FIGURE 1

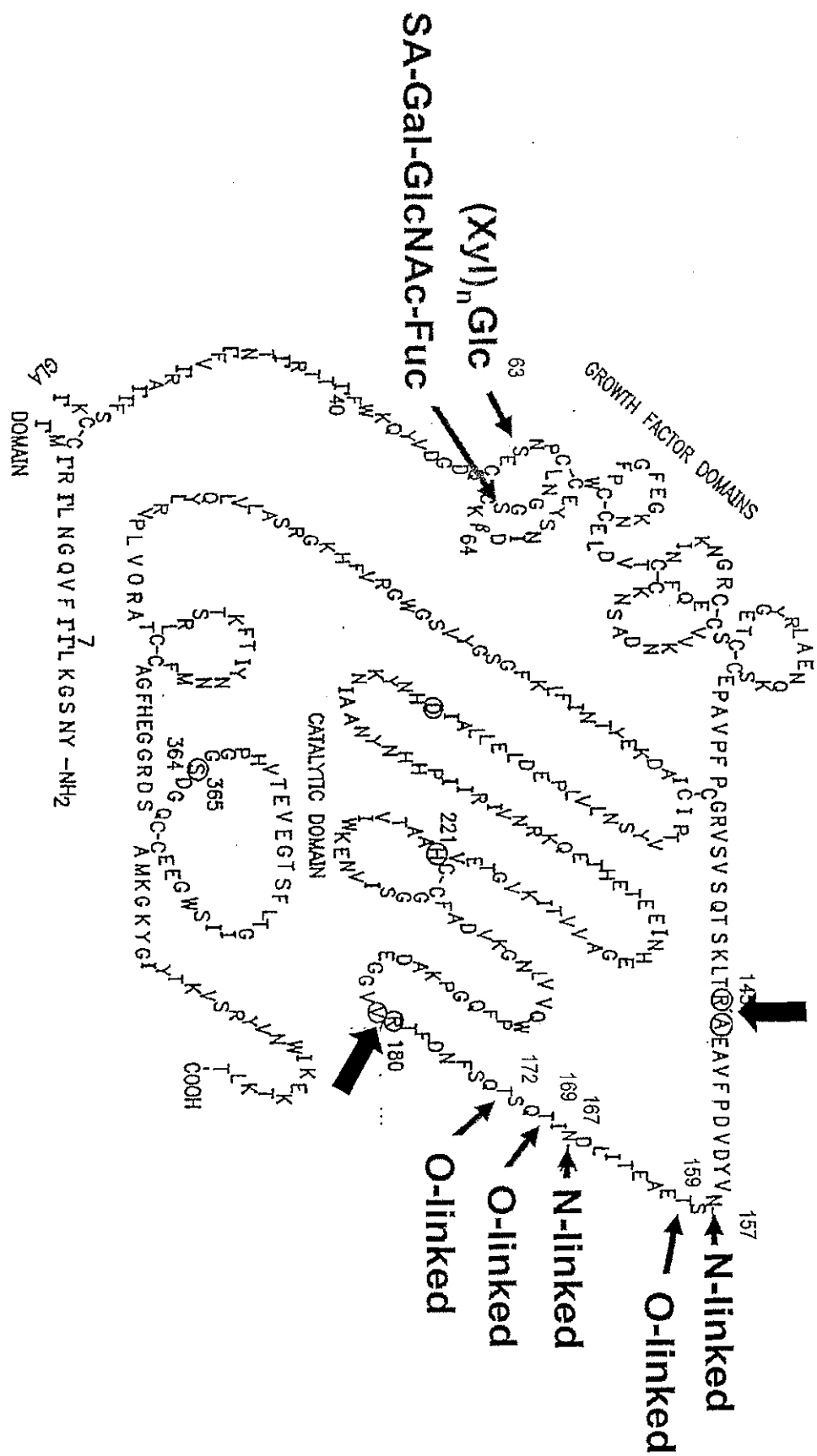

(Xyl)n Glc
SA-Gal-GlcNAc-Fuc
GROWTH FACTOR DOMAINS
CATALYTIC DOMAIN
DOMAIN
COOH
O-linked
O-linked
N-linked
O-linked
N-linked

FIGURE 2A

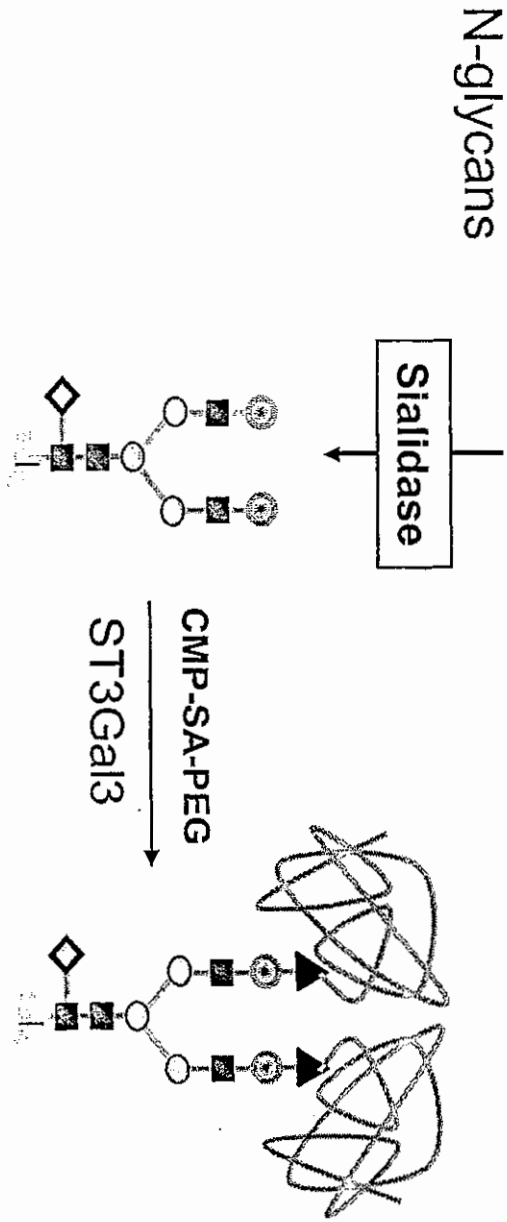


FIGURE 2B

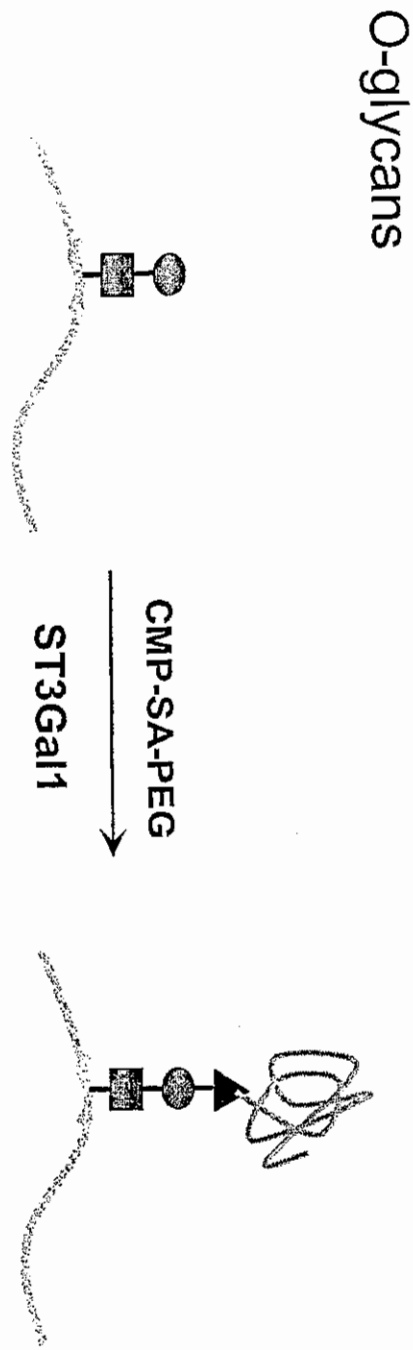


FIGURE 2C

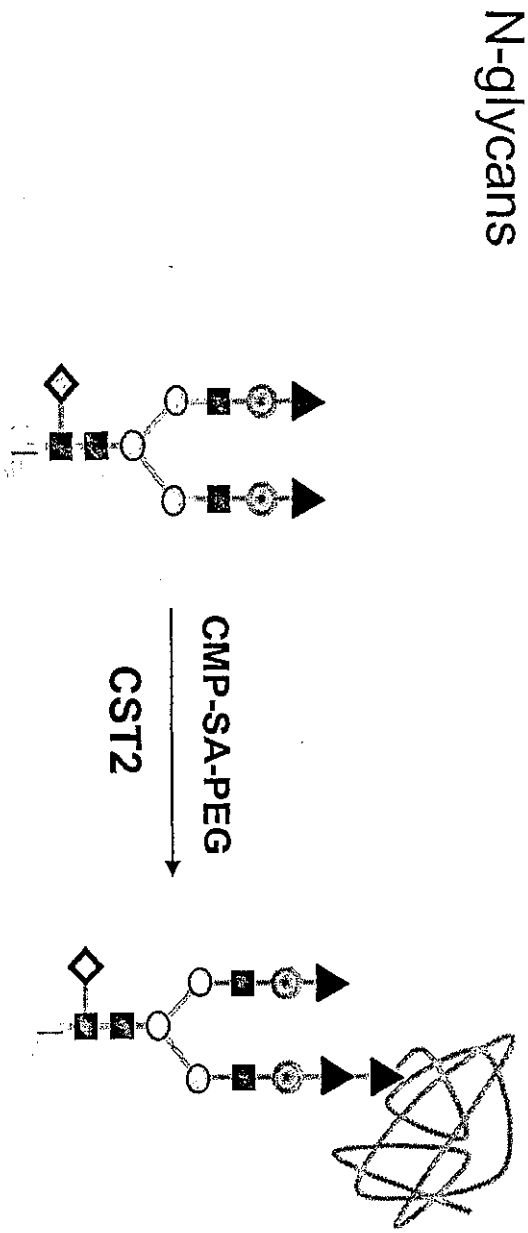


FIGURE 2D

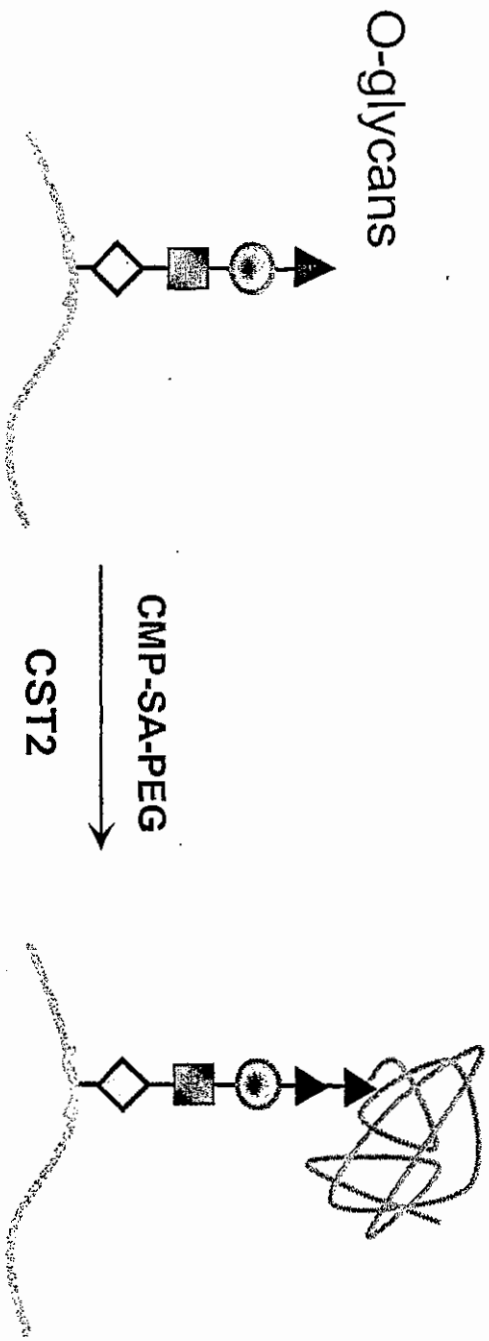


FIGURE 2E

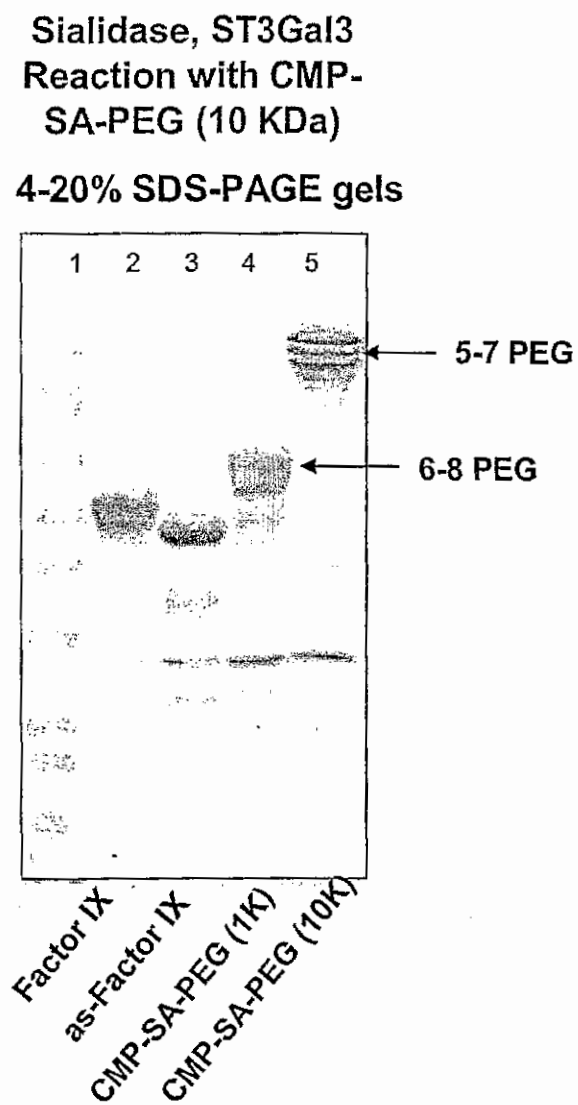

Sialidase, ST3Gal3
Reaction with CMP-
SA-PEG (10 KDa)
4-20% SDS-PAGE gels
1 2 3 4 5
5-7 PEG
6-8 PEG
Factor IX
as-Factor IX
CMP-SA-PEG (1K)
CMP-SA-PEG (10K)

FIGURE 2F

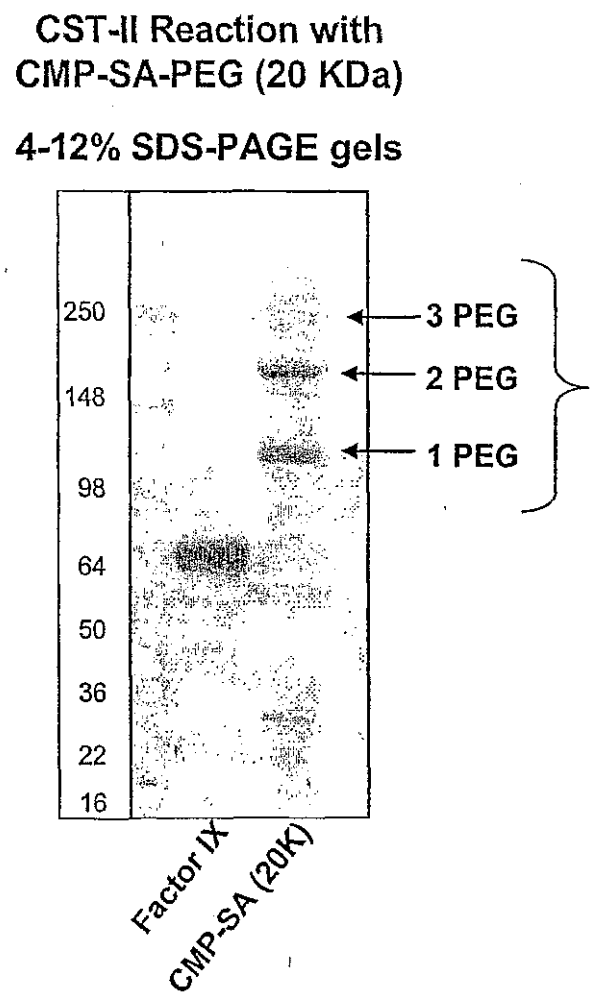

CST-II Reaction with
CMP-SA-PEG (20 KDa)
4-12% SDS-PAGE gels
250
148
98
64
50
36
22
16
3 PEG
2 PEG
1 PEG
Factor IX
CMP-SA (20K)

FIGURE 3

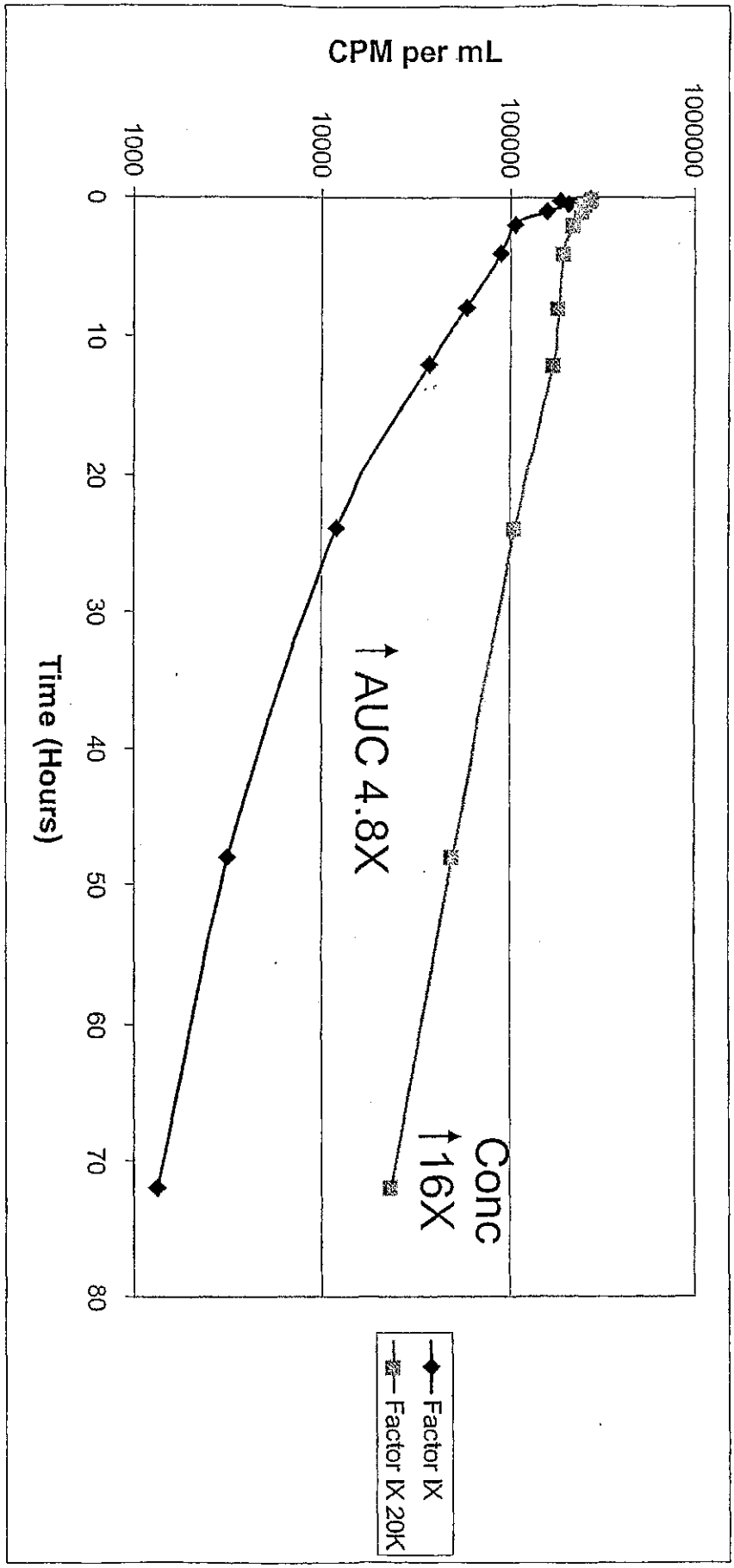

CPM per mL
1000
10000
100000
1000000
0
10
20
30
40
50
60
70
80
Time (Hours)
AUC 4.8X
Conc
16X
Factor IX
Factor IX 20K

FIGURE 4

Enzyme	Specificity and Linkage	Reaction Conditions	PEG Incorporation (size)	Clotting Activity (Chromogenic Assay)
Factor IX (unmodified)	NA	NA	>90% sialylation	110-139%
ST3Gal3	N-linked glycans	CMP-SA	None. (incorporated 2-3% SA)	Full Activity
ST3Gal3	***N-linked glycans***	***1. Sialidase 2. CMP-SA-PEG (1 & 10KDa)***	***(1 K); 6-8 PEG (10 K); 5-7 PEG***	***PEG (1 KDa); 52%. PEG (10 KDa); 54%.***
ST3Gal3	N-linked glycans	CMP-SA-PEG (10 KDa); no sialidase.	1-6 PEG (10 KDa);	1-2 PEG's (27 hrs); 164% ~3 PEG's (119 or 27 hrs); 97% 4-5 PEG's (119 hrs);
CST-II (Campylobacter)		***CMP-SA-PEG (20 KDa); no sialidase***	***2-3 PEG (20 KDa)***	***96%***

FIGURE 5

YNSGKLEEFVQGNLERECMEEKCSFEEAREVFENTERTTEFWKQYVDGDQ
CESNPCLNGGSCKDDINSYECWCPFGFEGKNCELDVTCNIKNGRCEQFCK
NSADNKVVCSCTEGYRLAENQKSCEPAVPFPCGRVSVSQTSKLTRAEAVFP
DVDYVNSTEAETILDNITQSTQSFNDFTRVVGGEDAKPGQFPWQVVLNGKV
DAFCGGSIVNEKWIVTAAHCVETGVKITVVAGEHNIEETEHTEQKRNVIRIIPH
HNYNAAINKYNHDIALLELDEPLVLNSYVTPICIADKEYTNIFLKFGSGYVSGWGRVFHK
GRSALVLQYLRVPLVDRATCLRSTKFTIYNNMFCAGFHEGGRDSCQGDSGGPHVTEVE
GTSFLTGIISWGEECAMKGKYGIYTKVSRYVNWIKEKTKLT

FIGURE 6

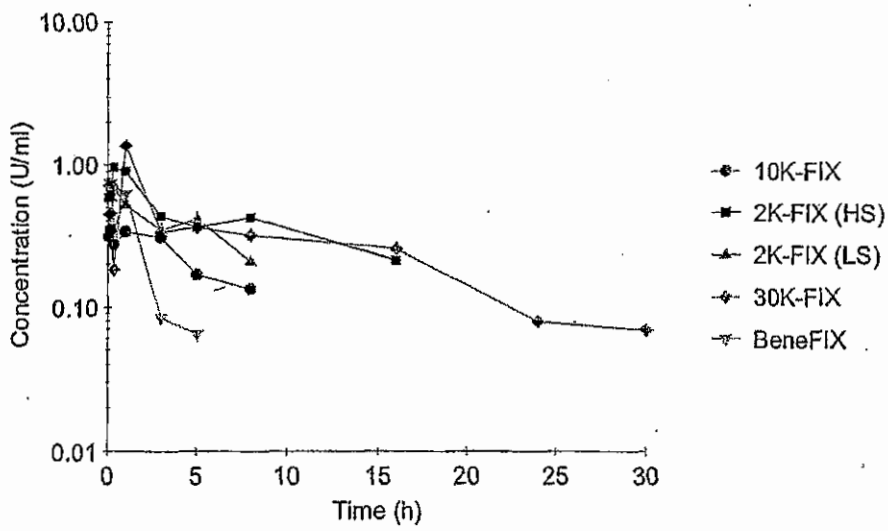
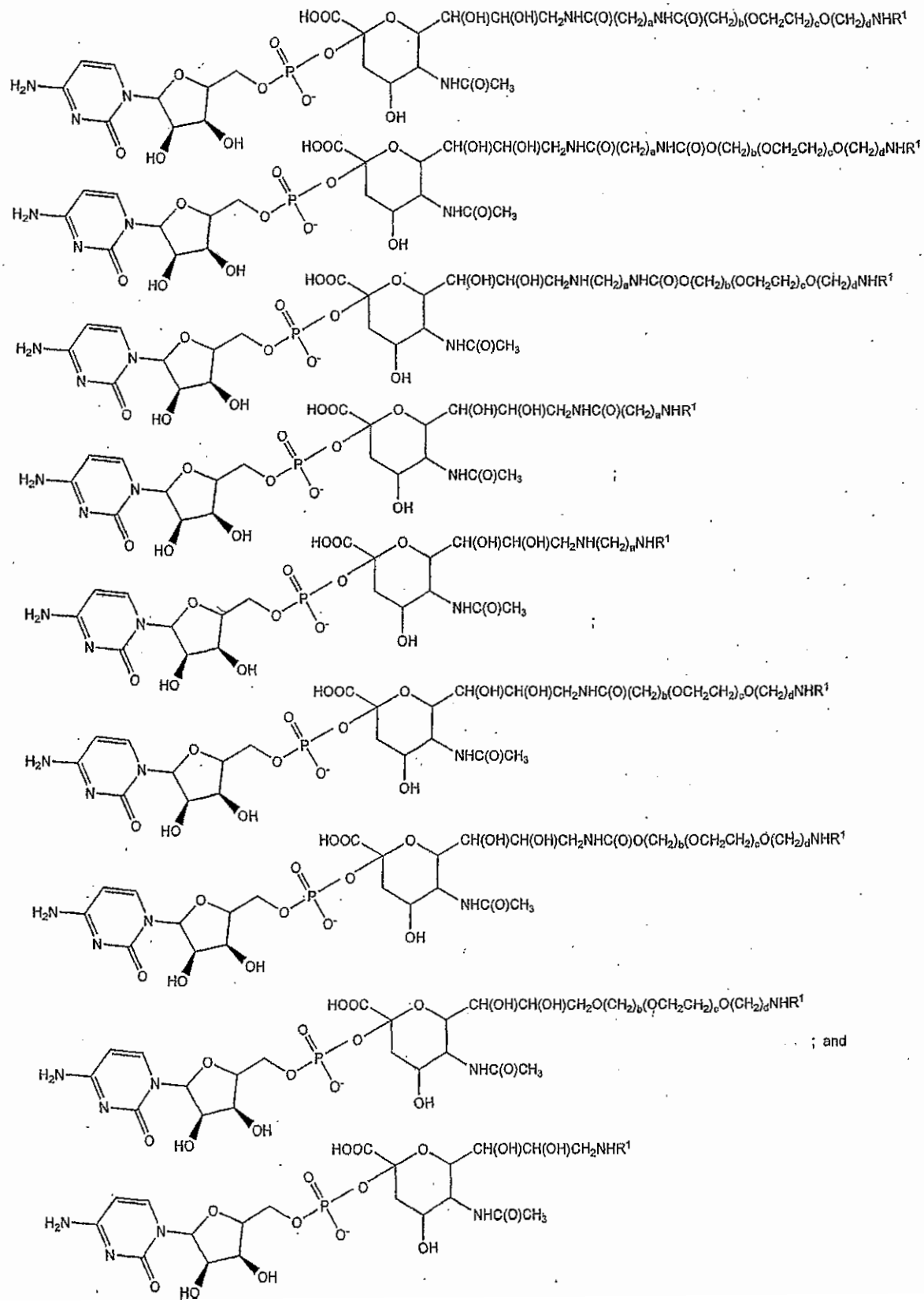
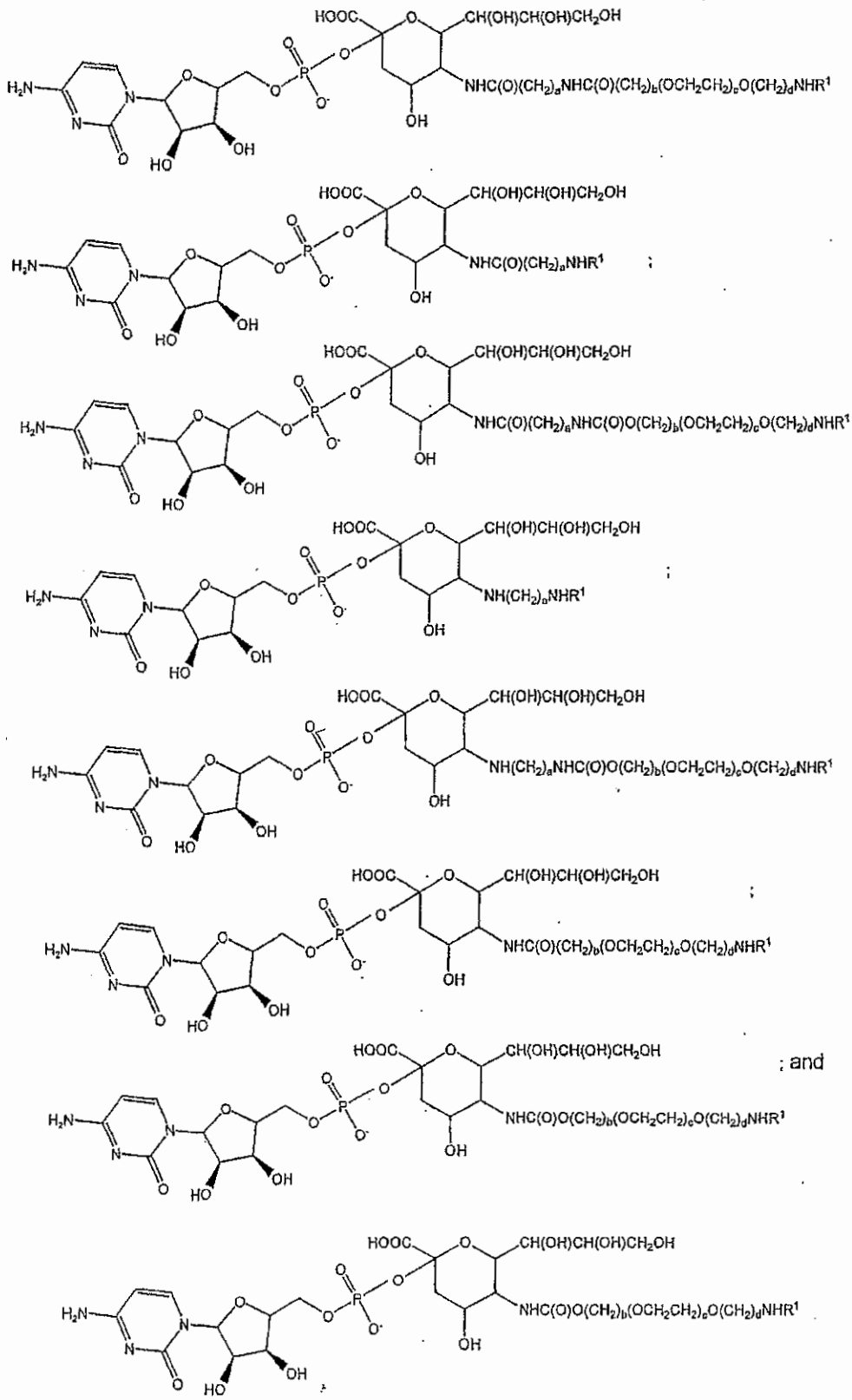

10.00
1.00
0.10
0.01
Concentration (U/ml)
0
5
10
15
20
25
30
Time (h)
10K-FIX
2K-FIX (HS)
2K-FIX (LS)
30K-FIX
BeneFIX

FIGURE 7


HOOC
CH(OH)CH(OH)CH2NHC(O)(CH2)aNHC(O)(CH2)b(OCH2CH2)cO(CH2)dNHR1
NHC(O)CH3
OH
H2N
HO
CH(OH)CH(OH)CH2NHC(O)(CH2)aNHC(O)O(CH2)b(OCH2CH2)cO(CH2)dNHR1
CH(OH)CH(OH)CH2NH(CH2)aNHC(O)O(CH2)b(OCH2CH2)cO(CH2)dNHR1
CH(OH)CH(OH)CH2NHC(O)(CH2)aNHR1
CH(OH)CH(OH)CH2NH(CH2)nNHR1
CH(OH)CH(OH)CH2NHC(O)(CH2)b(OCH2CH2)cO(CH2)dNHR1
CH(OH)CH(OH)CH2NHC(O)O(CH2)b(OCH2CH2)cO(CH2)dNHR1
CH(OH)CH(OH)CH2O(CH2)b(OCH2CH2)cO(CH2)dNHR1
CH(OH)CH(OH)CH2NHR1

; and

FIGURE 8


HOOC
CH(OH)CH(OH)CH2OH
NHC(O)(CH2)aNHC(O)(CH2)b(OCH2CH2)cO(CH2)dNHR1
NHC(O)(CH2)aNHR1
NHC(O)(CH2)aNHC(O)O(CH2)b(OCH2CH2)cO(CH2)dNHR1
NH(CH2)aNHR1
NH(CH2)aNHC(O)O(CH2)b(OCH2CH2)cO(CH2)dNHR1
NHC(O)(CH2)b(OCH2CH2)cO(CH2)dNHR1
NHC(O)O(CH2)b(OCH2CH2)cO(CH2)dNHR1
H2N
HO
OH

; and

FIGURE 9A

Protein	Organism	EC#	GenBank / GenPept	SwissProt	PDB 3D
At1g08280	*Arabidopsis thaliana*	n.d.	AC011438 AAF18241.1 BT004583 AAO42829.1 NC_003070 **NP_172305.1**	Q84W00 Q9SGD2	
At1g08660/F22O13.14	*Arabidopsis thaliana*	n.d.	AC003981 AAF99778.1 AY064135 **AAL36042.1** AY124807 AAM70516.1 NC_003070 NP_172342.1 NM_180609 NP_850940.1	Q8VZJ0 Q9FRR9	
At3g48820/T21J18_90	*Arabidopsis thaliana*	n.d.	AY080589 AAL85966.1 AY133816 AAM91750.1 AL132963 **CAB87910.1** NM_114741 NP_190451.1	Q8RY00 Q9M301	
α-2,3-sialyltransferase (ST3GAL-IV)	*Bos taurus*	n.d.	AJ584673 **CAE48298.1**		
α-2,3-sialyltransferase (St3Gal-V)	*Bos taurus*	n.d.	AJ585768 **CAE51392.1**		
α-2,6-sialyltransferase (Siat7b)	*Bos taurus*	n.d.	AJ620651 **CAF05850.1**		
α-2,8-sialyltransferase (SIAT8A)	*Bos taurus*	2.4.99.8	AJ699418 **CAG27880.1**		
α-2,8-sialyltransferase (Siat8D)	*Bos taurus*	n.d.	AJ699421 **CAG27883.1**		
α-2,8-sialyltransferase ST8Siα-III (Siat8C)	*Bos taurus*	n.d.	AJ704563 **CAG28696.1**		
CMP α-2,6-sialyltransferase (ST6Gal I)	*Bos taurus*	2.4.99.1	Y15111 CAA75385.1 NM_177517 NP_803483.1	**O18974**	
sialyltransferase 8 (fragment)	*Bos taurus*	n.d.	AF450088 **AAL47018.1**	Q8WN13	
sialyltransferase ST3Gal-II (Siat4B)	*Bos taurus*	n.d.	AJ748841 **CAG44450.1**		
sialyltransferase ST3Gal-III (Siat6)	*Bos taurus*	n.d.	AJ748842 **CAG44451.1**		
sialyltransferase ST3Gal-VI (Siat10)	*Bos taurus*	n.d.	AJ748843 **CAG44452.1**		
ST3Gal I	*Bos taurus*	n.d.	AJ305086 **CAC24698.1**	Q9BEG4	
St6GalNAc-VI	*Bos taurus*	n.d.	AJ620949 **CAF06586.1**		
CDS4	*Branchiostoma floridae*	n.d.	AF391289 **AAM18873.1**	Q8T771	
polysialyltransferase (PST) (fragment) ST8Sia IV	*Cercopithecus aethiops*	2.4.99.-	AF210729 **AAF17105.1**	Q9TT09	
polysialyltransferase (STX) (fragment) ST8Sia II	*Cercopithecus aethiops*	2.4.99.-	AF210318 **AAF17104.1**	Q9TT10	
α-2,3-sialyltransferase ST3Gal I (Siat4)	*Ciona intestinalis*	n.d.	AJ626815 **CAF25173.1**		
α-2,3-sialyltransferase ST3Gal I (Siat4)	*Ciona savignyi*	n.d.	AJ626814 **CAF25172.1**		
α-2,8-polysialyltransferase ST8Sia IV	*Cricetulus griseus*	2.4.99.-	-AAE28634 Z46801 CAA86822.1	**Q64690**	
Gal β-1,3/4-GlcNAc α-2,3-sialyltransferase St3Gal I	*Cricetulus griseus*	n.d.	AY266675 **AAP22942.1**	Q80WL0	
Gal β-1,3/4-GlcNAc α-2,3-sialyltransferase St3Gal II (fragment)	*Cricetulus griseus*	n.d.	AY266676 **AAP22943.1**	Q80WK9	
α-2,3-sialyltransferase ST3Gal I (Siat4)	*Danio rerio*	n.d.	AJ783740 **CAH04017.1**		
α-2,3-sialyltransferase ST3Gal II (Siat5)	*Danio rerio*	n.d.	AJ783741 **CAH04018.1**		
α-2,3-sialyltransferase ST3Gal III (Siat6)	*Danio rerio*	n.d.	AJ626821 **CAF25179.1**		
α-2,3-sialyltransferase ST3Gal IV (Siat4c)	*Danio rerio*	n.d.	AJ744809 **CAG32845.1**		

FIGURE 9B

Protein	Organism	EC#	GenBank / GenPept	SwissProt	PDB 3D
α-2,3-sialyltransferase ST3Gal V-r (Siat5-related)	*Danio rerio*	n.d.	AJ783742 **CAH04019.1**		
α-2,6-sialyltransferase ST6Gal I (Siat1)	*Danio rerio*	n.d.	AJ744801 **CAG32837.1**		
α-2,6-sialyltransferase ST6GalNAc II (Siat7B)	*Danio rerio*	n.d.	AJ634459 **CAG25680.1**		
α-2,6-sialyltransferase ST6GalNAc V (Siat7E) (fragment)	*Danio rerio*	n.d.	AJ646874 **CAG26703.1**		
α-2,6-sialyltransferase ST6GalNAc VI (Siat7F) (fragment)	*Danio rerio*	n.d.	AJ646883 **CAG26712.1**		
α-2,8-sialyltransferase ST8Sia I (Siat 8A) (fragment)	*Danio rerio*	n.d.	AJ715535 **CAG29374.1**		
α-2,8-sialyltransferase ST8Sia III (Siat 8C) (fragment)	*Danio rerio*	n.d.	AJ715543 **CAG29382.1**		
α-2,8-sialyltransferase ST8Sia IV (Siat 8D) (fragment)	*Danio rerio*	n.d.	AJ715545 **CAG29384.1**		
α-2,8-sialyltransferase ST8Sia V (Siat 8E) (fragment)	*Danio rerio*	n.d.	AJ715546 **CAG29385.1**		
α-2,8-sialyltransferase ST8Sia VI (Siat 8F) (fragment)	*Danio rerio*	n.d.	AJ715551 **CAG29390.1**		
β-galactosamide α-2,6-sialyltransferase II (ST6Gal II)	*Danio rerio*	n.d.	AJ627627 **CAF29495.1**		
N-glycan α-2,8-sialyltransferase	*Danio rerio.*	n.d.	BC050483 AAH50483.1 AY055462 **AAL17875.1** NM_153662 NP_705948.1	Q7ZU51 Q8QH83	
ST3Gal III-related (siat6r)	*Danio rerio*	n.d.	BC053179 AAH53179.1 AJ626820 **CAF25178.1** NM_200355 NP_956649.1	Q7T3B9	
St3Gal-V	*Danio rerio*	n.d.	AJ619960 **CAF04061.1**		
st6GalNAc-VI	*Danio rerio*	n.d.	BC060932 **AAH60932.1** AJ620947 CAF06584.1		
α-2,6-sialyltransferase (CG4871) ST6Gal I	*Drosophila melanogaster*	2.4.99.1	AE003465 AAF47256.1 AF218237 **AAG13185.1** AF397532 AAK92126.1 AE003465 AAM70791.1 NM_079129 NP_523853.1 NM_166684 NP_726474.1	Q9GU23 Q9W121	
α-2,3-sialyltransferase (ST3Gal-VI)	*Gallus gallus*	n.d.	AJ585767 **CAE51391.1** AJ627204 CAF25503.1		
α-2,3-sialyltransferase ST3Gal I	*Gallus gallus*	2.4.99.4	X80503 **CAA56666.1** NM_205217 NP_990548.1	Q11200	
α-2,3-sialyltransferase ST3Gal IV (fragment)	*Gallus gallus*	2.4.99.-	AF035250 AAC14163.1	**O73724**	
α-2,3-sialytransferase (ST3GAL-II)	*Gallus gallus*	n.d.	AJ585761 **CAE51385.2**		
α-2,6-sialyltransferase (Siat7b)	*Gallus gallus*	n.d.	AJ620653 **CAF05852.1**		
α-2,6-sialyltransferase ST6Gal I	*Gallus gallus*	2.4.99.1	X75558 **CAA53235.1** NM_205241 NP_990572.1	Q92182	
α-2,6-sialyltransferase ST6GalNAc I	*Gallus gallus*	2.4.99.3	-AAE68028.1 -AAE68029.1 X74946 **CAA52902.1** NM_205240 NP_990571.1	Q92183	
α-2,6-sialyltransferase ST6GalNAc II	*Gallus gallus*	2.4.99.-	X77775 AAE68030.1 NM_205233 **CAA54813.1** NP_990564.1	Q92184	
α-2,6-sialyltransferase ST6GalNAc III (SIAT7C)	*Gallus gallus*	n.d.	AJ634455 **CAG25677.1**		

FIGURE 9C

Protein	Organism	EC#	GenBank / GenPept	SwissProt	PDB 3D
(fragment)					
α-2,6-sialyltransferase ST6GalNAc V (SIAT7E) (fragment)	*Gallus gallus*	n.d.	AJ646877 **CAG26706.1**		
α-2,8-sialyltransferase (GD3 Synthase) ST8Sia I	*Gallus gallus*	2.4.99.-	U73176 AAC28888.1	**P79783**	
α-2,8-sialyltransferase (SIAT8B)	*Gallus gallus*	n.d.	AJ699419 **CAG27881.1**		
α-2,8-sialyltransferase (SIAT8C)	*Gallus gallus*	n.d.	AJ699420 **CAG27882.1**		
α-2,8-sialyltransferase (SIAT8F)	*Gallus gallus*	n.d.	AJ699424 **CAG27886.1**		
α-2,8-syalyltransferase ST8Siα-V (SIAT8C)	*Gallus gallus*	n.d.	AJ704564 **CAG28697.1**		
β-galactosamide α-2,6-sialyltransferase II (ST6Gal II)	*Gallus gallus*	n.d.	AJ627629 **CAF29497.1**		
GM3 synthase (SIAT9)	*Gallus gallus*	2.4.99.9	AY515255 **AAS83519.1**		
polysialyltransferase ST8Sia IV	*Gallus gallus*	2.4.99.-	AF008194 AAB95120.1	**O42399**	
α-2,3-sialyltransferase ST3Gal I	*Homo sapiens*	2.4.99.4	L29555 AAA36612.1	Q11201	
			AF059321 AAC17874.1	O60677	
			L13972 AAC37574.1	Q9UN51	
			AF155238 AAD39238.1		
			AF186191 **AAG29876.1**		
			BC018357 AAH18357.1		
			NM_003033 NP_003024.1		
			NM_173344 NP_775479.1		
α-2,3-sialyltransferase ST3Gal II	*Homo sapiens*	2.4.99.4	U63090 AAB40389.1	Q16842	
			BC036777 **AAH36777.1**	O00654	
			X96667 CAA65447.1		
			NM_006927 NP_008858.1		
α-2,3-sialyltransferase ST3Gal III (SiaT6)	*Homo sapiens*	2.4.99.6	L23768 AAA35778.1	Q11203	
			BC050380 AAH50380.1	Q86UR6	
			AF425851 AAO13859.1	Q86UR7	
			AF425852 AAO13860.1	Q86UR8	
			AF425853 AAO13861.1	Q86UR9	
			AF425854 AAO13862.1	Q86US0	
			AF425855 AAO13863.1	Q86US1	
			AF425856 AAO13864.1	Q86US2	
			AF425857 AAO13865.1	Q8IX43	
			AF425858 AAO13866.1	Q8IX44	
			AF425859 AAO13867.1	Q8IX45	
			AF425860 AAO13868.1	Q8IX46	
			AF425861 AAO13869.1	Q8IX47	
			AF425862 **AAO13870.1**	Q8IX48	
			AF425863 AAO13871.1	Q8IX49	
			AF425864 AAO13872.1	Q8IX50	
			AF425865 AAO13873.1	Q8IX51	
			AF425866 AAO13874.1	Q8IX52	
			AF425867 AAO13875.1	Q8IX53	
			AY167992 AAO38806.1	Q8IX54	
			AY167993 AAO38807.1	Q8IX55	
			AY167994 AAO38808.1	Q8IX56	
			AY167995 AAO38809.1	Q8IX57	
			AY167996 AAO38810.1	Q8IX58	
			AY167997 AAO38811.1		
			AY167998 AAO38812.1		
			NM_006279 NP_006270.1		
			NM_174964 NP_777624.1		
			NM_174965 NP_777625.1		
			NM_174966 NP_777626.1		
			NM_174967 NP_777627.1		
			NM_174969 NP_777629.1		
			NM_174970 NP_777630.1		
			NM_174972 NP_777632.1		

FIGURE 9D

Protein	Organism	EC	GenBank / GenPept	SwissProt / PDB
α-2,3-sialyltransferase ST3Gal IV	*Homo sapiens*	2.4.99.-	L23767 AAA16460.1 AF035249 AAC14162.1 BC010645 **AAH10645.1** AY040826 AAK93790.1 AF516602 AAM66431.1 AF516603 AAM66432.1 AF516604 AAM66433.1 AF525084 AAM81378.1 X74570 CAA52662.1 CR456858 CAG33139.1 NM_006278 NP_006269.1	Q11206 O60497 Q96QQ9 Q8N6A6 Q8N6A7 Q8NFD3 Q8NFG7
α-2,3-sialyltransferase ST3Gal VI	*Homo sapiens*	2.4.99.4	AF119391 **AAD39131.1** BC023312 AAH23312.1 AB022918 BAA77609.1 AX877828 CAE89895.1 AX886023 CAF00161.1 NM_006100 NP_006091.1	Q9Y274
α-2,6-sialyltransferase (ST6Gal II ; KIAA1877)	*Homo sapiens*	n.d.	BC008680 AAH08680.1 AB058780 **BAB47506.1** AB059555 BAC24793.1 AJ512141 CAD54408.1 AX795193 CAE48260.1 AX795193 CAE48261.1 NM_032528 NP_115917.1	Q86Y44 Q8IUG7 Q96HE4 Q96JF0
α-2,6-sialyltransferase (ST6GALNAC III)	*Homo sapiens*	n.d.	BC059363 AAH59363.1 AY358540 AAQ88904.1 AK091215 BAC03611.1 AJ507291 **CAD45371.1** NM_152996 NP_694541.1	Q8N259 Q8NDV1
α-2,6-sialyltransferase (ST6GalNAc V)	*Homo sapiens*	n.d.	BC001201 **AAH01201.1** AK056241 BAB71127.1 AL035409 CAB72344.1 AJ507292 CAD45372.1 NM_030965 NP_112227.1	Q9BVH7
α-2,6-sialyltransferase (SThM) ST6GalNAc II	*Homo sapiens*	2.4.99.-	U14550 AAA52228.1 BC040455 **AAH40455.1** AJ251053 CAB61434.1 NM_006456 NP_006447.1	Q9UJ37 Q12971
α-2,6-sialyltransferase ST6Gal I	*Homo sapiens*	2.4.99.1	BC031476 AAH31476.1 BC040009 AAH40009.1 A17362 **CAA01327.1** A23699 CAA01686.1 X17247 CAA35111.1 X54363 CAA38246.1 X62822 CAA44634.1 NM_003032 NP_003023.1 NM_173216 NP_775323.1	P15907
α-2,6-sialyltransferase ST6GalNAc I	*Homo sapiens*	2.4.99.3	BC022462 AAH22462.1 AY096001 AAM22800.1 AY358918 AAQ89277.1 AK000113 BAA90953.1 Y11339 **CAA72179.2** NM_018414 NP_060884.1	Q8TBJ6 Q9NSC7 Q9NXQ7
α-2,8-polysialyltransferase ST8Sia IV	*Homo sapiens*	2.4.99.-	L41680 AAC41775.1 BC027866 AAH27866.1 BC053657 **AAH53657.1** NM_005668 NP_005659.1	Q8N1F4 Q92187 Q92693
α-2,8-sialyltransferase (GD3 synthase) ST8Sia I	*Homo sapiens*	2.4.99.8	L32867 AAA62366.1 L43494 **AAC37586.1** BC046158 AAH46158.1 -AAQ53140.1 AY569975 AAS75783.1 D26360 BAA05391.1 X77922 CAA54891.1 NM_003034 NP_003025.1	Q86X71 Q92185 Q93064
α-2,8-sialyltransferase	*Homo sapiens*	2.4.99.-	L29556 AAA36613.1	Q92186

FIGURE 9E

Protein	Organism	EC#	GenBank / GenPept	SwissProt	PDB 3D
ST8Sia II			U82762 AAB51242.1	Q92470	
			U33551 **AAC24458.1**	Q92746	
			BC069584 AAH69584.1		
			NM_006011 NP_006002.1		
α-2,8-sialyltransferase ST8Sia III	*Homo sapiens*	2.4.99.-	AF004668 **AAB87642.1**	O43173	
			AF003092 AAC15901.2	Q9NS41	
			NM_015879 NP_056963.1		
α-2,8-sialyltransferase ST8Sia V	*Homo sapiens*	2.4.99.-	U91641 **AAC51727.1**	O15466	
			CR457037 CAG33318.1		
			NM_013305 NP_037437.1		
ENSP00000020221 (fragment)		n.d.	AC023295 -		
lactosylceramide α-2,3-sialyltransferase (ST3Gal V)	*Homo sapiens*	2.4.99.9	AF105026 **AAD14634.1**	Q9UNP4	
			AF119415 AAF66146.1	O94902	
			BC065936 AAH65936.1		
			AY152815 AAO16866.1		
			AAP65066 AAP65066.1		
			AY359105 AAQ89463.1		
			AB018356 BAA33950.1		
			AX876536 CAE89320.1		
			NM_003896 NP_003887.2		
N-acetylgalactosaminide α-2,6-sialyltransferase (ST6GalNAc VI)	*Homo sapiens*	2.4.99.-	BC006564 AAH06564.1	Q969X2	
			BC007802 AAH07802.1	Q9H8A2	
			BC016299 AAH16299.1	Q9ULB8	
			AY358672 AAQ89035.1		
			AB035173 **BAA87035.1**		
			AK023900 BAB14715.1		
			AJ507293 CAD45373.1		
			AX880950 CAE91145.1		
			CR457318 CAG33599.1		
			NM_013443 NP_038471.2		
N-acetylgalactosaminide α-2,6-sialyltransferase IV (ST6GalNAc IV)	*Homo sapiens*	2.4.99.-	AF127142 AAF00102.1	Q9H4F1	
			BC036705 AAH36705.1	Q9NWU6	
			- AAP63349.1	Q9UKU1	
			AB035172 **BAA87034.1**	Q9ULB9	
			AK000600 BAA91281.1	Q9Y3G3	
			Y17461 CAB44354.1	Q9Y3G4	
			AJ271734 CAC07404.1		
			AX061620 CAC24981.1		
			AX068265 CAC27250.1		
			AX969252 CAF14360.1		
			NM_014403 NP_055218.3		
			NM_175039 NP_778204.1		
ST8SIA-VI (fragment)	*Homo sapiens*	n.d.	AJ621583 **CAF21722.1**		
			XM_291725 XP_291725.2		
unnamed protein product	*Homo sapiens*	n.d.	AK021929 **BAB13940.1**	Q9HAA9	
			AX881696 CAE91353.1		
Gal β-1,3/4-GlcNAc α-2,3-sialyltransferase (ST3Gal III)	*Mesocricetus auratus*	2.4.99.6	AJ245699 **CAB53394.1**	Q9QXF6	

FIGURE 9F

Protein		Organism	EC#	GenBank / GenPept	SwissProt	PDB 3D
Gal β-1,3/4-GlcNAc α-2,3-sialyltransferase (ST3Gal IV)		*Mesocricetus auratus*	2.4.99.6	AJ245700 **CAB53395.1**	Q9QXF5	
GD3 synthase (fragment) ST8Sia I		*Mesocricetus auratus*	n.d.	AF141657 **AAD33879.1**	Q9WUL1	
polysialyltransferase (ST8Sia IV)		*Mesocricetus auratus*	2.4.99.-	AJ245701 **CAB53396.1**	Q9QXF4	
α-2,3-sialyltransferase ST3Gal I	*St3gal1*	*Mus musculus*	2.4.99.4	AF214028 AAF60973.1 AK031344 BAC27356.1 AK078469 BAC37290.1 X73523 CAA51919.1 NM_009177 NP_033203.1	**P54751** **Q11202** Q9JL30	
α-2,3-sialyltransferase ST3Gal II	*St3gal2*	*Mus musculus*	2.4.99.4	BC015264 **AAH15264.1** BC066064 AAH66064.1 AK034554 BAC28752.1 AK034863 BAC28859.1 AK053827 BAC35543.1 X76989 CAA54294.1 NM_009179 NP_033205.1 NM_178048 NP_835149.1	Q11204 Q8BPL0 Q8BSA0 Q8BSE9 Q91WH6	
α-2,3-sialyltransferase ST3Gal III	*St3gal3*	*Mus musculus*	2.4.99.-	BC006710 AAH06710.1 AK005053 **BAB23779.1** AK013016 BAB28598.1 X84234 CAA59013.1 NM_009176 NP_033202.2	P97325 Q922X5 Q9CZ48 Q9DBB6	
α-2,3-sialyltransferase ST3Gal IV	*St3gal4*	*Mus musculus*	2.4.99.4	BC011121 AAH11121.1 BC050773 **AAH50773.1** D28941 BAA06068.1 AK008543 BAB25732.1 AB061305 BAB47508.1 X95809 CAA65076.1 NM_009178 NP_033204.2	P97354 Q61325 Q91Y74 Q921R5 Q9CVE8	
α-2,3-sialyltransferase ST3Gal VI	*St3gal6*	*Mus musculus*	2.4.99.4	AF119390 AAD39130.1 BC052338 AAH52338.1 AB063326 **BAB79494.1** AK033562 BAC28360.1 AK041173 BAC30851.1 NM_018784 NP_061254	Q80UR7 Q8BLV1 Q8VIB3 Q9WVG2	
α-2,6-sialyltransferase ST6GalNAc II	*St6galnac2*	*Mus musculus*	2.4.99.-	NM_009180 6677963 BC010208 AAH10208.1 AB027198 BAB00637.1 AK004613 BAB23410.1 X93999 CAA63821.1 X94000 CAA63822.1 NM_009180 NP_033206.2	**P70277** Q9DC24 Q9JJM5	
α-2,6-sialyltransferase ST6Gal I	*St6gal1*	*Mus musculus*	2.4.99.1	-AAE68031.1 BC027833 AAH27833.1 D16106 BAA03680.1 AK034768 BAC28828.1 AK084124 BAC39120.1 NM_145933 NP_666045.1	**Q64685** Q8BM62 Q8K1L1	
α-2,6-sialyltransferase ST6Gal II	*St6gal2*	*Mus musculus*	n.d.	AK082566 BAC38534.1 AB095093 **BAC87752.1** AK129462 BAC98272.1 NM_172829 NP_766417.1	Q8BUU4	
α-2,6-sialyltransferase ST6GalNAc I	*St6galnac1*	*Mus musculus*	2.4.99.3	Y11274 **CAA72137.1** NM_011371 NP_035501.1	Q9QZ39 Q9JJP5	
α-2,6-sialyltransferase ST6GalNAc III	*St6galnac3*	*Mus musculus*	n.d.	BC058387 AAH58387.1 AK034804 BAC28836.1 Y11342 CAA72181.2 Y11343 **CAB95031.1** NM_011372 NP_035502	Q9WUV2 Q9JHP5	
α-2,6-sialyltransferase ST6GalNAc IV	*St6galnac4*	*Mus musculus*	2.4.99.7	BC056451 AAH56451.1 AK085730 BAC39523.1 AJ007310 CAA07446.1 Y15779 CAB43507.1	Q8C3J2 Q9JHP2 Q9R2B6 O88725	

FIGURE 9G

Protein		Organism	EC#	GenBank / GenPept	SwissProt	PDB ID
				Y15780 **CAB43514.1**	Q9JHP0	
				Y19055 CAB93946.1	Q9QUP9	
				Y19057 CAB93948.1	Q9R2B5	
				NM_011373 NP_035503.1		
α-2,8-sialyltransferase (GD3 synthase) ST8Sia I	*St8sia1*	*Mus musculus*	2.4.99.8	L38677 AAA91869.1	Q64468	
				BC024821 **AAH24821.1**	Q64687	
				AK046188 BAC32625.1	Q8BL76	
				AK052444 BAC34994.1	Q8BWI0	
				X84235 CAA59014.1	Q8K1C1	
				AJ401102 CAC20706.1	Q9EPK0	
				NM_011374 NP_035504.1		
α-2,8-sialyltransferase (ST8Sia VI)	*St8sia6*	*Mus musculus*	n.d.	AB059554 **BAC01265.1**	Q8BI43	
				AK085105 BAC39367.1	Q8K4T1	
				NM_145838 NP_665837.1		
α-2,8-sialyltransferase ST8Sia II	*St8sia2*	*Mus musculus*	2.4.99.-	X83562 CAA58548.1	**O35696**	
				X99646 CAA67965.1		
				X99647 CAA67965.1		
				X99648 CAA67965.1		
				X99649 CAA67965.1		
				X99650 CAA67965.1		
				X99651 CAA67965.1		
				NM_009181 NP_033207.1		
α-2,8-sialyltransferase ST8Sia IV	*St8sia4*	*Mus musculus*	2.4.99.8	BC060112 AAH60112.1	Q64692	
				AK003690 BAB22941.1	Q8BY70	
				AK041723 BAC31044.1		
				AJ223956 **CAA11685.1**		
				X86000 CAA59992.1		
				Y09484 CAA70692.1		
				NM_009183 NP_033209.1		
α-2,8-sialyltransferase ST8Sia V	*St8sia5*	*Mus musculus*	2.4.99.-	BC034855 AAH34855.1	P70126	
				AK078670 BAC37354.1	P70127	
				X98014 **CAA66642.1**	P70128	
				X98014 CAA66643.1	Q8BJW0	
				X98014 CAA66644.1	Q8JZQ3	
				NM_013666 NP_038694.1		
				NM_153124 NP_694764.1		
				NM_177416 NP_803135.1		
α-2,8-sialytransferase ST8Sia III	*St8sia3*	*Mus musculus*	2.4.99.-	BC075645 AAH75645.1	**Q64689**	
				AK015874 BAB30012.1	Q9CUJ6	
				X80502 CAA56665.1		
				NM_009182 NP_033208.1		
GD1 synthase (ST6GalNAc V)	*St6galnac5*	*Mus musculus*	n.d.	BC055737 **AAH55737.1**	Q8CAM7	
				AB030836 BAA85747.1	Q8CBX1	
				AB028840 BAA89292.1	Q9QYJ1	
				AK034387 BAC28693.1	Q9R0K6	
				AK038434 BAC29997.1		
				AK042683 BAC31331.1		
				NM_012028 NP_036158.2		
GM3 synthase (α-2,3-sialyltransferase) ST3Gal V	*St3gal5*	*Mus musculus*	2.4.99.9	AF119416 **AAF66147.1**	O88829	
				- AAP65063.1	Q9CZ65	
				AB018048 BAA33491.1	Q9QWF9	
				AB013302 BAA76467.1		
				AK012961 BAB28571.1		
				Y15003 CAA75235.1		
				NM_011375 NP_035505.1		
N-acetylgalactosaminide α-2,6-sialyltransferase (ST6GalNAc VI)	*St6galnac6*	*Mus musculus*	2.4.99.-	BC036985 **AAH36985.1**	Q8CDC3	
				AB035174 BAA87036.1	Q8JZW3	
				AB035123 BAA95940.1	Q9JM95	
				AK030648 BAC27064.1	Q9R0G9	
				NM_016973 NP_058669.1		
M138L		*Myxoma virus*	n.d.	U46578 AAD00069.1		
				AF170726 AAE61323.1		
				NC_001132 AAE61326.1		
				AAF15026.1		
				NP_051852.1		
α-2,3-sialyltransferase		*Oncorhynchus*	n.d.	AJ585760 **CAE51384.1**		

FIGURE 9H

Protein	Organism	EC#	GenBank / GenPept	SwissProt	PDB 3D
(St3Gal-I)	*mykiss*				
α-2,6-sialyltransferase (Siat1)	*Oncorhynchus mykiss*	n.d.	AJ620649 **CAF05848.1**		
α-2,8-polysialyltransferase IV (ST8Sia IV)	*Oncorhynchus mykiss*	n.d.	AB094402 **BAC77411.1**	Q7T2X5	
GalNAc α-2,6-sialyltransferase (RtST6GalNAc)	*Oncorhynchus mykiss*	n.d.	AB097943 **BAC77520.1**	Q7T2X4	
α-2,3-sialyltransferase ST3Gal IV	*Oryctolagus cuniculus*	2.4.99.-	AF121967 **AAF28871.1**	Q9N257	
OJ1217_F02.7	*Oryza sativa (japonica cultivar-group)*	n.d.	AP004084 **BAD07616.1**		
OSJNBa0043L24.2 or OSJNBb0002J11.9	*Oryza sativa (japonica cultivar-group)*	n.d.	AL731626 **CAD41185.1** AL662969 CAE04714.1		
P0683f02.18 or P0489B03.1	*Oryza sativa (japonica cultivar-group)*	n.d.	AP003289 **BAB63715.1** AP003794 BAB90552.1		
α-2,6-sialyltransferase ST6GalNAc V (Siat7E) (fragment)	*Oryzias latipes*	n.d.	AJ646876 **CAG26705.1**		
α-2,3-sialyltransferase ST3Gal I (Siat4)	*Pan troglodytes*	n.d.	AJ744803 **CAG32839.1**		
α-2,3-sialyltransferase ST3Gal II (Siat5)	*Pan troglodytes*	n.d.	AJ744804 **CAG32840.1**		
α-2,3-sialyltransferase ST3Gal III (Siat6)	*Pan troglodytes*	n.d.	AJ626819 **CAF25177.1**		
α-2,3-sialyltransferase ST3Gal IV (Siat4c)	*Pan troglodytes*	n.d.	AJ626824 **CAF25182.1**		
α-2,3-sialyltransferase ST3Gal VI (Siat10)	*Pan troglodytes*	n.d.	AJ744808 **CAG32844.1**		
α-2,6-sialyltransferase (Sia7A)	*Pan troglodytes*	n.d.	AJ748740 **CAG38615.1**		
α-2,6-sialyltransferase (Sia7B)	*Pan troglodytes*	n.d.	AJ748741 **CAG38616.1**		
α-2,6-sialyltransferase ST6GalNAc III (Siat7C)	*Pan troglodytes*	n.d.	AJ634454 **CAG25676.1**		
α-2,6-sialyltransferase ST6GalNAc IV (Siat7D) (fragment)	*Pan troglodytes*	n.d.	AJ646870 **CAG26699.1**		
α-2,6-sialyltransferase ST6GalNAc V (Siat7E)	*Pan troglodytes*	n.d.	AJ646875 **CAG26704.1**		
α-2,6-sialyltransferase ST6GalNAc VI (Siat7F) (fragment)	*Pan troglodytes*	n.d.	AJ646882 **CAG26711.1**		
α-2,8-sialyltransferase 8A (Siat8A)	*Pan troglodytes*	2.4.99.8	AJ697658 **CAG26896.1**		
α-2,8-sialyltransferase 8B (Siat8B)	*Pan troglodytes*	n.d.	AJ697659 **CAG26897.1**		
α-2,8-sialyltransferase 8C (Siat8C)	*Pan troglodytes*	n.d.	AJ697660 **CAG26898.1**		
α-2,8-sialyltransferase 8D (Siat8D)	*Pan troglodytes*	n.d.	AJ697661 **CAG26899.1**		
α-2,8-sialyltransferase 8E (Siat8E)	*Pan troglodytes*	n.d.	AJ697662 **CAG26900.1**		
α-2,8-sialyltransferase 8F (Siat8F)	*Pan troglodytes*	n.d.	AJ697663 **CAG26901.1**		
β-galactosamide α-2,6-sialyltransferase I (ST6Gal I; Siat1)	*Pan troglodytes*	2.4.99.1	AJ627624 **CAF29492.1**		
β-galactosamide α-2,6-sialyltransferase II (ST6Gal II)	*Pan troglodytes*	n.d.	AJ627625 **CAF29493.1**		
GM3 synthase ST3Gal V	*Pan troglodytes*	n.d.	AJ744807 **CAG32843.1**		

FIGURE 9I

Protein	Organism	EC#	GenBank / GenPept	SwissProt	PDB 3D
(Siat9) S138L	*Rabbit fibroma virus Kasza*	n.d.	NC_001266 **NP_052025**		
α-2,3-sialyltransferase ST3Gal III	*Rattus norvegicus*	2.4.99.6	M97754 AAA42146.1 NM_031697 NP_113885.1	**Q02734**	
α-2,3-sialyltransferase ST3Gal IV (Siat4c)	*Rattus norvegicus*	n.d.	AJ626825 **CAF25183.1**		
α-2,3-sialyltransferase ST3Gal Vi	*Rattus norvegicus*	n.d.	AJ626743 **CAF25053.1**		
α-2,6-sialyltransferase ST3Gal II	*Rattus norvegicus*	2.4.99.-	X76988 CAA54293.1 NM_031695 NP_113883.1	**Q11205**	
α-2,6-sialyltransferase ST6Gal I	*Rattus norvegicus*	2.4.99.1	M18769 AAA41196.1 M83143 AAB07233.1	**P13721**	
α-2,6-sialyltransferase ST6GalNAc I (Siat7A)	*Rattus norvegicus*	n.d.	AJ634458 **CAG25684.1**		
α-2,6-sialyltransferase ST6GalNAc II (Siat7B)	*Rattus norvegicus*	n.d.	AJ634457 **CAG25679.1**		
α-2,6-sialyltransferase ST6GalNAc III	*Rattus norvegicus*	2.4.99.-	L29554 AAC42086.1 BC072501 AAH72501.1 NM_019123 NP_061996.1	**Q64686**	
α-2,6-sialyltransferase ST6GalNAc IV (Siat7D) (fragment)	*Rattus norvegicus*	n.d.	AJ646871 **CAG26700.1**		
α-2,6-sialyltransferase ST6GalNAc V (Siat7E)	*Rattus norvegicus*	n.d.	AJ646872 **CAG26701.1**		
α-2,6-sialyltransferase ST6GalNAc VI (Siat7F) (fragment)	*Rattus norvegicus*	n.d.	AJ646881 **CAG26710.1**		
α-2,8-sialyltransferase (GD3 synthase) ST8Sia I	*Rattus norvegicus*	2.4.99.-	U53883 AAC27541.1 D45255 BAA08213.1	**P70554** **P97713**	
α-2,8-sialyltransferase (SIAT8E)	*Rattus norvegicus*	n.d.	AJ699422 **CAG27884.1**		
α-2,8-sialyltransferase (SIAT8F)	*Rattus norvegicus*	n.d.	AJ699423 **CAG27885.1**		
α-2,8-sialyltransferase ST8Sia II	*Rattus norvegicus*	2.4.99.-	L13445 AAA42147.1 NM_057156 NP_476497.1	**Q07977** **Q64688**	
α-2,8-sialyltransferase ST8Sia III	*Rattus norvegicus*	2.4.99.-	U55938 AAB50061.1 NM_013029 NP_037161.1	**P97877**	
α-2,8-sialyltransferase ST8Sia IV	*Rattus norvegicus*	2.4.99.-	U90215 AAB49989.1	**O08563**	
β-galactosamide α-2,6-sialyltransferase II (ST6Gal II)	*Rattus norvegicus*	n.d.	AJ627626 **CAF29494.1**		
GM3 synthase ST3Gal V	*Rattus norvegicus*	n.d.	AB018049 BAA33492.1 NM_031337 NP_112627.1	**O88830**	
sialyltransferase ST3Gal-I (Siat4A)	*Rattus norvegicus*	n.d.	AJ748840 **CAG44449.1**		
α-2,3-sialyltransferase (St3Gal-II)	*Silurana tropicalis*	n.d.	AJ585763 **CAE51387.1**		
α-2,6-sialyltransferase (Siat7b)	*Silurana tropicalis*	n.d.	AJ620650 **CAF05849.1**		
α-2,6-sialyltransferase (St6galnac)	*Strongylocentrotus purpuratus*	n.d.	AJ699425 **CAG27887.1**		
α-2,3-sialyltransferase (ST3GAL-III)	*Sus scrofa*	n.d.	AJ585765 **CAE51389.1**		
α-2,3-sialyltransferase (ST3GAL-IV)	*Sus scrofa*	n.d.	AJ584674 **CAE48299.1**		
α-2,3-sialyltransferase ST3Gal I	*Sus scrofa*	2.4.99.4	M97753 AAA31125.1	**Q02745**	
α-2,6-sialyltransferase (fragment) ST6Gal I	*Sus scrofa*	2.4.99.1	AF136746 **AAD33059.1**	Q9XSG8	
β-galactosamide α-2,6-sialyltransferase (ST6GalNAc-V)	*Sus scrofa*	n.d.	AJ620948 **CAF06585.2**		
sialyltransferase (fragment) ST6Gal I	*sus scrofa*	n.d.	AF041031 **AAC15633.1**	O62717	

FIGURE 9J

Protein	Organism	EC	GenBank / GenPept	SwissProt	PDB / 3D
ST6GALNAC-V	*Sus scrofa*	n.d.	AJ620948 CAF06585.1		
α-2,3-sialyltransferase (Siat5-r)	*Takifugu rubripes*	n.d.	AJ744805 CAG32841.1		
α-2,3-sialyltransferase ST3Gal I (Siat4)	*Takifugu rubripes*	n.d.	AJ626816 CAF25174.1		
α-2,3-sialyltransferase ST3Gal II (Siat5) (fragment)	*Takifugu rubripes*	n.d.	AJ626817 CAF25175.1		
α-2,3-sialyltransferase ST3Gal III (Siat6)	*Takifugu rubripes*	n.d.	AJ626818 CAF25176.1		
α-2,6-sialyltransferase ST6Gal I (Siat1)	*Takifugu rubripes*	n.d.	AJ744800 CAG32836.1		
α-2,6-sialyltransferase ST6GalNAc II (Siat7B)	*Takifugu rubripes*	n.d.	AJ634460 CAG25681.1		
α-2,6-sialyltransferase ST6GalNAc II B (Siat7B-related)	*Takifugu rubripes*	n.d.	AJ634461 CAG25682.1		
α-2,6-sialyltransferase ST6GalNAc III (Siat7C) (fragment)	*Takifugu rubripes*	n.d.	AJ634456 CAG25678.1		
α-2,6-sialyltransferase ST6GalNAc IV (siat7D) (fragment)	*Takifugu rubripes*	2.4.99.3	Y17466 CAB44338.1 AJ646869 CAG26698.1	Q9W6U6	
α-2,6-sialyltransferase ST6GalNAc V (Siat7E) (fragment)	*Takifugu rubripes*	n.d.	AJ646873 CAG26702.1		
α-2,6-sialyltransferase ST6GalNAc VI (Siat7F) (fragment)	*Takifugu rubripes*	n.d.	AJ646880 CAG26709.1		
α-2,8-sialyltransferase ST8Sia I (Siat 8A) (fragment)	*Takifugu rubripes*	n.d.	AJ715534 CAG29373.1		
α-2,8-sialyltransferase ST8Sia II (Siat 8B) (fragment)	*Takifugu rubripes*	n.d.	AJ715538 CAG29377.1		
α-2,8-sialyltransferase ST8Sia III (Siat 8C) (fragment)	*Takifugu rubripes*	n.d.	AJ715541 CAG29380.1		
α-2,8-sialyltransferase ST8Sia IIIr (Siat 8Cr)	*Takifugu rubripes*	n.d.	AJ715542 CAG29381.1		
α-2,8-sialyltransferase ST8Sia V (Siat 8E) (fragment)	*Takifugu rubripes*	n.d.	AJ715547 CAG29386.1		
α-2,8-sialyltransferase ST8Sia VI (Siat 8F) (fragment)	*Takifugu rubripes*	n.d.	AJ715549 CAG29388.1		
α-2,8-sialyltransferase ST8Sia VIr (Siat 8Fr)	*Takifugu rubripes*	n.d.	AJ715550 CAG29389.1		
α-2,3-sialyltransferase (Siat5-r)	*Tetraodon nigroviridis*	n.d.	AJ744806 CAG32842.1		
α-2,3-sialyltransferase ST3Gal I (Siat4)	*Tetraodon nigroviridis*	n.d.	AJ744802 CAG32838.1		
α-2,3-sialyltransferase ST3Gal III (Siat6)	*Tetraodon nigroviridis*	n.d.	AJ626822 CAF25180.1		
α-2,6-sialyltransferase ST6GalNAc II (Siat7B)	*Tetraodon nigroviridis*	n.d.	AJ634462 CAG25683.1		
α-2,6-sialyltransferase ST6GalNAc V (Siat7E) (fragment)	*Tetraodon nigroviridis*	n.d.	AJ646879 CAG26708.1		
α-2,8-sialyltransferase ST8Sia I (Siat 8A) (fragment)	*Tetraodon nigroviridis*	n.d.	AJ715536 CAG29375.1		
α-2,8-sialyltransferase ST8Sia II (Siat 8B) (fragment)	*Tetraodon nigroviridis*	n.d.	AJ715537 CAG29376.1		
α-2,8-sialyltransferase	*Tetraodon*	n.d.	AJ715539 CAG29378.1		

FIGURE 9K

Protein	Organism	EC#	GenBank / GenPept	SwissProt	PDB 3D
ST8Sia III (Siat 8C) (fragment)	*nigroviridis*				
α-2,8-sialyltransferase ST8Sia IIIr (Siat 8Cr) (fragment)	*Tetraodon nigroviridis*	n.d.	AJ715540 **CAG29379.1**		
α-2,8-sialyltransferase ST8Sia V (Siat 8E) (fragment)	*Tetraodon nigroviridis*	n.d.	AJ715548 **CAG29387.1**		
α-2,3-sialyltransferase (St3Gal-II)	*Xenopus laevis*	n.d.	AJ585762 **CAE51386.1**		
α-2,3-sialyltransferase (St3Gal-VI)	*Xenopus laevis*	n.d.	AJ585766 **CAE51390.1**		
α-2,3-sialyltransferase St3Gal-III (Siat6)	*Xenopus laevis*	n.d.	AJ585764 **CAE51388.1** AJ626823 CAF25181.1		
α-2,8-polysialyltransferase	*Xenopus laevis*	2.4.99.-	AB007468 BAA32617.1	O93234	
α-2,8-sialyltransferase ST8Siα-I (Siat8A;GD3 synthase)	*Xenopus laevis*	n.d.	AY272056 AAQ16162.1 AY272057 AAQ16163.1 AJ704562 **CAG28695.1**		
Unknown (protein for MGC:81265)	*Xenopus laevis*	n.d.	BC068760 **AAH68760.1**		
α-2,3-sialyltransferase (3Gal-VI)	*Xenopus tropicalis*	n.d.	AJ626744 **CAF25054.1**		
α-2,3-sialyltransferase (Siat4c)	*Xenopus tropicalis*	n.d.	AJ622908 **CAF22058.1**		
α-2,6-sialyltransferase ST6GalNAc V (Siat7E) (fragment)	*Xenopus tropicalis*	n.d.	AJ646878 **CAG26707.1**		
α-2,8-sialyltransferase ST8Sia III (Siat 8C) (fragment)	*Xenopus tropicalis*	n.d.	AJ715544 **CAG29383.1**		
β-galactosamide α-2,6-sialyltransferase II (ST6Gal II)	*Xenopus tropicalis*	n.d.	AJ627628 **CAF29496.1**		
sialytransferase St8Sial	*Xenopus tropicalis*	n.d.	AY652775 **AAT67042**		
poly-α-2,8-sialosyl sialyltransferase (NeuS)	*Escherichia coli K1*	2.4.-.-	M76370 AAA24213.1 X60598 CAA43053.1	**Q57269**	
polysialyltransferase	*Escherichia coli K92*	2.4.-.-	M88479 **AAA24215.1**	Q47404	
α-2,8 polysialyltransferase SiaD	*Neisseria meningitidis B1940*	2.4.-.-	M95053 AAA20478.1 X78068 CAA54985.1	**Q51281** **Q51145**	
SynE	*Neisseria meningitidis FAM18*	n.d.	U75650 **AAB53842.1**	O06435	
polysialyltransferase (SiaD)(fragment)	*Neisseria meningitidis M1019*	n.d.	AY234192 **AAO85290.1**		
SiaD (fragment)	*Neisseria meningitidis M209*	n.d.	AY281046 **AAP34769.1**		
SiaD (fragment)	*Neisseria meningitidis M3045*	n.d.	AY281044 **AAP34767.1**		
polysialyltransferase (SiaD)(fragment)	*Neisseria meningitidis M3315*	n.d.	AY234191 **AAO85289.1**		
SiaD (fragment)	*Neisseria meningitidis M3515*	n.d.	AY281047 **AAP34770.1**		
polysialyltransferase (SiaD)(fragment)	*Neisseria meningitidis M4211*	n.d.	AY234190 **AAO85288.1**		
SiaD (fragment)	*Neisseria meningitidis M4642*	n.d.	AY281048 **AAP34771.1**		
polysialyltransferase (SiaD)(fragment)	*Neisseria meningitidis M5177*	n.d.	AY234193 **AAO85291.1**		
SiaD	*Neisseria meningitidis M5178*	n.d.	AY281043 **AAP34766.1**		
SiaD (fragment)	*Neisseria meningitidis M980*	n.d.	AY281045 **AAP34768.1**		
NMB0067	*Neisseria meningitidis MC58*	n.d.	NC_003112 **NP_273131**		

FIGURE 9L

Protein	Organism	EC#	GenBank / GenPept	SwissProt	PDB 3D
Lst	*Aeromonas punctata Sch3*	n.d.	AF126256 **AAS66624.1**		
ORF2	*Haemophilus influenzae A2*	n.d.	M94855 **AAA24979.1**		
HI1699	*Haemophilus influenzae Rd*	n.d.	U32842 **AAC23345.1** NC_000907 NP_439841.1	Q48211	
α-2,3-sialyltransferase	*Neisseria gonorrhoeae F62*	2.4.99.4	U60664 **AAC44539.1** AAE67205.1	P72074	
α-2,3-sialyltransferase	*Neisseria meningitidis 126E, NRCC 4010*	2.4.99.4	U60662 **AAC44544.2**		
α-2,3-sialyltransferase	*Neisseria meningitidis 406Y, NRCC 4030*	2.4.99.4	U60661 **AAC44543.1**		
α-2,3-sialyltransferase (NMB0922)	*Neisseria meningitidis MC58*	2.4.99.4	U60660 AAC44541.1 AE002443 AAF41330.1 NC_003112 **NP_273962.1**	P72097	
NMA1118	*Neisseria meningitidis Z2491*	n.d.	AL162755 CAB84380.1 NC_003116 NP_283887.1	**Q9JUV5**	
PM0508	*Pasteurella multocida PM70*	n.d.	AE006086 **AAK02592.1** NC_002663 NP_245445.1	Q9CNC4	
WaaH	*Salmonella enterica SARB25*	n.d.	AF519787 **AAM82550.1**	Q8KS93	
WaaH	*Salmonella enterica SARB3*	n.d.	AF519788 **AAM82551.1**	Q8KS92	
WaaH	*Salmonella enterica SARB39*	n.d.	AF519789 **AAM82552.1**		
WaaH	*Salmonella enterica SARB53*	n.d.	AF519790 **AAM82553.1**		
WaaH	*Salmonella enterica SARB57*	n.d.	AF519791 **AAM82554.1**	Q8KS91	
WaaH	*Salmonella enterica SARB71*	n.d.	AF519793 **AAM82556.1**	Q8KS89	
WaaH	*Salmonella enterica SARB8*	n.d.	AF519792 **AAM82555.1**	Q8KS90	
WaaH	*Salmonella enterica SARC10V*	n.d.	AF519779 **AAM88840.1**	Q8KS99	
WaaH (fragment)	*Salmonella enterica SARC12*	n.d.	AF519781 **AAM88842.1**		
WaaH (fragment)	*Salmonella enterica SARC13I*	n.d.	AF519782 **AAM88843.1**	Q8KS98	
WaaH (fragment)	*Salmonella enterica SARC14I*	n.d.	AF519783 **AAM88844.1**	Q8KS97	
WaaH	*Salmonella enterica SARC15II*	n.d.	AF519784 **AAM88845.1**	Q8KS96	
WaaH	*Salmonella enterica SARC16II*	n.d.	AF519785 **AAM88846.1**	Q8KS95	

FIGURE 9M

Protein	Organism	EC#	GenBank / GenRept	SwissProt	PDB 3D
WaaH (fragment)	*Salmonella enterica SARC3I*	n.d.	AF519772 **AAM88834.1**	Q8KSA4	
WaaH (fragment)	*Salmonella enterica SARC4I*	n.d.	AF519773 **AAM88835.1**	Q8KSA3	
WaaH	*Salmonella enterica SARC5IIa*	n.d.	AF519774 **AAM88836.1**		
WaaH	*Salmonella enterica SARC6IIa*	n.d.	AF519775 **AAM88837.1**	Q8KSA2	
WaaH	*Salmonella enterica SARC8*	n.d.	AF519777 **AAM88838.1**	Q8KSA1	
WaaH	*Salmonella enterica SARC9V*	n.d.	AF519778 **AAM88839.1**	Q8KSA0	
UDP-glucose: α-1,2-glucosyltransferase (WaaH)	*Salmonella enterica subsp. arizonae SARC 5*	2.4.1.-	AF511116 **AAM48166.1**		
bifunctional α-2,3/-2,8-sialyltransferase (Cst-II)	*Campylobacter jejuni ATCC 43449*	n.d.	AF401529 **AAL06004.1**	Q93CZ5	
Cst	*Campylobacter jejuni 81-176*	n.d.	AF305571 **AAL09368.1**		
α-2,3-sialyltransferase (Cst-III)	*Campylobacter jejuni ATCC 43429*	2.4.99.-	AY044156 **AAK73183.1**		
α-2,3-sialyltransferase (Cst-III)	*Campylobacter jejuni ATCC 43430*	2.4.99.-	AF400047 **AAK85419.1**		
α-2,3-sialyltransferase (Cst-II)	*Campylobacter jejuni ATCC 43432*	2.4.99.-	AF215659 **AAG43979.1**	Q9F0M9	
α-2,3/8-sialyltransferase (CstII)	*Campylobacter jejuni ATCC 43438*	n.d.	AF400048 **AAK91725.1**	Q93MQ0	
α-2,3-sialyltransferase cst-II	*Campylobacter jejuni ATCC 43446*	2.4.99.-	AF167344 **AAF34137.1**		
α-2,3-sialyltransferase (Cst-II)	*Campylobacter jejuni ATCC 43456*	2.4.99.-	AF401528 **AAL05990.1**	Q93D05	
α-2,3-/α-2,8-sialyltransferase (CstII)	*Campylobacter jejuni ATCC 43460*	2.4.99.-	AY044868 **AAK96001.1**	Q938X6	
α-2,3/8-sialyltransferase (Cst-II)	*Campylobacter jejuni ATCC 700297*	n.d.	AF216647 **AAL36462.1**		
ORF	*Campylobacter jejuni GB11*	n.d.	AY422197 **AAR82875.1**		
α-2,3-sialyltransferase cstIII	*Campylobacter jejuni MSC57360*	2.4.99.-	AF195055 **AAG29922.1**		
α-2,3-sialyltransferase cstIII Cj1140	*Campylobacter jejuni NCTC 11168*	2.4.99.-	AL139077 **CAB73395.1** NC_002163 NP_282288.1	Q9PNF4	
α-2,3/α-2,8-sialyltransferase II (cstII)	*Campylobacter jejuni O:10*	n.d.	-AAO96669.1 AX934427 **CAF04167.1**		
α-2,3/α-2,8-sialyltransferase II (CstII)	*Campylobacter jejuni O:19*	n.d.	AX934431 **CAF04169.1**		
α-2,3/α-2,8-sialyltransferase II (CstII)	*Campylobacter jejuni O:36*	n.d.	AX934436 **CAF04171.1**		
α-2,3/α-2,8-sialyltransferase II (CstII)	*Campylobacter jejuni O:4*	n.d.	AX934434 **CAF04170.1**		
α-2,3/α-2,8-sialyltransferase II (CstII)	*Campylobacter jejuni O:41*	n.d.	-AAO96670.1 -AAT17967.1 AX934429 **CAF04168.1**		
α-2,3-sialyltransferase cst-I	*Campylobacter jejuni OH4384*	2.4.99.-	AF130466 **AAF13495.1** -AAS36261.1	Q9RGF1	
bifunctional α-2,3/-2,8-sialyltransferase (Cst-II)	*Campylobacter jejuni OH4384*	2.4.99.-	AF130984 **AAF31771.1** AX934425 CAF04166.1		1RO7 C 1RO8 A

FIGURE 9N

Protein	Organism	EC#	GenBank / GenPept	SwissProt	PDB 3D
HI0352 (fragment)	*Haemophilus influenzae Rd*	n.d.	U32720 AAC22013.1 X57315 CAA40567.1 NC_000907 NP_438516.1	P24324	
PM1174	*Pasteurella multocida PM70*	n.d.	AE006157 AAK03258.1 NC_002663 NP_246111.1	Q9CLP3	
Sequence 10 from patent US 6503744	*Unknown.*	n.d.	-AAO96672.1		
Sequence 10 from patent US 6699705	*Unknown.*	n.d.	-AAT17969.1		
Sequence 12 from patent US 6699705	*Unknown.*	n.d.	-AAT17970.1		
Sequence 2 from patent US 6709834	*Unknown.*	n.d.	-AAT23232.1		
Sequence 3 from patent US 6503744	*Unknown.*	n.d.	-AAO96668.1		
Sequence 3 from patent US 6699705	*Unknown.*	n.d.	-AAT17965.1		
Sequence 34 from patent US 6503744	*Unknown.*	n.d.	-AAO96684.1		
Sequence 35 from patent US 6503744 (fragment)	*Unknown.*	n.d.	-AAO96685.1 -AAS36262.1		
Sequence 48 from patent US 6699705	*Unknown.*	n.d.	-AAT17988.1		
Sequence 5 from patent US 6699705	*Unknown.*	n.d.	-AAT17966.1		
Sequence 9 from patent US 6503744	*Unknown.*	n.d.	-AAO96671.1		