr	Gln	Phe	Leu	Thr	Pro	Pro	Ser	Gln	His
	2450					2455					2460			
Ser	Tyr	Ser	Ser	Ser	Pro	Val	Asp	Asn	Thr	Pro	Ser	His	Gln	Leu
	2465					2470					2475			
Gln	Val	Pro	Glu	His	Pro	Phe	Leu	Thr	Pro	Ser	Pro	Glu	Ser	Pro
	2480					2485					2490			
Asp	Gln	Trp	Ser	Ser	Ser	Ser	Pro	His	Ser	Asn	Ile	Ser	Asp	Trp
	2495					2500					2505			
Ser	Glu	Gly	Ile	Ser	Ser	Pro	Pro	Thr	Thr	Met	Pro	Ser	Gln	Ile
	2510					2515					2520			
Thr	His	Ile	Pro	Glu	Ala	Phe	Lys							
	2525					2530								

<210> 2
 <211> 23
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> Synthetic construct

 <400> 2

X20729 Sequence Listing_ST25

Met	Pro	Arg	Leu	Leu	Thr	Pro	Leu	Leu	Cys	Leu	Thr	Leu	Leu	Pro	Ala
1				5					10					15	

Leu	Ala	Ala	Arg	Gly	Leu	Arg
			20			