Accession number	Description
IPI00000024	Isoform 1 of Protocadherin-1 precursor
IPI00000024	Isoform 1 of Protocadherin-1 precursor
IPI00000137	N-acetylglucosamine-1-phosphotransferase subunit gamma
IPI00000137	N-acetylglucosamine-1-phosphotransferase subunit gamma
IPI00000137	N-acetylglucosamine-1-phosphotransferase subunit gamma
IPI00000137	N-acetylglucosamine-1-phosphotransferase subunit gamma
IPI00000137	N-acetylglucosamine-1-phosphotransferase subunit gamma
IPI00000137	N-acetylglucosamine-1-phosphotransferase subunit gamma
IPI00000138	Alpha-1,3-mannosyl-glycoprotein 2-beta-N-acetylglucosami
IPI00000138	Alpha-1,3-mannosyl-glycoprotein 2-beta-N-acetylglucosami
IPI00000138	Alpha-1,3-mannosyl-glycoprotein 2-beta-N-acetylglucosami
IPI00000138	Alpha-1,3-mannosyl-glycoprotein 2-beta-N-acetylglucosami
IPI00000190	CD81 antigen
IPI00000190	CD81 antigen
IPI00000265	hypothetical protein LOC221061
IPI00000265	hypothetical protein LOC221061
IPI00000779	Isoform 1 of ADAM 22 precursor
IPI00000779	Isoform 1 of ADAM 22 precursor
IPI00000779	Isoform 1 of ADAM 22 precursor
IPI00000779	Isoform 1 of ADAM 22 precursor
IPI00000779	Isoform 1 of ADAM 22 precursor
IPI00000816	14-3-3 protein epsilon
IPI00000816	14-3-3 protein epsilon
IPI00000816	14-3-3 protein epsilon
IPI00000824	Isoform A of NT-3 growth factor receptor precursor
IPI00000874	Peroxiredoxin-1
IPI00000874	Peroxiredoxin-1
IPI00000874	Peroxiredoxin-1
IPI00000874	Peroxiredoxin-1
IPI00000874	Peroxiredoxin-1
IPI00000874	Peroxiredoxin-1
IPI00000874	Peroxiredoxin-1
IPI00000949	Mu-crystallin homolog
IPI00000949	Mu-crystallin homolog
IPI00000949	Mu-crystallin homolog
IPI00001516	Isoform Long of Protocadherin alpha C2 precursor
IPI00001516	Isoform Long of Protocadherin alpha C2 precursor
IPI00001516	Isoform Long of Protocadherin alpha C2 precursor
IPI00001516	Isoform Long of Protocadherin alpha C2 precursor
IPI00001516	Isoform Long of Protocadherin alpha C2 precursor
IPI00001516	Isoform Long of Protocadherin alpha C2 precursor
IPI00001516	Isoform Long of Protocadherin alpha C2 precursor
IPI00001516	Isoform Long of Protocadherin alpha C2 precursor
IPI00001516	Isoform Long of Protocadherin alpha C2 precursor
IPI00001611	Isoform 1 of Insulin-like growth factor II precursor
IPI00001611	Isoform 1 of Insulin-like growth factor II precursor
IPI00001611	Isoform 1 of Insulin-like growth factor II precursor
IPI00001662	Opioid-binding protein/cell adhesion molecule precursor
IPI00001662	Opioid-binding protein/cell adhesion molecule precursor
IPI00001662	Opioid-binding protein/cell adhesion molecule precursor
IPI00001662	Opioid-binding protein/cell adhesion molecule precursor
IPI00001662	Opioid-binding protein/cell adhesion molecule precursor
IPI00001662	Opioid-binding protein/cell adhesion molecule precursor
IPI00001662	Opioid-binding protein/cell adhesion molecule precursor
IPI00001734	Isoform 1 of Phosphoserine aminotransferase
IPI00001734	Isoform 1 of Phosphoserine aminotransferase

IPI00001734	Isoform 1 of Phosphoserine aminotransferase
IPI00001734	Isoform 1 of Phosphoserine aminotransferase
IPI00001734	Isoform 1 of Phosphoserine aminotransferase
IPI00001753	Myosin-4
IPI00001753	Myosin-4
IPI00001893	Isoform A of Protocadherin-7 precursor
IPI00001893	Isoform A of Protocadherin-7 precursor
IPI00001895	Isoform 1 of Protocadherin-8 precursor
IPI00001952	Endonuclease domain-containing 1 protein precursor
IPI00001952	Endonuclease domain-containing 1 protein precursor
IPI00001952	Endonuclease domain-containing 1 protein precursor
IPI00001952	Endonuclease domain-containing 1 protein precursor
IPI00001952	Endonuclease domain-containing 1 protein precursor
IPI00001952	Endonuclease domain-containing 1 protein precursor
IPI00001952	Endonuclease domain-containing 1 protein precursor
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IPI00001952	Endonuclease domain-containing 1 protein precursor
IPI00001952	Endonuclease domain-containing 1 protein precursor
IPI00001952	Endonuclease domain-containing 1 protein precursor
IPI00001952	Endonuclease domain-containing 1 protein precursor
IPI00001952	Endonuclease domain-containing 1 protein precursor
IPI00001952	Endonuclease domain-containing 1 protein precursor
IPI00002142	Protocadherin-10 precursor
IPI00002142	Protocadherin-10 precursor
IPI00002142	Protocadherin-10 precursor
IPI00002142	Protocadherin-10 precursor
IPI00002147	Chitinase-3-like protein 1 precursor
IPI00002147	Chitinase-3-like protein 1 precursor
IPI00002147	Chitinase-3-like protein 1 precursor
IPI00002147	Chitinase-3-like protein 1 precursor
IPI00002147	Chitinase-3-like protein 1 precursor
IPI00002147	Chitinase-3-like protein 1 precursor
IPI00002147	Chitinase-3-like protein 1 precursor
IPI00002147	Chitinase-3-like protein 1 precursor
IPI00002147	Chitinase-3-like protein 1 precursor
IPI00002147	Chitinase-3-like protein 1 precursor
IPI00002147	Chitinase-3-like protein 1 precursor
IPI00002147	Chitinase-3-like protein 1 precursor
IPI00002147	Chitinase-3-like protein 1 precursor
IPI00002147	Chitinase-3-like protein 1 precursor
IPI00002147	Chitinase-3-like protein 1 precursor
IPI00002147	Chitinase-3-like protein 1 precursor
IPI00002147	Chitinase-3-like protein 1 precursor
IPI00002147	Chitinase-3-like protein 1 precursor
IPI00002147	Chitinase-3-like protein 1 precursor
IPI00002147	Chitinase-3-like protein 1 precursor
IPI00002211	Isoform 2 of Semaphorin-6A precursor
IPI00002236	Lactadherin precursor
IPI00002236	Lactadherin precursor
IPI00002236	Lactadherin precursor
IPI00002236	Lactadherin precursor
IPI00002236	Lactadherin precursor
IPI00002236	Lactadherin precursor
IPI00002280	proprotein convertase subtilisin/kexin type 1 inhibitor precu

IPI00002280	proprotein convertase subtilisin/kexin type 1 inhibitor precur
IPI00002280	proprotein convertase subtilisin/kexin type 1 inhibitor precur
IPI00002280	proprotein convertase subtilisin/kexin type 1 inhibitor precur
IPI00002280	proprotein convertase subtilisin/kexin type 1 inhibitor precur
IPI00002280	proprotein convertase subtilisin/kexin type 1 inhibitor precur
IPI00002280	proprotein convertase subtilisin/kexin type 1 inhibitor precur
IPI00002280	proprotein convertase subtilisin/kexin type 1 inhibitor precur
IPI00002525	Neudesin precursor
IPI00002525	Neudesin precursor
IPI00002714	Dickkopf-related protein 3 precursor
IPI00002714	Dickkopf-related protein 3 precursor
IPI00002714	Dickkopf-related protein 3 precursor
IPI00002714	Dickkopf-related protein 3 precursor
IPI00002714	Dickkopf-related protein 3 precursor
IPI00002714	Dickkopf-related protein 3 precursor
IPI00002714	Dickkopf-related protein 3 precursor
IPI00002714	Dickkopf-related protein 3 precursor
IPI00002714	Dickkopf-related protein 3 precursor
IPI00002714	Dickkopf-related protein 3 precursor
IPI00002714	Dickkopf-related protein 3 precursor
IPI00002732	Exostosin-like 2
IPI00002732	Exostosin-like 2
IPI00002732	Exostosin-like 2
IPI00002732	Exostosin-like 2
IPI00002732	Exostosin-like 2
IPI00002732	Exostosin-like 2
IPI00002732	Exostosin-like 2
IPI00002732	Exostosin-like 2
IPI00002745	Cathepsin Z precursor
IPI00002745	Cathepsin Z precursor
IPI00002745	Cathepsin Z precursor
IPI00002745	Cathepsin Z precursor
IPI00002745	Cathepsin Z precursor
IPI00002745	Cathepsin Z precursor
IPI00002816	Cathepsin F precursor
IPI00002816	Cathepsin F precursor
IPI00002816	Cathepsin F precursor
IPI00002818	Isoform 1 of Kallikrein-11 precursor
IPI00002818	Isoform 1 of Kallikrein-11 precursor
IPI00002925	Cocaine- and amphetamine-regulated transcript protein pre
IPI00002925	Cocaine- and amphetamine-regulated transcript protein pre
IPI00003102	Ciliary neurotrophic factor receptor alpha precursor
IPI00003176	Serine protease HTRA1 precursor
IPI00003176	Serine protease HTRA1 precursor
IPI00003176	Serine protease HTRA1 precursor
IPI00003351	Extracellular matrix protein 1 precursor
IPI00003351	Extracellular matrix protein 1 precursor
IPI00003351	Extracellular matrix protein 1 precursor
IPI00003351	Extracellular matrix protein 1 precursor
IPI00003351	Extracellular matrix protein 1 precursor
IPI00003351	Extracellular matrix protein 1 precursor
IPI00003351	Extracellular matrix protein 1 precursor
IPI00003351	Extracellular matrix protein 1 precursor
IPI00003351	Extracellular matrix protein 1 precursor
IPI00003351	Extracellular matrix protein 1 precursor

IPI00003351	Extracellular matrix protein 1 precursor
IPI00003362	Hypothetical protein
IPI00003362	Hypothetical protein
IPI00003362	Hypothetical protein
IPI00003362	Hypothetical protein
IPI00003362	Hypothetical protein
IPI00003362	Hypothetical protein
IPI00003362	Hypothetical protein
IPI00003366	Isoform TrkB of BDNF/NT-3 growth factors receptor precurs
IPI00003366	Isoform TrkB of BDNF/NT-3 growth factors receptor precurs
IPI00003366	Isoform TrkB of BDNF/NT-3 growth factors receptor precurs
IPI00003448	Melanoma-derived growth regulatory protein precursor
IPI00003448	Melanoma-derived growth regulatory protein precursor
IPI00003469	Ig kappa chain V-I region WEA
IPI00003469	Ig kappa chain V-I region WEA
IPI00003590	quiescin Q6 isoform a
IPI00003590	quiescin Q6 isoform a
IPI00003590	quiescin Q6 isoform a
IPI00003590	quiescin Q6 isoform a
IPI00003590	quiescin Q6 isoform a
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IPI00003590	quiescin Q6 isoform a
IPI00003590	quiescin Q6 isoform a
IPI00003590	quiescin Q6 isoform a
IPI00003802	Alpha-mannosidase 2
IPI00003802	Alpha-mannosidase 2
IPI00003802	Alpha-mannosidase 2
IPI00003802	Alpha-mannosidase 2
IPI00003802	Alpha-mannosidase 2
IPI00003802	Alpha-mannosidase 2
IPI00003802	Alpha-mannosidase 2
IPI00003802	Alpha-mannosidase 2
IPI00003802	Alpha-mannosidase 2
IPI00003802	Alpha-mannosidase 2
IPI00003813	Nectin-like protein 2
IPI00003815	Rho GDP-dissociation inhibitor 1
IPI00003865	Isoform 1 of Heat shock cognate 71 kDa protein
IPI00003865	Isoform 1 of Heat shock cognate 71 kDa protein
IPI00003907	Isoform 1 of Protocadherin gamma C5 precursor
IPI00003907	Isoform 1 of Protocadherin gamma C5 precursor
IPI00003907	Isoform 1 of Protocadherin gamma C5 precursor
IPI00003907	Isoform 1 of Protocadherin gamma C5 precursor
IPI00003907	Isoform 1 of Protocadherin gamma C5 precursor
IPI00003919	Glutaminyl-peptide cyclotransferase precursor
IPI00003919	Glutaminyl-peptide cyclotransferase precursor
IPI00003919	Glutaminyl-peptide cyclotransferase precursor
IPI00003919	Glutaminyl-peptide cyclotransferase precursor
IPI00003919	Glutaminyl-peptide cyclotransferase precursor
IPI00003919	Glutaminyl-peptide cyclotransferase precursor
IPI00004114	Ribonuclease K6 precursor
IPI00004433	Contactin-6 precursor
IPI00004433	Contactin-6 precursor
IPI00004433	Contactin-6 precursor

IPI00004433	Contactin-6 precursor
IPI00004433	Contactin-6 precursor
IPI00004433	Contactin-6 precursor
IPI00004433	Contactin-6 precursor
IPI00004440	Receptor-type tyrosine-protein phosphatase-like N precursc
IPI00004440	Receptor-type tyrosine-protein phosphatase-like N precursc
IPI00004440	Receptor-type tyrosine-protein phosphatase-like N precursc
IPI00004440	Receptor-type tyrosine-protein phosphatase-like N precursc
IPI00004503	LAMP1 protein
IPI00004503	LAMP1 protein
IPI00004503	LAMP1 protein
IPI00004503	LAMP1 protein
IPI00004573	Polymeric-immunoglobulin receptor precursor
IPI00004573	Polymeric-immunoglobulin receptor precursor
IPI00004656	Beta-2-microglobulin precursor
IPI00004656	Beta-2-microglobulin precursor
IPI00004656	Beta-2-microglobulin precursor
IPI00004656	Beta-2-microglobulin precursor
IPI00004656	Beta-2-microglobulin precursor
IPI00004656	Beta-2-microglobulin precursor
IPI00004656	Beta-2-microglobulin precursor
IPI00004669	Polypeptide N-acetylgalactosaminyltransferase 2
IPI00004669	Polypeptide N-acetylgalactosaminyltransferase 2
IPI00004946	chemokine (C-X-C motif) ligand 16
IPI00004946	chemokine (C-X-C motif) ligand 16
IPI00004946	chemokine (C-X-C motif) ligand 16
IPI00005038	Ribonuclease UK114
IPI00005038	Ribonuclease UK114
IPI00005038	Ribonuclease UK114
IPI00005126	Ephrin-B2 precursor
IPI00005126	Ephrin-B2 precursor
IPI00005126	Ephrin-B2 precursor
IPI00005142	Isoform 1 of Basic fibroblast growth factor receptor 1 precu
IPI00005142	Isoform 1 of Basic fibroblast growth factor receptor 1 precu
IPI00005292	Testican-1 precursor
IPI00005292	Testican-1 precursor
IPI00005292	Testican-1 precursor
IPI00005292	Testican-1 precursor
IPI00005292	Testican-1 precursor
IPI00005491	Calsyntenin-2 precursor
IPI00005491	Calsyntenin-2 precursor
IPI00005491	Calsyntenin-2 precursor
IPI00005491	Calsyntenin-2 precursor
IPI00005707	Macrophage mannose receptor 2 precursor
IPI00005707	Macrophage mannose receptor 2 precursor
IPI00005707	Macrophage mannose receptor 2 precursor
IPI00005794	Blood plasma glutamate carboxypeptidase precursor
IPI00005794	Blood plasma glutamate carboxypeptidase precursor
IPI00005794	Blood plasma glutamate carboxypeptidase precursor
IPI00005794	Blood plasma glutamate carboxypeptidase precursor
IPI00005794	Blood plasma glutamate carboxypeptidase precursor
IPI00005794	Blood plasma glutamate carboxypeptidase precursor
IPI00005837	Angiopoietin-related protein 1 precursor
IPI00005908	ADAMTS-1 precursor
IPI00005908	ADAMTS-1 precursor
IPI00006009	Isoform 2 of Pleckstrin homology domain-containing family

IPI00006009	Isoform 2 of Pleckstrin homology domain-containing family
IPI00006114	Pigment epithelium-derived factor precursor
IPI00006114	Pigment epithelium-derived factor precursor
IPI00006114	Pigment epithelium-derived factor precursor
IPI00006114	Pigment epithelium-derived factor precursor
IPI00006114	Pigment epithelium-derived factor precursor
IPI00006114	Pigment epithelium-derived factor precursor
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IPI00006114	Pigment epithelium-derived factor precursor
IPI00006114	Pigment epithelium-derived factor precursor
IPI00006114	Pigment epithelium-derived factor precursor
IPI00006128	Testican-2 precursor
IPI00006128	Testican-2 precursor
IPI00006154	Isoform Long of Complement factor H-related protein 2 prec
IPI00006154	Isoform Long of Complement factor H-related protein 2 prec
IPI00006524	KIAA0319
IPI00006601	Secretogranin-1 precursor
IPI00006601	Secretogranin-1 precursor
IPI00006601	Secretogranin-1 precursor
IPI00006601	Secretogranin-1 precursor
IPI00006601	Secretogranin-1 precursor
IPI00006601	Secretogranin-1 precursor
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IPI00006601	Secretogranin-1 precursor
IPI00006601	Secretogranin-1 precursor
IPI00006601	Secretogranin-1 precursor
IPI00006601	Secretogranin-1 precursor
IPI00006608	Isoform APP770 of Amyloid beta A4 protein precursor (Frag
IPI00006608	Isoform APP770 of Amyloid beta A4 protein precursor (Frag
IPI00006608	Isoform APP770 of Amyloid beta A4 protein precursor (Frag
IPI00006608	Isoform APP770 of Amyloid beta A4 protein precursor (Frag

IPI00006608	Isoform APP770 of Amyloid beta A4 protein precursor (Frag
IPI00006608	Isoform APP770 of Amyloid beta A4 protein precursor (Frag
IPI00006608	Isoform APP770 of Amyloid beta A4 protein precursor (Frag
IPI00006608	Isoform APP770 of Amyloid beta A4 protein precursor (Frag
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IPI00006608	Isoform APP770 of Amyloid beta A4 protein precursor (Frag
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IPI00006608	Isoform APP770 of Amyloid beta A4 protein precursor (Frag
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IPI00006608	Isoform APP770 of Amyloid beta A4 protein precursor (Frag
IPI00006608	Isoform APP770 of Amyloid beta A4 protein precursor (Frag
IPI00006608	Isoform APP770 of Amyloid beta A4 protein precursor (Frag
IPI00006608	Isoform APP770 of Amyloid beta A4 protein precursor (Frag
IPI00006644	Isoform 2 of Plexin-B1 precursor
IPI00006644	Isoform 2 of Plexin-B1 precursor
IPI00006662	Apolipoprotein D precursor
IPI00006662	Apolipoprotein D precursor
IPI00006662	Apolipoprotein D precursor
IPI00006662	Apolipoprotein D precursor
IPI00006662	Apolipoprotein D precursor
IPI00006662	Apolipoprotein D precursor
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IPI00006662	Apolipoprotein D precursor
IPI00006662	Apolipoprotein D precursor
IPI00006662	Apolipoprotein D precursor
IPI00006662	Apolipoprotein D precursor
IPI00006967	Protocadherin-9 precursor
IPI00006967	Protocadherin-9 precursor
IPI00006967	Protocadherin-9 precursor
IPI00006967	Protocadherin-9 precursor
IPI00007047	Protein S100-A8
IPI00007047	Protein S100-A8
IPI00007102	CGI-150 protein
IPI00007102	CGI-150 protein
IPI00007102	CGI-150 protein
IPI00007221	Plasma serine protease inhibitor precursor
IPI00007221	Plasma serine protease inhibitor precursor
IPI00007221	Plasma serine protease inhibitor precursor
IPI00007221	Plasma serine protease inhibitor precursor
IPI00007221	Plasma serine protease inhibitor precursor
IPI00007221	Plasma serine protease inhibitor precursor
IPI00007221	Plasma serine protease inhibitor precursor
IPI00007221	Plasma serine protease inhibitor precursor
IPI00007221	Plasma serine protease inhibitor precursor
IPI00007221	Plasma serine protease inhibitor precursor
IPI00007236	Isoform 2 of Neuroligin-1 precursor
IPI00007236	Isoform 2 of Neuroligin-1 precursor

IPI00007236	Isoform 2 of Neuroligin-1 precursor
IPI00007257	calsyntenin 1 isoform 2
IPI00007257	calsyntenin 1 isoform 2
IPI00007257	calsyntenin 1 isoform 2
IPI00007257	calsyntenin 1 isoform 2
IPI00007257	calsyntenin 1 isoform 2
IPI00007257	calsyntenin 1 isoform 2
IPI00007257	calsyntenin 1 isoform 2
IPI00007257	calsyntenin 1 isoform 2
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IPI00007257	calsyntenin 1 isoform 2
IPI00007257	calsyntenin 1 isoform 2
IPI00007257	calsyntenin 1 isoform 2
IPI00007257	calsyntenin 1 isoform 2
IPI00007257	calsyntenin 1 isoform 2
IPI00007632	Glutamate receptor 4 precursor
IPI00007632	Glutamate receptor 4 precursor
IPI00007632	Glutamate receptor 4 precursor
IPI00007778	Di-N-acetylchitobiase precursor
IPI00007778	Di-N-acetylchitobiase precursor
IPI00007778	Di-N-acetylchitobiase precursor
IPI00007778	Di-N-acetylchitobiase precursor
IPI00007797	Fatty acid-binding protein, epidermal
IPI00007797	Fatty acid-binding protein, epidermal
IPI00007797	Fatty acid-binding protein, epidermal
IPI00007798	Thyrotropin-releasing hormone degrading ectoenzyme
IPI00007798	Thyrotropin-releasing hormone degrading ectoenzyme
IPI00007798	Thyrotropin-releasing hormone degrading ectoenzyme
IPI00007798	Thyrotropin-releasing hormone degrading ectoenzyme
IPI00007800	Angiopoietin-related protein 2 precursor
IPI00007800	Angiopoietin-related protein 2 precursor
IPI00007899	Single chain Fv (Fragment)
IPI00007921	Isoform 1 of Neurexin-2-alpha precursor
IPI00007921	Isoform 1 of Neurexin-2-alpha precursor
IPI00007921	Isoform 1 of Neurexin-2-alpha precursor
IPI00007921	Isoform 1 of Neurexin-2-alpha precursor
IPI00007921	Isoform 1 of Neurexin-2-alpha precursor
IPI00007921	Isoform 1 of Neurexin-2-alpha precursor
IPI00007921	Isoform 1 of Neurexin-2-alpha precursor
IPI00007921	Isoform 1 of Neurexin-2-alpha precursor
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IPI00007921	Isoform 1 of Neurexin-2-alpha precursor
IPI00007921	Isoform 1 of Neurexin-2-alpha precursor
IPI00007921	Isoform 1 of Neurexin-2-alpha precursor
IPI00007921	Isoform 1 of Neurexin-2-alpha precursor
IPI00007921	Isoform 1 of Neurexin-2-alpha precursor

IPI00007921	Isoform 1 of Neurexin-2-alpha precursor
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IPI00007921	Isoform 1 of Neurexin-2-alpha precursor
IPI00007921	Isoform 1 of Neurexin-2-alpha precursor
IPI00007921	Isoform 1 of Neurexin-2-alpha precursor
IPI00007921	Isoform 1 of Neurexin-2-alpha precursor
IPI00007921	Isoform 1 of Neurexin-2-alpha precursor
IPI00007921	Isoform 1 of Neurexin-2-alpha precursor
IPI00007921	Isoform 1 of Neurexin-2-alpha precursor
IPI00007921	Isoform 1 of Neurexin-2-alpha precursor
IPI00007921	Isoform 1 of Neurexin-2-alpha precursor
IPI00007921	Isoform 1 of Neurexin-2-alpha precursor
IPI00008087	Follistatin-related protein 5 precursor
IPI00008087	Follistatin-related protein 5 precursor
IPI00008087	Follistatin-related protein 5 precursor
IPI00008087	Follistatin-related protein 5 precursor
IPI00008087	Follistatin-related protein 5 precursor
IPI00008087	Follistatin-related protein 5 precursor
IPI00008087	Follistatin-related protein 5 precursor
IPI00008207	Endoplasmic reticulum mannosyl-oligosaccharide 1,2-alpha
IPI00008207	Endoplasmic reticulum mannosyl-oligosaccharide 1,2-alpha
IPI00008290	Isoform 1 of Ephrin type-A receptor 5 precursor
IPI00008290	Isoform 1 of Ephrin type-A receptor 5 precursor
IPI00008290	Isoform 1 of Ephrin type-A receptor 5 precursor
IPI00008290	Isoform 1 of Ephrin type-A receptor 5 precursor
IPI00008318	Ephrin type-A receptor 4 precursor
IPI00008318	Ephrin type-A receptor 4 precursor
IPI00008318	Ephrin type-A receptor 4 precursor
IPI00008318	Ephrin type-A receptor 4 precursor
IPI00008318	Ephrin type-A receptor 4 precursor
IPI00008318	Ephrin type-A receptor 4 precursor
IPI00008318	Ephrin type-A receptor 4 precursor
IPI00008554	Angiogenin precursor
IPI00008554	Angiogenin precursor
IPI00008558	Plasma kallikrein precursor
IPI00008558	Plasma kallikrein precursor
IPI00008558	Plasma kallikrein precursor
IPI00008586	chondroitin sulfate proteoglycan 5-III
IPI00008787	Alpha-N-acetylglucosaminidase precursor
IPI00008787	Alpha-N-acetylglucosaminidase precursor
IPI00008787	Alpha-N-acetylglucosaminidase precursor
IPI00008787	Alpha-N-acetylglucosaminidase precursor
IPI00008787	Alpha-N-acetylglucosaminidase precursor
IPI00008787	Alpha-N-acetylglucosaminidase precursor
IPI00008787	Alpha-N-acetylglucosaminidase precursor
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IPI00008787	Alpha-N-acetylglucosaminidase precursor
IPI00008787	Alpha-N-acetylglucosaminidase precursor
IPI00008787	Alpha-N-acetylglucosaminidase precursor
IPI00008787	Alpha-N-acetylglucosaminidase precursor
IPI00008787	Alpha-N-acetylglucosaminidase precursor
IPI00008860	Isoform 1 of Complement C1q tumor necrosis factor-relatec

IPI00008860	Isoform 1 of Complement C1q tumor necrosis factor-relatec
IPI00008944	Isoform 1 of Neuroendocrine protein 7B2 precursor
IPI00008944	Isoform 1 of Neuroendocrine protein 7B2 precursor
IPI00008944	Isoform 1 of Neuroendocrine protein 7B2 precursor
IPI00008944	Isoform 1 of Neuroendocrine protein 7B2 precursor
IPI00008994	Isoform 1 of Protein NDRG2
IPI00008994	Isoform 1 of Protein NDRG2
IPI00008994	Isoform 1 of Protein NDRG2
IPI00009028	Tetranectin precursor
IPI00009028	Tetranectin precursor
IPI00009028	Tetranectin precursor
IPI00009028	Tetranectin precursor
IPI00009028	Tetranectin precursor
IPI00009028	Tetranectin precursor
IPI00009028	Tetranectin precursor
IPI00009028	Tetranectin precursor
IPI00009028	Tetranectin precursor
IPI00009028	Tetranectin precursor
IPI00009028	Tetranectin precursor
IPI00009028	Tetranectin precursor
IPI00009028	Tetranectin precursor
IPI00009030	Isoform LAMP-2A of Lysosome-associated membrane glyc
IPI00009030	Isoform LAMP-2A of Lysosome-associated membrane glyc
IPI00009030	Isoform LAMP-2A of Lysosome-associated membrane glyc
IPI00009362	Secretogranin-2 precursor
IPI00009362	Secretogranin-2 precursor
IPI00009362	Secretogranin-2 precursor
IPI00009362	Secretogranin-2 precursor
IPI00009362	Secretogranin-2 precursor
IPI00009362	Secretogranin-2 precursor
IPI00009362	Secretogranin-2 precursor
IPI00009362	Secretogranin-2 precursor
IPI00009362	Secretogranin-2 precursor
IPI00009362	Secretogranin-2 precursor
IPI00009362	Secretogranin-2 precursor
IPI00009619	Isoform 2 of Immunoglobulin superfamily member 4B precu
IPI00009619	Isoform 2 of Immunoglobulin superfamily member 4B precu
IPI00009619	Isoform 2 of Immunoglobulin superfamily member 4B precu
IPI00009619	Isoform 2 of Immunoglobulin superfamily member 4B precu
IPI00009619	Isoform 2 of Immunoglobulin superfamily member 4B precu
IPI00009619	Isoform 2 of Immunoglobulin superfamily member 4B precu
IPI00009619	Isoform 2 of Immunoglobulin superfamily member 4B precu
IPI00009619	Isoform 2 of Immunoglobulin superfamily member 4B precu
IPI00009793	Complement component 1, r subcomponent-like variant (Fr
IPI00009793	Complement component 1, r subcomponent-like variant (Fr
IPI00009794	Calcium binding protein Cab45
IPI00009802	Isoform V0 of Versican core protein precursor
IPI00009802	Isoform V0 of Versican core protein precursor
IPI00009802	Isoform V0 of Versican core protein precursor
IPI00009802	Isoform V0 of Versican core protein precursor
IPI00009802	Isoform V0 of Versican core protein precursor
IPI00009802	Isoform V0 of Versican core protein precursor
IPI00009890	Glia-derived nexin precursor
IPI00009890	Glia-derived nexin precursor
IPI00009890	Glia-derived nexin precursor

IPI00009890	Glia-derived nexin precursor
IPI00009890	Glia-derived nexin precursor
IPI00009890	Glia-derived nexin precursor
IPI00009920	Complement component C6 precursor
IPI00009920	Complement component C6 precursor
IPI00009920	Complement component C6 precursor
IPI00009920	Complement component C6 precursor
IPI00009920	Complement component C6 precursor
IPI00009920	Complement component C6 precursor
IPI00009920	Complement component C6 precursor
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IPI00009920	Complement component C6 precursor
IPI00009920	Complement component C6 precursor
IPI00009920	Complement component C6 precursor
IPI00009920	Complement component C6 precursor
IPI00009920	Complement component C6 precursor
IPI00009950	Vesicular integral-membrane protein VIP36 precursor
IPI00009950	Vesicular integral-membrane protein VIP36 precursor
IPI00009950	Vesicular integral-membrane protein VIP36 precursor
IPI00009950	Vesicular integral-membrane protein VIP36 precursor
IPI00009950	Vesicular integral-membrane protein VIP36 precursor
IPI00009950	Vesicular integral-membrane protein VIP36 precursor
IPI00009950	Vesicular integral-membrane protein VIP36 precursor
IPI00009950	Vesicular integral-membrane protein VIP36 precursor
IPI00009950	Vesicular integral-membrane protein VIP36 precursor
IPI00009997	N-acetyllactosaminide beta-1,3-N-acetylglucosaminyltransfe
IPI00009997	N-acetyllactosaminide beta-1,3-N-acetylglucosaminyltransfe
IPI00009997	N-acetyllactosaminide beta-1,3-N-acetylglucosaminyltransfe
IPI00009997	N-acetyllactosaminide beta-1,3-N-acetylglucosaminyltransfe
IPI00009997	N-acetyllactosaminide beta-1,3-N-acetylglucosaminyltransfe
IPI00009997	N-acetyllactosaminide beta-1,3-N-acetylglucosaminyltransfe
IPI00009997	N-acetyllactosaminide beta-1,3-N-acetylglucosaminyltransfe
IPI00009997	N-acetyllactosaminide beta-1,3-N-acetylglucosaminyltransfe
IPI00009997	N-acetyllactosaminide beta-1,3-N-acetylglucosaminyltransfe
IPI00009997	N-acetyllactosaminide beta-1,3-N-acetylglucosaminyltransfe
IPI00009997	N-acetyllactosaminide beta-1,3-N-acetylglucosaminyltransfe
IPI00009997	N-acetyllactosaminide beta-1,3-N-acetylglucosaminyltransfe
IPI00009997	N-acetyllactosaminide beta-1,3-N-acetylglucosaminyltransfe
IPI00010154	Rab GDP dissociation inhibitor alpha
IPI00010154	Rab GDP dissociation inhibitor alpha
IPI00010154	Rab GDP dissociation inhibitor alpha
IPI00010295	Carboxypeptidase N catalytic chain precursor
IPI00010381	VPS10 domain-containing receptor SorCS3 precursor
IPI00010381	VPS10 domain-containing receptor SorCS3 precursor
IPI00010381	VPS10 domain-containing receptor SorCS3 precursor
IPI00010402	Hypothetical protein
IPI00010471	Plastin-2
IPI00010471	Plastin-2
IPI00010471	Plastin-2

IPI00010471	Plastin-2
IPI00010471	Plastin-2
IPI00010680	Isoform 2 of Fibroblast growth factor receptor 2 precursor
IPI00010680	Isoform 2 of Fibroblast growth factor receptor 2 precursor
IPI00010796	Protein disulfide-isomerase precursor
IPI00010796	Protein disulfide-isomerase precursor
IPI00010849	Beta-casein precursor
IPI00010849	Beta-casein precursor
IPI00010849	Beta-casein precursor
IPI00010849	Beta-casein precursor
IPI00010849	Beta-casein precursor
IPI00010849	Beta-casein precursor
IPI00010896	Chloride intracellular channel protein 1
IPI00010949	Isoform 1 of Sialate O-acetylesterase precursor
IPI00010949	Isoform 1 of Sialate O-acetylesterase precursor
IPI00010949	Isoform 1 of Sialate O-acetylesterase precursor
IPI00011094	Complement C1q tumor necrosis factor-related protein 4 pr
IPI00011094	Complement C1q tumor necrosis factor-related protein 4 pr
IPI00011094	Complement C1q tumor necrosis factor-related protein 4 pr
IPI00011094	Complement C1q tumor necrosis factor-related protein 4 pr
IPI00011094	Complement C1q tumor necrosis factor-related protein 4 pr
IPI00011140	Protein NOV homolog precursor
IPI00011140	Protein NOV homolog precursor
IPI00011140	Protein NOV homolog precursor
IPI00011140	Protein NOV homolog precursor
IPI00011140	Protein NOV homolog precursor
IPI00011218	Macrophage colony-stimulating factor 1 receptor precursor
IPI00011218	Macrophage colony-stimulating factor 1 receptor precursor
IPI00011218	Macrophage colony-stimulating factor 1 receptor precursor
IPI00011218	Macrophage colony-stimulating factor 1 receptor precursor
IPI00011218	Macrophage colony-stimulating factor 1 receptor precursor
IPI00011218	Macrophage colony-stimulating factor 1 receptor precursor
IPI00011229	Cathepsin D precursor
IPI00011229	Cathepsin D precursor
IPI00011229	Cathepsin D precursor
IPI00011229	Cathepsin D precursor
IPI00011229	Cathepsin D precursor
IPI00011229	Cathepsin D precursor
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IPI00011229	Cathepsin D precursor
IPI00011229	Cathepsin D precursor
IPI00011229	Cathepsin D precursor
IPI00011229	Cathepsin D precursor
IPI00011252	Complement component C8 alpha chain precursor
IPI00011252	Complement component C8 alpha chain precursor
IPI00011252	Complement component C8 alpha chain precursor
IPI00011252	Complement component C8 alpha chain precursor
IPI00011261	Complement component C8 gamma chain precursor
IPI00011261	Complement component C8 gamma chain precursor
IPI00011261	Complement component C8 gamma chain precursor
IPI00011261	Complement component C8 gamma chain precursor
IPI00011261	Complement component C8 gamma chain precursor

IPI00011261	Complement component C8 gamma chain precursor
IPI00011261	Complement component C8 gamma chain precursor
IPI00011261	Complement component C8 gamma chain precursor
IPI00011261	Complement component C8 gamma chain precursor
IPI00011264	Complement factor H-related protein 1 precursor
IPI00011264	Complement factor H-related protein 1 precursor
IPI00011264	Complement factor H-related protein 1 precursor
IPI00011302	CD59 glycoprotein precursor
IPI00011302	CD59 glycoprotein precursor
IPI00011302	CD59 glycoprotein precursor
IPI00011605	Cerebellin precursor
IPI00011605	Cerebellin precursor
IPI00011605	Cerebellin precursor
IPI00011605	Cerebellin precursor
IPI00011605	Cerebellin precursor
IPI00011605	Cerebellin precursor
IPI00011642	Isoform 1 of Receptor-type tyrosine-protein phosphatase de
IPI00011642	Isoform 1 of Receptor-type tyrosine-protein phosphatase de
IPI00011642	Isoform 1 of Receptor-type tyrosine-protein phosphatase de
IPI00011642	Isoform 1 of Receptor-type tyrosine-protein phosphatase de
IPI00011642	Isoform 1 of Receptor-type tyrosine-protein phosphatase de
IPI00011642	Isoform 1 of Receptor-type tyrosine-protein phosphatase de
IPI00011642	Isoform 1 of Receptor-type tyrosine-protein phosphatase de
IPI00011642	Isoform 1 of Receptor-type tyrosine-protein phosphatase de
IPI00011642	Isoform 1 of Receptor-type tyrosine-protein phosphatase de
IPI00011642	Isoform 1 of Receptor-type tyrosine-protein phosphatase de
IPI00011651	Receptor-type tyrosine-protein phosphatase gamma precur
IPI00011651	Receptor-type tyrosine-protein phosphatase gamma precur
IPI00011651	Receptor-type tyrosine-protein phosphatase gamma precur
IPI00011651	Receptor-type tyrosine-protein phosphatase gamma precur
IPI00011662	Kunitz-type protease inhibitor 2 precursor
IPI00011732	Isoform 1 of GDNF family receptor alpha-2 precursor
IPI00011732	Isoform 1 of GDNF family receptor alpha-2 precursor
IPI00011732	Isoform 1 of GDNF family receptor alpha-2 precursor
IPI00011732	Isoform 1 of GDNF family receptor alpha-2 precursor
IPI00011994	Ectonucleotide pyrophosphatase/phosphodiesterase 5 prec
IPI00011994	Ectonucleotide pyrophosphatase/phosphodiesterase 5 prec
IPI00012011	Cofilin-1
IPI00012011	Cofilin-1
IPI00012011	Cofilin-1
IPI00012011	Cofilin-1
IPI00012011	Cofilin-1
IPI00012119	Isoform A of Decorin precursor
IPI00012119	Isoform A of Decorin precursor
IPI00012119	Isoform A of Decorin precursor
IPI00012119	Isoform A of Decorin precursor
IPI00012386	Cochlin precursor
IPI00012386	Cochlin precursor
IPI00012440	Plasma alpha-L-fucosidase precursor
IPI00012440	Plasma alpha-L-fucosidase precursor
IPI00012440	Plasma alpha-L-fucosidase precursor
IPI00012440	Plasma alpha-L-fucosidase precursor
IPI00012440	Plasma alpha-L-fucosidase precursor
IPI00012503	Isoform Sap-mu-0 of Proactivator polypeptide precursor
IPI00012503	Isoform Sap-mu-0 of Proactivator polypeptide precursor

IPI00012503	Isoform Sap-mu-0 of Proactivator polypeptide precursor
IPI00012503	Isoform Sap-mu-0 of Proactivator polypeptide precursor
IPI00012503	Isoform Sap-mu-0 of Proactivator polypeptide precursor
IPI00012503	Isoform Sap-mu-0 of Proactivator polypeptide precursor
IPI00012503	Isoform Sap-mu-0 of Proactivator polypeptide precursor
IPI00012503	Isoform Sap-mu-0 of Proactivator polypeptide precursor
IPI00012887	Cathepsin L precursor
IPI00012887	Cathepsin L precursor
IPI00012887	Cathepsin L precursor
IPI00012887	Cathepsin L precursor
IPI00012887	Cathepsin L precursor
IPI00012887	Cathepsin L precursor
IPI00013162	Isoform 1 of OX-2 membrane glycoprotein precursor
IPI00013162	Isoform 1 of OX-2 membrane glycoprotein precursor
IPI00013179	Prostaglandin-H2 D-isomerase precursor
IPI00013179	Prostaglandin-H2 D-isomerase precursor
IPI00013179	Prostaglandin-H2 D-isomerase precursor
IPI00013179	Prostaglandin-H2 D-isomerase precursor
IPI00013179	Prostaglandin-H2 D-isomerase precursor
IPI00013179	Prostaglandin-H2 D-isomerase precursor
IPI00013179	Prostaglandin-H2 D-isomerase precursor
IPI00013179	Prostaglandin-H2 D-isomerase precursor
IPI00013179	Prostaglandin-H2 D-isomerase precursor
IPI00013179	Prostaglandin-H2 D-isomerase precursor
IPI00013179	Prostaglandin-H2 D-isomerase precursor
IPI00013299	Neuroblastoma, suppression of tumorigenicity 1
IPI00013299	Neuroblastoma, suppression of tumorigenicity 1
IPI00013299	Neuroblastoma, suppression of tumorigenicity 1
IPI00013299	Neuroblastoma, suppression of tumorigenicity 1
IPI00013303	Limbic system-associated membrane protein precursor
IPI00013303	Limbic system-associated membrane protein precursor
IPI00013303	Limbic system-associated membrane protein precursor
IPI00013303	Limbic system-associated membrane protein precursor
IPI00013303	Limbic system-associated membrane protein precursor
IPI00013303	Limbic system-associated membrane protein precursor
IPI00013303	Limbic system-associated membrane protein precursor
IPI00013682	Isoform 3 of Ecto-ADP-ribosyltransferase 3 precursor
IPI00013682	Isoform 3 of Ecto-ADP-ribosyltransferase 3 precursor
IPI00013682	Isoform 3 of Ecto-ADP-ribosyltransferase 3 precursor
IPI00013682	Isoform 3 of Ecto-ADP-ribosyltransferase 3 precursor
IPI00013682	Isoform 3 of Ecto-ADP-ribosyltransferase 3 precursor
IPI00013682	Isoform 3 of Ecto-ADP-ribosyltransferase 3 precursor
IPI00013682	Isoform 3 of Ecto-ADP-ribosyltransferase 3 precursor
IPI00013682	Isoform 3 of Ecto-ADP-ribosyltransferase 3 precursor
IPI00013698	Acid ceramidase precursor
IPI00013698	Acid ceramidase precursor
IPI00013933	Isoform DPI of Desmoplakin
IPI00013933	Isoform DPI of Desmoplakin
IPI00013933	Isoform DPI of Desmoplakin
IPI00013933	Isoform DPI of Desmoplakin
IPI00013933	Isoform DPI of Desmoplakin
IPI00013976	Laminin beta-1 chain precursor
IPI00013976	Laminin beta-1 chain precursor
IPI00013976	Laminin beta-1 chain precursor
IPI00013976	Laminin beta-1 chain precursor

IPI00013976	Laminin beta-1 chain precursor
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IPI00013976	Laminin beta-1 chain precursor
IPI00013976	Laminin beta-1 chain precursor
IPI00013976	Laminin beta-1 chain precursor
IPI00014048	Ribonuclease pancreatic precursor
IPI00014048	Ribonuclease pancreatic precursor
IPI00014048	Ribonuclease pancreatic precursor
IPI00014048	Ribonuclease pancreatic precursor
IPI00014048	Ribonuclease pancreatic precursor
IPI00014048	Ribonuclease pancreatic precursor
IPI00014048	Ribonuclease pancreatic precursor
IPI00014439	Dihydropteridine reductase
IPI00014439	Dihydropteridine reductase
IPI00014439	Dihydropteridine reductase
IPI00014439	Dihydropteridine reductase
IPI00014439	Dihydropteridine reductase
IPI00014439	Dihydropteridine reductase
IPI00014572	SPARC precursor
IPI00014572	SPARC precursor
IPI00014572	SPARC precursor
IPI00014572	SPARC precursor
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IPI00014572	SPARC precursor
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IPI00014572	SPARC precursor
IPI00014572	SPARC precursor
IPI00014592	Chondroadherin precursor
IPI00014592	Chondroadherin precursor
IPI00014592	Chondroadherin precursor
IPI00014592	Chondroadherin precursor
IPI00014592	Chondroadherin precursor
IPI00014964	Lymphocyte antigen Ly-6H precursor
IPI00014964	Lymphocyte antigen Ly-6H precursor
IPI00014964	Lymphocyte antigen Ly-6H precursor
IPI00015049	Isoform 2 of Repulsive guidance molecule A precursor
IPI00015049	Isoform 2 of Repulsive guidance molecule A precursor
IPI00015049	Isoform 2 of Repulsive guidance molecule A precursor
IPI00015049	Isoform 2 of Repulsive guidance molecule A precursor
IPI00015049	Isoform 2 of Repulsive guidance molecule A precursor
IPI00015102	CD166 antigen precursor
IPI00015102	CD166 antigen precursor
IPI00015102	CD166 antigen precursor
IPI00015102	CD166 antigen precursor
IPI00015102	CD166 antigen precursor
IPI00015102	CD166 antigen precursor
IPI00015260	Protein kinase C-binding protein NELL2 precursor
IPI00015260	Protein kinase C-binding protein NELL2 precursor
IPI00015260	Protein kinase C-binding protein NELL2 precursor
IPI00015260	Protein kinase C-binding protein NELL2 precursor
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IPI00015260	Protein kinase C-binding protein NELL2 precursor
IPI00015260	Protein kinase C-binding protein NELL2 precursor
IPI00015260	Protein kinase C-binding protein NELL2 precursor
IPI00015260	Protein kinase C-binding protein NELL2 precursor
IPI00015315	Extracellular matrix protein 2 precursor
IPI00015315	Extracellular matrix protein 2 precursor
IPI00015346	Cadherin EGF LAG seven-pass G-type receptor 2 precurso
IPI00015346	Cadherin EGF LAG seven-pass G-type receptor 2 precurso
IPI00015346	Cadherin EGF LAG seven-pass G-type receptor 2 precurso
IPI00015346	Cadherin EGF LAG seven-pass G-type receptor 2 precurso
IPI00015351	AD039
IPI00015351	AD039
IPI00015351	AD039
IPI00015525	Multimerin-2 precursor
IPI00015525	Multimerin-2 precursor
IPI00015525	Multimerin-2 precursor
IPI00015525	Multimerin-2 precursor
IPI00015688	Glypican-1 precursor
IPI00015688	Glypican-1 precursor
IPI00015756	Receptor-type tyrosine-protein phosphatase kappa precurso
IPI00015756	Receptor-type tyrosine-protein phosphatase kappa precurso
IPI00015834	Myo-inositol monophosphatase A3
IPI00015834	Myo-inositol monophosphatase A3
IPI00015834	Myo-inositol monophosphatase A3
IPI00015834	Myo-inositol monophosphatase A3
IPI00015834	Myo-inositol monophosphatase A3
IPI00015834	Myo-inositol monophosphatase A3
IPI00015881	Isoform 1 of Macrophage colony-stimulating factor 1 precur
IPI00015881	Isoform 1 of Macrophage colony-stimulating factor 1 precur
IPI00015911	Dihydrolipoyl dehydrogenase, mitochondrial precursor
IPI00016150	Neuroserpin precursor
IPI00016150	Neuroserpin precursor
IPI00016150	Neuroserpin precursor
IPI00016150	Neuroserpin precursor
IPI00016150	Neuroserpin precursor

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IPI00016150	Neuroserpin precursor
IPI00016150	Neuroserpin precursor
IPI00016150	Neuroserpin precursor
IPI00016150	Neuroserpin precursor
IPI00016334	Isoform 1 of Cell surface glycoprotein MUC18 precursor
IPI00016334	Isoform 1 of Cell surface glycoprotein MUC18 precursor
IPI00016334	Isoform 1 of Cell surface glycoprotein MUC18 precursor
IPI00016334	Isoform 1 of Cell surface glycoprotein MUC18 precursor
IPI00016334	Isoform 1 of Cell surface glycoprotein MUC18 precursor
IPI00016334	Isoform 1 of Cell surface glycoprotein MUC18 precursor
IPI00016334	Isoform 1 of Cell surface glycoprotein MUC18 precursor
IPI00016334	Isoform 1 of Cell surface glycoprotein MUC18 precursor
IPI00016334	Isoform 1 of Cell surface glycoprotein MUC18 precursor
IPI00016334	Isoform 1 of Cell surface glycoprotein MUC18 precursor
IPI00016334	Isoform 1 of Cell surface glycoprotein MUC18 precursor
IPI00016371	Isoform JM-A of Receptor tyrosine-protein kinase erbB-4 pr
IPI00016371	Isoform JM-A of Receptor tyrosine-protein kinase erbB-4 pr
IPI00016422	Netrin receptor DCC precursor
IPI00016422	Netrin receptor DCC precursor
IPI00016422	Netrin receptor DCC precursor
IPI00016422	Netrin receptor DCC precursor
IPI00016422	Netrin receptor DCC precursor
IPI00016422	Netrin receptor DCC precursor
IPI00016422	Netrin receptor DCC precursor
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IPI00016422	Netrin receptor DCC precursor
IPI00016422	Netrin receptor DCC precursor
IPI00016422	Netrin receptor DCC precursor
IPI00016422	Netrin receptor DCC precursor
IPI00016915	Insulin-like growth factor-binding protein 7 precursor
IPI00016915	Insulin-like growth factor-binding protein 7 precursor
IPI00016915	Insulin-like growth factor-binding protein 7 precursor
IPI00016915	Insulin-like growth factor-binding protein 7 precursor
IPI00016915	Insulin-like growth factor-binding protein 7 precursor
IPI00016915	Insulin-like growth factor-binding protein 7 precursor
IPI00016915	Insulin-like growth factor-binding protein 7 precursor
IPI00016915	Insulin-like growth factor-binding protein 7 precursor
IPI00016915	Insulin-like growth factor-binding protein 7 precursor
IPI00016915	Insulin-like growth factor-binding protein 7 precursor
IPI00016915	Insulin-like growth factor-binding protein 7 precursor
IPI00016915	Insulin-like growth factor-binding protein 7 precursor
IPI00016915	Insulin-like growth factor-binding protein 7 precursor
IPI00017257	Cathepsin O precursor
IPI00017257	Cathepsin O precursor
IPI00017257	Cathepsin O precursor

IPI00017257	Cathepsin O precursor
IPI00017601	Ceruloplasmin precursor
IPI00017601	Ceruloplasmin precursor
IPI00017601	Ceruloplasmin precursor
IPI00017601	Ceruloplasmin precursor
IPI00017601	Ceruloplasmin precursor
IPI00017601	Ceruloplasmin precursor
IPI00017601	Ceruloplasmin precursor
IPI00017601	Ceruloplasmin precursor
IPI00017601	Ceruloplasmin precursor
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IPI00017601	Ceruloplasmin precursor
IPI00017696	Complement C1s subcomponent precursor
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IPI00017696	Complement C1s subcomponent precursor
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IPI00017696	Complement C1s subcomponent precursor
IPI00017696	Complement C1s subcomponent precursor
IPI00017696	Complement C1s subcomponent precursor
IPI00017704	Coactosin-like protein
IPI00017704	Coactosin-like protein
IPI00017704	Coactosin-like protein
IPI00017704	Coactosin-like protein
IPI00017704	Coactosin-like protein
IPI00017745	Metalloproteinase inhibitor 4 precursor
IPI00017745	Metalloproteinase inhibitor 4 precursor
IPI00018136	Isoform 1 of Vascular cell adhesion protein 1 precursor
IPI00018136	Isoform 1 of Vascular cell adhesion protein 1 precursor
IPI00018136	Isoform 1 of Vascular cell adhesion protein 1 precursor
IPI00018136	Isoform 1 of Vascular cell adhesion protein 1 precursor
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IPI00018136	Isoform 1 of Vascular cell adhesion protein 1 precursor
IPI00018136	Isoform 1 of Vascular cell adhesion protein 1 precursor
IPI00018206	Aspartate aminotransferase, mitochondrial precursor
IPI00018206	Aspartate aminotransferase, mitochondrial precursor
IPI00018206	Aspartate aminotransferase, mitochondrial precursor
IPI00018206	Aspartate aminotransferase, mitochondrial precursor
IPI00018206	Aspartate aminotransferase, mitochondrial precursor
IPI00018206	Aspartate aminotransferase, mitochondrial precursor
IPI00018206	Aspartate aminotransferase, mitochondrial precursor
IPI00018206	Aspartate aminotransferase, mitochondrial precursor
IPI00018219	Transforming growth factor-beta-induced protein ig-h3 prec
IPI00018219	Transforming growth factor-beta-induced protein ig-h3 prec
IPI00018219	Transforming growth factor-beta-induced protein ig-h3 prec
IPI00018219	Transforming growth factor-beta-induced protein ig-h3 prec
IPI00018219	Transforming growth factor-beta-induced protein ig-h3 prec
IPI00018219	Transforming growth factor-beta-induced protein ig-h3 prec
IPI00018236	Ganglioside GM2 activator precursor
IPI00018236	Ganglioside GM2 activator precursor
IPI00018236	Ganglioside GM2 activator precursor

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IPI00018236	Ganglioside GM2 activator precursor
IPI00018236	Ganglioside GM2 activator precursor
IPI00018236	Ganglioside GM2 activator precursor
IPI00018274	Isoform 1 of Epidermal growth factor receptor precursor
IPI00018274	Isoform 1 of Epidermal growth factor receptor precursor
IPI00018276	Type I transmembrane receptor precursor
IPI00018276	Type I transmembrane receptor precursor
IPI00018276	Type I transmembrane receptor precursor
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IPI00018276	Type I transmembrane receptor precursor
IPI00018276	Type I transmembrane receptor precursor
IPI00018396	Cerebellin-4 precursor
IPI00018769	Thrombospondin-2 precursor
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IPI00018769	Thrombospondin-2 precursor
IPI00019038	Lysozyme C precursor
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IPI00019038	Lysozyme C precursor
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IPI00019038	Lysozyme C precursor
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IPI00019157	Chondroitin sulfate proteoglycan 4 precursor
IPI00019157	Chondroitin sulfate proteoglycan 4 precursor
IPI00019157	Chondroitin sulfate proteoglycan 4 precursor
IPI00019157	Chondroitin sulfate proteoglycan 4 precursor
IPI00019176	Retinoic acid receptor responder protein 2 precursor
IPI00019176	Retinoic acid receptor responder protein 2 precursor
IPI00019176	Retinoic acid receptor responder protein 2 precursor
IPI00019399	Serum amyloid A-4 protein precursor
IPI00019399	Serum amyloid A-4 protein precursor
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IPI00019399	Serum amyloid A-4 protein precursor
IPI00019524	Alpha-lactalbumin precursor
IPI00019524	Alpha-lactalbumin precursor
IPI00019524	Alpha-lactalbumin precursor
IPI00019533	Chitinase 3-like protein 2 precursor
IPI00019533	Chitinase 3-like protein 2 precursor
IPI00019568	Prothrombin precursor (Fragment)
IPI00019568	Prothrombin precursor (Fragment)
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IPI00019576	Coagulation factor X precursor
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IPI00019576	Coagulation factor X precursor
IPI00019579	Complement factor D precursor
IPI00019579	Complement factor D precursor
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IPI00019579	Complement factor D precursor
IPI00019579	Complement factor D precursor
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IPI00019581	Coagulation factor XII precursor
IPI00019581	Coagulation factor XII precursor
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IPI00019591	Isoform 1 of Complement factor B precursor (Fragment)
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IPI00019943	Afamin precursor
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IPI00019943	Afamin precursor
IPI00019954	Cystatin M precursor
IPI00019954	Cystatin M precursor
IPI00020012	Amyloid-like protein 1 precursor
IPI00020012	Amyloid-like protein 1 precursor
IPI00020012	Amyloid-like protein 1 precursor
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IPI00020012	Amyloid-like protein 1 precursor
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IPI00020012	Amyloid-like protein 1 precursor
IPI00020012	Amyloid-like protein 1 precursor
IPI00020012	Amyloid-like protein 1 precursor
IPI00020091	Alpha-1-acid glycoprotein 2 precursor
IPI00020091	Alpha-1-acid glycoprotein 2 precursor
IPI00020091	Alpha-1-acid glycoprotein 2 precursor
IPI00020091	Alpha-1-acid glycoprotein 2 precursor
IPI00020091	Alpha-1-acid glycoprotein 2 precursor
IPI00020091	Alpha-1-acid glycoprotein 2 precursor
IPI00020091	Alpha-1-acid glycoprotein 2 precursor
IPI00020430	Vacuolar ATP synthase subunit S1 precursor
IPI00020430	Vacuolar ATP synthase subunit S1 precursor
IPI00020430	Vacuolar ATP synthase subunit S1 precursor
IPI00020430	Vacuolar ATP synthase subunit S1 precursor
IPI00020430	Vacuolar ATP synthase subunit S1 precursor
IPI00020430	Vacuolar ATP synthase subunit S1 precursor
IPI00020557	Low-density lipoprotein receptor-related protein 1 precursor
IPI00020557	Low-density lipoprotein receptor-related protein 1 precursor

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IPI00020599	Calreticulin precursor
IPI00020599	Calreticulin precursor
IPI00020599	Calreticulin precursor
IPI00020599	Calreticulin precursor
IPI00020665	Latent TGF-beta binding protein-4
IPI00020977	Isoform 1 of Connective tissue growth factor precursor
IPI00020977	Isoform 1 of Connective tissue growth factor precursor
IPI00020977	Isoform 1 of Connective tissue growth factor precursor
IPI00020977	Isoform 1 of Connective tissue growth factor precursor
IPI00020986	Lumican precursor
IPI00020986	Lumican precursor
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IPI00020986	Lumican precursor
IPI00020986	Lumican precursor
IPI00020986	Lumican precursor
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IPI00020987	Prolargin precursor
IPI00020987	Prolargin precursor

IPI00020987	Prolargin precursor
IPI00020990	Osteomodulin precursor
IPI00020990	Osteomodulin precursor
IPI00020996	Insulin-like growth factor-binding protein complex acid labile
IPI00020996	Insulin-like growth factor-binding protein complex acid labile
IPI00020996	Insulin-like growth factor-binding protein complex acid labile
IPI00020996	Insulin-like growth factor-binding protein complex acid labile
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IPI00020996	Insulin-like growth factor-binding protein complex acid labile
IPI00020996	Insulin-like growth factor-binding protein complex acid labile
IPI00020996	Insulin-like growth factor-binding protein complex acid labile
IPI00021000	Isoform A of Osteopontin precursor
IPI00021000	Isoform A of Osteopontin precursor
IPI00021000	Isoform A of Osteopontin precursor
IPI00021000	Isoform A of Osteopontin precursor
IPI00021000	Isoform A of Osteopontin precursor
IPI00021000	Isoform A of Osteopontin precursor
IPI00021000	Isoform A of Osteopontin precursor
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IPI00021000	Isoform A of Osteopontin precursor
IPI00021000	Isoform A of Osteopontin precursor
IPI00021000	Isoform A of Osteopontin precursor
IPI00021033	Collagen alpha-1(III) chain precursor
IPI00021033	Collagen alpha-1(III) chain precursor
IPI00021263	14-3-3 protein zeta/delta
IPI00021263	14-3-3 protein zeta/delta
IPI00021263	14-3-3 protein zeta/delta
IPI00021263	14-3-3 protein zeta/delta
IPI00021263	14-3-3 protein zeta/delta
IPI00021439	Actin, cytoplasmic 1
IPI00021439	Actin, cytoplasmic 1
IPI00021439	Actin, cytoplasmic 1
IPI00021439	Actin, cytoplasmic 1
IPI00021439	Actin, cytoplasmic 1
IPI00021439	Actin, cytoplasmic 1
IPI00021439	Actin, cytoplasmic 1
IPI00021439	Actin, cytoplasmic 1
IPI00021439	Actin, cytoplasmic 1
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IPI00021439	Actin, cytoplasmic 1
IPI00021439	Actin, cytoplasmic 1
IPI00021439	Actin, cytoplasmic 1
IPI00021439	Actin, cytoplasmic 1
IPI00021485	Leucine-rich repeats neuronal protein 1 precursor
IPI00021841	Apolipoprotein A-I precursor
IPI00021841	Apolipoprotein A-I precursor
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IPI00021854	Apolipoprotein A-II precursor
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IPI00021855	Apolipoprotein C-I precursor
IPI00021855	Apolipoprotein C-I precursor
IPI00021855	Apolipoprotein C-I precursor
IPI00021855	Apolipoprotein C-I precursor
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IPI00021856	Apolipoprotein C-II precursor
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IPI00021900	Tumor necrosis factor ligand superfamily member 12
IPI00021900	Tumor necrosis factor ligand superfamily member 12
IPI00021923	Protein FAM3C precursor
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IPI00022284	Major prion protein precursor
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IPI00022284	Major prion protein precursor
IPI00022296	Mast/stem cell growth factor receptor precursor
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IPI00022314	Superoxide dismutase [Mn], mitochondrial precursor
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IPI00022331	Phosphatidylcholine-sterol acyltransferase precursor
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IPI00022333	Brain-specific angiogenesis inhibitor 1 precursor
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IPI00022337	Interphotoreceptor retinoid-binding protein precursor
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IPI00022371	Histidine-rich glycoprotein precursor
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IPI00022389	Isoform 1 of C-reactive protein precursor
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IPI00022392	Complement C1q subcomponent subunit A precursor
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IPI00022608	Sortilin-related receptor precursor

IPI00022733	45 kDa protein
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IPI00022792	Microfibril-associated glycoprotein 4 precursor
IPI00022792	Microfibril-associated glycoprotein 4 precursor
IPI00022792	Microfibril-associated glycoprotein 4 precursor
IPI00022822	Isoform Long of Collagen alpha-1(XVIII) chain precursor
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IPI00022822	Isoform Long of Collagen alpha-1(XVIII) chain precursor
IPI00022822	Isoform Long of Collagen alpha-1(XVIII) chain precursor
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IPI00022892	Thy-1 membrane glycoprotein precursor
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IPI00022895	Alpha-1B-glycoprotein precursor
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IPI00022937	Coagulation factor V
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IPI00022937	Coagulation factor V
IPI00022937	Coagulation factor V

IPI00022937	Coagulation factor V
IPI00022937	Coagulation factor V
IPI00022937	Coagulation factor V
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IPI00022937	Coagulation factor V
IPI00022937	Coagulation factor V
IPI00022937	Coagulation factor V
IPI00022937	Coagulation factor V
IPI00022977	Creatine kinase B-type
IPI00022977	Creatine kinase B-type
IPI00022977	Creatine kinase B-type
IPI00022977	Creatine kinase B-type
IPI00022977	Creatine kinase B-type
IPI00023014	Von Willebrand factor precursor
IPI00023014	Von Willebrand factor precursor
IPI00023014	Von Willebrand factor precursor
IPI00023014	Von Willebrand factor precursor
IPI00023014	Von Willebrand factor precursor
IPI00023014	Von Willebrand factor precursor
IPI00023019	Isoform 1 of Sex hormone-binding globulin precursor
IPI00023019	Isoform 1 of Sex hormone-binding globulin precursor
IPI00023019	Isoform 1 of Sex hormone-binding globulin precursor
IPI00023019	Isoform 1 of Sex hormone-binding globulin precursor
IPI00023019	Isoform 1 of Sex hormone-binding globulin precursor
IPI00023019	Isoform 1 of Sex hormone-binding globulin precursor
IPI00023019	Isoform 1 of Sex hormone-binding globulin precursor
IPI00023648	immunoglobulin superfamily containing leucine-rich repeat
IPI00023648	immunoglobulin superfamily containing leucine-rich repeat
IPI00023648	immunoglobulin superfamily containing leucine-rich repeat
IPI00023648	immunoglobulin superfamily containing leucine-rich repeat
IPI00023648	immunoglobulin superfamily containing leucine-rich repeat
IPI00023648	immunoglobulin superfamily containing leucine-rich repeat
IPI00023673	Galectin-3-binding protein precursor
IPI00023673	Galectin-3-binding protein precursor
IPI00023673	Galectin-3-binding protein precursor
IPI00023673	Galectin-3-binding protein precursor
IPI00023673	Galectin-3-binding protein precursor
IPI00023673	Galectin-3-binding protein precursor
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IPI00023673	Galectin-3-binding protein precursor
IPI00023673	Galectin-3-binding protein precursor
IPI00023673	Galectin-3-binding protein precursor
IPI00023728	Gamma-glutamyl hydrolase precursor
IPI00023728	Gamma-glutamyl hydrolase precursor
IPI00023728	Gamma-glutamyl hydrolase precursor
IPI00023728	Gamma-glutamyl hydrolase precursor
IPI00023728	Gamma-glutamyl hydrolase precursor
IPI00023728	Gamma-glutamyl hydrolase precursor

IPI00023728	Gamma-glutamyl hydrolase precursor
IPI00023751	Growth/differentiation factor 8 precursor
IPI00023751	Growth/differentiation factor 8 precursor
IPI00023751	Growth/differentiation factor 8 precursor
IPI00023751	Growth/differentiation factor 8 precursor
IPI00023814	Isoform 1 of Neogenin precursor
IPI00023814	Isoform 1 of Neogenin precursor
IPI00023814	Isoform 1 of Neogenin precursor
IPI00023814	Isoform 1 of Neogenin precursor
IPI00023814	Isoform 1 of Neogenin precursor
IPI00023814	Isoform 1 of Neogenin precursor
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IPI00023814	Isoform 1 of Neogenin precursor
IPI00023814	Isoform 1 of Neogenin precursor
IPI00023814	Isoform 1 of Neogenin precursor
IPI00023824	Fibulin-2 precursor
IPI00023824	Fibulin-2 precursor
IPI00023824	Fibulin-2 precursor
IPI00023824	Fibulin-2 precursor
IPI00023824	Fibulin-2 precursor
IPI00023824	Fibulin-2 precursor
IPI00023824	Fibulin-2 precursor
IPI00023824	Fibulin-2 precursor
IPI00023824	Fibulin-2 precursor
IPI00023824	Fibulin-2 precursor
IPI00023824	Fibulin-2 precursor
IPI00023845	Kallikrein-6 precursor
IPI00023845	Kallikrein-6 precursor
IPI00023845	Kallikrein-6 precursor
IPI00023845	Kallikrein-6 precursor
IPI00023845	Kallikrein-6 precursor
IPI00023845	Kallikrein-6 precursor
IPI00023845	Kallikrein-6 precursor
IPI00023845	Kallikrein-6 precursor
IPI00023845	Kallikrein-6 precursor
IPI00023845	Kallikrein-6 precursor
IPI00023845	Kallikrein-6 precursor
IPI00024034	Cadherin-4 precursor
IPI00024034	Cadherin-4 precursor
IPI00024034	Cadherin-4 precursor
IPI00024034	Cadherin-4 precursor
IPI00024035	Isoform 1 of Cadherin-6 precursor
IPI00024035	Isoform 1 of Cadherin-6 precursor
IPI00024035	Isoform 1 of Cadherin-6 precursor
IPI00024046	Cadherin-13 precursor
IPI00024046	Cadherin-13 precursor
IPI00024046	Cadherin-13 precursor

IPI00024046	Cadherin-13 precursor
IPI00024046	Cadherin-13 precursor
IPI00024046	Cadherin-13 precursor
IPI00024046	Cadherin-13 precursor
IPI00024046	Cadherin-13 precursor
IPI00024048	Muscle-cadherin precursor
IPI00024048	Muscle-cadherin precursor
IPI00024048	Muscle-cadherin precursor
IPI00024048	Muscle-cadherin precursor
IPI00024048	Muscle-cadherin precursor
IPI00024105	Complement C1q tumor necrosis factor-related protein 5 pr
IPI00024105	Complement C1q tumor necrosis factor-related protein 5 pr
IPI00024105	Complement C1q tumor necrosis factor-related protein 5 pr
IPI00024129	Peptidyl-prolyl cis-trans isomerase C
IPI00024129	Peptidyl-prolyl cis-trans isomerase C
IPI00024129	Peptidyl-prolyl cis-trans isomerase C
IPI00024138	Ig kappa chain V-III region VH precursor (Fragment)
IPI00024138	Ig kappa chain V-III region VH precursor (Fragment)
IPI00024248	Sodium/iodide cotransporter
IPI00024248	Sodium/iodide cotransporter
IPI00024273	Isoform Long of Very low-density lipoprotein receptor precu
IPI00024284	Basement membrane-specific heparan sulfate proteoglycar
IPI00024284	Basement membrane-specific heparan sulfate proteoglycar
IPI00024284	Basement membrane-specific heparan sulfate proteoglycar
IPI00024284	Basement membrane-specific heparan sulfate proteoglycar
IPI00024284	Basement membrane-specific heparan sulfate proteoglycar
IPI00024284	Basement membrane-specific heparan sulfate proteoglycar
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IPI00024284	Basement membrane-specific heparan sulfate proteoglycar
IPI00024284	Basement membrane-specific heparan sulfate proteoglycar
IPI00024284	Basement membrane-specific heparan sulfate proteoglycar
IPI00024284	Basement membrane-specific heparan sulfate proteoglycar
IPI00024284	Basement membrane-specific heparan sulfate proteoglycar
IPI00024289	KIAA0387 protein (Fragment)
IPI00024289	KIAA0387 protein (Fragment)
IPI00024570	Semaphorin sem2
IPI00024570	Semaphorin sem2
IPI00024570	Semaphorin sem2
IPI00024601	Carbonic anhydrase-related protein 10
IPI00024601	Carbonic anhydrase-related protein 10
IPI00024621	Isoform 1 of Olfactomedin-like protein 3 precursor
IPI00024621	Isoform 1 of Olfactomedin-like protein 3 precursor
IPI00024621	Isoform 1 of Olfactomedin-like protein 3 precursor
IPI00024621	Isoform 1 of Olfactomedin-like protein 3 precursor
IPI00024621	Isoform 1 of Olfactomedin-like protein 3 precursor
IPI00024825	Isoform A of Proteoglycan-4 precursor
IPI00024825	Isoform A of Proteoglycan-4 precursor
IPI00024825	Isoform A of Proteoglycan-4 precursor
IPI00024825	Isoform A of Proteoglycan-4 precursor
IPI00024825	Isoform A of Proteoglycan-4 precursor
IPI00024825	Isoform A of Proteoglycan-4 precursor
IPI00024825	Isoform A of Proteoglycan-4 precursor
IPI00024825	Isoform A of Proteoglycan-4 precursor
IPI00024825	Isoform A of Proteoglycan-4 precursor
IPI00024887	Bone morphogenetic protein 6 precursor
IPI00024966	Contactin-2 precursor
IPI00024966	Contactin-2 precursor
IPI00024966	Contactin-2 precursor
IPI00024966	Contactin-2 precursor
IPI00024966	Contactin-2 precursor
IPI00024966	Contactin-2 precursor
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IPI00024966	Contactin-2 precursor
IPI00024966	Contactin-2 precursor

IPI00024966	Contactin-2 precursor
IPI00024966	Contactin-2 precursor
IPI00024989	Protein-L-isoaspartate (D-aspartate) O-methyltransferase
IPI00024989	Protein-L-isoaspartate (D-aspartate) O-methyltransferase
IPI00025257	Semaphorin-7A precursor
IPI00025257	Semaphorin-7A precursor
IPI00025257	Semaphorin-7A precursor
IPI00025257	Semaphorin-7A precursor
IPI00025257	Semaphorin-7A precursor
IPI00025257	Semaphorin-7A precursor
IPI00025257	Semaphorin-7A precursor
IPI00025257	Semaphorin-7A precursor
IPI00025257	Semaphorin-7A precursor
IPI00025257	Semaphorin-7A precursor
IPI00025257	Semaphorin-7A precursor
IPI00025257	Semaphorin-7A precursor
IPI00025257	Semaphorin-7A precursor
IPI00025276	Isoform XB of Tenascin-X precursor
IPI00025276	Isoform XB of Tenascin-X precursor
IPI00025276	Isoform XB of Tenascin-X precursor
IPI00025276	Isoform XB of Tenascin-X precursor
IPI00025276	Isoform XB of Tenascin-X precursor
IPI00025276	Isoform XB of Tenascin-X precursor
IPI00025276	Isoform XB of Tenascin-X precursor
IPI00025276	Isoform XB of Tenascin-X precursor
IPI00025276	Isoform XB of Tenascin-X precursor
IPI00025276	Isoform XB of Tenascin-X precursor
IPI00025276	Isoform XB of Tenascin-X precursor
IPI00025276	Isoform XB of Tenascin-X precursor
IPI00025276	Isoform XB of Tenascin-X precursor
IPI00025276	Isoform XB of Tenascin-X precursor
IPI00025276	Isoform XB of Tenascin-X precursor
IPI00025276	Isoform XB of Tenascin-X precursor
IPI00025276	Isoform XB of Tenascin-X precursor
IPI00025276	Isoform XB of Tenascin-X precursor
IPI00025276	Isoform XB of Tenascin-X precursor
IPI00025318	SH3 domain-binding glutamic acid-rich-like protein
IPI00025318	SH3 domain-binding glutamic acid-rich-like protein
IPI00025318	SH3 domain-binding glutamic acid-rich-like protein
IPI00025318	SH3 domain-binding glutamic acid-rich-like protein
IPI00025318	SH3 domain-binding glutamic acid-rich-like protein
IPI00025426	Pregnancy zone protein precursor
IPI00025426	Pregnancy zone protein precursor
IPI00025426	Pregnancy zone protein precursor
IPI00025426	Pregnancy zone protein precursor
IPI00025426	Pregnancy zone protein precursor
IPI00025465	Mimecan precursor
IPI00025465	Mimecan precursor
IPI00025465	Mimecan precursor
IPI00025465	Mimecan precursor
IPI00025465	Mimecan precursor
IPI00025465	Mimecan precursor
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IPI00025465	Mimecan precursor
IPI00025465	Mimecan precursor
IPI00025465	Mimecan precursor
IPI00025465	Mimecan precursor
IPI00025465	Mimecan precursor
IPI00025686	Latent transforming growth factor-beta binding protein 4L
IPI00025686	Latent transforming growth factor-beta binding protein 4L
IPI00025686	Latent transforming growth factor-beta binding protein 4L
IPI00025686	Latent transforming growth factor-beta binding protein 4L
IPI00025686	Latent transforming growth factor-beta binding protein 4L
IPI00025686	Latent transforming growth factor-beta binding protein 4L
IPI00025686	Latent transforming growth factor-beta binding protein 4L
IPI00025686	Latent transforming growth factor-beta binding protein 4L
IPI00025812	Carbonic anhydrase-related protein 2 precursor
IPI00025812	Carbonic anhydrase-related protein 2 precursor
IPI00025840	Isoform 1 of Ephrin-A1 precursor
IPI00025846	Isoform 2A of Desmocollin-2 precursor
IPI00025846	Isoform 2A of Desmocollin-2 precursor
IPI00025861	Epithelial-cadherin precursor
IPI00025861	Epithelial-cadherin precursor
IPI00025864	Cholinesterase precursor
IPI00025864	Cholinesterase precursor
IPI00026104	Isoform Long of Iduronate 2-sulfatase precursor
IPI00026104	Isoform Long of Iduronate 2-sulfatase precursor
IPI00026104	Isoform Long of Iduronate 2-sulfatase precursor
IPI00026104	Isoform Long of Iduronate 2-sulfatase precursor
IPI00026104	Isoform Long of Iduronate 2-sulfatase precursor
IPI00026104	Isoform Long of Iduronate 2-sulfatase precursor
IPI00026104	Isoform Long of Iduronate 2-sulfatase precursor
IPI00026104	Isoform Long of Iduronate 2-sulfatase precursor
IPI00026125	Deoxyribonuclease I-like 1 precursor
IPI00026154	Glucosidase 2 subunit beta precursor
IPI00026154	Glucosidase 2 subunit beta precursor
IPI00026154	Glucosidase 2 subunit beta precursor
IPI00026199	Glutathione peroxidase 3 precursor
IPI00026199	Glutathione peroxidase 3 precursor
IPI00026199	Glutathione peroxidase 3 precursor
IPI00026199	Glutathione peroxidase 3 precursor
IPI00026199	Glutathione peroxidase 3 precursor
IPI00026199	Glutathione peroxidase 3 precursor
IPI00026199	Glutathione peroxidase 3 precursor
IPI00026216	Puromycin-sensitive aminopeptidase
IPI00026216	Puromycin-sensitive aminopeptidase
IPI00026216	Puromycin-sensitive aminopeptidase
IPI00026216	Puromycin-sensitive aminopeptidase
IPI00026216	Puromycin-sensitive aminopeptidase
IPI00026216	Puromycin-sensitive aminopeptidase
IPI00026216	Puromycin-sensitive aminopeptidase
IPI00026259	N(4)-(beta-N-acetylglucosaminyl)-L-asparaginase precurso
IPI00026259	N(4)-(beta-N-acetylglucosaminyl)-L-asparaginase precurso
IPI00026259	N(4)-(beta-N-acetylglucosaminyl)-L-asparaginase precurso
IPI00026259	N(4)-(beta-N-acetylglucosaminyl)-L-asparaginase precurso
IPI00026259	N(4)-(beta-N-acetylglucosaminyl)-L-asparaginase precurso
IPI00026260	Nucleoside diphosphate kinase B
IPI00026314	Isoform 1 of Gelsolin precursor

IPI00026314	Isoform 1 of Gelsolin precursor
IPI00026314	Isoform 1 of Gelsolin precursor
IPI00026314	Isoform 1 of Gelsolin precursor
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IPI00026314	Isoform 1 of Gelsolin precursor
IPI00026314	Isoform 1 of Gelsolin precursor
IPI00026314	Isoform 1 of Gelsolin precursor
IPI00026358	Gamma-aminobutyric acid receptor-associated protein-like
IPI00026800	Scrapie-responsive protein 1 precursor
IPI00026800	Scrapie-responsive protein 1 precursor
IPI00026800	Scrapie-responsive protein 1 precursor
IPI00026800	Scrapie-responsive protein 1 precursor
IPI00026944	Isoform 1 of Nidogen-1 precursor
IPI00026944	Isoform 1 of Nidogen-1 precursor
IPI00026944	Isoform 1 of Nidogen-1 precursor
IPI00026944	Isoform 1 of Nidogen-1 precursor

IPI00026944	Isoform 1 of Nidogen-1 precursor
IPI00026944	Isoform 1 of Nidogen-1 precursor
IPI00026944	Isoform 1 of Nidogen-1 precursor
IPI00026944	Isoform 1 of Nidogen-1 precursor
IPI00026944	Isoform 1 of Nidogen-1 precursor
IPI00026944	Isoform 1 of Nidogen-1 precursor
IPI00026944	Isoform 1 of Nidogen-1 precursor
IPI00026944	Isoform 1 of Nidogen-1 precursor
IPI00026946	Neuronal pentraxin-2 precursor
IPI00026946	Neuronal pentraxin-2 precursor
IPI00026946	Neuronal pentraxin-2 precursor
IPI00026946	Neuronal pentraxin-2 precursor
IPI00026991	Polypeptide N-acetylgalactosaminyltransferase 6
IPI00026991	Polypeptide N-acetylgalactosaminyltransferase 6
IPI00026991	Polypeptide N-acetylgalactosaminyltransferase 6
IPI00027038	Isoform 1 of V-set and immunoglobulin domain-containing p
IPI00027038	Isoform 1 of V-set and immunoglobulin domain-containing p
IPI00027038	Isoform 1 of V-set and immunoglobulin domain-containing p
IPI00027087	Isoform 1 of Neural cell adhesion molecule L1 precursor
IPI00027087	Isoform 1 of Neural cell adhesion molecule L1 precursor
IPI00027087	Isoform 1 of Neural cell adhesion molecule L1 precursor
IPI00027087	Isoform 1 of Neural cell adhesion molecule L1 precursor
IPI00027087	Isoform 1 of Neural cell adhesion molecule L1 precursor
IPI00027087	Isoform 1 of Neural cell adhesion molecule L1 precursor
IPI00027087	Isoform 1 of Neural cell adhesion molecule L1 precursor
IPI00027087	Isoform 1 of Neural cell adhesion molecule L1 precursor
IPI00027166	Metalloproteinase inhibitor 2 precursor
IPI00027166	Metalloproteinase inhibitor 2 precursor
IPI00027166	Metalloproteinase inhibitor 2 precursor
IPI00027166	Metalloproteinase inhibitor 2 precursor
IPI00027166	Metalloproteinase inhibitor 2 precursor
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IPI00027166	Metalloproteinase inhibitor 2 precursor
IPI00027166	Metalloproteinase inhibitor 2 precursor
IPI00027166	Metalloproteinase inhibitor 2 precursor
IPI00027192	Procollagen-lysine,2-oxoglutarate 5-dioxygenase 1 precurso
IPI00027192	Procollagen-lysine,2-oxoglutarate 5-dioxygenase 1 precurso
IPI00027192	Procollagen-lysine,2-oxoglutarate 5-dioxygenase 1 precurso
IPI00027192	Procollagen-lysine,2-oxoglutarate 5-dioxygenase 1 precurso
IPI00027230	Endoplasmin precursor
IPI00027230	Endoplasmin precursor
IPI00027230	Endoplasmin precursor
IPI00027230	Endoplasmin precursor
IPI00027230	Endoplasmin precursor
IPI00027230	Endoplasmin precursor
IPI00027235	Isoform 1 of Attractin precursor
IPI00027235	Isoform 1 of Attractin precursor
IPI00027235	Isoform 1 of Attractin precursor
IPI00027235	Isoform 1 of Attractin precursor
IPI00027235	Isoform 1 of Attractin precursor
IPI00027235	Isoform 1 of Attractin precursor

IPI00027235	Isoform 1 of Attractin precursor
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IPI00027235	Isoform 1 of Attractin precursor
IPI00027235	Isoform 1 of Attractin precursor
IPI00027235	Isoform 1 of Attractin precursor
IPI00027310	Isoform 1 of Multiple epidermal growth factor-like domains 8
IPI00027310	Isoform 1 of Multiple epidermal growth factor-like domains 8
IPI00027310	Isoform 1 of Multiple epidermal growth factor-like domains 8
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IPI00027310	Isoform 1 of Multiple epidermal growth factor-like domains 8
IPI00027310	Isoform 1 of Multiple epidermal growth factor-like domains 8
IPI00027350	Peroxiredoxin-2
IPI00027350	Peroxiredoxin-2
IPI00027350	Peroxiredoxin-2
IPI00027350	Peroxiredoxin-2
IPI00027350	Peroxiredoxin-2
IPI00027444	Leukocyte elastase inhibitor
IPI00027444	Leukocyte elastase inhibitor
IPI00027444	Leukocyte elastase inhibitor
IPI00027444	Leukocyte elastase inhibitor
IPI00027466	Carbonic anhydrase 4 precursor
IPI00027466	Carbonic anhydrase 4 precursor
IPI00027466	Carbonic anhydrase 4 precursor
IPI00027466	Carbonic anhydrase 4 precursor
IPI00027466	Carbonic anhydrase 4 precursor
IPI00027466	Carbonic anhydrase 4 precursor
IPI00027482	Corticosteroid-binding globulin precursor
IPI00027482	Corticosteroid-binding globulin precursor

IPI00027482	Corticosteroid-binding globulin precursor
IPI00027482	Corticosteroid-binding globulin precursor
IPI00027482	Corticosteroid-binding globulin precursor
IPI00027482	Corticosteroid-binding globulin precursor
IPI00027482	Corticosteroid-binding globulin precursor
IPI00027493	4F2 cell-surface antigen heavy chain
IPI00027493	4F2 cell-surface antigen heavy chain
IPI00027547	Dermcidin precursor
IPI00027547	Dermcidin precursor
IPI00027703	Isoform Long of Alpha-mannosidase IIx
IPI00027703	Isoform Long of Alpha-mannosidase IIx
IPI00027703	Isoform Long of Alpha-mannosidase IIx
IPI00027703	Isoform Long of Alpha-mannosidase IIx
IPI00027703	Isoform Long of Alpha-mannosidase IIx
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IPI00027703	Isoform Long of Alpha-mannosidase IIx
IPI00027703	Isoform Long of Alpha-mannosidase IIx
IPI00027703	Isoform Long of Alpha-mannosidase IIx
IPI00027703	Isoform Long of Alpha-mannosidase IIx
IPI00027780	72 kDa type IV collagenase precursor
IPI00027780	72 kDa type IV collagenase precursor
IPI00027780	72 kDa type IV collagenase precursor
IPI00027780	72 kDa type IV collagenase precursor
IPI00027780	72 kDa type IV collagenase precursor
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IPI00027780	72 kDa type IV collagenase precursor
IPI00027780	72 kDa type IV collagenase precursor
IPI00027780	72 kDa type IV collagenase precursor
IPI00027780	72 kDa type IV collagenase precursor
IPI00027827	Extracellular superoxide dismutase [Cu-Zn] precursor
IPI00027827	Extracellular superoxide dismutase [Cu-Zn] precursor
IPI00027827	Extracellular superoxide dismutase [Cu-Zn] precursor
IPI00027827	Extracellular superoxide dismutase [Cu-Zn] precursor
IPI00027827	Extracellular superoxide dismutase [Cu-Zn] precursor
IPI00027827	Extracellular superoxide dismutase [Cu-Zn] precursor
IPI00027848	Macrophage mannose receptor 1 precursor
IPI00027848	Macrophage mannose receptor 1 precursor
IPI00027848	Macrophage mannose receptor 1 precursor
IPI00027848	Macrophage mannose receptor 1 precursor
IPI00027848	Macrophage mannose receptor 1 precursor
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IPI00027848	Macrophage mannose receptor 1 precursor
IPI00027848	Macrophage mannose receptor 1 precursor
IPI00027848	Macrophage mannose receptor 1 precursor
IPI00027851	Beta-hexosaminidase alpha chain precursor
IPI00027851	Beta-hexosaminidase alpha chain precursor
IPI00027851	Beta-hexosaminidase alpha chain precursor
IPI00027851	Beta-hexosaminidase alpha chain precursor
IPI00027851	Beta-hexosaminidase alpha chain precursor
IPI00027851	Beta-hexosaminidase alpha chain precursor
IPI00028030	Cartilage oligomeric matrix protein precursor
IPI00028413	Inter-alpha-trypsin inhibitor heavy chain H3 precursor
IPI00028413	Inter-alpha-trypsin inhibitor heavy chain H3 precursor

IPI00028413	Inter-alpha-trypsin inhibitor heavy chain H3 precursor
IPI00028413	Inter-alpha-trypsin inhibitor heavy chain H3 precursor
IPI00028413	Inter-alpha-trypsin inhibitor heavy chain H3 precursor
IPI00028414	glia maturation factor, gamma
IPI00028448	Brain-specific angiogenesis inhibitor 3 precursor
IPI00028448	Brain-specific angiogenesis inhibitor 3 precursor
IPI00028714	Matrix Gla-protein precursor
IPI00028908	Nidogen-2 precursor
IPI00028908	Nidogen-2 precursor
IPI00028908	Nidogen-2 precursor
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IPI00028908	Nidogen-2 precursor
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IPI00028908	Nidogen-2 precursor
IPI00028911	Dystroglycan precursor
IPI00028911	Dystroglycan precursor
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IPI00028911	Dystroglycan precursor
IPI00028911	Dystroglycan precursor
IPI00028931	Desmoglein 2
IPI00028931	Desmoglein 2
IPI00028931	Desmoglein 2
IPI00029046	Protein KIAA0152 precursor
IPI00029046	Protein KIAA0152 precursor
IPI00029061	Selenoprotein P precursor
IPI00029061	Selenoprotein P precursor
IPI00029061	Selenoprotein P precursor
IPI00029061	Selenoprotein P precursor
IPI00029193	Hepatocyte growth factor activator precursor
IPI00029193	Hepatocyte growth factor activator precursor
IPI00029193	Hepatocyte growth factor activator precursor
IPI00029193	Hepatocyte growth factor activator precursor
IPI00029235	Insulin-like growth factor-binding protein 6 precursor
IPI00029235	Insulin-like growth factor-binding protein 6 precursor
IPI00029235	Insulin-like growth factor-binding protein 6 precursor
IPI00029235	Insulin-like growth factor-binding protein 6 precursor
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IPI00029235	Insulin-like growth factor-binding protein 6 precursor
IPI00029236	Insulin-like growth factor-binding protein 5 precursor
IPI00029236	Insulin-like growth factor-binding protein 5 precursor
IPI00029260	Monocyte differentiation antigen CD14 precursor
IPI00029260	Monocyte differentiation antigen CD14 precursor
IPI00029260	Monocyte differentiation antigen CD14 precursor
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IPI00029343	Isoform 1 of Contactin-associated protein-like 2 precursor
IPI00029343	Isoform 1 of Contactin-associated protein-like 2 precursor
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IPI00029343	Isoform 1 of Contactin-associated protein-like 2 precursor
IPI00029343	Isoform 1 of Contactin-associated protein-like 2 precursor
IPI00029658	Isoform 1 of EGF-containing fibulin-like extracellular matrix
IPI00029658	Isoform 1 of EGF-containing fibulin-like extracellular matrix
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IPI00029658	Isoform 1 of EGF-containing fibulin-like extracellular matrix
IPI00029658	Isoform 1 of EGF-containing fibulin-like extracellular matrix
IPI00029658	Isoform 1 of EGF-containing fibulin-like extracellular matrix
IPI00029693	Isoform A22 of Neuropilin-2 precursor
IPI00029693	Isoform A22 of Neuropilin-2 precursor
IPI00029693	Isoform A22 of Neuropilin-2 precursor
IPI00029693	Isoform A22 of Neuropilin-2 precursor
IPI00029699	Ribonuclease 4 precursor
IPI00029699	Ribonuclease 4 precursor
IPI00029699	Ribonuclease 4 precursor
IPI00029699	Ribonuclease 4 precursor
IPI00029723	Follistatin-related protein 1 precursor
IPI00029723	Follistatin-related protein 1 precursor
IPI00029723	Follistatin-related protein 1 precursor
IPI00029723	Follistatin-related protein 1 precursor
IPI00029723	Follistatin-related protein 1 precursor
IPI00029739	Isoform 1 of Complement factor H precursor
IPI00029739	Isoform 1 of Complement factor H precursor
IPI00029739	Isoform 1 of Complement factor H precursor
IPI00029739	Isoform 1 of Complement factor H precursor
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IPI00029739	Isoform 1 of Complement factor H precursor
IPI00029751	Isoform 1 of Contactin-1 precursor
IPI00029751	Isoform 1 of Contactin-1 precursor
IPI00029751	Isoform 1 of Contactin-1 precursor
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IPI00029751	Isoform 1 of Contactin-1 precursor
IPI00029751	Isoform 1 of Contactin-1 precursor
IPI00029751	Isoform 1 of Contactin-1 precursor
IPI00029863	Alpha-2-antiplasmin precursor
IPI00029863	Alpha-2-antiplasmin precursor
IPI00029863	Alpha-2-antiplasmin precursor
IPI00029863	Alpha-2-antiplasmin precursor
IPI00029863	Alpha-2-antiplasmin precursor
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IPI00029863	Alpha-2-antiplasmin precursor
IPI00029863	Alpha-2-antiplasmin precursor
IPI00029863	Alpha-2-antiplasmin precursor
IPI00029863	Alpha-2-antiplasmin precursor
IPI00030111	Growth/differentiation factor 11 precursor
IPI00030111	Growth/differentiation factor 11 precursor
IPI00030111	Growth/differentiation factor 11 precursor
IPI00030255	Procollagen-lysine,2-oxoglutarate 5-dioxygenase 3 precurs
IPI00030255	Procollagen-lysine,2-oxoglutarate 5-dioxygenase 3 precurs
IPI00030385	CDNA FLJ13813 fis, clone THYRO1000358, moderately sir
IPI00030385	CDNA FLJ13813 fis, clone THYRO1000358, moderately sir
IPI00030385	CDNA FLJ13813 fis, clone THYRO1000358, moderately sir
IPI00030385	CDNA FLJ13813 fis, clone THYRO1000358, moderately sir
IPI00030385	CDNA FLJ13813 fis, clone THYRO1000358, moderately sir
IPI00030385	CDNA FLJ13813 fis, clone THYRO1000358, moderately sir
IPI00030739	Apolipoprotein M
IPI00030739	Apolipoprotein M
IPI00030739	Apolipoprotein M
IPI00030877	15 kDa selenoprotein isoform 1 precursor
IPI00030877	15 kDa selenoprotein isoform 1 precursor

IPI00031030	Isoform 1 of Amyloid-like protein 2 precursor
IPI00031030	Isoform 1 of Amyloid-like protein 2 precursor
IPI00031030	Isoform 1 of Amyloid-like protein 2 precursor
IPI00031030	Isoform 1 of Amyloid-like protein 2 precursor
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IPI00031030	Isoform 1 of Amyloid-like protein 2 precursor
IPI00031030	Isoform 1 of Amyloid-like protein 2 precursor
IPI00031030	Isoform 1 of Amyloid-like protein 2 precursor
IPI00031121	Carboxypeptidase E precursor
IPI00031121	Carboxypeptidase E precursor
IPI00031121	Carboxypeptidase E precursor
IPI00031121	Carboxypeptidase E precursor
IPI00031121	Carboxypeptidase E precursor
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IPI00031121	Carboxypeptidase E precursor
IPI00031121	Carboxypeptidase E precursor
IPI00031121	Carboxypeptidase E precursor
IPI00031289	Neuronal pentraxin receptor
IPI00031289	Neuronal pentraxin receptor
IPI00031289	Neuronal pentraxin receptor
IPI00031411	Cadherin-related tumor suppressor homolog precursor
IPI00031461	Rab GDP dissociation inhibitor beta
IPI00031461	Rab GDP dissociation inhibitor beta
IPI00031461	Rab GDP dissociation inhibitor beta
IPI00031461	Rab GDP dissociation inhibitor beta
IPI00031549	Isoform 3A of Desmocollin-3 precursor
IPI00031549	Isoform 3A of Desmocollin-3 precursor
IPI00031549	Isoform 3A of Desmocollin-3 precursor
IPI00031564	Protein C7orf24
IPI00031564	Protein C7orf24
IPI00031564	Protein C7orf24
IPI00031821	Integral membrane protein 2B
IPI00031821	Integral membrane protein 2B
IPI00031821	Integral membrane protein 2B
IPI00032179	Antithrombin III variant
IPI00032179	Antithrombin III variant
IPI00032179	Antithrombin III variant
IPI00032179	Antithrombin III variant
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IPI00032179	Antithrombin III variant
IPI00032179	Antithrombin III variant
IPI00032179	Antithrombin III variant
IPI00032179	Antithrombin III variant
IPI00032220	Angiotensinogen precursor
IPI00032220	Angiotensinogen precursor
IPI00032220	Angiotensinogen precursor
IPI00032220	Angiotensinogen precursor
IPI00032220	Angiotensinogen precursor
IPI00032220	Angiotensinogen precursor
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IPI00032220	Angiotensinogen precursor
IPI00032220	Angiotensinogen precursor
IPI00032258	Complement C4-A precursor
IPI00032258	Complement C4-A precursor
IPI00032288	MANSC domain-containing protein 1 precursor
IPI00032288	MANSC domain-containing protein 1 precursor
IPI00032291	Complement C5 precursor
IPI00032291	Complement C5 precursor
IPI00032291	Complement C5 precursor
IPI00032291	Complement C5 precursor
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IPI00032291	Complement C5 precursor
IPI00032291	Complement C5 precursor
IPI00032291	Complement C5 precursor
IPI00032292	Metalloproteinase inhibitor 1 precursor
IPI00032292	Metalloproteinase inhibitor 1 precursor
IPI00032292	Metalloproteinase inhibitor 1 precursor
IPI00032292	Metalloproteinase inhibitor 1 precursor
IPI00032292	Metalloproteinase inhibitor 1 precursor
IPI00032292	Metalloproteinase inhibitor 1 precursor
IPI00032292	Metalloproteinase inhibitor 1 precursor
IPI00032292	Metalloproteinase inhibitor 1 precursor
IPI00032293	Cystatin C precursor
IPI00032293	Cystatin C precursor
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IPI00032293	Cystatin C precursor
IPI00032293	Cystatin C precursor
IPI00032293	Cystatin C precursor
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IPI00032293	Cystatin C precursor
IPI00032293	Cystatin C precursor
IPI00032293	Cystatin C precursor
IPI00032311	Lipopolysaccharide-binding protein precursor
IPI00032311	Lipopolysaccharide-binding protein precursor
IPI00032311	Lipopolysaccharide-binding protein precursor
IPI00032328	Isoform HMW of Kininogen-1 precursor
IPI00032328	Isoform HMW of Kininogen-1 precursor
IPI00032328	Isoform HMW of Kininogen-1 precursor
IPI00032328	Isoform HMW of Kininogen-1 precursor
IPI00032328	Isoform HMW of Kininogen-1 precursor
IPI00032328	Isoform HMW of Kininogen-1 precursor
IPI00032876	Cytokine-like protein 1 precursor
IPI00032876	Cytokine-like protein 1 precursor
IPI00033466	C-type lectin domain family 11 member A precursor
IPI00033466	C-type lectin domain family 11 member A precursor
IPI00033466	C-type lectin domain family 11 member A precursor
IPI00034319	Isoform A of Protein CutA precursor
IPI00034319	Isoform A of Protein CutA precursor
IPI00034319	Isoform A of Protein CutA precursor
IPI00034319	Isoform A of Protein CutA precursor
IPI00042514	Abhydrolase domain containing 14A
IPI00043215	immunoglobulin superfamily, member 1 isoform 1
IPI00044369	Isoform 1 of Plexin domain-containing protein 2 precursor
IPI00044369	Isoform 1 of Plexin domain-containing protein 2 precursor
IPI00044369	Isoform 1 of Plexin domain-containing protein 2 precursor
IPI00044369	Isoform 1 of Plexin domain-containing protein 2 precursor
IPI00044369	Isoform 1 of Plexin domain-containing protein 2 precursor
IPI00044369	Isoform 1 of Plexin domain-containing protein 2 precursor
IPI00044369	Isoform 1 of Plexin domain-containing protein 2 precursor
IPI00044369	Isoform 1 of Plexin domain-containing protein 2 precursor
IPI00056357	Uncharacterized protein C19orf10 precursor
IPI00056357	Uncharacterized protein C19orf10 precursor
IPI00056357	Uncharacterized protein C19orf10 precursor

IPI00056357	Uncharacterized protein C19orf10 precursor
IPI00056478	Isoform 1 of Immunoglobulin superfamily member 8 precurs
IPI00056478	Isoform 1 of Immunoglobulin superfamily member 8 precurs
IPI00056478	Isoform 1 of Immunoglobulin superfamily member 8 precurs
IPI00056478	Isoform 1 of Immunoglobulin superfamily member 8 precurs
IPI00056478	Isoform 1 of Immunoglobulin superfamily member 8 precurs
IPI00056478	Isoform 1 of Immunoglobulin superfamily member 8 precurs
IPI00060310	phospholipase D family, member 4
IPI00060310	phospholipase D family, member 4
IPI00060310	phospholipase D family, member 4
IPI00060310	phospholipase D family, member 4
IPI00060715	Potassium channel tetramerization domain-containing prote
IPI00060715	Potassium channel tetramerization domain-containing prote
IPI00060715	Potassium channel tetramerization domain-containing prote
IPI00063827	Isoform 1 of Abhydrolase domain-containing protein 14B
IPI00063827	Isoform 1 of Abhydrolase domain-containing protein 14B
IPI00064607	Hypothetical protein DKFZp781K1852
IPI00064607	Hypothetical protein DKFZp781K1852
IPI00064607	Hypothetical protein DKFZp781K1852
IPI00064607	Hypothetical protein DKFZp781K1852
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IPI00064607	Hypothetical protein DKFZp781K1852
IPI00064607	Hypothetical protein DKFZp781K1852
IPI00064607	Hypothetical protein DKFZp781K1852
IPI00064667	Glutamate carboxypeptidase-like protein 2 precursor
IPI00064667	Glutamate carboxypeptidase-like protein 2 precursor
IPI00064667	Glutamate carboxypeptidase-like protein 2 precursor
IPI00064667	Glutamate carboxypeptidase-like protein 2 precursor
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IPI00064667	Glutamate carboxypeptidase-like protein 2 precursor
IPI00064667	Glutamate carboxypeptidase-like protein 2 precursor
IPI00069058	VGF nerve growth factor inducible precursor
IPI00069058	VGF nerve growth factor inducible precursor
IPI00069058	VGF nerve growth factor inducible precursor
IPI00069058	VGF nerve growth factor inducible precursor
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IPI00069058	VGF nerve growth factor inducible precursor
IPI00069058	VGF nerve growth factor inducible precursor
IPI00069058	VGF nerve growth factor inducible precursor
IPI00072917	alpha 3 type VI collagen isoform 3 precursor

IPI00072917	alpha 3 type VI collagen isoform 3 precursor
IPI00072917	alpha 3 type VI collagen isoform 3 precursor
IPI00072917	alpha 3 type VI collagen isoform 3 precursor
IPI00072917	alpha 3 type VI collagen isoform 3 precursor
IPI00075013	Complement C1q tumor necrosis factor-related protein 1 pr
IPI00075248	Calmodulin
IPI00075248	Calmodulin
IPI00075248	Calmodulin
IPI00075248	Calmodulin
IPI00099670	Carboxyl ester lipase
IPI00099670	Carboxyl ester lipase
IPI00099670	Carboxyl ester lipase
IPI00099670	Carboxyl ester lipase
IPI00099670	Carboxyl ester lipase
IPI00099670	Carboxyl ester lipase
IPI00101405	Farnesyl diphosphate synthase
IPI00102543	SLIT and NTRK-like protein 1 precursor
IPI00102543	SLIT and NTRK-like protein 1 precursor
IPI00102543	SLIT and NTRK-like protein 1 precursor
IPI00102543	SLIT and NTRK-like protein 1 precursor
IPI00102543	SLIT and NTRK-like protein 1 precursor
IPI00102543	SLIT and NTRK-like protein 1 precursor
IPI00102543	SLIT and NTRK-like protein 1 precursor
IPI00102543	SLIT and NTRK-like protein 1 precursor
IPI00102543	SLIT and NTRK-like protein 1 precursor
IPI00103175	Isoform 1 of Soluble calcium-activated nucleotidase 1
IPI00103175	Isoform 1 of Soluble calcium-activated nucleotidase 1
IPI00103175	Isoform 1 of Soluble calcium-activated nucleotidase 1
IPI00103175	Isoform 1 of Soluble calcium-activated nucleotidase 1
IPI00103175	Isoform 1 of Soluble calcium-activated nucleotidase 1
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IPI00103175	Isoform 1 of Soluble calcium-activated nucleotidase 1
IPI00103175	Isoform 1 of Soluble calcium-activated nucleotidase 1
IPI00103471	Selenoprotein M precursor
IPI00103471	Selenoprotein M precursor
IPI00103597	Isoform 1 of VPS10 domain-containing receptor SorCS1 pr
IPI00103597	Isoform 1 of VPS10 domain-containing receptor SorCS1 pr
IPI00103597	Isoform 1 of VPS10 domain-containing receptor SorCS1 pr
IPI00103597	Isoform 1 of VPS10 domain-containing receptor SorCS1 pr
IPI00103597	Isoform 1 of VPS10 domain-containing receptor SorCS1 pr
IPI00103597	Isoform 1 of VPS10 domain-containing receptor SorCS1 pr
IPI00103636	Isoform 2 of WAP four-disulfide core domain protein 2 prec
IPI00103636	Isoform 2 of WAP four-disulfide core domain protein 2 prec
IPI00104074	Isoform 1 of Scavenger receptor cysteine-rich type 1 protein
IPI00104074	Isoform 1 of Scavenger receptor cysteine-rich type 1 protein
IPI00104074	Isoform 1 of Scavenger receptor cysteine-rich type 1 protein
IPI00104074	Isoform 1 of Scavenger receptor cysteine-rich type 1 protein
IPI00104074	Isoform 1 of Scavenger receptor cysteine-rich type 1 protein
IPI00104074	Isoform 1 of Scavenger receptor cysteine-rich type 1 protein
IPI00104074	Isoform 1 of Scavenger receptor cysteine-rich type 1 protein
IPI00104074	Isoform 1 of Scavenger receptor cysteine-rich type 1 protein
IPI00104074	Isoform 1 of Scavenger receptor cysteine-rich type 1 protein
IPI00104074	Isoform 1 of Scavenger receptor cysteine-rich type 1 protein
IPI00104074	Isoform 1 of Scavenger receptor cysteine-rich type 1 protein

IPI00107555	Isoform IIb of Profilin-2
IPI00107831	Receptor-type tyrosine-protein phosphatase F precursor
IPI00107831	Receptor-type tyrosine-protein phosphatase F precursor
IPI00107831	Receptor-type tyrosine-protein phosphatase F precursor
IPI00107831	Receptor-type tyrosine-protein phosphatase F precursor
IPI00107831	Receptor-type tyrosine-protein phosphatase F precursor
IPI00107831	Receptor-type tyrosine-protein phosphatase F precursor
IPI00107831	Receptor-type tyrosine-protein phosphatase F precursor
IPI00107831	Receptor-type tyrosine-protein phosphatase F precursor
IPI00107831	Receptor-type tyrosine-protein phosphatase F precursor
IPI00152189	Hypothetical protein (Fragment)
IPI00152540	OTTHUMP00000016748
IPI00152540	OTTHUMP00000016748
IPI00152540	OTTHUMP00000016748
IPI00152540	OTTHUMP00000016748
IPI00152540	OTTHUMP00000016748
IPI00152540	OTTHUMP00000016748
IPI00152540	OTTHUMP00000016748
IPI00152540	OTTHUMP00000016748
IPI00152789	SNED1 protein
IPI00152847	WFIKKN2 protein
IPI00152847	WFIKKN2 protein
IPI00152847	WFIKKN2 protein
IPI00152847	WFIKKN2 protein
IPI00152847	WFIKKN2 protein
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IPI00152847	WFIKKN2 protein
IPI00152850	junctional adhesion molecule 3 precursor
IPI00152850	junctional adhesion molecule 3 precursor
IPI00152861	HBE120
IPI00152861	HBE120
IPI00152861	HBE120
IPI00153049	Novel protein
IPI00153049	Novel protein
IPI00154734	seizure related 6 homolog
IPI00154734	seizure related 6 homolog
IPI00154734	seizure related 6 homolog
IPI00154742	IGLC1 protein
IPI00154742	IGLC1 protein
IPI00154742	IGLC1 protein
IPI00154742	IGLC1 protein
IPI00154742	IGLC1 protein
IPI00154742	IGLC1 protein
IPI00154742	IGLC1 protein
IPI00154742	IGLC1 protein
IPI00156171	Isoform 1 of Ectonucleotide pyrophosphatase/phosphodiest
IPI00156171	Isoform 1 of Ectonucleotide pyrophosphatase/phosphodiest
IPI00156171	Isoform 1 of Ectonucleotide pyrophosphatase/phosphodiest
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IPI00156171	Isoform 1 of Ectonucleotide pyrophosphatase/phosphodiest
IPI00157417	Isoform 4 of Seizure 6-like protein precursor
IPI00157417	Isoform 4 of Seizure 6-like protein precursor
IPI00157417	Isoform 4 of Seizure 6-like protein precursor
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IPI00159927	Neurocan core protein precursor
IPI00159927	Neurocan core protein precursor
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IPI00160552	Isoform 1 of Tenascin-R precursor
IPI00160552	Isoform 1 of Tenascin-R precursor
IPI00160552	Isoform 1 of Tenascin-R precursor
IPI00160552	Isoform 1 of Tenascin-R precursor
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IPI00160552	Isoform 1 of Tenascin-R precursor
IPI00162547	Isoform 2 of Latrophilin-3 precursor
IPI00162547	Isoform 2 of Latrophilin-3 precursor
IPI00163207	Isoform 1 of N-acetylmuramoyl-L-alanine amidase precurso
IPI00163207	Isoform 1 of N-acetylmuramoyl-L-alanine amidase precurso
IPI00163207	Isoform 1 of N-acetylmuramoyl-L-alanine amidase precurso
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IPI00163446	IGHD protein
IPI00163446	IGHD protein
IPI00163446	IGHD protein
IPI00163563	Cousin-of-RKIP 1 protein
IPI00163563	Cousin-of-RKIP 1 protein
IPI00163563	Cousin-of-RKIP 1 protein
IPI00163563	Cousin-of-RKIP 1 protein
IPI00163563	Cousin-of-RKIP 1 protein
IPI00163563	Cousin-of-RKIP 1 protein
IPI00163563	Cousin-of-RKIP 1 protein
IPI00164623	Complement C3 precursor
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IPI00164755	alpha 2 type I collagen
IPI00164755	alpha 2 type I collagen
IPI00164755	alpha 2 type I collagen
IPI00164755	alpha 2 type I collagen
IPI00164755	alpha 2 type I collagen
IPI00164755	alpha 2 type I collagen
IPI00164755	alpha 2 type I collagen
IPI00165421	SERPINC1 protein
IPI00165421	SERPINC1 protein
IPI00165421	SERPINC1 protein
IPI00165438	Muscle type neuropilin 1
IPI00165438	Muscle type neuropilin 1
IPI00165438	Muscle type neuropilin 1
IPI00165438	Muscle type neuropilin 1
IPI00165438	Muscle type neuropilin 1
IPI00165949	type 1 tumor necrosis factor receptor shedding aminopeptic
IPI00165949	type 1 tumor necrosis factor receptor shedding aminopeptic
IPI00165949	type 1 tumor necrosis factor receptor shedding aminopeptic
IPI00165949	type 1 tumor necrosis factor receptor shedding aminopeptic
IPI00165949	type 1 tumor necrosis factor receptor shedding aminopeptic
IPI00165949	type 1 tumor necrosis factor receptor shedding aminopeptic

IPI00166339	Isoform 1 of Ephrin type-A receptor 10 precursor
IPI00166339	Isoform 1 of Ephrin type-A receptor 10 precursor
IPI00166339	Isoform 1 of Ephrin type-A receptor 10 precursor
IPI00166622	PREDICTED: similar to CG14446-PA
IPI00166622	PREDICTED: similar to CG14446-PA
IPI00166622	PREDICTED: similar to CG14446-PA
IPI00166729	alpha-2-glycoprotein 1, zinc
IPI00166729	alpha-2-glycoprotein 1, zinc
IPI00166729	alpha-2-glycoprotein 1, zinc
IPI00166729	alpha-2-glycoprotein 1, zinc
IPI00166729	alpha-2-glycoprotein 1, zinc
IPI00166729	alpha-2-glycoprotein 1, zinc
IPI00166729	alpha-2-glycoprotein 1, zinc
IPI00166729	alpha-2-glycoprotein 1, zinc
IPI00166729	alpha-2-glycoprotein 1, zinc
IPI00166729	alpha-2-glycoprotein 1, zinc
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IPI00166729	alpha-2-glycoprotein 1, zinc
IPI00166729	alpha-2-glycoprotein 1, zinc
IPI00166729	alpha-2-glycoprotein 1, zinc
IPI00166729	alpha-2-glycoprotein 1, zinc
IPI00166729	alpha-2-glycoprotein 1, zinc
IPI00166766	CDNA FLJ90761 fis, clone THYRO1000099
IPI00166766	CDNA FLJ90761 fis, clone THYRO1000099
IPI00166766	CDNA FLJ90761 fis, clone THYRO1000099
IPI00166766	CDNA FLJ90761 fis, clone THYRO1000099
IPI00166766	CDNA FLJ90761 fis, clone THYRO1000099
IPI00166766	CDNA FLJ90761 fis, clone THYRO1000099
IPI00166766	CDNA FLJ90761 fis, clone THYRO1000099
IPI00166766	CDNA FLJ90761 fis, clone THYRO1000099
IPI00166766	CDNA FLJ90761 fis, clone THYRO1000099
IPI00166766	CDNA FLJ90761 fis, clone THYRO1000099
IPI00166766	CDNA FLJ90761 fis, clone THYRO1000099
IPI00166766	CDNA FLJ90761 fis, clone THYRO1000099
IPI00166768	TUBA6 protein
IPI00168479	apolipoprotein A-I binding protein precursor
IPI00168479	apolipoprotein A-I binding protein precursor
IPI00168520	Isoform 2 of Matrilin-2 precursor
IPI00168812	PTK7 protein tyrosine kinase 7 isoform d precursor
IPI00168866	Isoform 1 of MAM domain-containing glycosylphosphatidylii
IPI00168866	Isoform 1 of MAM domain-containing glycosylphosphatidylii
IPI00168866	Isoform 1 of MAM domain-containing glycosylphosphatidylii
IPI00168884	Renin receptor precursor
IPI00168884	Renin receptor precursor
IPI00168884	Renin receptor precursor
IPI00168884	Renin receptor precursor
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IPI00168884	Renin receptor precursor
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IPI00168884	Renin receptor precursor
IPI00169383	Phosphoglycerate kinase 1

IPI00169383	Phosphoglycerate kinase 1
IPI00169383	Phosphoglycerate kinase 1
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IPI00169383	Phosphoglycerate kinase 1
IPI00169383	Phosphoglycerate kinase 1
IPI00170551	Isoform 2 of Semaphorin-6D precursor
IPI00170551	Isoform 2 of Semaphorin-6D precursor
IPI00170551	Isoform 2 of Semaphorin-6D precursor
IPI00170551	Isoform 2 of Semaphorin-6D precursor
IPI00170551	Isoform 2 of Semaphorin-6D precursor
IPI00170551	Isoform 2 of Semaphorin-6D precursor
IPI00170551	Isoform 2 of Semaphorin-6D precursor
IPI00171411	Golgi phosphoprotein 2
IPI00171411	Golgi phosphoprotein 2
IPI00171411	Golgi phosphoprotein 2
IPI00171411	Golgi phosphoprotein 2
IPI00171473	Spondin-1 precursor
IPI00171473	Spondin-1 precursor
IPI00171473	Spondin-1 precursor
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IPI00171473	Spondin-1 precursor
IPI00171473	Spondin-1 precursor
IPI00171473	Spondin-1 precursor
IPI00171473	Spondin-1 precursor
IPI00172450	Isoform 4 of Calcium/calmodulin-dependent protein kinase
IPI00172450	Isoform 4 of Calcium/calmodulin-dependent protein kinase
IPI00172636	calcium/calmodulin-dependent protein kinase II delta isofor
IPI00172636	calcium/calmodulin-dependent protein kinase II delta isofor
IPI00176193	Isoform 1 of Collagen alpha-1(XIV) chain precursor
IPI00176193	Isoform 1 of Collagen alpha-1(XIV) chain precursor
IPI00176221	Neuronal growth regulator 1 precursor
IPI00176221	Neuronal growth regulator 1 precursor
IPI00176221	Neuronal growth regulator 1 precursor
IPI00176221	Neuronal growth regulator 1 precursor
IPI00176221	Neuronal growth regulator 1 precursor
IPI00176221	Neuronal growth regulator 1 precursor
IPI00176221	Neuronal growth regulator 1 precursor
IPI00176221	Neuronal growth regulator 1 precursor
IPI00176221	Neuronal growth regulator 1 precursor
IPI00176221	Neuronal growth regulator 1 precursor
IPI00176221	Neuronal growth regulator 1 precursor
IPI00176424	Neuroligin-2 precursor
IPI00176424	Neuroligin-2 precursor
IPI00176424	Neuroligin-2 precursor
IPI00176427	TSLC1-like 2
IPI00176427	TSLC1-like 2
IPI00176427	TSLC1-like 2
IPI00176427	TSLC1-like 2
IPI00176427	TSLC1-like 2
IPI00176427	TSLC1-like 2
IPI00176427	TSLC1-like 2

IPI00176427	TSLC1-like 2
IPI00176427	TSLC1-like 2
IPI00176458	protocadherin 1 isoform 2 precursor
IPI00176458	protocadherin 1 isoform 2 precursor
IPI00176458	protocadherin 1 isoform 2 precursor
IPI00176458	protocadherin 1 isoform 2 precursor
IPI00176458	protocadherin 1 isoform 2 precursor
IPI00176458	protocadherin 1 isoform 2 precursor
IPI00176482	CDNA FLJ42404 fis, clone ASTRO3000301, highly similar t
IPI00176482	CDNA FLJ42404 fis, clone ASTRO3000301, highly similar t
IPI00177543	Isoform 1 of Peptidyl-glycine alpha-amidating monooxygena
IPI00177543	Isoform 1 of Peptidyl-glycine alpha-amidating monooxygena
IPI00177543	Isoform 1 of Peptidyl-glycine alpha-amidating monooxygena
IPI00177543	Isoform 1 of Peptidyl-glycine alpha-amidating monooxygena
IPI00177543	Isoform 1 of Peptidyl-glycine alpha-amidating monooxygena
IPI00177543	Isoform 1 of Peptidyl-glycine alpha-amidating monooxygena
IPI00177543	Isoform 1 of Peptidyl-glycine alpha-amidating monooxygena
IPI00177543	Isoform 1 of Peptidyl-glycine alpha-amidating monooxygena
IPI00177543	Isoform 1 of Peptidyl-glycine alpha-amidating monooxygena
IPI00177543	Isoform 1 of Peptidyl-glycine alpha-amidating monooxygena
IPI00177543	Isoform 1 of Peptidyl-glycine alpha-amidating monooxygena
IPI00177543	Isoform 1 of Peptidyl-glycine alpha-amidating monooxygena
IPI00177543	Isoform 1 of Peptidyl-glycine alpha-amidating monooxygena
IPI00177543	Isoform 1 of Peptidyl-glycine alpha-amidating monooxygena
IPI00177543	Isoform 1 of Peptidyl-glycine alpha-amidating monooxygena
IPI00177543	Isoform 1 of Peptidyl-glycine alpha-amidating monooxygena
IPI00177543	Isoform 1 of Peptidyl-glycine alpha-amidating monooxygena
IPI00177543	Isoform 1 of Peptidyl-glycine alpha-amidating monooxygena
IPI00177543	Isoform 1 of Peptidyl-glycine alpha-amidating monooxygena
IPI00177543	Isoform 1 of Peptidyl-glycine alpha-amidating monooxygena
IPI00177543	Isoform 1 of Peptidyl-glycine alpha-amidating monooxygena
IPI00177543	Isoform 1 of Peptidyl-glycine alpha-amidating monooxygena
IPI00177543	Isoform 1 of Peptidyl-glycine alpha-amidating monooxygena
IPI00177543	Isoform 1 of Peptidyl-glycine alpha-amidating monooxygena
IPI00178854	Isoform 1 of Contactin-4 precursor
IPI00178854	Isoform 1 of Contactin-4 precursor
IPI00178854	Isoform 1 of Contactin-4 precursor
IPI00178854	Isoform 1 of Contactin-4 precursor
IPI00178854	Isoform 1 of Contactin-4 precursor
IPI00178854	Isoform 1 of Contactin-4 precursor
IPI00178854	Isoform 1 of Contactin-4 precursor
IPI00178854	Isoform 1 of Contactin-4 precursor
IPI00178854	Isoform 1 of Contactin-4 precursor
IPI00178854	Isoform 1 of Contactin-4 precursor
IPI00178854	Isoform 1 of Contactin-4 precursor
IPI00178854	Isoform 1 of Contactin-4 precursor
IPI00178926	immunoglobulin J chain
IPI00178926	immunoglobulin J chain
IPI00178926	immunoglobulin J chain
IPI00179330	ubiquitin and ribosomal protein S27a precursor
IPI00179330	ubiquitin and ribosomal protein S27a precursor
IPI00179330	ubiquitin and ribosomal protein S27a precursor

IPI00183445	Isoform 1 of Latrophilin-1 precursor
IPI00183445	Isoform 1 of Latrophilin-1 precursor
IPI00183445	Isoform 1 of Latrophilin-1 precursor
IPI00183445	Isoform 1 of Latrophilin-1 precursor
IPI00183445	Isoform 1 of Latrophilin-1 precursor
IPI00183445	Isoform 1 of Latrophilin-1 precursor
IPI00183445	Isoform 1 of Latrophilin-1 precursor
IPI00183445	Isoform 1 of Latrophilin-1 precursor
IPI00183445	Isoform 1 of Latrophilin-1 precursor
IPI00183487	Xylosyltransferase 1
IPI00183487	Xylosyltransferase 1
IPI00184988	CDNA FLJ31966 fis, clone NT2RP7007925, weakly similar
IPI00184988	CDNA FLJ31966 fis, clone NT2RP7007925, weakly similar
IPI00185362	Neural cell adhesion molecule 1
IPI00185362	Neural cell adhesion molecule 1
IPI00185362	Neural cell adhesion molecule 1
IPI00185362	Neural cell adhesion molecule 1
IPI00185362	Neural cell adhesion molecule 1
IPI00185362	Neural cell adhesion molecule 1
IPI00185362	Neural cell adhesion molecule 1
IPI00185362	Neural cell adhesion molecule 1
IPI00185362	Neural cell adhesion molecule 1
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IPI00185362	Neural cell adhesion molecule 1
IPI00185362	Neural cell adhesion molecule 1
IPI00185362	Neural cell adhesion molecule 1
IPI00185362	Neural cell adhesion molecule 1
IPI00185362	Neural cell adhesion molecule 1
IPI00185362	Neural cell adhesion molecule 1
IPI00185362	Neural cell adhesion molecule 1
IPI00186903	Isoform 2 of Apolipoprotein-L1 precursor
IPI00186903	Isoform 2 of Apolipoprotein-L1 precursor
IPI00186903	Isoform 2 of Apolipoprotein-L1 precursor
IPI00186903	Isoform 2 of Apolipoprotein-L1 precursor
IPI00186903	Isoform 2 of Apolipoprotein-L1 precursor
IPI00186903	Isoform 2 of Apolipoprotein-L1 precursor
IPI00215715	Isoform A of Calcium/calmodulin-dependent protein kinase
IPI00215715	Isoform A of Calcium/calmodulin-dependent protein kinase
IPI00215715	Isoform A of Calcium/calmodulin-dependent protein kinase
IPI00215894	Isoform LMW of Kininogen-1 precursor
IPI00215894	Isoform LMW of Kininogen-1 precursor
IPI00215894	Isoform LMW of Kininogen-1 precursor
IPI00215894	Isoform LMW of Kininogen-1 precursor
IPI00215894	Isoform LMW of Kininogen-1 precursor
IPI00215894	Isoform LMW of Kininogen-1 precursor
IPI00215894	Isoform LMW of Kininogen-1 precursor
IPI00215894	Isoform LMW of Kininogen-1 precursor
IPI00215899	Isoform 2 of Sushi repeat-containing protein SRPX precurs
IPI00215899	Isoform 2 of Sushi repeat-containing protein SRPX precurs
IPI00215983	Carbonic anhydrase 1
IPI00215983	Carbonic anhydrase 1
IPI00215983	Carbonic anhydrase 1
IPI00215983	Carbonic anhydrase 1
IPI00216138	Transgelin
IPI00216138	Transgelin

IPI00216138	Transgelin
IPI00216138	Transgelin
IPI00216138	Transgelin
IPI00216138	Transgelin
IPI00216138	Transgelin
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IPI00216138	Transgelin
IPI00216138	Transgelin
IPI00216138	Transgelin
IPI00216138	Transgelin
IPI00216138	Transgelin
IPI00216171	Gamma-enolase
IPI00216171	Gamma-enolase
IPI00216171	Gamma-enolase
IPI00216171	Gamma-enolase
IPI00216171	Gamma-enolase
IPI00216171	Gamma-enolase
IPI00216171	Gamma-enolase
IPI00216171	Gamma-enolase
IPI00216171	Gamma-enolase
IPI00216171	Gamma-enolase
IPI00216171	Gamma-enolase
IPI00216250	cell recognition protein CASPR4 isoform 1
IPI00216250	cell recognition protein CASPR4 isoform 1
IPI00216250	cell recognition protein CASPR4 isoform 1
IPI00216250	cell recognition protein CASPR4 isoform 1
IPI00216250	cell recognition protein CASPR4 isoform 1
IPI00216250	cell recognition protein CASPR4 isoform 1
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IPI00216250	cell recognition protein CASPR4 isoform 1
IPI00216250	cell recognition protein CASPR4 isoform 1
IPI00216250	cell recognition protein CASPR4 isoform 1
IPI00216283	Isoform Short of Receptor-type tyrosine-protein phosphatas
IPI00216283	Isoform Short of Receptor-type tyrosine-protein phosphatas
IPI00216283	Isoform Short of Receptor-type tyrosine-protein phosphatas
IPI00216283	Isoform Short of Receptor-type tyrosine-protein phosphatas
IPI00216298	Thioredoxin
IPI00216298	Thioredoxin
IPI00216298	Thioredoxin
IPI00216298	Thioredoxin
IPI00216298	Thioredoxin
IPI00216318	tyrosine 3-monooxygenase/tryptophan 5-monooxygenase a
IPI00216318	tyrosine 3-monooxygenase/tryptophan 5-monooxygenase a
IPI00216318	tyrosine 3-monooxygenase/tryptophan 5-monooxygenase a
IPI00216318	tyrosine 3-monooxygenase/tryptophan 5-monooxygenase a

IPI00216461	Acylphosphatase, muscle type isozyme
IPI00216461	Acylphosphatase, muscle type isozyme
IPI00216461	Acylphosphatase, muscle type isozyme
IPI00216691	Profilin-1
IPI00216691	Profilin-1
IPI00216691	Profilin-1
IPI00216691	Profilin-1
IPI00216691	Profilin-1
IPI00216723	Neurexin 3-beta (Fragment)
IPI00216723	Neurexin 3-beta (Fragment)
IPI00216723	Neurexin 3-beta (Fragment)
IPI00216728	Neurexin 3-alpha
IPI00216728	Neurexin 3-alpha
IPI00216728	Neurexin 3-alpha
IPI00216728	Neurexin 3-alpha
IPI00216728	Neurexin 3-alpha
IPI00216728	Neurexin 3-alpha
IPI00216728	Neurexin 3-alpha
IPI00216728	Neurexin 3-alpha
IPI00216728	Neurexin 3-alpha
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IPI00216728	Neurexin 3-alpha
IPI00216728	Neurexin 3-alpha
IPI00216728	Neurexin 3-alpha
IPI00216728	Neurexin 3-alpha
IPI00216728	Neurexin 3-alpha
IPI00216882	mannan-binding lectin serine protease 1 isoform 3
IPI00217146	SLIT and NTRK-like protein 4 precursor
IPI00217146	SLIT and NTRK-like protein 4 precursor
IPI00217146	SLIT and NTRK-like protein 4 precursor
IPI00217146	SLIT and NTRK-like protein 4 precursor
IPI00217146	SLIT and NTRK-like protein 4 precursor
IPI00217146	SLIT and NTRK-like protein 4 precursor
IPI00217146	SLIT and NTRK-like protein 4 precursor
IPI00217423	Hyaluronan binding protein (Fragment)
IPI00217423	Hyaluronan binding protein (Fragment)
IPI00217493	Myoglobin
IPI00217493	Myoglobin
IPI00217561	Isoform Beta-1C of Integrin beta-1 precursor
IPI00217778	Isoform 2 of Phospholipid transfer protein precursor
IPI00217778	Isoform 2 of Phospholipid transfer protein precursor
IPI00217882	Sortilin precursor
IPI00217882	Sortilin precursor
IPI00217882	Sortilin precursor
IPI00217882	Sortilin precursor
IPI00217966	lactate dehydrogenase A
IPI00217966	lactate dehydrogenase A
IPI00217966	lactate dehydrogenase A
IPI00217966	lactate dehydrogenase A
IPI00217966	lactate dehydrogenase A

IPI00217966	lactate dehydrogenase A
IPI00217966	lactate dehydrogenase A
IPI00217966	lactate dehydrogenase A
IPI00218046	Heparan-sulfate 6-O-sulfotransferase 3
IPI00218046	Heparan-sulfate 6-O-sulfotransferase 3
IPI00218192	Isoform 2 of Inter-alpha-trypsin inhibitor heavy chain H4 pre
IPI00218192	Isoform 2 of Inter-alpha-trypsin inhibitor heavy chain H4 pre
IPI00218192	Isoform 2 of Inter-alpha-trypsin inhibitor heavy chain H4 pre
IPI00218192	Isoform 2 of Inter-alpha-trypsin inhibitor heavy chain H4 pre
IPI00218192	Isoform 2 of Inter-alpha-trypsin inhibitor heavy chain H4 pre
IPI00218192	Isoform 2 of Inter-alpha-trypsin inhibitor heavy chain H4 pre
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IPI00218192	Isoform 2 of Inter-alpha-trypsin inhibitor heavy chain H4 pre
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IPI00218192	Isoform 2 of Inter-alpha-trypsin inhibitor heavy chain H4 pre
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IPI00218192	Isoform 2 of Inter-alpha-trypsin inhibitor heavy chain H4 pre
IPI00218192	Isoform 2 of Inter-alpha-trypsin inhibitor heavy chain H4 pre
IPI00218192	Isoform 2 of Inter-alpha-trypsin inhibitor heavy chain H4 pre
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IPI00218192	Isoform 2 of Inter-alpha-trypsin inhibitor heavy chain H4 pre
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IPI00218192	Isoform 2 of Inter-alpha-trypsin inhibitor heavy chain H4 pre
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IPI00218192	Isoform 2 of Inter-alpha-trypsin inhibitor heavy chain H4 pre
IPI00218192	Isoform 2 of Inter-alpha-trypsin inhibitor heavy chain H4 pre
IPI00218192	Isoform 2 of Inter-alpha-trypsin inhibitor heavy chain H4 pre
IPI00218192	Isoform 2 of Inter-alpha-trypsin inhibitor heavy chain H4 pre
IPI00218192	Isoform 2 of Inter-alpha-trypsin inhibitor heavy chain H4 pre
IPI00218192	Isoform 2 of Inter-alpha-trypsin inhibitor heavy chain H4 pre
IPI00218413	biotinidase precursor
IPI00218413	biotinidase precursor
IPI00218413	biotinidase precursor
IPI00218413	biotinidase precursor
IPI00218413	biotinidase precursor
IPI00218413	biotinidase precursor
IPI00218493	Hypoxanthine-guanine phosphoribosyltransferase
IPI00218493	Hypoxanthine-guanine phosphoribosyltransferase
IPI00218493	Hypoxanthine-guanine phosphoribosyltransferase
IPI00218493	Hypoxanthine-guanine phosphoribosyltransferase
IPI00218725	laminin alpha 2 subunit precursor
IPI00218725	laminin alpha 2 subunit precursor
IPI00218725	laminin alpha 2 subunit precursor
IPI00218725	laminin alpha 2 subunit precursor
IPI00218725	laminin alpha 2 subunit precursor
IPI00218725	laminin alpha 2 subunit precursor

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IPI00218725	laminin alpha 2 subunit precursor
IPI00218725	laminin alpha 2 subunit precursor
IPI00218725	laminin alpha 2 subunit precursor
IPI00218725	laminin alpha 2 subunit precursor
IPI00218725	laminin alpha 2 subunit precursor
IPI00218732	Serum paraoxonase/arylesterase 1
IPI00218732	Serum paraoxonase/arylesterase 1
IPI00218732	Serum paraoxonase/arylesterase 1
IPI00218732	Serum paraoxonase/arylesterase 1
IPI00218732	Serum paraoxonase/arylesterase 1
IPI00218732	Serum paraoxonase/arylesterase 1
IPI00218732	Serum paraoxonase/arylesterase 1
IPI00218732	Serum paraoxonase/arylesterase 1
IPI00218732	Serum paraoxonase/arylesterase 1
IPI00218733	Superoxide dismutase
IPI00218733	Superoxide dismutase
IPI00218733	Superoxide dismutase
IPI00218733	Superoxide dismutase
IPI00218733	Superoxide dismutase
IPI00218733	Superoxide dismutase
IPI00218733	Superoxide dismutase
IPI00218733	Superoxide dismutase
IPI00218733	Superoxide dismutase
IPI00218746	Complement Component 1, q subComponent, beta polypep
IPI00218746	Complement Component 1, q subComponent, beta polypep
IPI00218746	Complement Component 1, q subComponent, beta polypep
IPI00218803	Isoform B of Fibulin-1 precursor
IPI00218803	Isoform B of Fibulin-1 precursor
IPI00218803	Isoform B of Fibulin-1 precursor
IPI00218803	Isoform B of Fibulin-1 precursor
IPI00218803	Isoform B of Fibulin-1 precursor
IPI00219018	Glyceraldehyde-3-phosphate dehydrogenase
IPI00219018	Glyceraldehyde-3-phosphate dehydrogenase
IPI00219018	Glyceraldehyde-3-phosphate dehydrogenase
IPI00219018	Glyceraldehyde-3-phosphate dehydrogenase
IPI00219018	Glyceraldehyde-3-phosphate dehydrogenase
IPI00219018	Glyceraldehyde-3-phosphate dehydrogenase
IPI00219018	Glyceraldehyde-3-phosphate dehydrogenase
IPI00219018	Glyceraldehyde-3-phosphate dehydrogenase
IPI00219018	Glyceraldehyde-3-phosphate dehydrogenase
IPI00219018	Glyceraldehyde-3-phosphate dehydrogenase

IPI00219029	Aspartate aminotransferase, cytoplasmic
IPI00219029	Aspartate aminotransferase, cytoplasmic
IPI00219029	Aspartate aminotransferase, cytoplasmic
IPI00219029	Aspartate aminotransferase, cytoplasmic
IPI00219029	Aspartate aminotransferase, cytoplasmic
IPI00219029	Aspartate aminotransferase, cytoplasmic
IPI00219029	Aspartate aminotransferase, cytoplasmic
IPI00219029	Aspartate aminotransferase, cytoplasmic
IPI00219029	Aspartate aminotransferase, cytoplasmic
IPI00219029	Aspartate aminotransferase, cytoplasmic
IPI00219029	Aspartate aminotransferase, cytoplasmic
IPI00219029	Aspartate aminotransferase, cytoplasmic
IPI00219029	Aspartate aminotransferase, cytoplasmic
IPI00219029	Aspartate aminotransferase, cytoplasmic
IPI00219029	Aspartate aminotransferase, cytoplasmic
IPI00219029	Aspartate aminotransferase, cytoplasmic
IPI00219029	Aspartate aminotransferase, cytoplasmic
IPI00219029	Aspartate aminotransferase, cytoplasmic
IPI00219029	Aspartate aminotransferase, cytoplasmic
IPI00219131	Isoform 1 of ICOS ligand precursor
IPI00219131	Isoform 1 of ICOS ligand precursor
IPI00219217	L-lactate dehydrogenase B chain
IPI00219217	L-lactate dehydrogenase B chain
IPI00219217	L-lactate dehydrogenase B chain
IPI00219217	L-lactate dehydrogenase B chain
IPI00219217	L-lactate dehydrogenase B chain
IPI00219217	L-lactate dehydrogenase B chain
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IPI00219217	L-lactate dehydrogenase B chain
IPI00219217	L-lactate dehydrogenase B chain
IPI00219217	L-lactate dehydrogenase B chain
IPI00219217	L-lactate dehydrogenase B chain
IPI00219217	L-lactate dehydrogenase B chain
IPI00219219	Galectin-1
IPI00219219	Galectin-1
IPI00219219	Galectin-1
IPI00219219	Galectin-1
IPI00219219	Galectin-1
IPI00219219	Galectin-1
IPI00219219	Galectin-1
IPI00219365	Moesin
IPI00219365	Moesin
IPI00219365	Moesin
IPI00219446	Phosphatidylethanolamine-binding protein 1
IPI00219446	Phosphatidylethanolamine-binding protein 1
IPI00219446	Phosphatidylethanolamine-binding protein 1
IPI00219446	Phosphatidylethanolamine-binding protein 1
IPI00219446	Phosphatidylethanolamine-binding protein 1
IPI00219446	Phosphatidylethanolamine-binding protein 1
IPI00219446	Phosphatidylethanolamine-binding protein 1
IPI00219446	Phosphatidylethanolamine-binding protein 1
IPI00219446	Phosphatidylethanolamine-binding protein 1

IPI00219446	Phosphatidylethanolamine-binding protein 1
IPI00219446	Phosphatidylethanolamine-binding protein 1
IPI00219446	Phosphatidylethanolamine-binding protein 1
IPI00219465	Transcobalamin-2 precursor
IPI00219465	Transcobalamin-2 precursor
IPI00219465	Transcobalamin-2 precursor
IPI00219465	Transcobalamin-2 precursor
IPI00219465	Transcobalamin-2 precursor
IPI00219575	bleomycin hydrolase
IPI00219664	Isoform 2 of Myelin-oligodendrocyte glycoprotein precursor
IPI00219664	Isoform 2 of Myelin-oligodendrocyte glycoprotein precursor
IPI00219664	Isoform 2 of Myelin-oligodendrocyte glycoprotein precursor
IPI00219703	Parvalbumin alpha
IPI00219703	Parvalbumin alpha
IPI00219703	Parvalbumin alpha
IPI00219703	Parvalbumin alpha
IPI00219703	Parvalbumin alpha
IPI00219757	Glutathione S-transferase P
IPI00219757	Glutathione S-transferase P
IPI00219757	Glutathione S-transferase P
IPI00219757	Glutathione S-transferase P
IPI00219757	Glutathione S-transferase P
IPI00219757	Glutathione S-transferase P
IPI00219757	Glutathione S-transferase P
IPI00219757	Glutathione S-transferase P
IPI00219798	Isoform 1 of Roundabout homolog 1 precursor
IPI00219798	Isoform 1 of Roundabout homolog 1 precursor
IPI00219798	Isoform 1 of Roundabout homolog 1 precursor
IPI00219798	Isoform 1 of Roundabout homolog 1 precursor
IPI00219798	Isoform 1 of Roundabout homolog 1 precursor
IPI00219798	Isoform 1 of Roundabout homolog 1 precursor
IPI00219798	Isoform 1 of Roundabout homolog 1 precursor
IPI00219798	Isoform 1 of Roundabout homolog 1 precursor
IPI00219798	Isoform 1 of Roundabout homolog 1 precursor
IPI00219798	Isoform 1 of Roundabout homolog 1 precursor
IPI00219798	Isoform 1 of Roundabout homolog 1 precursor
IPI00219798	Isoform 1 of Roundabout homolog 1 precursor
IPI00219798	Isoform 1 of Roundabout homolog 1 precursor
IPI00219798	Isoform 1 of Roundabout homolog 1 precursor
IPI00219930	Cellular retinoic acid-binding protein 1
IPI00219930	Cellular retinoic acid-binding protein 1
IPI00219930	Cellular retinoic acid-binding protein 1
IPI00219930	Cellular retinoic acid-binding protein 1
IPI00219930	Cellular retinoic acid-binding protein 1
IPI00220249	Latent-transforming growth factor beta-binding protein, isof
IPI00220249	Latent-transforming growth factor beta-binding protein, isof
IPI00220303	mannosidase, alpha, class 2A, member 2
IPI00220303	mannosidase, alpha, class 2A, member 2
IPI00220303	mannosidase, alpha, class 2A, member 2
IPI00220303	mannosidase, alpha, class 2A, member 2
IPI00220303	mannosidase, alpha, class 2A, member 2
IPI00220303	mannosidase, alpha, class 2A, member 2
IPI00220303	mannosidase, alpha, class 2A, member 2
IPI00220303	mannosidase, alpha, class 2A, member 2
IPI00220303	mannosidase, alpha, class 2A, member 2
IPI00220303	mannosidase, alpha, class 2A, member 2

IPI00220342	NG,NG-dimethylarginine dimethylaminohydrolase 1
IPI00220342	NG,NG-dimethylarginine dimethylaminohydrolase 1
IPI00220342	NG,NG-dimethylarginine dimethylaminohydrolase 1
IPI00220342	NG,NG-dimethylarginine dimethylaminohydrolase 1
IPI00220342	NG,NG-dimethylarginine dimethylaminohydrolase 1
IPI00220342	NG,NG-dimethylarginine dimethylaminohydrolase 1
IPI00220342	NG,NG-dimethylarginine dimethylaminohydrolase 1
IPI00220361	Calbindin
IPI00220361	Calbindin
IPI00220361	Calbindin
IPI00220361	Calbindin
IPI00220562	Neuronal pentraxin-1 precursor
IPI00220562	Neuronal pentraxin-1 precursor
IPI00220562	Neuronal pentraxin-1 precursor
IPI00220562	Neuronal pentraxin-1 precursor
IPI00220562	Neuronal pentraxin-1 precursor
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IPI00220562	Neuronal pentraxin-1 precursor
IPI00220562	Neuronal pentraxin-1 precursor
IPI00220562	Neuronal pentraxin-1 precursor
IPI00220642	14-3-3 protein gamma
IPI00220642	14-3-3 protein gamma
IPI00220642	14-3-3 protein gamma
IPI00220644	Isoform M1 of Pyruvate kinase isozymes M1/M2
IPI00220644	Isoform M1 of Pyruvate kinase isozymes M1/M2
IPI00220644	Isoform M1 of Pyruvate kinase isozymes M1/M2
IPI00220644	Isoform M1 of Pyruvate kinase isozymes M1/M2
IPI00220644	Isoform M1 of Pyruvate kinase isozymes M1/M2
IPI00220644	Isoform M1 of Pyruvate kinase isozymes M1/M2
IPI00220644	Isoform M1 of Pyruvate kinase isozymes M1/M2
IPI00220644	Isoform M1 of Pyruvate kinase isozymes M1/M2
IPI00220644	Isoform M1 of Pyruvate kinase isozymes M1/M2
IPI00220644	Isoform M1 of Pyruvate kinase isozymes M1/M2
IPI00220644	Isoform M1 of Pyruvate kinase isozymes M1/M2
IPI00220644	Isoform M1 of Pyruvate kinase isozymes M1/M2
IPI00220644	Isoform M1 of Pyruvate kinase isozymes M1/M2
IPI00220644	Isoform M1 of Pyruvate kinase isozymes M1/M2
IPI00220644	Isoform M1 of Pyruvate kinase isozymes M1/M2
IPI00220701	Isoform 2 of Collagen alpha-3(VI) chain precursor
IPI00220701	Isoform 2 of Collagen alpha-3(VI) chain precursor
IPI00220701	Isoform 2 of Collagen alpha-3(VI) chain precursor
IPI00220701	Isoform 2 of Collagen alpha-3(VI) chain precursor
IPI00220701	Isoform 2 of Collagen alpha-3(VI) chain precursor
IPI00220701	Isoform 2 of Collagen alpha-3(VI) chain precursor
IPI00220701	Isoform 2 of Collagen alpha-3(VI) chain precursor
IPI00220737	Isoform N-CAM 120 of Neural cell adhesion molecule 1, 12
IPI00220737	Isoform N-CAM 120 of Neural cell adhesion molecule 1, 12
IPI00220737	Isoform N-CAM 120 of Neural cell adhesion molecule 1, 12
IPI00220737	Isoform N-CAM 120 of Neural cell adhesion molecule 1, 12
IPI00220737	Isoform N-CAM 120 of Neural cell adhesion molecule 1, 12
IPI00220737	Isoform N-CAM 120 of Neural cell adhesion molecule 1, 12
IPI00220737	Isoform N-CAM 120 of Neural cell adhesion molecule 1, 12

IPI00220737	Isoform N-CAM 120 of Neural cell adhesion molecule 1, 12
IPI00220737	Isoform N-CAM 120 of Neural cell adhesion molecule 1, 12
IPI00220991	Isoform 2 of AP-2 complex subunit beta-1
IPI00220991	Isoform 2 of AP-2 complex subunit beta-1
IPI00220991	Isoform 2 of AP-2 complex subunit beta-1
IPI00220991	Isoform 2 of AP-2 complex subunit beta-1
IPI00221224	Aminopeptidase N
IPI00221224	Aminopeptidase N
IPI00240988	PREDICTED: 6720455I24Rik homolog
IPI00240988	PREDICTED: 6720455I24Rik homolog
IPI00241562	Isoform 2 of Reelin precursor
IPI00241562	Isoform 2 of Reelin precursor
IPI00241562	Isoform 2 of Reelin precursor
IPI00241562	Isoform 2 of Reelin precursor
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IPI00241562	Isoform 2 of Reelin precursor
IPI00241562	Isoform 2 of Reelin precursor
IPI00241562	Isoform 2 of Reelin precursor
IPI00242956	Fc fragment of IgG binding protein
IPI00242956	Fc fragment of IgG binding protein
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IPI00242956	Fc fragment of IgG binding protein
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IPI00242956	Fc fragment of IgG binding protein
IPI00242956	Fc fragment of IgG binding protein
IPI00242956	Fc fragment of IgG binding protein
IPI00242956	Fc fragment of IgG binding protein
IPI00247243	PREDICTED: hypothetical protein XP_292785
IPI00247243	PREDICTED: hypothetical protein XP_292785
IPI00247243	PREDICTED: hypothetical protein XP_292785
IPI00247616	Scavenger receptor with C-type lectin type I
IPI00247616	Scavenger receptor with C-type lectin type I
IPI00247616	Scavenger receptor with C-type lectin type I
IPI00252731	Isoform DPPX-S of Dipeptidyl aminopeptidase-like protein 6
IPI00252731	Isoform DPPX-S of Dipeptidyl aminopeptidase-like protein 6
IPI00252731	Isoform DPPX-S of Dipeptidyl aminopeptidase-like protein 6
IPI00252731	Isoform DPPX-S of Dipeptidyl aminopeptidase-like protein 6
IPI00252731	Isoform DPPX-S of Dipeptidyl aminopeptidase-like protein 6
IPI00257882	Xaa-Pro dipeptidase
IPI00259102	Mammalian ependymin-related protein 1 precursor
IPI00259102	Mammalian ependymin-related protein 1 precursor
IPI00259102	Mammalian ependymin-related protein 1 precursor
IPI00259102	Mammalian ependymin-related protein 1 precursor
IPI00289058	Ly-6/neurotoxin-like protein 1 precursor
IPI00289058	Ly-6/neurotoxin-like protein 1 precursor
IPI00289083	Novel protein
IPI00289083	Novel protein
IPI00289204	Reticulon-4 receptor precursor

IPI00289204	Reticulon-4 receptor precursor
IPI00289204	Reticulon-4 receptor precursor
IPI00289204	Reticulon-4 receptor precursor
IPI00289204	Reticulon-4 receptor precursor
IPI00289819	Cation-independent mannose-6-phosphate receptor precurs
IPI00289819	Cation-independent mannose-6-phosphate receptor precurs
IPI00289819	Cation-independent mannose-6-phosphate receptor precurs
IPI00289831	Isoform PTPS of Receptor-type tyrosine-protein phosphatas
IPI00289831	Isoform PTPS of Receptor-type tyrosine-protein phosphatas
IPI00289831	Isoform PTPS of Receptor-type tyrosine-protein phosphatas
IPI00289831	Isoform PTPS of Receptor-type tyrosine-protein phosphatas
IPI00289831	Isoform PTPS of Receptor-type tyrosine-protein phosphatas
IPI00289831	Isoform PTPS of Receptor-type tyrosine-protein phosphatas
IPI00289831	Isoform PTPS of Receptor-type tyrosine-protein phosphatas
IPI00289831	Isoform PTPS of Receptor-type tyrosine-protein phosphatas
IPI00289831	Isoform PTPS of Receptor-type tyrosine-protein phosphatas
IPI00289924	Alpha-2,8-sialyltransferase 8E
IPI00289926	Isoform 1 of Leukocyte immunoglobulin-like receptor subfar
IPI00289926	Isoform 1 of Leukocyte immunoglobulin-like receptor subfar
IPI00289926	Isoform 1 of Leukocyte immunoglobulin-like receptor subfar
IPI00290085	Neural-cadherin precursor
IPI00290085	Neural-cadherin precursor
IPI00290085	Neural-cadherin precursor
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IPI00290085	Neural-cadherin precursor
IPI00290085	Neural-cadherin precursor
IPI00290085	Neural-cadherin precursor
IPI00290283	mannan-binding lectin serine protease 1 isoform 2 precursc
IPI00290283	mannan-binding lectin serine protease 1 isoform 2 precursc
IPI00290283	mannan-binding lectin serine protease 1 isoform 2 precursc
IPI00290315	Chromogranin A precursor
IPI00290315	Chromogranin A precursor
IPI00290315	Chromogranin A precursor
IPI00290315	Chromogranin A precursor
IPI00290315	Chromogranin A precursor
IPI00290411	Immunoglobulin-like domain protein MGC33530 precursor
IPI00290411	Immunoglobulin-like domain protein MGC33530 precursor
IPI00290411	Immunoglobulin-like domain protein MGC33530 precursor
IPI00290411	Immunoglobulin-like domain protein MGC33530 precursor
IPI00290456	Intercellular adhesion molecule 5 precursor
IPI00290456	Intercellular adhesion molecule 5 precursor
IPI00290456	Intercellular adhesion molecule 5 precursor
IPI00290456	Intercellular adhesion molecule 5 precursor
IPI00290456	Intercellular adhesion molecule 5 precursor
IPI00290456	Intercellular adhesion molecule 5 precursor
IPI00290856	extracellular link domain containing 1
IPI00290856	extracellular link domain containing 1
IPI00291005	Malate dehydrogenase, cytoplasmic
IPI00291005	Malate dehydrogenase, cytoplasmic

IPI00291005	Malate dehydrogenase, cytoplasmic
IPI00291005	Malate dehydrogenase, cytoplasmic
IPI00291005	Malate dehydrogenase, cytoplasmic
IPI00291005	Malate dehydrogenase, cytoplasmic
IPI00291005	Malate dehydrogenase, cytoplasmic
IPI00291005	Malate dehydrogenase, cytoplasmic
IPI00291005	Malate dehydrogenase, cytoplasmic
IPI00291005	Malate dehydrogenase, cytoplasmic
IPI00291005	Malate dehydrogenase, cytoplasmic
IPI00291005	Malate dehydrogenase, cytoplasmic
IPI00291006	Malate dehydrogenase, mitochondrial precursor
IPI00291006	Malate dehydrogenase, mitochondrial precursor
IPI00291006	Malate dehydrogenase, mitochondrial precursor
IPI00291006	Malate dehydrogenase, mitochondrial precursor
IPI00291099	Splice Isoform Long of Receptor-type tyrosine-protein phos
IPI00291099	Splice Isoform Long of Receptor-type tyrosine-protein phos
IPI00291099	Splice Isoform Long of Receptor-type tyrosine-protein phos
IPI00291099	Splice Isoform Long of Receptor-type tyrosine-protein phos
IPI00291099	Splice Isoform Long of Receptor-type tyrosine-protein phos
IPI00291099	Splice Isoform Long of Receptor-type tyrosine-protein phos
IPI00291099	Splice Isoform Long of Receptor-type tyrosine-protein phos
IPI00291099	Splice Isoform Long of Receptor-type tyrosine-protein phos
IPI00291136	Collagen alpha-1(VI) chain precursor
IPI00291136	Collagen alpha-1(VI) chain precursor
IPI00291136	Collagen alpha-1(VI) chain precursor
IPI00291136	Collagen alpha-1(VI) chain precursor
IPI00291136	Collagen alpha-1(VI) chain precursor
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IPI00291136	Collagen alpha-1(VI) chain precursor
IPI00291136	Collagen alpha-1(VI) chain precursor
IPI00291136	Collagen alpha-1(VI) chain precursor
IPI00291136	Collagen alpha-1(VI) chain precursor
IPI00291170	Isoform 2 of Protein KIAA1199 precursor
IPI00291170	Isoform 2 of Protein KIAA1199 precursor
IPI00291262	Clusterin precursor
IPI00291262	Clusterin precursor
IPI00291262	Clusterin precursor
IPI00291262	Clusterin precursor
IPI00291262	Clusterin precursor
IPI00291262	Clusterin precursor
IPI00291262	Clusterin precursor
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IPI00291262	Clusterin precursor
IPI00291262	Clusterin precursor
IPI00291641	Mannosyl-oligosaccharide 1,2-alpha-mannosidase IA

IPI00291641	Mannosyl-oligosaccharide 1,2-alpha-mannosidase IA
IPI00291866	Plasma protease C1 inhibitor precursor
IPI00291866	Plasma protease C1 inhibitor precursor
IPI00291866	Plasma protease C1 inhibitor precursor
IPI00291866	Plasma protease C1 inhibitor precursor
IPI00291866	Plasma protease C1 inhibitor precursor
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IPI00291866	Plasma protease C1 inhibitor precursor
IPI00291866	Plasma protease C1 inhibitor precursor
IPI00291867	Complement factor I precursor
IPI00291867	Complement factor I precursor
IPI00291867	Complement factor I precursor
IPI00291867	Complement factor I precursor
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IPI00291867	Complement factor I precursor
IPI00291867	Complement factor I precursor
IPI00291867	Complement factor I precursor
IPI00292071	Secretogranin-3 precursor
IPI00292071	Secretogranin-3 precursor
IPI00292071	Secretogranin-3 precursor
IPI00292071	Secretogranin-3 precursor
IPI00292071	Secretogranin-3 precursor
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IPI00292071	Secretogranin-3 precursor
IPI00292071	Secretogranin-3 precursor
IPI00292130	Dermatopontin precursor
IPI00292130	Dermatopontin precursor
IPI00292150	Latent-transforming growth factor beta-binding protein 2 pre
IPI00292150	Latent-transforming growth factor beta-binding protein 2 pre
IPI00292150	Latent-transforming growth factor beta-binding protein 2 pre

IPI00292150	Latent-transforming growth factor beta-binding protein 2 pre
IPI00292150	Latent-transforming growth factor beta-binding protein 2 pre
IPI00292150	Latent-transforming growth factor beta-binding protein 2 pre
IPI00292300	contactin associated protein-like 5 isoform 1
IPI00292300	contactin associated protein-like 5 isoform 1
IPI00292530	Inter-alpha-trypsin inhibitor heavy chain H1 precursor
IPI00292530	Inter-alpha-trypsin inhibitor heavy chain H1 precursor
IPI00292530	Inter-alpha-trypsin inhibitor heavy chain H1 precursor
IPI00292530	Inter-alpha-trypsin inhibitor heavy chain H1 precursor
IPI00292530	Inter-alpha-trypsin inhibitor heavy chain H1 precursor
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IPI00292530	Inter-alpha-trypsin inhibitor heavy chain H1 precursor
IPI00292530	Inter-alpha-trypsin inhibitor heavy chain H1 precursor
IPI00292530	Inter-alpha-trypsin inhibitor heavy chain H1 precursor
IPI00292791	Contactin-3 precursor
IPI00292791	Contactin-3 precursor
IPI00292946	Thyroxine-binding globulin precursor
IPI00292946	Thyroxine-binding globulin precursor
IPI00292946	Thyroxine-binding globulin precursor
IPI00292946	Thyroxine-binding globulin precursor
IPI00292946	Thyroxine-binding globulin precursor
IPI00292946	Thyroxine-binding globulin precursor
IPI00292950	Heparin cofactor 2 precursor
IPI00292950	Heparin cofactor 2 precursor
IPI00292950	Heparin cofactor 2 precursor
IPI00292950	Heparin cofactor 2 precursor
IPI00292950	Heparin cofactor 2 precursor
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IPI00292950	Heparin cofactor 2 precursor
IPI00292950	Heparin cofactor 2 precursor
IPI00292950	Heparin cofactor 2 precursor
IPI00293088	Lysosomal alpha-glucosidase precursor
IPI00293088	Lysosomal alpha-glucosidase precursor
IPI00293088	Lysosomal alpha-glucosidase precursor
IPI00293276	Macrophage migration inhibitory factor
IPI00293276	Macrophage migration inhibitory factor
IPI00293276	Macrophage migration inhibitory factor
IPI00293748	Isoform 1 of Multiple inositol polyphosphate phosphatase 1

IPI00293748	Isoform 1 of Multiple inositol polyphosphate phosphatase 1
IPI00293748	Isoform 1 of Multiple inositol polyphosphate phosphatase 1
IPI00293757	Isoform 1 of Netrin receptor UNC5C precursor
IPI00293836	immunoglobulin superfamily, member 4D
IPI00293836	immunoglobulin superfamily, member 4D
IPI00293836	immunoglobulin superfamily, member 4D
IPI00293849	Receptor-type tyrosine-protein phosphatase mu precursor
IPI00294004	Vitamin K-dependent protein S precursor
IPI00294004	Vitamin K-dependent protein S precursor
IPI00294004	Vitamin K-dependent protein S precursor
IPI00294004	Vitamin K-dependent protein S precursor
IPI00294004	Vitamin K-dependent protein S precursor
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IPI00294004	Vitamin K-dependent protein S precursor
IPI00294004	Vitamin K-dependent protein S precursor
IPI00294004	Vitamin K-dependent protein S precursor
IPI00294395	Complement component C8 beta chain precursor
IPI00294395	Complement component C8 beta chain precursor
IPI00294395	Complement component C8 beta chain precursor
IPI00294395	Complement component C8 beta chain precursor
IPI00294395	Complement component C8 beta chain precursor
IPI00294395	Complement component C8 beta chain precursor
IPI00294615	Fibulin-5 precursor
IPI00294615	Fibulin-5 precursor
IPI00294615	Fibulin-5 precursor
IPI00294615	Fibulin-5 precursor
IPI00294650	Secreted frizzled-related protein 3 precursor
IPI00294650	Secreted frizzled-related protein 3 precursor
IPI00294650	Secreted frizzled-related protein 3 precursor
IPI00294705	Hypothetical protein
IPI00294705	Hypothetical protein
IPI00295386	Carbonyl reductase [NADPH] 1
IPI00295386	Carbonyl reductase [NADPH] 1
IPI00295386	Carbonyl reductase [NADPH] 1
IPI00295414	Collagen alpha-1(XV) chain precursor
IPI00295414	Collagen alpha-1(XV) chain precursor
IPI00295414	Collagen alpha-1(XV) chain precursor
IPI00295414	Collagen alpha-1(XV) chain precursor
IPI00295414	Collagen alpha-1(XV) chain precursor
IPI00295542	Nucleobindin-1 precursor
IPI00295542	Nucleobindin-1 precursor
IPI00295542	Nucleobindin-1 precursor
IPI00295741	Cathepsin B precursor
IPI00295741	Cathepsin B precursor
IPI00295741	Cathepsin B precursor
IPI00295741	Cathepsin B precursor
IPI00295741	Cathepsin B precursor
IPI00295832	Oligodendrocyte-myelin glycoprotein precursor
IPI00295832	Oligodendrocyte-myelin glycoprotein precursor

IPI00295832	Oligodendrocyte-myelin glycoprotein precursor
IPI00296058	EGF-containing fibulin-like extracellular matrix protein 2 pre
IPI00296058	EGF-containing fibulin-like extracellular matrix protein 2 pre
IPI00296141	Dipeptidyl-peptidase 2 precursor
IPI00296141	Dipeptidyl-peptidase 2 precursor
IPI00296165	Complement C1r subcomponent precursor
IPI00296165	Complement C1r subcomponent precursor
IPI00296165	Complement C1r subcomponent precursor
IPI00296165	Complement C1r subcomponent precursor
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IPI00296165	Complement C1r subcomponent precursor
IPI00296165	Complement C1r subcomponent precursor
IPI00296165	Complement C1r subcomponent precursor
IPI00296259	Transmembrane emp24 domain-containing protein 4 precu
IPI00296259	Transmembrane emp24 domain-containing protein 4 precu
IPI00296534	Isoform D of Fibulin-1 precursor
IPI00296534	Isoform D of Fibulin-1 precursor
IPI00296537	Isoform C of Fibulin-1 precursor
IPI00296537	Isoform C of Fibulin-1 precursor
IPI00296537	Isoform C of Fibulin-1 precursor
IPI00296537	Isoform C of Fibulin-1 precursor
IPI00296537	Isoform C of Fibulin-1 precursor
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IPI00296537	Isoform C of Fibulin-1 precursor
IPI00296537	Isoform C of Fibulin-1 precursor
IPI00296537	Isoform C of Fibulin-1 precursor
IPI00296537	Isoform C of Fibulin-1 precursor
IPI00296537	Isoform C of Fibulin-1 precursor

IPI00296537	Isoform C of Fibulin-1 precursor
IPI00296537	Isoform C of Fibulin-1 precursor
IPI00296608	Complement component C7 precursor
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IPI00296608	Complement component C7 precursor
IPI00296777	SPARC-like protein 1 precursor
IPI00296777	SPARC-like protein 1 precursor
IPI00296777	SPARC-like protein 1 precursor
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IPI00296777	SPARC-like protein 1 precursor
IPI00296777	SPARC-like protein 1 precursor
IPI00296922	Laminin beta-2 chain precursor
IPI00296922	Laminin beta-2 chain precursor
IPI00296922	Laminin beta-2 chain precursor

IPI00296922	Laminin beta-2 chain precursor
IPI00296922	Laminin beta-2 chain precursor
IPI00296922	Laminin beta-2 chain precursor
IPI00296992	AXL receptor tyrosine kinase isoform 1
IPI00296992	AXL receptor tyrosine kinase isoform 1
IPI00296992	AXL receptor tyrosine kinase isoform 1
IPI00297124	Isoform 1 of Interleukin-6 receptor beta chain precursor
IPI00297124	Isoform 1 of Interleukin-6 receptor beta chain precursor
IPI00297124	Isoform 1 of Interleukin-6 receptor beta chain precursor
IPI00297124	Isoform 1 of Interleukin-6 receptor beta chain precursor
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IPI00297124	Isoform 1 of Interleukin-6 receptor beta chain precursor
IPI00297124	Isoform 1 of Interleukin-6 receptor beta chain precursor
IPI00297124	Isoform 1 of Interleukin-6 receptor beta chain precursor
IPI00297160	CD44 antigen isoform 4 precursor
IPI00297160	CD44 antigen isoform 4 precursor
IPI00297188	Brain-specific angiogenesis inhibitor 2 precursor
IPI00297188	Brain-specific angiogenesis inhibitor 2 precursor
IPI00297224	PREDICTED: hypothetical protein LOC26032
IPI00297224	PREDICTED: hypothetical protein LOC26032
IPI00297263	PREDICTED: HEG homolog 1
IPI00297263	PREDICTED: HEG homolog 1
IPI00297284	Insulin-like growth factor-binding protein 2 precursor
IPI00297284	Insulin-like growth factor-binding protein 2 precursor
IPI00297284	Insulin-like growth factor-binding protein 2 precursor
IPI00297284	Insulin-like growth factor-binding protein 2 precursor
IPI00297284	Insulin-like growth factor-binding protein 2 precursor
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IPI00297284	Insulin-like growth factor-binding protein 2 precursor
IPI00297284	Insulin-like growth factor-binding protein 2 precursor
IPI00297284	Insulin-like growth factor-binding protein 2 precursor
IPI00297487	Cathepsin H precursor
IPI00297487	Cathepsin H precursor
IPI00297487	Cathepsin H precursor
IPI00297646	Collagen alpha-1(I) chain precursor
IPI00297646	Collagen alpha-1(I) chain precursor
IPI00297646	Collagen alpha-1(I) chain precursor
IPI00297646	Collagen alpha-1(I) chain precursor
IPI00297646	Collagen alpha-1(I) chain precursor
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IPI00297646	Collagen alpha-1(I) chain precursor
IPI00297646	Collagen alpha-1(I) chain precursor
IPI00297646	Collagen alpha-1(I) chain precursor
IPI00298237	Isoform 1 of Tripeptidyl-peptidase 1 precursor
IPI00298237	Isoform 1 of Tripeptidyl-peptidase 1 precursor
IPI00298237	Isoform 1 of Tripeptidyl-peptidase 1 precursor

IPI00298237	Isoform 1 of Tripeptidyl-peptidase 1 precursor
IPI00298237	Isoform 1 of Tripeptidyl-peptidase 1 precursor
IPI00298281	Laminin gamma-1 chain precursor
IPI00298281	Laminin gamma-1 chain precursor
IPI00298281	Laminin gamma-1 chain precursor
IPI00298281	Laminin gamma-1 chain precursor
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IPI00298281	Laminin gamma-1 chain precursor
IPI00298281	Laminin gamma-1 chain precursor
IPI00298388	HGFL protein
IPI00298388	HGFL protein
IPI00298388	HGFL protein
IPI00298497	Fibrinogen beta chain precursor
IPI00298497	Fibrinogen beta chain precursor
IPI00298497	Fibrinogen beta chain precursor
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IPI00298497	Fibrinogen beta chain precursor
IPI00298497	Fibrinogen beta chain precursor
IPI00298547	Protein DJ-1
IPI00298547	Protein DJ-1
IPI00298547	Protein DJ-1
IPI00298547	Protein DJ-1
IPI00298547	Protein DJ-1
IPI00298547	Protein DJ-1
IPI00298547	Protein DJ-1
IPI00298547	Protein DJ-1
IPI00298547	Protein DJ-1
IPI00298793	Beta-mannosidase precursor
IPI00298793	Beta-mannosidase precursor
IPI00298793	Beta-mannosidase precursor
IPI00298793	Beta-mannosidase precursor
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IPI00298793	Beta-mannosidase precursor
IPI00298793	Beta-mannosidase precursor
IPI00298828	Beta-2-glycoprotein 1 precursor
IPI00298828	Beta-2-glycoprotein 1 precursor
IPI00298828	Beta-2-glycoprotein 1 precursor
IPI00298828	Beta-2-glycoprotein 1 precursor
IPI00298828	Beta-2-glycoprotein 1 precursor
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IPI00298828	Beta-2-glycoprotein 1 precursor
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IPI00298828	Beta-2-glycoprotein 1 precursor
IPI00298860	Growth-inhibiting protein 12
IPI00298860	Growth-inhibiting protein 12
IPI00298860	Growth-inhibiting protein 12
IPI00298860	Growth-inhibiting protein 12
IPI00298860	Growth-inhibiting protein 12
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IPI00298860	Growth-inhibiting protein 12
IPI00298860	Growth-inhibiting protein 12
IPI00298956	Hypothetical protein DKFZp686E04229
IPI00298956	Hypothetical protein DKFZp686E04229
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IPI00298956	Hypothetical protein DKFZp686E04229
IPI00298971	Vitronectin precursor
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IPI00298971	Vitronectin precursor
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IPI00298971	Vitronectin precursor
IPI00298994	Talin-1
IPI00298994	Talin-1
IPI00299026	Alpha-L-fucosidase
IPI00299059	cell adhesion molecule with homology to L1CAM precursor
IPI00299059	cell adhesion molecule with homology to L1CAM precursor
IPI00299059	cell adhesion molecule with homology to L1CAM precursor
IPI00299059	cell adhesion molecule with homology to L1CAM precursor
IPI00299059	cell adhesion molecule with homology to L1CAM precursor
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IPI00299059	cell adhesion molecule with homology to L1CAM precursor
IPI00299059	cell adhesion molecule with homology to L1CAM precursor
IPI00299083	Junctional adhesion molecule B precursor
IPI00299083	Junctional adhesion molecule B precursor
IPI00299150	Cathepsin S precursor
IPI00299503	Phosphatidylinositol-glycan-specific phospholipase D 1 prec
IPI00299503	Phosphatidylinositol-glycan-specific phospholipase D 1 prec
IPI00299503	Phosphatidylinositol-glycan-specific phospholipase D 1 prec
IPI00299503	Phosphatidylinositol-glycan-specific phospholipase D 1 prec
IPI00299503	Phosphatidylinositol-glycan-specific phospholipase D 1 prec
IPI00299503	Phosphatidylinositol-glycan-specific phospholipase D 1 prec
IPI00299503	Phosphatidylinositol-glycan-specific phospholipase D 1 prec
IPI00299503	Phosphatidylinositol-glycan-specific phospholipase D 1 prec
IPI00299547	Neutrophil gelatinase-associated lipocalin precursor
IPI00299547	Neutrophil gelatinase-associated lipocalin precursor
IPI00299669	Mannosyl-oligosaccharide 1,2-alpha-mannosidase IC
IPI00299699	Neural proliferation differentiation and control protein 1 prec
IPI00299699	Neural proliferation differentiation and control protein 1 prec
IPI00299738	Procollagen C-endopeptidase enhancer 1 precursor
IPI00299738	Procollagen C-endopeptidase enhancer 1 precursor
IPI00299738	Procollagen C-endopeptidase enhancer 1 precursor
IPI00299738	Procollagen C-endopeptidase enhancer 1 precursor
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IPI00299738	Procollagen C-endopeptidase enhancer 1 precursor
IPI00300241	Leucine-rich repeat-containing protein 4B precursor
IPI00300241	Leucine-rich repeat-containing protein 4B precursor
IPI00300241	Leucine-rich repeat-containing protein 4B precursor
IPI00300241	Leucine-rich repeat-containing protein 4B precursor
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IPI00300241	Leucine-rich repeat-containing protein 4B precursor
IPI00300241	Leucine-rich repeat-containing protein 4B precursor
IPI00300376	Protein-glutamine gamma-glutamyltransferase E precursor
IPI00301143	protease inhibitor 16 precursor
IPI00301143	protease inhibitor 16 precursor
IPI00301255	Immunoglobulin superfamily member 21 precursor
IPI00301255	Immunoglobulin superfamily member 21 precursor
IPI00301459	1-O-acylceramide synthase precursor
IPI00301459	1-O-acylceramide synthase precursor
IPI00301579	Epididymal secretory protein E1 precursor
IPI00301579	Epididymal secretory protein E1 precursor
IPI00301579	Epididymal secretory protein E1 precursor
IPI00301579	Epididymal secretory protein E1 precursor
IPI00301579	Epididymal secretory protein E1 precursor
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IPI00301579	Epididymal secretory protein E1 precursor
IPI00301579	Epididymal secretory protein E1 precursor
IPI00301865	transmembrane protein 132A isoform b
IPI00301865	transmembrane protein 132A isoform b
IPI00301865	transmembrane protein 132A isoform b
IPI00301865	transmembrane protein 132A isoform b
IPI00301865	transmembrane protein 132A isoform b
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IPI00301865	transmembrane protein 132A isoform b
IPI00301865	transmembrane protein 132A isoform b
IPI00302181	calcium channel, voltage-dependent, alpha 2/delta 3 subun
IPI00302181	calcium channel, voltage-dependent, alpha 2/delta 3 subun
IPI00302181	calcium channel, voltage-dependent, alpha 2/delta 3 subun
IPI00302181	calcium channel, voltage-dependent, alpha 2/delta 3 subun
IPI00302181	calcium channel, voltage-dependent, alpha 2/delta 3 subun
IPI00302181	calcium channel, voltage-dependent, alpha 2/delta 3 subun
IPI00302592	Filamin A, alpha
IPI00302592	Filamin A, alpha
IPI00302592	Filamin A, alpha
IPI00302592	Filamin A, alpha
IPI00302592	Filamin A, alpha
IPI00302592	Filamin A, alpha
IPI00302592	Filamin A, alpha
IPI00302641	Protocadherin Fat 2 precursor
IPI00302641	Protocadherin Fat 2 precursor
IPI00302641	Protocadherin Fat 2 precursor
IPI00302641	Protocadherin Fat 2 precursor

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IPI00302641	Protocadherin Fat 2 precursor
IPI00302641	Protocadherin Fat 2 precursor
IPI00302641	Protocadherin Fat 2 precursor
IPI00302840	Sodium/potassium-transporting ATPase alpha-3 chain
IPI00303161	Endothelial cell-selective adhesion molecule precursor
IPI00303161	Endothelial cell-selective adhesion molecule precursor
IPI00303318	Protein FAM49B
IPI00303318	Protein FAM49B
IPI00303963	Complement C2 precursor (Fragment)
IPI00303963	Complement C2 precursor (Fragment)
IPI00303963	Complement C2 precursor (Fragment)
IPI00303963	Complement C2 precursor (Fragment)
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IPI00303963	Complement C2 precursor (Fragment)
IPI00303963	Complement C2 precursor (Fragment)
IPI00304273	Apolipoprotein A-IV precursor
IPI00304273	Apolipoprotein A-IV precursor
IPI00304273	Apolipoprotein A-IV precursor
IPI00304273	Apolipoprotein A-IV precursor
IPI00304273	Apolipoprotein A-IV precursor
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IPI00304273	Apolipoprotein A-IV precursor
IPI00304273	Apolipoprotein A-IV precursor
IPI00304273	Apolipoprotein A-IV precursor
IPI00304865	transforming growth factor, beta receptor III
IPI00304865	transforming growth factor, beta receptor III
IPI00304865	transforming growth factor, beta receptor III
IPI00304865	transforming growth factor, beta receptor III

IPI00304865	transforming growth factor, beta receptor III
IPI00305380	Insulin-like growth factor-binding protein 4 precursor
IPI00305380	Insulin-like growth factor-binding protein 4 precursor
IPI00305380	Insulin-like growth factor-binding protein 4 precursor
IPI00305461	Inter-alpha-trypsin inhibitor heavy chain H2 precursor
IPI00305461	Inter-alpha-trypsin inhibitor heavy chain H2 precursor
IPI00305461	Inter-alpha-trypsin inhibitor heavy chain H2 precursor
IPI00305461	Inter-alpha-trypsin inhibitor heavy chain H2 precursor
IPI00305461	Inter-alpha-trypsin inhibitor heavy chain H2 precursor
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IPI00305461	Inter-alpha-trypsin inhibitor heavy chain H2 precursor
IPI00305719	selenium binding protein 1
IPI00305719	selenium binding protein 1
IPI00305719	selenium binding protein 1
IPI00305719	selenium binding protein 1
IPI00305719	selenium binding protein 1
IPI00305719	selenium binding protein 1
IPI00307276	ADAMTS-4
IPI00328113	Fibrillin-1 precursor
IPI00328113	Fibrillin-1 precursor
IPI00328243	Phospholipase D3, isoform 1
IPI00328243	Phospholipase D3, isoform 1
IPI00328243	Phospholipase D3, isoform 1
IPI00328391	N-acetylgalactosaminyltransferase 7
IPI00328550	Thrombospondin-4 precursor
IPI00328550	Thrombospondin-4 precursor
IPI00328550	Thrombospondin-4 precursor
IPI00328550	Thrombospondin-4 precursor
IPI00328609	Kallistatin precursor
IPI00328609	Kallistatin precursor
IPI00328609	Kallistatin precursor
IPI00328609	Kallistatin precursor
IPI00328609	Kallistatin precursor

IPI00328609	Kallistatin precursor
IPI00328609	Kallistatin precursor
IPI00328609	Kallistatin precursor
IPI00328609	Kallistatin precursor
IPI00328609	Kallistatin precursor
IPI00328703	Hypothetical protein OAF
IPI00328703	Hypothetical protein OAF
IPI00328703	Hypothetical protein OAF
IPI00328703	Hypothetical protein OAF
IPI00328703	Hypothetical protein OAF
IPI00328703	Hypothetical protein OAF
IPI00328703	Hypothetical protein OAF
IPI00328703	Hypothetical protein OAF
IPI00328746	Reticulon-4 receptor-like 2 precursor
IPI00328746	Reticulon-4 receptor-like 2 precursor
IPI00328746	Reticulon-4 receptor-like 2 precursor
IPI00328829	inter-alpha trypsin inhibitor heavy chain precursor 5 isoform
IPI00328829	inter-alpha trypsin inhibitor heavy chain precursor 5 isoform
IPI00328829	inter-alpha trypsin inhibitor heavy chain precursor 5 isoform
IPI00328829	inter-alpha trypsin inhibitor heavy chain precursor 5 isoform
IPI00328829	inter-alpha trypsin inhibitor heavy chain precursor 5 isoform
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IPI00328829	inter-alpha trypsin inhibitor heavy chain precursor 5 isoform
IPI00328829	inter-alpha trypsin inhibitor heavy chain precursor 5 isoform
IPI00328829	inter-alpha trypsin inhibitor heavy chain precursor 5 isoform
IPI00329332	Syntaxin-12
IPI00329332	Syntaxin-12
IPI00329482	Isoform 1 of Laminin alpha-4 chain precursor
IPI00329482	Isoform 1 of Laminin alpha-4 chain precursor
IPI00329482	Isoform 1 of Laminin alpha-4 chain precursor
IPI00329775	Isoform 1 of Carboxypeptidase B2 precursor
IPI00329775	Isoform 1 of Carboxypeptidase B2 precursor
IPI00329801	Annexin A5
IPI00329801	Annexin A5
IPI00332887	signal-regulatory protein alpha precursor
IPI00332887	signal-regulatory protein alpha precursor
IPI00332887	signal-regulatory protein alpha precursor
IPI00332887	signal-regulatory protein alpha precursor
IPI00332887	signal-regulatory protein alpha precursor
IPI00332887	signal-regulatory protein alpha precursor
IPI00332887	signal-regulatory protein alpha precursor
IPI00332887	signal-regulatory protein alpha precursor
IPI00332887	signal-regulatory protein alpha precursor
IPI00332887	signal-regulatory protein alpha precursor
IPI00333140	Delta-notch-like EGF repeat-containing transmembrane
IPI00333140	Delta-notch-like EGF repeat-containing transmembrane
IPI00333776	Isoform 1 of Neuronal cell adhesion molecule precursor
IPI00333776	Isoform 1 of Neuronal cell adhesion molecule precursor
IPI00333776	Isoform 1 of Neuronal cell adhesion molecule precursor

IPI00333776	Isoform 1 of Neuronal cell adhesion molecule precursor
IPI00333776	Isoform 1 of Neuronal cell adhesion molecule precursor
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IPI00333776	Isoform 1 of Neuronal cell adhesion molecule precursor
IPI00333776	Isoform 1 of Neuronal cell adhesion molecule precursor
IPI00334238	neuronal pentraxin receptor isoform 1
IPI00334238	neuronal pentraxin receptor isoform 1
IPI00334238	neuronal pentraxin receptor isoform 1
IPI00334238	neuronal pentraxin receptor isoform 1
IPI00334238	neuronal pentraxin receptor isoform 1
IPI00334238	neuronal pentraxin receptor isoform 1
IPI00334238	neuronal pentraxin receptor isoform 1
IPI00334238	neuronal pentraxin receptor isoform 1
IPI00335946	KIAA1838 protein
IPI00337351	MAM domain-containing protein 1 precursor
IPI00337351	MAM domain-containing protein 1 precursor
IPI00337548	Cell growth regulator with EF hand domain 1
IPI00337548	Cell growth regulator with EF hand domain 1
IPI00337548	Cell growth regulator with EF hand domain 1
IPI00339225	Isoform 5 of Fibronectin precursor
IPI00339225	Isoform 5 of Fibronectin precursor
IPI00339225	Isoform 5 of Fibronectin precursor

IPI00339225	Isoform 5 of Fibronectin precursor
IPI00339225	Isoform 5 of Fibronectin precursor
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IPI00339225	Isoform 5 of Fibronectin precursor
IPI00339225	Isoform 5 of Fibronectin precursor
IPI00339225	Isoform 5 of Fibronectin precursor
IPI00374563	Agrin precursor
IPI00374563	Agrin precursor
IPI00374563	Agrin precursor
IPI00374563	Agrin precursor
IPI00374563	Agrin precursor
IPI00374563	Agrin precursor
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IPI00374563	Agrin precursor
IPI00374563	Agrin precursor
IPI00374563	Agrin precursor
IPI00374563	Agrin precursor
IPI00374914	PREDICTED: hypothetical protein XP_379250
IPI00375364	Isoform 3 of Chitotriosidase-1 precursor
IPI00375364	Isoform 3 of Chitotriosidase-1 precursor
IPI00375676	Ferritin light polypeptide variant
IPI00375676	Ferritin light polypeptide variant
IPI00376394	Similar to quiescin Q6-like 1
IPI00376394	Similar to quiescin Q6-like 1
IPI00376394	Similar to quiescin Q6-like 1
IPI00376394	Similar to quiescin Q6-like 1
IPI00376394	Similar to quiescin Q6-like 1
IPI00376427	Neural cell adhesion molecule 2 precursor
IPI00376427	Neural cell adhesion molecule 2 precursor
IPI00376427	Neural cell adhesion molecule 2 precursor
IPI00376427	Neural cell adhesion molecule 2 precursor
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IPI00376427	Neural cell adhesion molecule 2 precursor
IPI00376427	Neural cell adhesion molecule 2 precursor
IPI00376427	Neural cell adhesion molecule 2 precursor
IPI00382428	Full-length cDNA clone CS0DI085YI08 of Placenta of Hom
IPI00382428	Full-length cDNA clone CS0DI085YI08 of Placenta of Hom
IPI00382428	Full-length cDNA clone CS0DI085YI08 of Placenta of Hom
IPI00382436	Ig lambda chain V-III region SH
IPI00382436	Ig lambda chain V-III region SH
IPI00382436	Ig lambda chain V-III region SH
IPI00382440	Ig lambda chain V-IV region Hil
IPI00382470	heat shock 90kDa protein 1, alpha isoform 1
IPI00382478	Ig heavy chain V-III region TIL
IPI00382481	Ig heavy chain V-III region BUT
IPI00382481	Ig heavy chain V-III region BUT
IPI00382483	Ig heavy chain V-III region GA
IPI00382483	Ig heavy chain V-III region GA
IPI00382483	Ig heavy chain V-III region GA
IPI00382500	Ig heavy chain V-III region GAL
IPI00382539	Ig heavy chain V-II region WAH
IPI00382950	Beta-globin gene from a thalassemia patient, complete cds
IPI00382950	Beta-globin gene from a thalassemia patient, complete cds
IPI00383814	transmembrane protein 132A isoform a
IPI00383814	transmembrane protein 132A isoform a
IPI00383814	transmembrane protein 132A isoform a
IPI00383814	transmembrane protein 132A isoform a
IPI00383814	transmembrane protein 132A isoform a
IPI00383814	transmembrane protein 132A isoform a
IPI00383975	Full-length cDNA clone CS0DN001YP04 of Adult brain of H
IPI00383975	Full-length cDNA clone CS0DN001YP04 of Adult brain of H
IPI00383975	Full-length cDNA clone CS0DN001YP04 of Adult brain of H
IPI00384293	SPARC-like protein 1 precursor
IPI00384293	SPARC-like protein 1 precursor
IPI00384293	SPARC-like protein 1 precursor
IPI00384293	SPARC-like protein 1 precursor
IPI00384293	SPARC-like protein 1 precursor
IPI00384293	SPARC-like protein 1 precursor
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IPI00384293	SPARC-like protein 1 precursor
IPI00384293	SPARC-like protein 1 precursor
IPI00384293	SPARC-like protein 1 precursor
IPI00384293	SPARC-like protein 1 precursor
IPI00384400	Myosin-reactive immunoglobulin heavy chain variable regio
IPI00384401	Myosin-reactive immunoglobulin kappa chain variable regio
IPI00384406	Myosin-reactive immunoglobulin heavy chain variable regio
IPI00384542	Isoform 2 of Nidogen-1 precursor
IPI00384542	Isoform 2 of Nidogen-1 precursor
IPI00384697	ALB protein
IPI00384697	ALB protein
IPI00384998	Isoform 7 of Neurofascin precursor

IPI00384998	Isoform 7 of Neurofascin precursor
IPI00384998	Isoform 7 of Neurofascin precursor
IPI00384998	Isoform 7 of Neurofascin precursor
IPI00384998	Isoform 7 of Neurofascin precursor
IPI00384998	Isoform 7 of Neurofascin precursor
IPI00384998	Isoform 7 of Neurofascin precursor
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IPI00384998	Isoform 7 of Neurofascin precursor
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IPI00384998	Isoform 7 of Neurofascin precursor
IPI00384998	Isoform 7 of Neurofascin precursor
IPI00384998	Isoform 7 of Neurofascin precursor
IPI00384998	Isoform 7 of Neurofascin precursor
IPI00384998	Isoform 7 of Neurofascin precursor
IPI00384998	Isoform 7 of Neurofascin precursor
IPI00384998	Isoform 7 of Neurofascin precursor
IPI00384998	Isoform 7 of Neurofascin precursor
IPI00385252	Ig kappa chain V-III region GOL
IPI00385252	Ig kappa chain V-III region GOL
IPI00385555	Ig kappa chain V-I region BAN
IPI00385980	ROBO2 isoform a
IPI00385980	ROBO2 isoform a
IPI00385980	ROBO2 isoform a
IPI00385980	ROBO2 isoform a
IPI00386142	Ig heavy chain V-II region ARH-77 precursor
IPI00386575	Ig lambda chain V-I region EPS
IPI00386630	TCN2 protein
IPI00386630	TCN2 protein
IPI00386630	TCN2 protein
IPI00386630	TCN2 protein
IPI00386630	TCN2 protein
IPI00387025	Ig kappa chain V-I region DEE
IPI00387025	Ig kappa chain V-I region DEE
IPI00387025	Ig kappa chain V-I region DEE
IPI00387025	Ig kappa chain V-I region DEE
IPI00387113	Ig kappa chain V-III region B6
IPI00387113	Ig kappa chain V-III region B6
IPI00387113	Ig kappa chain V-III region B6
IPI00387113	Ig kappa chain V-III region B6
IPI00387115	Ig kappa chain V-III region SIE
IPI00387115	Ig kappa chain V-III region SIE
IPI00387120	Ig kappa chain V-IV region Len
IPI00387120	Ig kappa chain V-IV region Len
IPI00387120	Ig kappa chain V-IV region Len
IPI00394655	Isoform 4 of Neurofascin precursor
IPI00394655	Isoform 4 of Neurofascin precursor
IPI00394655	Isoform 4 of Neurofascin precursor
IPI00394655	Isoform 4 of Neurofascin precursor
IPI00394655	Isoform 4 of Neurofascin precursor
IPI00395488	Vasorin precursor
IPI00395488	Vasorin precursor
IPI00395488	Vasorin precursor
IPI00395488	Vasorin precursor
IPI00395488	Vasorin precursor

IPI00396423	Alcadein beta
IPI00396423	Alcadein beta
IPI00396423	Alcadein beta
IPI00397828	20 kDa protein
IPI00397828	20 kDa protein
IPI00398435	PREDICTED: plexin B2
IPI00398435	PREDICTED: plexin B2
IPI00398435	PREDICTED: plexin B2
IPI00398435	PREDICTED: plexin B2
IPI00398435	PREDICTED: plexin B2
IPI00398435	PREDICTED: plexin B2
IPI00398435	PREDICTED: plexin B2
IPI00398435	PREDICTED: plexin B2
IPI00398435	PREDICTED: plexin B2
IPI00398435	PREDICTED: plexin B2
IPI00399007	Hypothetical protein DKFZp686I04196 (Fragment)
IPI00399007	Hypothetical protein DKFZp686I04196 (Fragment)
IPI00399007	Hypothetical protein DKFZp686I04196 (Fragment)
IPI00399007	Hypothetical protein DKFZp686I04196 (Fragment)
IPI00399007	Hypothetical protein DKFZp686I04196 (Fragment)
IPI00399007	Hypothetical protein DKFZp686I04196 (Fragment)
IPI00399007	Hypothetical protein DKFZp686I04196 (Fragment)
IPI00401283	Multiple epidermal growth factor-like domains 9 precursor
IPI00401283	Multiple epidermal growth factor-like domains 9 precursor
IPI00402157	Cerebellin-3 precursor
IPI00402157	Cerebellin-3 precursor
IPI00402157	Cerebellin-3 precursor
IPI00402157	Cerebellin-3 precursor
IPI00410312	Isoform 3 of Latrophilin-3 precursor
IPI00410312	Isoform 3 of Latrophilin-3 precursor
IPI00410312	Isoform 3 of Latrophilin-3 precursor
IPI00410312	Isoform 3 of Latrophilin-3 precursor
IPI00410312	Isoform 3 of Latrophilin-3 precursor
IPI00410600	Calcium channel, alpha 2/delta subunit 2
IPI00410600	Calcium channel, alpha 2/delta subunit 2
IPI00410600	Calcium channel, alpha 2/delta subunit 2
IPI00410600	Calcium channel, alpha 2/delta subunit 2
IPI00410600	Calcium channel, alpha 2/delta subunit 2
IPI00410600	Calcium channel, alpha 2/delta subunit 2
IPI00410714	Hemoglobin subunit alpha
IPI00410714	Hemoglobin subunit alpha
IPI00410714	Hemoglobin subunit alpha
IPI00410714	Hemoglobin subunit alpha
IPI00410714	Hemoglobin subunit alpha
IPI00410714	Hemoglobin subunit alpha
IPI00410714	Hemoglobin subunit alpha
IPI00410714	Hemoglobin subunit alpha
IPI00410714	Hemoglobin subunit alpha
IPI00410714	Hemoglobin subunit alpha
IPI00412264	Pleiotrophin precursor
IPI00412264	Pleiotrophin precursor
IPI00412264	Pleiotrophin precursor
IPI00412264	Pleiotrophin precursor
IPI00412987	GMFB protein
IPI00412987	GMFB protein
IPI00412987	GMFB protein

IPI00412988	Isoform 1 of Netrin G1 precursor
IPI00412988	Isoform 1 of Netrin G1 precursor
IPI00413273	Nebulette
IPI00413344	Isoform CFL2b of Cofilin-2
IPI00413344	Isoform CFL2b of Cofilin-2
IPI00413451	Hypothetical protein DKFZp686I04222
IPI00413451	Hypothetical protein DKFZp686I04222
IPI00413451	Hypothetical protein DKFZp686I04222
IPI00413451	Hypothetical protein DKFZp686I04222
IPI00413451	Hypothetical protein DKFZp686I04222
IPI00413641	Aldose reductase
IPI00413641	Aldose reductase
IPI00413641	Aldose reductase
IPI00413778	FK506 binding protein12
IPI00413959	Calsyntenin-1 precursor
IPI00413959	Calsyntenin-1 precursor
IPI00413959	Calsyntenin-1 precursor
IPI00413959	Calsyntenin-1 precursor
IPI00413959	Calsyntenin-1 precursor
IPI00413959	Calsyntenin-1 precursor
IPI00413959	Calsyntenin-1 precursor
IPI00413959	Calsyntenin-1 precursor
IPI00414283	fibronectin 1 isoform 4 preproprotein
IPI00414283	fibronectin 1 isoform 4 preproprotein
IPI00414283	fibronectin 1 isoform 4 preproprotein
IPI00414283	fibronectin 1 isoform 4 preproprotein
IPI00414283	fibronectin 1 isoform 4 preproprotein
IPI00414283	fibronectin 1 isoform 4 preproprotein
IPI00414283	fibronectin 1 isoform 4 preproprotein
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IPI00414283	fibronectin 1 isoform 4 preproprotein
IPI00414283	fibronectin 1 isoform 4 preproprotein
IPI00414283	fibronectin 1 isoform 4 preproprotein
IPI00414896	Isoform 1 of Ribonuclease T2 precursor
IPI00414896	Isoform 1 of Ribonuclease T2 precursor
IPI00414896	Isoform 1 of Ribonuclease T2 precursor
IPI00414896	Isoform 1 of Ribonuclease T2 precursor
IPI00414896	Isoform 1 of Ribonuclease T2 precursor
IPI00414984	Isoform SGCE-1 of Epsilon-sarcoglycan precursor
IPI00414984	Isoform SGCE-1 of Epsilon-sarcoglycan precursor
IPI00414984	Isoform SGCE-1 of Epsilon-sarcoglycan precursor

IPI00414984	Isoform SGCE-1 of Epsilon-sarcoglycan precursor
IPI00414984	Isoform SGCE-1 of Epsilon-sarcoglycan precursor
IPI00418163	complement component 4B preproprotein
IPI00418163	complement component 4B preproprotein
IPI00418163	complement component 4B preproprotein
IPI00418163	complement component 4B preproprotein
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IPI00418163	complement component 4B preproprotein
IPI00418163	complement component 4B preproprotein
IPI00418262	Fructose-bisphosphate aldolase C
IPI00418262	Fructose-bisphosphate aldolase C
IPI00418262	Fructose-bisphosphate aldolase C
IPI00418262	Fructose-bisphosphate aldolase C
IPI00418262	Fructose-bisphosphate aldolase C
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IPI00418262	Fructose-bisphosphate aldolase C
IPI00418262	Fructose-bisphosphate aldolase C
IPI00418262	Fructose-bisphosphate aldolase C
IPI00418262	Fructose-bisphosphate aldolase C
IPI00418262	Fructose-bisphosphate aldolase C
IPI00418262	Fructose-bisphosphate aldolase C
IPI00418262	Fructose-bisphosphate aldolase C
IPI00418262	Fructose-bisphosphate aldolase C
IPI00418262	Fructose-bisphosphate aldolase C
IPI00418531	gliomedin
IPI00418531	gliomedin
IPI00418531	gliomedin
IPI00418531	gliomedin
IPI00418531	gliomedin
IPI00418531	gliomedin
IPI00419424	IGKV1-5 protein
IPI00419424	IGKV1-5 protein

IPI00419424	IGKV1-5 protein
IPI00419442	IGLV6-57 protein
IPI00419442	IGLV6-57 protein
IPI00419453	PREDICTED: similar to Ig kappa chain V-III region VG prec
IPI00419453	PREDICTED: similar to Ig kappa chain V-III region VG prec
IPI00419585	Peptidyl-prolyl cis-trans isomerase A
IPI00419585	Peptidyl-prolyl cis-trans isomerase A
IPI00419585	Peptidyl-prolyl cis-trans isomerase A
IPI00419585	Peptidyl-prolyl cis-trans isomerase A
IPI00419585	Peptidyl-prolyl cis-trans isomerase A
IPI00419585	Peptidyl-prolyl cis-trans isomerase A
IPI00419585	Peptidyl-prolyl cis-trans isomerase A
IPI00419585	Peptidyl-prolyl cis-trans isomerase A
IPI00419590	Isoform 2 of Testican-3 precursor
IPI00419590	Isoform 2 of Testican-3 precursor
IPI00419595	Podocalyxin-like protein
IPI00419595	Podocalyxin-like protein
IPI00419630	Carnosinase 1
IPI00419630	Carnosinase 1
IPI00419630	Carnosinase 1
IPI00419630	Carnosinase 1
IPI00419630	Carnosinase 1
IPI00419630	Carnosinase 1
IPI00419630	Carnosinase 1
IPI00419630	Carnosinase 1
IPI00419630	Carnosinase 1
IPI00419724	semaphorin 4B precursor
IPI00419724	semaphorin 4B precursor
IPI00419724	semaphorin 4B precursor
IPI00419966	NeshBP
IPI00426056	Hypothetical protein DKFZp686L19235
IPI00430808	Hypothetical protein
IPI00430808	Hypothetical protein
IPI00430808	Hypothetical protein
IPI00430808	Hypothetical protein
IPI00430808	Hypothetical protein
IPI00430808	Hypothetical protein
IPI00430808	Hypothetical protein
IPI00430808	Hypothetical protein
IPI00430808	Hypothetical protein
IPI00430808	Hypothetical protein
IPI00430808	Hypothetical protein
IPI00430842	IGHA1 protein
IPI00430842	IGHA1 protein
IPI00431645	HP protein
IPI00431645	HP protein
IPI00431645	HP protein
IPI00431645	HP protein
IPI00431645	HP protein
IPI00431645	HP protein
IPI00431645	HP protein
IPI00432525	PREDICTED: similar to Sialic acid-binding Ig-like lectin 5 pı
IPI00432525	PREDICTED: similar to Sialic acid-binding Ig-like lectin 5 pı
IPI00432525	PREDICTED: similar to Sialic acid-binding Ig-like lectin 5 pı
IPI00435020	Neural cell adhesion molecule 1, 140 kDa isoform precurso
IPI00435020	Neural cell adhesion molecule 1, 140 kDa isoform precurso

IPI00435020	Neural cell adhesion molecule 1, 140 kDa isoform precurso
IPI00441043	Hypothetical protein
IPI00441043	Hypothetical protein
IPI00441515	Neurexin-3-alpha precursor
IPI00441515	Neurexin-3-alpha precursor
IPI00441515	Neurexin-3-alpha precursor
IPI00441515	Neurexin-3-alpha precursor
IPI00441515	Neurexin-3-alpha precursor
IPI00441515	Neurexin-3-alpha precursor
IPI00441515	Neurexin-3-alpha precursor
IPI00441515	Neurexin-3-alpha precursor
IPI00441515	Neurexin-3-alpha precursor
IPI00441515	Neurexin-3-alpha precursor
IPI00441515	Neurexin-3-alpha precursor
IPI00441515	Neurexin-3-alpha precursor
IPI00441515	Neurexin-3-alpha precursor
IPI00441515	Neurexin-3-alpha precursor
IPI00442294	Isoform 1 of Neurotrimin precursor
IPI00442294	Isoform 1 of Neurotrimin precursor
IPI00442294	Isoform 1 of Neurotrimin precursor
IPI00442294	Isoform 1 of Neurotrimin precursor
IPI00442294	Isoform 1 of Neurotrimin precursor
IPI00442294	Isoform 1 of Neurotrimin precursor
IPI00442294	Isoform 1 of Neurotrimin precursor
IPI00442294	Isoform 1 of Neurotrimin precursor
IPI00442299	Isoform 1 of Neurexin-1-alpha precursor
IPI00442299	Isoform 1 of Neurexin-1-alpha precursor
IPI00442299	Isoform 1 of Neurexin-1-alpha precursor
IPI00442299	Isoform 1 of Neurexin-1-alpha precursor
IPI00442299	Isoform 1 of Neurexin-1-alpha precursor
IPI00442299	Isoform 1 of Neurexin-1-alpha precursor
IPI00442299	Isoform 1 of Neurexin-1-alpha precursor
IPI00442299	Isoform 1 of Neurexin-1-alpha precursor
IPI00442299	Isoform 1 of Neurexin-1-alpha precursor
IPI00442299	Isoform 1 of Neurexin-1-alpha precursor
IPI00442299	Isoform 1 of Neurexin-1-alpha precursor
IPI00442299	Isoform 1 of Neurexin-1-alpha precursor
IPI00442299	Isoform 1 of Neurexin-1-alpha precursor
IPI00442299	Isoform 1 of Neurexin-1-alpha precursor
IPI00442299	Isoform 1 of Neurexin-1-alpha precursor
IPI00444378	CDNA FLJ45634 fis, clone CHONS2002829, moderately si
IPI00444378	CDNA FLJ45634 fis, clone CHONS2002829, moderately si
IPI00451624	Isoform 1 of Cartilage acidic protein 1 precursor
IPI00451624	Isoform 1 of Cartilage acidic protein 1 precursor
IPI00451624	Isoform 1 of Cartilage acidic protein 1 precursor
IPI00451624	Isoform 1 of Cartilage acidic protein 1 precursor
IPI00451624	Isoform 1 of Cartilage acidic protein 1 precursor
IPI00451624	Isoform 1 of Cartilage acidic protein 1 precursor
IPI00451624	Isoform 1 of Cartilage acidic protein 1 precursor
IPI00451624	Isoform 1 of Cartilage acidic protein 1 precursor
IPI00451624	Isoform 1 of Cartilage acidic protein 1 precursor
IPI00451624	Isoform 1 of Cartilage acidic protein 1 precursor
IPI00451624	Isoform 1 of Cartilage acidic protein 1 precursor
IPI00451624	Isoform 1 of Cartilage acidic protein 1 precursor

IPI00453476	PREDICTED: similar to Phosphoglycerate mutase 1 (Phosp
IPI00453476	PREDICTED: similar to Phosphoglycerate mutase 1 (Phosp
IPI00455667	PREDICTED: similar to opioid binding protein/cell adhesion
IPI00456623	Isoform 1 of Brevican core protein precursor
IPI00456623	Isoform 1 of Brevican core protein precursor
IPI00456623	Isoform 1 of Brevican core protein precursor
IPI00456623	Isoform 1 of Brevican core protein precursor
IPI00456623	Isoform 1 of Brevican core protein precursor
IPI00456623	Isoform 1 of Brevican core protein precursor
IPI00456623	Isoform 1 of Brevican core protein precursor
IPI00456623	Isoform 1 of Brevican core protein precursor
IPI00456623	Isoform 1 of Brevican core protein precursor
IPI00456623	Isoform 1 of Brevican core protein precursor
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IPI00456623	Isoform 1 of Brevican core protein precursor
IPI00456623	Isoform 1 of Brevican core protein precursor
IPI00456623	Isoform 1 of Brevican core protein precursor
IPI00456623	Isoform 1 of Brevican core protein precursor
IPI00456736	Isoform 1 of RGM domain family member B precursor
IPI00456736	Isoform 1 of RGM domain family member B precursor
IPI00456736	Isoform 1 of RGM domain family member B precursor
IPI00456736	Isoform 1 of RGM domain family member B precursor
IPI00456736	Isoform 1 of RGM domain family member B precursor
IPI00456736	Isoform 1 of RGM domain family member B precursor
IPI00465028	Triosephosphate isomerase
IPI00465028	Triosephosphate isomerase
IPI00465028	Triosephosphate isomerase
IPI00465028	Triosephosphate isomerase
IPI00465028	Triosephosphate isomerase
IPI00465028	Triosephosphate isomerase
IPI00465028	Triosephosphate isomerase
IPI00465028	Triosephosphate isomerase
IPI00465028	Triosephosphate isomerase
IPI00465028	Triosephosphate isomerase
IPI00465028	Triosephosphate isomerase
IPI00465028	Triosephosphate isomerase
IPI00465028	Triosephosphate isomerase
IPI00465028	Triosephosphate isomerase
IPI00465184	Guanine deaminase
IPI00465184	Guanine deaminase
IPI00465184	Guanine deaminase
IPI00465184	Guanine deaminase
IPI00465184	Guanine deaminase
IPI00465184	Guanine deaminase
IPI00465248	enolase 1
IPI00465248	enolase 1
IPI00465248	enolase 1
IPI00465248	enolase 1
IPI00465248	enolase 1
IPI00465248	enolase 1
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IPI00465248	enolase 1
IPI00465248	enolase 1
IPI00465248	enolase 1
IPI00465315	Cytochrome c
IPI00465315	Cytochrome c
IPI00465315	Cytochrome c
IPI00465315	Cytochrome c
IPI00465315	Cytochrome c
IPI00465315	Cytochrome c
IPI00465325	leucine-rich repeat neuronal 6A
IPI00465325	leucine-rich repeat neuronal 6A
IPI00465325	leucine-rich repeat neuronal 6A
IPI00465325	leucine-rich repeat neuronal 6A
IPI00465325	leucine-rich repeat neuronal 6A
IPI00465439	Fructose-bisphosphate aldolase A
IPI00465439	Fructose-bisphosphate aldolase A
IPI00465439	Fructose-bisphosphate aldolase A
IPI00465439	Fructose-bisphosphate aldolase A
IPI00465439	Fructose-bisphosphate aldolase A
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IPI00465439	Fructose-bisphosphate aldolase A
IPI00465439	Fructose-bisphosphate aldolase A
IPI00465439	Fructose-bisphosphate aldolase A
IPI00470535	Dihydropyridine receptor alpha 2 subunit
IPI00470535	Dihydropyridine receptor alpha 2 subunit
IPI00470535	Dihydropyridine receptor alpha 2 subunit
IPI00470535	Dihydropyridine receptor alpha 2 subunit
IPI00470535	Dihydropyridine receptor alpha 2 subunit
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IPI00470535	Dihydropyridine receptor alpha 2 subunit
IPI00470535	Dihydropyridine receptor alpha 2 subunit
IPI00470625	neuritin precursor
IPI00470625	neuritin precursor
IPI00470652	single-chain Fv fragment
IPI00470653	PREDICTED: similar to Ig heavy chain V-III region VH26 pr
IPI00470653	PREDICTED: similar to Ig heavy chain V-III region VH26 pr
IPI00472011	154 kDa protein
IPI00472011	154 kDa protein
IPI00472011	154 kDa protein
IPI00472011	154 kDa protein
IPI00472011	154 kDa protein
IPI00472011	154 kDa protein
IPI00472011	154 kDa protein
IPI00472013	HLA class I histocompatibility antigen, A-33 alpha chain pre
IPI00472013	HLA class I histocompatibility antigen, A-33 alpha chain pre
IPI00472610	IGHM protein
IPI00472610	IGHM protein
IPI00472610	IGHM protein
IPI00472610	IGHM protein
IPI00472610	IGHM protein
IPI00472610	IGHM protein
IPI00472610	IGHM protein
IPI00472610	IGHM protein
IPI00472610	IGHM protein
IPI00472610	IGHM protein
IPI00472610	IGHM protein
IPI00472610	IGHM protein
IPI00472610	IGHM protein
IPI00472610	IGHM protein
IPI00472762	IGHG1 protein
IPI00472762	IGHG1 protein
IPI00472762	IGHG1 protein
IPI00472882	HLA class I histocompatibility antigen, A-68 alpha chain pre
IPI00472882	HLA class I histocompatibility antigen, A-68 alpha chain pre
IPI00473011	Hemoglobin subunit delta
IPI00473011	Hemoglobin subunit delta
IPI00473011	Hemoglobin subunit delta
IPI00473011	Hemoglobin subunit delta
IPI00473011	Hemoglobin subunit delta
IPI00473011	Hemoglobin subunit delta
IPI00473011	Hemoglobin subunit delta
IPI00473011	Hemoglobin subunit delta
IPI00473014	Destrin
IPI00473014	Destrin
IPI00477069	10 kDa protein
IPI00477090	IGHM protein
IPI00477090	IGHM protein
IPI00477597	Isoform 1 of Haptoglobin-related protein precursor
IPI00477597	Isoform 1 of Haptoglobin-related protein precursor
IPI00477644	26 kDa protein
IPI00477644	26 kDa protein
IPI00477747	follistatin-like 4
IPI00477747	follistatin-like 4
IPI00477747	follistatin-like 4

IPI00477747	follistatin-like 4
IPI00477747	follistatin-like 4
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IPI00477747	follistatin-like 4
IPI00477747	follistatin-like 4
IPI00477747	follistatin-like 4
IPI00477747	follistatin-like 4
IPI00477992	complement component 1, q subcomponent, B chain precu
IPI00477992	complement component 1, q subcomponent, B chain precu
IPI00477992	complement component 1, q subcomponent, B chain precu
IPI00477992	complement component 1, q subcomponent, B chain precu
IPI00478003	Alpha-2-macroglobulin precursor
IPI00478003	Alpha-2-macroglobulin precursor
IPI00478003	Alpha-2-macroglobulin precursor
IPI00478003	Alpha-2-macroglobulin precursor
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IPI00478003	Alpha-2-macroglobulin precursor
IPI00478003	Alpha-2-macroglobulin precursor
IPI00478003	Alpha-2-macroglobulin precursor
IPI00478414	Ventroptin (Fragment)
IPI00478414	Ventroptin (Fragment)
IPI00478414	Ventroptin (Fragment)
IPI00478600	Hypothetical protein
IPI00478600	Hypothetical protein
IPI00478809	Coagulation factor V precursor
IPI00478809	Coagulation factor V precursor
IPI00478809	Coagulation factor V precursor
IPI00478809	Coagulation factor V precursor
IPI00478809	Coagulation factor V precursor
IPI00478809	Coagulation factor V precursor
IPI00478809	Coagulation factor V precursor
IPI00478809	Coagulation factor V precursor
IPI00479016	Isoform 3 of Contactin-4 precursor
IPI00479016	Isoform 3 of Contactin-4 precursor
IPI00479016	Isoform 3 of Contactin-4 precursor
IPI00479016	Isoform 3 of Contactin-4 precursor
IPI00479018	Syntenin iSoform 3
IPI00479018	Syntenin iSoform 3
IPI00479018	Syntenin iSoform 3
IPI00479169	Full-length cDNA clone CS0DD006YL02 of Neuroblastoma
IPI00479169	Full-length cDNA clone CS0DD006YL02 of Neuroblastoma
IPI00479169	Full-length cDNA clone CS0DD006YL02 of Neuroblastoma
IPI00479169	Full-length cDNA clone CS0DD006YL02 of Neuroblastoma

IPI00479708	IGHM protein
IPI00479708	IGHM protein
IPI00479708	IGHM protein
IPI00479708	IGHM protein
IPI00479708	IGHM protein
IPI00479805	Apolipoprotein A-IV precursor
IPI00479805	Apolipoprotein A-IV precursor
IPI00479805	Apolipoprotein A-IV precursor
IPI00479805	Apolipoprotein A-IV precursor
IPI00479805	Apolipoprotein A-IV precursor
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IPI00479805	Apolipoprotein A-IV precursor
IPI00479805	Apolipoprotein A-IV precursor
IPI00479805	Apolipoprotein A-IV precursor
IPI00479805	Apolipoprotein A-IV precursor
IPI00479981	26 kDa protein
IPI00479981	26 kDa protein
IPI00480183	LAR splice variant 1
IPI00514402	Hypothetical protein DKFZp547N203
IPI00514402	Hypothetical protein DKFZp547N203
IPI00514402	Hypothetical protein DKFZp547N203
IPI00514402	Hypothetical protein DKFZp547N203
IPI00514982	26 kDa protein
IPI00514982	26 kDa protein
IPI00549747	Hypothetical protein
IPI00550162	IGLV3-25 protein
IPI00550162	IGLV3-25 protein
IPI00550162	IGLV3-25 protein
IPI00550363	Transgelin-2
IPI00550363	Transgelin-2
IPI00550488	TALDO1 protein
IPI00550488	TALDO1 protein
IPI00550488	TALDO1 protein
IPI00550558	Protein O-linked-mannose beta-1,2-N-acetylglucosaminyltra
IPI00550558	Protein O-linked-mannose beta-1,2-N-acetylglucosaminyltra
IPI00550640	IGHG4 protein
IPI00550640	IGHG4 protein
IPI00550640	IGHG4 protein
IPI00550640	IGHG4 protein
IPI00550991	Isoform 1 of Alpha-1-antichymotrypsin precursor
IPI00550991	Isoform 1 of Alpha-1-antichymotrypsin precursor
IPI00550991	Isoform 1 of Alpha-1-antichymotrypsin precursor
IPI00550991	Isoform 1 of Alpha-1-antichymotrypsin precursor
IPI00550991	Isoform 1 of Alpha-1-antichymotrypsin precursor
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IPI00550991	Isoform 1 of Alpha-1-antichymotrypsin precursor

IPI00550991	Isoform 1 of Alpha-1-antichymotrypsin precursor
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IPI00550991	Isoform 1 of Alpha-1-antichymotrypsin precursor
IPI00550991	Isoform 1 of Alpha-1-antichymotrypsin precursor
IPI00550991	Isoform 1 of Alpha-1-antichymotrypsin precursor
IPI00550991	Isoform 1 of Alpha-1-antichymotrypsin precursor
IPI00550991	Isoform 1 of Alpha-1-antichymotrypsin precursor
IPI00553132	SNC73 protein
IPI00553132	SNC73 protein
IPI00553177	Alpha-1-antitrypsin precursor
IPI00553177	Alpha-1-antitrypsin precursor
IPI00553177	Alpha-1-antitrypsin precursor
IPI00553177	Alpha-1-antitrypsin precursor
IPI00553177	Alpha-1-antitrypsin precursor
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IPI00553177	Alpha-1-antitrypsin precursor
IPI00553177	Alpha-1-antitrypsin precursor
IPI00553177	Alpha-1-antitrypsin precursor
IPI00553177	Alpha-1-antitrypsin precursor
IPI00554811	Actin-related protein 2/3 complex subunit 4
IPI00554811	Actin-related protein 2/3 complex subunit 4
IPI00555812	Vitamin D-binding protein precursor
IPI00555812	Vitamin D-binding protein precursor
IPI00555812	Vitamin D-binding protein precursor
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IPI00555812	Vitamin D-binding protein precursor
IPI00555812	Vitamin D-binding protein precursor
IPI00555945	IGLC2 protein
IPI00555945	IGLC2 protein
IPI00555945	IGLC2 protein
IPI00555945	IGLC2 protein
IPI00607648	Isoform 2 of Neurexin-1-alpha precursor
IPI00607648	Isoform 2 of Neurexin-1-alpha precursor
IPI00607648	Isoform 2 of Neurexin-1-alpha precursor
IPI00607648	Isoform 2 of Neurexin-1-alpha precursor
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IPI00607648	Isoform 2 of Neurexin-1-alpha precursor
IPI00607648	Isoform 2 of Neurexin-1-alpha precursor
IPI00641737	Haptoglobin precursor
IPI00641737	Haptoglobin precursor
IPI00641737	Haptoglobin precursor
IPI00641737	Haptoglobin precursor
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IPI00641737	Haptoglobin precursor

IPI00641737	Haptoglobin precursor
IPI00641737	Haptoglobin precursor
IPI00641737	Haptoglobin precursor
IPI00641961	Collagen, type XII, alpha 1
IPI00641961	Collagen, type XII, alpha 1
IPI00641961	Collagen, type XII, alpha 1
IPI00641961	Collagen, type XII, alpha 1
IPI00642019	HLA class I histocompatibility antigen, B-40 alpha chain pre
IPI00642019	HLA class I histocompatibility antigen, B-40 alpha chain pre
IPI00642632	Ig-like protein
IPI00642632	Ig-like protein
IPI00643034	Isoform 1 of Phospholipid transfer protein precursor
IPI00643034	Isoform 1 of Phospholipid transfer protein precursor
IPI00643034	Isoform 1 of Phospholipid transfer protein precursor
IPI00643034	Isoform 1 of Phospholipid transfer protein precursor
IPI00643034	Isoform 1 of Phospholipid transfer protein precursor
IPI00643525	Complement component 4A
IPI00643525	Complement component 4A
IPI00643525	Complement component 4A
IPI00643914	Tenascin XB
IPI00643914	Tenascin XB
IPI00643914	Tenascin XB
IPI00643914	Tenascin XB
IPI00643914	Tenascin XB
IPI00643914	Tenascin XB
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IPI00643914	Tenascin XB
IPI00643914	Tenascin XB
IPI00643914	Tenascin XB
IPI00643914	Tenascin XB
IPI00645206	protocadherin 17
IPI00645206	protocadherin 17
IPI00645206	protocadherin 17
IPI00645206	protocadherin 17
IPI00645206	protocadherin 17
IPI00645206	protocadherin 17
IPI00646304	peptidylprolyl isomerase B precursor
IPI00646304	peptidylprolyl isomerase B precursor
IPI00646304	peptidylprolyl isomerase B precursor
IPI00646304	peptidylprolyl isomerase B precursor
IPI00646304	peptidylprolyl isomerase B precursor
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IPI00646304	peptidylprolyl isomerase B precursor
IPI00646304	peptidylprolyl isomerase B precursor
IPI00646304	peptidylprolyl isomerase B precursor
IPI00654755	Hemoglobin subunit beta
IPI00654755	Hemoglobin subunit beta
IPI00654755	Hemoglobin subunit beta
IPI00654755	Hemoglobin subunit beta
IPI00654755	Hemoglobin subunit beta
IPI00654755	Hemoglobin subunit beta
IPI00654755	Hemoglobin subunit beta
IPI00654875	Complement C4-B precursor

IPI00654875	Complement C4-B precursor
IPI00654875	Complement C4-B precursor
IPI00654888	Plasma kallikrein precursor
IPI00654888	Plasma kallikrein precursor
IPI00654888	Plasma kallikrein precursor
IPI00654888	Plasma kallikrein precursor
IPI00654888	Plasma kallikrein precursor
IPI00654888	Plasma kallikrein precursor
IPI00719088	collagen, type VI, alpha 1 precursor
IPI00719088	collagen, type VI, alpha 1 precursor
IPI00719088	collagen, type VI, alpha 1 precursor
IPI00719088	collagen, type VI, alpha 1 precursor
IPI00719088	collagen, type VI, alpha 1 precursor
IPI00719088	collagen, type VI, alpha 1 precursor
IPI00719088	collagen, type VI, alpha 1 precursor
IPI00719088	collagen, type VI, alpha 1 precursor
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IPI00719088	collagen, type VI, alpha 1 precursor
IPI00719088	collagen, type VI, alpha 1 precursor
IPI00719088	collagen, type VI, alpha 1 precursor
IPI00719088	collagen, type VI, alpha 1 precursor
IPI00719088	collagen, type VI, alpha 1 precursor
IPI00719088	collagen, type VI, alpha 1 precursor
IPI00719233	IGHA1
IPI00719233	IGHA1
IPI00719233	IGHA1
IPI00719233	IGHA1
IPI00719233	IGHA1
IPI00719233	IGHA1
IPI00719233	IGHA1
IPI00719233	IGHA1
IPI00735451	PREDICTED: similar to Ig heavy chain V region 102 precur
IPI00735451	PREDICTED: similar to Ig heavy chain V region 102 precur
IPI00735451	PREDICTED: similar to Ig heavy chain V region 102 precur
IPI00738433	PREDICTED: similar to Carboxypeptidase N subunit 2 prec
IPI00738433	PREDICTED: similar to Carboxypeptidase N subunit 2 prec
IPI00738433	PREDICTED: similar to Carboxypeptidase N subunit 2 prec
IPI00740550	PREDICTED: similar to Ig kappa chain V-I region HK102 pr
IPI00740550	PREDICTED: similar to Ig kappa chain V-I region HK102 pr
IPI00743517	protein tyrosine phosphatase, receptor type, sigma isoform
IPI00743517	protein tyrosine phosphatase, receptor type, sigma isoform
IPI00743589	PREDICTED: similar to Ig gamma-1 chain C region
IPI00743589	PREDICTED: similar to Ig gamma-1 chain C region
IPI00743589	PREDICTED: similar to Ig gamma-1 chain C region
IPI00743589	PREDICTED: similar to Ig gamma-1 chain C region
IPI00744023	Imunoglobulin heavy chain
IPI00744023	Imunoglobulin heavy chain
IPI00744023	Imunoglobulin heavy chain
IPI00744109	91 kDa protein
IPI00744109	91 kDa protein
IPI00744109	91 kDa protein
IPI00744109	91 kDa protein
IPI00744476	IGLC2 protein
IPI00744476	IGLC2 protein

IPI00744630	Isoform 2 of Protein-L-isoaspartate(D-aspartate) O-methyltr
IPI00744630	Isoform 2 of Protein-L-isoaspartate(D-aspartate) O-methyltr
IPI00744630	Isoform 2 of Protein-L-isoaspartate(D-aspartate) O-methyltr
IPI00745082	calcium/calmodulin-dependent protein kinase IIA isoform 2
IPI00745082	calcium/calmodulin-dependent protein kinase IIA isoform 2
IPI00745313	adipocyte enhancer binding protein 1 precursor
IPI00745313	adipocyte enhancer binding protein 1 precursor
IPI00745363	PREDICTED: similar to Ig heavy chain V-III region VH26 pr
IPI00745363	PREDICTED: similar to Ig heavy chain V-III region VH26 pr
IPI00745872	Serum albumin precursor
IPI00745872	Serum albumin precursor
IPI00745872	Serum albumin precursor
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IPI00745872	Serum albumin precursor
IPI00745872	Serum albumin precursor
IPI00746686	Hypothetical protein DKFZp686O0186 (Fragment)
IPI00746686	Hypothetical protein DKFZp686O0186 (Fragment)
IPI00746686	Hypothetical protein DKFZp686O0186 (Fragment)
IPI00746686	Hypothetical protein DKFZp686O0186 (Fragment)
IPI00746686	Hypothetical protein DKFZp686O0186 (Fragment)
IPI00746686	Hypothetical protein DKFZp686O0186 (Fragment)
IPI00746813	chromogranin A precursor
IPI00746813	chromogranin A precursor
IPI00746813	chromogranin A precursor
IPI00746813	chromogranin A precursor
IPI00748036	94 kDa protein
IPI00748036	94 kDa protein

IPI00748036	94 kDa protein
IPI00749009	Immunoglobulin-like domain containing protein
IPI00749009	Immunoglobulin-like domain containing protein
IPI00760824	IGHG1 protein
IPI00760824	IGHG1 protein
IPI00760824	IGHG1 protein

Sequence	Modifications	MASC(	Peptide length	Number of peptides per protein
AQVVVTVK		30	8	2
EQQSTYTFQLK		51	11	2
CFSLVESTYK		33	10	6
QWDQVEQDLADELITPQGHEK		30	21	6
RDPSPVSGPVHLFR		65	14	6
TLFEDAGYLK		45	10	6
TPEENEPTQLEGGPDSLGFETLENCR		89	26	6
VVEEPNAFGVNNPFLPQASR		57	20	6
AFWDDWMR		31	8	4
GIVTFQFR		37	8	4
LNQQFVHFTQLDLSYLQR		33	18	4
WALGQVFR		48	8	4
NNLCPSGSNIISNLFK		63	16	2
QFYDQALQQAVVDDDANNAK		89	20	2
HNAYVAAWR		39	9	2
LGSQLIVTASK		88	11	2
FAISENPLITLR		31	12	5
HDALDTR		31	7	5
LFEFSLDDLPSEFQQVNITPSK		70	22	5
QSIVPLR		35	7	5
SGGEDESRHDALDTR		34	15	5
AAFDDAIAELDTLSEESYK		90	19	3
LAEQAER		48	7	3
LICCDILDVLDK		70	12	3
VVSLEEPELR		45	10	1
ATAVMPDGQFK		34	11	7
DISLSDYK		41	8	7
GLFIIDDK		44	8	7
LVQAFQFTDK		57	10	7
QITVNDLPVGR		45	11	7
SVDETLR		27	7	7
TIAQDYGVLK		30	10	7
EAVLYVDSQEAALK		45	14	3
ESGDVLLSGAEIFAELGEVIK		153	21	3
VPAFLSAAEVEEHLR		65	15	3
ALGLELR		51	7	8
FPRPNYQLQVSESVAPGAR		30	19	8
QLFSIDASTGEVR		72	13	8
VLDTNDNSPAFDQSTYR		105	17	8
VLELVLR		50	7	8
VSLGLEATLPFR		55	12	8
YSLSSYTSDR		61	10	8
YSVPEEQAPGALVGNVAR		43	18	8
FFQYDTWK		33	8	3
GIVEECCFR		48	9	3
SCDLALLETYCATPAK		69	16	3
DQSGEYECSALNDVAAPDVR		89	20	7
EGQGFVSEDEYLEISDIK		92	18	7
GILSCEASAVPMAEFQWFK		45	19	7
ITVNYPPYISK		60	11	7
LATGLDGMR		45	9	7
MSTLTFFNVSEK		31	12	7
STILYAGNDK		49	10	7
ALELNMLSLK		34	10	5
ASLYNAVTIEDVQK		79	14	5

DDLLGFALR	43	9	5
ELLAVPDNYK	33	10	5
IINNTENLVR	41	10	5
LETDISQIQGEMEDIVQEAR	111	20	2
MQGTLEDQIISANPLLEAFGN/ 1Met(ox)	40	23	2
LDETSGWLSVLHR	36	13	2
LFEIDPTSGVVSLVGK	67	16	2
AQIPVEVSEGAAVGTR	69	16	1
ALTPQCGSGEDLYILTGTVPSDYR	52	24	16
APRPAPGGAEQR	45	12	16
DRIPVYSAFR	34	10	16
DSDIIEDVMVK	53	11	16
FATLYSTR	45	8	16
FFYAGTPPAGLAADSHVK	43	18	16
GQLYPFSLSSDVQVATFTLTN 1Met(ox)	43	32	16
ILEVVNQIQDEER	75	13	16
IPVYSAFR	39	8	16
KILEVVNQIQDEER	79	14	16
LLPFNPQLFQNNCGETEQDTEK	38	22	16
LVGEEEAGFGECDK	62	14	16
PAPGGAEQR	54	9	16
QALNTDYLDSDYQR	95	14	16
WLVEPQIDDPNSNLEEAINEAEAITSVNSLGS	79	33	16
WYVNLHSLMDR	47	11	16
FAELVLEKPLDR	41	12	4
FPLESAFDPDVGTNSLR	28	17	4
NYYTIVTEAPLDR	40	13	4
VLDANDNAPEISFSTVK	59	17	4
DKQHFTTLIK	59	10	20
EAGTLAYYEICDFLR	50	15	20
EGDGSCFPDALDR	53	13	20
FPLTNAIK	30	8	20
FSNTDYAVGYMLR	96	13	20
GNQWVGYDDQESVK	51	14	20
GQEDASPDRFSNTDYAVGYMLR	41	22	20
GTTGHHSPLFR	39	11	20
IASNTQSR	31	8	20
ILGQQVPYATK	50	11	20
ISQHLDFISIMTYDFHGAWR	41	20	20
LGAPASK	42	7	20
LVCYYTSWSQYR	67	12	20
LVMGIPTFGR	48	10	20
QLLLSAALSAGK	71	12	20
SFTLASSETGVGAPISGPGIPGR	92	23	20
SKVQYLK	29	7	20
THGFDGLDLAWLYPGR	90	16	20
TLLSVGGWNFGSQR	98	14	20
VTIDSSYDIAK	80	11	20
MDTLEPFGDEFSGMAR	34	16	1
AGMVNAWTPSSNDDNPWIQVNLLR	29	24	6
EVTGIITQGAR	34	11	6
FELLGCELNGCANPLGLK	71	18	6
NAVHVNLFETPVEAQYVR	34	18	6
NLFETPILAR	40	10	6
VTFLGLQHWVPELAR	45	15	6
ALAHLLEAER	54	10	8

ARAEAQEAEDQQAR	62	14	8
GEAAGAVQELAR	99	12	8
ILAGSADSEGVAAPR	104	15	8
LDPAALAAQLVPAPVPAAALR	106	21	8
NSDPALGLDDDPDAPAAQLAR	113	21	8
PRPPVYDDGPAGPDAEEAGDETPDVDPELL	71	31	8
VLAQLLR	37	7	8
ELEALDEVFTK	55	11	2
LFTEEELAR	51	9	2
DCQPGLCCAFQR	68	12	12
DQDGEILLPR	73	10	12
DSECCGDQLCVWGHCTK	59	17	12
EVEELMEDTQHK	58	12	12
EVPDEYEVGSFMEEVR	86	16	12
GLLFPVCTPLPVEGELCHDPASR	46	23	12
LLDLITWELEPDGALDR	89	17	12
PTFVGSR	43	7	12
QELEDLER	40	8	12
SAVEEMEAEEAAAK	104	14	12
SHECIIDEDCGPSMYCQFASFQYTCQPCR	54	29	12
SLTEEMALGEPAAAAAALLGGEEI	110	24	12
AEHALQR	47	7	8
APDELWNSLGPHPIPVIFK	32	19	8
ETNSGYSGMWHR	53	12	8
LVNIYDSMPLR	52	11	8
TSGIFVKPVNMDNLEK	40	16	8
VIVVWNNIGEK	38	11	8
YLELFQR	36	7	8
YSNIMISQFGFPYANYK	78	17	8
DQECDKFNQCGTCNEFK	34	17	6
GDGLAPLGR	31	9	6
IVTSTYKDGK	61	10	6
NVDGVNYASITR	82	12	6
VGDYGSLSGR	86	10	6
YNLAIEEHCTFGDPIV	30	16	6
AASFQAWGPPSPELLAPTR	60	19	3
TIYLNTLLR	56	9	3
TLLCSFQVLDELGR	84	14	3
CVTAGTSCLISGWGSTSSPQLR	119	22	2
LLCGATLIAPR	61	11	2
GTSCNSFLLK	50	10	2
YGQVPMCDAGEQCAVR	62	16	2
HSPQEAPHVQYER	44	13	1
LHRPPVIVLQR	38	11	3
VTAGISFAIPSDK	27	13	3
YNFIADVVEK	34	10	3
CCDLPFPEQACCAEEEK	72	17	11
DILTIDIGR	39	9	11
DPALCCYLSPGDEQVNCFNINYLR	61	24	11
ELLALIQLER	62	10	11
ELPSLQHPNEQK	47	12	11
EVGPPLPQEAVPLQK	39	15	11
HPPSPTR	32	7	11
LVWEEAMSR	50	9	11
NLPATDPLQR	36	10	11
NVALVSGDTENAK	71	13	11

QGETLNFLEIGYSR	95	14	11
DAGTIAGLNVMR	40	12	7
ITPSYVAFTPEGER	53	14	7
NELESYAYSLK	33	11	7
NQLTSNPENTVFDAK	64	15	7
SQIFSTASDNQPTVTIK	45	17	7
TFAPEEISAMVLTK	50	14	7
TWNDPSVQQDIK	33	12	7
IWCSDPSPGIVAFPR	37	15	3
PALQWFYNGAILNESK	75	16	3
SSPDTQDLYCLNESSK	51	16	3
GQVVYVFSK	62	9	2
TDKWDFYCQ	34	9	2
LIYGATSLQSGVPSR	29	15	2
RLIYGATSLQSGVPSR	27	16	2
AHFSPSNIILDFPAAGSAAR	48	20	12
DFNIPGFPTVR	35	11	12
FPVLEGQR	48	8	12
FVAVLAK	30	7	12
LAGAPSEDPQFPK	48	13	12
LDVPVWDVEATLNFLK	65	16	12
LEEIDGFFAR	57	10	12
NGSGAVFPVAGADVQTLR	55	18	12
NNEEYLALIFEK	74	12	12
SALYSPSDPLTLLQADTVR	45	19	12
SFYTAYLQR	48	9	12
VLNTEANVVR	64	10	12
FIWSEISYLSK	30	11	10
FLSSSLYTALTEAR	66	14	10
FSSPTLELQGEFSPLQSSLPCDIHLVNLR	43	29	10
GLEQGIQDNK	58	10	10
IQFGTLSDFFDALDK	69	15	10
IQFGTLSDFFDALDKADETQR	27	21	10
ITANLFR	43	7	10
LLAENNEIISNIR	95	13	10
VLLAPLGDDFR	28	11	10
WWDIIDIQK	34	9	10
SDDSVIQLLNPNR	33	13	1
AEEYEFLTPVEEAPK	39	15	1
CNEIINWLDK	58	10	2
QTQTFTTYSDNQPGVLIQVYEGER	48	24	2
ALLEDDSDTQQVVVLVR	73	17	5
NLFGLDPSSGAIHVLGPIDFEESR	32	24	5
VEVEILDLNDNSPSFATPER	82	20	5
VGIPENAPIGTLLLR	46	15	5
YSVVEESEPGTLVGNVAQDLGLK	98	23	5
LQAIEHELHELGLLK	73	15	6
MASTPHPPGAR	59	11	6
QIAEGTSISEMWQNDLQPLLIER	45	23	6
SFSNIISTLNPTAK	64	14	6
VFVGATDSAVPCAMMLELAR	35	20	6
YPGSPGSYAAR	44	11	6
LVPVHLDSIL	30	10	1
AGPDNNSPIQIFTIQTR	31	17	7
IFLLEDGSLK	51	10	7
PIFTQEPHDVIFPLDLSK	31	18	7

VLASAPDFSK		36	10	7
VLGYEVLYWTDDSK		79	14	7
VVAGNSIGIGEPSEPSELLR		56	20	7
YLCTVQTTLESLSAVADIIVR		31	21	7
LAAVLAGYGVELR		77	13	4
PAAEEYGYIVTDQKPLSLAAGVK		53	23	4
QLTPEQLSTLLTLLQLLPK		67	19	4
SELEAQTGLQILQTGVGQR		105	19	4
AFSVNIFK		32	8	4
ALQATVGNSYK		46	11	4
FFLQGIQLNTILPDAR		73	16	4
TVESITDIR		44	9	4
NADLQVLKPEPELVYEDLR		37	19	2
QSSGENCDVVVNTLGK		70	16	2
HPAENGK		36	7	7
IEKVEHSDLSFSK		33	13	7
IVKWDRDM	1Met(ox)	32	8	7
SNFLNCYVSGFHPSDIEVDLLK		41	22	7
TPKIQVYSR		32	9	7
VEHSDLSFSK		43	10	7
VNHVTLSQPK		57	10	7
EIILVDDYSNDPEDGALLGK		110	20	2
GGFDWNLVFK		51	10	2
DPWVQELMSCLDLK		55	14	3
FQLLSWSVCGGNK		50	13	3
ISSDSPPSVQFMNR		96	14	3
AAYQVAALPK		53	10	3
APGAIGPYSQAVLVDR		43	16	3
TTVLLADINDFNTVNEIYK		129	19	3
FLPGQGLVLYPQIGDK		56	16	3
FQEFSPNLWGLEFQK		86	15	3
VGQDASSAGSTR		88	12	3
IGPDNLPYVQILK		53	13	2
LRDDVQSINWLR		32	12	2
DSLGWMFNK		43	9	5
FDTSILPICK		41	10	5
SLATLCDGPCPCLPEPEPPK		62	20	5
SLLGAFIPR		49	9	5
VCVTQDYQTALCVSR		113	15	5
DVNEFAPTFK		37	10	4
EGLDINSLESLGQGIK		47	16	4
HGPSPGVR		49	8	4
NWRPASLEAR		33	10	4
QEVEELWIGLNDLK		76	14	3
TLGDQLSLLLGAR		74	13	3
TPLWIGLAGEEGSR		61	14	3
AIINLAVYGK		33	10	6
AIQIMYQNLQQDGLEK	1Met(ox)	71	16	6
LALLVDTVGPR		74	11	6
SVASFSIYSPHTGIQEYQDGVPK		43	23	6
TYPDTDSFNTVAEITGSK		76	18	6
VGALASLIR		71	9	6
DNSLELSQLENK		60	12	1
GAFYLLGEAYFIQPLPAASER		104	21	2
SGSETPLPETDLAHCFYSGTVNGDPSSAAAL		32	38	2
DGLLTVNLR		33	9	2

TALLEANSTPVR	83	12	2
ALYYDLISSPDIHGTYK	106	17	21
DTDTGALLFIGK	98	12	21
EIPDEISILLLGVAHFK	82	17	21
ELLDTVTAPQK	54	11	21
IAQLPLTGSMSIIFFLPLK	86	19	21
ITGKPIK	30	7	21
KTSLEDFYLDEER	89	13	21
LAAAVSNFGYDLYR	110	14	21
LDLQEINNWVQAQMK	87	15	21
LKLSYEGEVTK	57	11	21
LQSLFDSPDFSK	66	12	21
LSYEGEVTK	40	9	21
LTQVEHR	46	7	21
SSFVAPLEK	38	9	21
SSMSPTTNVLLSPLSVATALSALSLGAEQR	126	30	21
TESIIHR	40	7	21
TSLEDFYLDEER	100	12	21
TVQAVLTVPK	71	10	21
VLTGNPR	47	7	21
VPMMSDPK	30	8	21
YGLDSDLSCK	79	10	21
FRDEVEDDYIK	51	11	2
IQIQEAAK	42	8	2
ITCAEEGWSPTPK	60	13	2
TGDIVEFVCK	43	10	2
TYSNAVISPNLETTR	46	15	1
ADEPQWSLYPSDSQVSEEVK	57	20	25
ADQTVLTEDEK	44	11	25
ADQTVLTEDEKK	42	12	25
APRPQSEESWDEEDKR	51	16	25
ASEEEPEYGEEIK	50	13	25
CIIEVLSNALSK	71	12	25
DPADASEAHESSSR	73	14	25
EDEEEEEGENYQK	66	13	25
ELDRNYLNYGEEGAPGK	40	17	25
ELENLAAMDLELQK	72	14	25
ERADEPQWSLYPSDSQVSEEVK	53	22	25
FLGEGHHR	30	8	25
GEAGAPGEEDIQGPTK	87	16	25
GYPGVQAPEDLEWER	53	15	25
HLEEPGETQNAFLNER	103	16	25
KEELVAR	48	7	25
NYLNYGEEGAPGK	62	13	25
NYPSLELDK	29	9	25
SAEFPDFYDSEEPVSTHQEAENEKDR	33	26	25
SQREDEEEEEGENYQK	132	16	25
SSAPPITPECR	62	11	25
SSQGGSLPSEEK	44	12	25
VAQLDQLLHYR	44	11	25
WAEGGGHSR	28	9	25
WQQQGDLQDTK	64	11	25
AVIQHFQEK	36	9	25
CAPFFYGGCGGNR	47	13	25
CLVGEFVSDALLVPDK	78	16	25
EGILQYCQEVYPELQITNVVEANQPVTIQNW(	53	33	25

EQNYSDDVLANMISEPR		101	17	25
EVCSEQAETGPCR		86	13	25
EWEEAER		43	7	25
GLTTRPGSGLTNIK		60	14	25
ISYGNDALMPSLTETK	1Met(ox)	91	16	25
KAVIQHFQEK		59	10	25
LALENYITALQAVPPR		37	16	25
LALENYITALQAVPPRPR		59	18	25
LNMHMNVQNGK		51	11	25
LVFFAEDVGSNK		83	12	25
PGSGLTNIK		45	9	25
QQLVETHMAR		52	10	25
SQVMTHLR		45	8	25
STNLHDYGMLLPCGIDK		49	17	25
TEEISEVK		28	8	25
THPHFVIPYR		45	10	25
VEAMLNDR	1Met(ox)	54	8	25
VEAMLNDRR		28	9	25
VESLEQEAANER		75	12	25
WYFDVTEGK		42	9	25
YLETPGDENEHAHFQK		63	16	25
VYLGPGSDGHPYSTQSIQQGSAVSR		44	25	2
YGLIQAAAVATSR		53	13	2
CIQANYSLMENGK		41	13	12
CPNPPVQENFDVNK		75	14	12
IKVLNQELR		41	9	12
IPTTFENGR		50	9	12
KMTVTDQVNCPK		103	12	12
MTVTDQVNCPK		98	11	12
MTVTDQVNCPKLS		60	13	12
NILTSNNIDVK		66	11	12
NILTSNNIDVKK		66	12	12
NPNLPPETVDSLK		62	13	12
VLNQELR		42	7	12
WYEIEKIPTTFENGR		27	15	12
AVTLSILNDNDNFVLDPYSGVIK		71	23	4
AVYDNQYLLETSSLLDYEGTK		99	21	4
EELPENVPIGNIPK		48	14	4
FIFTVTAR		36	8	4
ALNSIIDVYHK		29	11	2
LLETECPQYIR		43	11	2
ILTPLVSLDTPGK		45	13	3
SLNYWCNLLGMK		42	12	3
VTLAVSDLQK		45	10	3
AAAATGTIFTFR		80	12	10
AVVEVDESGTR		69	11	10
DFTFDLYR		46	8	10
EDQYHYLLDR		45	10	10
FSIEGSYQLEK		54	11	10
GFQQLLQELNQPR		44	13	10
GTQEQDFYVTSETVVR		107	16	10
MQILEGLGLNLQK	1Met(ox)	63	13	10
MQQVENGLSEK		44	11	10
TLYLADTFPTNFR		49	13	10
AIAQSGTALSSWAVSFQPAK		77	20	3
LGVLGFLSTGDQAAK		78	15	3

WTSENIGFFGGDPLR		96	15	3
AASEFESSEGVFLFPELR		69	18	23
AMQHISYLNSR	1Met(ox)	45	11	23
ATEDVLVK		44	8	23
ATVIEGK		33	7	23
DYSFTIQAYDCGK		89	13	23
EGLDLQVLEDSGR		84	13	23
EKLDCELQK		37	9	23
ETILCSSDK		39	9	23
FAGEICGFK		57	9	23
GIEVSSSELGMTFTGVDTMAS	2Met(ox)	41	30	23
GNLAGLTLR		74	9	23
HHYSLYVHGCR		46	11	23
IHGQNVPFDAVVVDK		72	15	23
IISTITR		27	7	23
IPDGVVSVSPK		71	11	23
ISLSGVHHFAR		79	11	23
LDCELQK		47	7	23
QYDSILR		59	7	23
RATEDVLVK		42	9	23
STGEGVIR		47	8	23
VEAVDADCSPQFSQICSYEIITPDVPFTVDKD		45	36	23
VIDCLYTCK		52	9	23
YISNEFK		32	7	23
GVFAIFGLYDK		72	11	3
IQGLTGNVQFDHYGR		41	15	3
LQNILEQIVSVGK		70	13	3
ATYIQNYR		40	8	4
GIGMWNANCLDYSGDAVAK		70	19	4
HHPDFEVFVFDVGQK		33	15	4
SYDWSQITTVATFGK		74	15	4
ATVQQLEGR	1N-ac	70	9	3
ELGVGIALR		54	9	3
LVVECVMNNVTCTR		62	14	3
AGLIDDAFSLAR		72	12	4
LIEFYEDYFK		31	10	4
LISGVTEFLNTEGELK		77	16	4
NDLWNTLSEALK		48	12	4
IINQISTNEIQSDQNLK		73	17	2
LDGSVNFFR		40	9	2
EVQLVESGGGVVRPGGSLR		28	19	1
AIVADPVTFK		27	10	29
CEDVAALDPVTFESPEAFVALPR		109	23	29
DPGNVHTLK		40	9	29
DQGRPFQGQVSGLYYNGLK		28	19	29
FTLSCAEPATLQLDTPVADDR		105	21	29
GATADPLCAPAR		60	12	29
GELYIGGLSK		43	10	29
GGAGDVHQPTK		44	11	29
GKEEFVATFK		32	10	29
GLAEAQGAVGVAPFCSR		50	17	29
GLLANLK		31	7	29
GNEFFCYDLSHNPIQSSTDEITLAFR		45	26	29
GNSDKPVNDNQWHNVVVSR		35	19	29
LGERPPALLGSQGLR		55	15	29
LPDLIADALHR		56	11	29

LSALTLSTVK	63	10	29
LTVNLDCLR	54	9	29
NPCANGGLCTVLAPGEVGCDCSHTGFGGK	36	29	29
QLTIFNSQAAIK	36	12	29
SADYVNLSLK	39	10	29
TALAVDGEAR	54	10	29
TGSISLDFR	52	9	29
TTEPNGLLLFSQGR	57	14	29
VDLPLPPEVWTAALR	38	15	29
VLALAAESDPNVR	61	13	29
VNDGEWCHVDFQR	41	13	29
VVDEWLLDK	37	9	29
WAGAASSGELSFSLR	92	15	29
WHMVLLTR	39	8	29
AFQVIQLSLPEDQK	46	14	7
DLFDCTLYVLLK	52	12	7
DSLFILDGR	50	9	7
NEAGVDEDISSLFVEDSAR	109	19	7
VLIVDVQSQK	52	10	7
VVQAVSTDPVPVK	39	13	7
YFDADSNGLVDINELTQVIK	88	20	7
DVDVNLFESTIR	80	12	2
QETQLLEDYVEAIEGVR	100	17	2
DCNSLPGGLGTCK	35	13	4
IDTIAADESFTELDLGDR	83	18	4
TVMGDLGWIAFPK 1Met(ox)	37	13	4
VMEQNQNNWLLTSWISNEGASR	84	22	4
ETFNLYYYESDNDKER	59	16	7
GLNPLTSYVFHVR	65	13	7
IDTIAADESFTQVDIGDR	108	18	7
NLAQFPDTITGADTSSLVEVR	113	21	7
TYQVCNVMEPSQNNWLR 1Met(ox)	78	17	7
VYPANEVTLLDSR	66	13	7
YNPNPDQSVSVTVTTNQAAPSSIALVQAK	87	29	7
DINTFIHGNKR	39	11	2
NVVVACENGLPVHLDQSIFR	81	20	2
IAYGTQGSSGYSLR	82	14	3
TGAVSGHSLK	56	10	3
VAEYMDWILEK	79	11	3
EAGSAVEAEELVK	76	13	1
AAAVSEAEADFYEQNSR	46	17	15
AGGVLAYELLPALDEVLASDSR	60	22	15
DPVPDLAAWVTSFAAR	39	16	15
FLLGSWLEQAR	68	11	15
GSTGVAAAAGLHR	34	13	15
KDPVPDLAAWVTSFAAR	61	17	15
LLGPGPAADFSVSVER	37	16	15
LLLTSAPSLATSPAFR	35	16	15
LLVLDLFAESQPVYTR	54	16	15
NVFQLEQAFVLSK	58	13	15
QAVQELVSLYYEEAR	72	15	15
QLAGLVANYYTPR	49	13	15
SDVFEAWR	45	8	15
YDLLDLTR	35	8	15
YQLTLWGPEGNILDYANK	63	18	15
FSTFAGFLLFETK	53	13	2

MGNGALHGDHQR	43	12	2
DFSEDQGYPDPPNPCPVGK	87	19	4
LLHGVMEQLGIAR	27	13	4
TADDGCLENTPDTAEFSR	94	18	4
VSEADIQR	63	8	4
AELQEVQITEEKPLLPGQTPE/ 1N-ac	58	24	3
NIITHAPNLDNIELYWNSYNNR	43	22	3
TASLTSAASVDGNR	67	14	3
CFLAFTQTK	39	9	14
CRDQLPYICQFGIV	30	14	14
DQLPYICQFGIV	41	12	14
EQQALQTVCLK	60	11	14
GGTLSTPQTGSENDALYEYLR	77	21	14
KDVVNTK	53	7	14
LDTLAQEVALLK	102	12	14
LDTLAQEVALLKEQQALQTVCLK	73	23	14
NWETEITAQPDGGK	78	14	14
NWETEITAQPDGGKTENCAVLSGAANGK	34	28	14
QSVGNEAEIWLGLNDMAAEG 2Met(ox)	53	29	14
SRLDTLAQEVALLK	72	14	14
TENCAVLSGAANGK	105	14	14
TFHEASEDCISR	48	12	14
GILTVDELLAIR	74	12	3
IPLNDLFR	61	8	3
YLDFVFAVK	43	9	3
ALEYIENLR	58	9	11
ANNIAYEDVVGGEDWNPVEEK	116	21	11
DSLSEEDWMR	38	10	11
ENKPYALNSEK	38	11	11
FPSPEMIR	33	8	11
IESQTQEEVR	66	10	11
IILEALR	44	7	11
LFEKPLDSQSIYQLIEISR	57	19	11
LYTDDEDDIYK	43	11	11
SGQLGIQEEDLR	56	12	11
VLEYLNQEK	51	9	11
DHEDSSLQWSNPAQQTLYFGEK	77	22	8
EDDGASIVCSVNHESLK	50	17	8
EKDTATLNCQSSGSKPAAR	37	19	8
GNPVPQQYLWEK	29	12	8
KGDQELHGEPTR	67	12	8
LLLHCEGR	47	8	8
SLVTVLGIPQKPIITGYK	33	18	8
TFTVSSSVTFQVTR	90	14	8
LGNFPWQAFTSIHGR	49	15	2
VLSYVDWIK	27	9	2
DLGGFDEDAEPR	71	12	1
ITEEFLGK	27	8	6
LATVGELQAAWR	96	12	6
LDISEIK	30	7	6
LLASDAGLYR	53	10	6
QEIESETTSEEQIQEEK	70	17	6
YTLNFEAAQK	54	10	6
ASAATTAILIAR	65	12	6
DIVTVANAVFVK	54	12	6
DMIDNLLSPDLIDGVLTR	63	18	6

FTAVAQTDLKEPLK	73	14	6
NVNFEDPASACDSINAWVK	63	19	6
VLGITDMFDSSK	56	12	6
ALNHLPLEYNSALYSR	27	16	17
CPINCLLGDFGPWSDCDPCIEK	59	22	17
DLHLSDVFLK	45	10	17
DLTSLGHNENQQGSFSSQGGSSFSVPIFYS	30	32	17
ENPAVIDFELAPIVDLVR	91	18	17
GEVLDNSFTGGICK	95	14	17
IGESIELTCPK	63	11	17
QAIQASHK	28	8	17
QLEWGLER	40	8	17
SENINHNSAFK	61	11	17
SEYGAALAWEK	51	11	17
TECIKPVVQEVLTITPFQR	39	19	17
TFSEWLESVK	54	10	17
TLNICEVGTIR	47	11	17
VPANLENVGFEVQTAEDDLK	74	20	17
VPANLENVGFEVQTAEDDLKTDFYK	38	25	17
YYQENFCEQICSK	74	13	17
DHDTFLAVR	29	9	10
DNFHGLAIFLDTYPNDETTER	54	21	10
DWEMHVHFK	27	9	10
LFQLMVEHTPDEESIDWTK	70	19	10
LPTGYYFGASAGTGDLSDNHDIISMK	75	26	10
LTVMTDLEDK	44	10	10
LTVMTDLEDKNEWK	70	14	10
NLHGDGIALWYTR	36	13	10
NRDHDTFLAVR	39	11	10
WTELAGCTADFR	90	12	10
EEAQLATVLAYALSSHCPDMR	68	21	15
EGANYALVIDVDMVPSEGLWI 1Met(ox)	74	21	15
EMLDQSNQWGGTALVVPAFEIR	102	22	15
EPGEFALLR	37	9	15
NELVQLYQVGEVR	89	13	15
PAYVVPWQDPWEPFYVAGGK	57	20	15
QFKQELK	28	7	15
SCQEVFDK	34	8	15
TALASGGVLDASGDYR	120	16	15
VAMHLVCPSR	42	10	15
VPTFDER	40	7	15
WEGPLSVSVFAATK	86	14	15
WVNLPEESLLR	51	11	15
WVNLPEESLLRPAYVVPWQDPWEPFYVAG	33	31	15
YEAAVPDPR	38	9	15
FDLGQDVIDFTGHALALYR	37	19	3
FQLLEGPPESMGR	49	13	3
TDDYLDQPCLETVNR	47	15	3
NFPDLNTYIYYNEK	38	14	1
AGPELLPQQGGGR	34	13	3
AVASQWPEELASAR	37	14	3
LYDFNLGSVTESSLWR	113	16	3
VYSTSVTGSR	48	10	1
ISFDEFIK	56	8	5
ISTSLPVLDLIDAIQPGSINYDLLK	38	25	5
NEALIALLR	40	9	5

VDTDGNGYISFNELNDLFK		61	19	5
VNDDIIVNWVNETLR		77	15	5
DAAVISWTK		67	9	2
TVLIGEYLQIK		44	11	2
TGPAATTLPDGAAAESLVESSEVAVIGFFK		49	30	2
VDATEESDLAQQYGVR		45	16	2
AKDTVYTK		38	8	6
HEDQQQGEDEHQDK		28	14	6
IYPSFQPQPLIYPFVEPIPYGFLPQNILPLAQP/		36	49	6
SPTIPFFDPQIPK		40	13	6
VKHEDQQQGEDEHQDK		59	16	6
VLPIPQQVVPYPQR		35	14	6
AEEQPQVELFVK		27	12	1
ELSNTAAYQSVR		51	12	3
FFPFGLVQLSSDLSK		48	15	3
ILSVSPIQAEQELEDLVAVDLQWSK		34	25	3
GPPAPPEPR		42	9	5
NRDEVQALAFDEQR		76	14	5
SLSVMLVR		43	8	5
TTPLEGTSEMAVTFDK		69	16	5
VPGAYFFSFTAGK		34	13	5
AVLDGCSCCLVCAR		90	14	5
CQLDVLLPEPNCPAPR		66	16	5
DGQIGCVPR		40	9	5
NNEAFLQELELK		64	12	5
TIQAEFQCSPGQIVK		68	15	5
ALTFELTLR		57	9	6
CDEAQVLQVWDDPYPEVLSQEPFHK		37	25	6
FIQSQDYQCSALMGGR	1Met(ox)	107	16	6
KVMSISIR		37	8	6
LAIPQQSDFHNNR		54	13	6
VIPGPPALTLVPAELVR		64	17	6
AIGAVPLIQGEYMIPCEK	1Met(ox)	60	18	15
AYWQVHLDQVEVASGLTLCK		32	20	15
DPDAQPGGELMLGGTDSK		86	18	15
EGCEAIVDTGTSLMVGPVDE\	1Met(ox)	118	22	15
FDGILGMAYPR		97	11	15
ISVNNVLPVFDNLMQQK		44	17	15
LLDIACWIHHK		47	11	15
LVDQNIFSFYLSR		93	13	15
QPGITFIAAK		27	10	15
RTMSEVGGSVEDLIAK		86	16	15
TMSEVGGSVEDLIAK		103	15	15
VGFAEAAR		51	8	15
VSTLPAITLK		28	10	15
YSQAVPAVTEGPIPEVLK		76	18	15
YYTVFDR		27	7	15
AIDEDCSQYEPIPGSQK		71	17	4
FGGTICSGDIWDQASCSSSTTCVR		141	24	4
HLVCNGDQDCLDGSDEDDCEDVR		43	23	4
LGSLGAACEQTQTEGAK		42	17	4
AEATTLHVAPQGTAMAVSTFR		52	21	9
ANFDAQQFAGTWLLVAVGSACR		95	22	9
FLQEQGHR		32	8	9
KLDGICWQVR		59	10	9
LDGICWQVR		60	9	9

QLYGDTGVLGR		61	11	9
SLPVSDSVLSGFEQR		82	15	9
VQEAHLTEDQIFYFPK		40	16	9
YGFCEAADQFHVLDEVR		60	17	9
INHGILYDEEK		66	11	3
ITCTEEGWSPTPK		50	13	3
YKPFSQVPTGEVFYYSCEYNFVSPSK		41	26	3
AGLQVYNK		47	8	3
FEHCNFNDVTTR		60	12	3
LRENELTYYCCK		59	12	3
CLVVCDSNPTSDPTGTALGISVR		120	23	6
EAASNGVLIQMEK		78	13	6
GNLMGGWK		30	8	6
STFIAPR		33	7	6
VAFSAIR		32	7	6
YSTFSGFLVFPL		39	12	6
AEPESETSILLSWTPPR		29	17	11
FEVIEFDDGSGSVLR		122	15	11
GFPTIDMGPQLK		53	12	11
GPPSEPVLTQTSEQAPSSAPR		114	21	11
NVLELNDVR		46	9	11
SDTIANYELVYK		62	12	11
SPQGLGASTAEISAR		92	15	11
SYSFVLTNR		45	9	11
TATMLCAASGNPDPEITWFK		71	20	11
YSAPANLYVR		42	10	11
YSVAGLSPYSDYEFR		62	15	11
DDYFVSGAGLPGR		41	13	4
DLLPASLGSYYR		57	12	4
DNSALDPIIHGLK		36	13	4
ETFLDPFVLR		42	10	4
NAADSSVPSAPR		75	12	1
DFTENPCLR		27	9	4
ECQAALEVLQESPLYDCR		114	18	4
LASIFSGTGADPVVSAK		81	17	4
MLFCSCQDQACAER		95	14	4
FWEEATPIWITNQR		62	14	2
VLLVSFDGFR		48	10	2
ASGVAVSDGVIK	1N-ac	50	12	5
AVLFCLSEDKK		44	11	5
EILVGDVGQTVDDPYATFVK		87	20	5
NIILEEGKEILVGDVGQTVDDPYATFVK		42	28	5
YALYDATYETK		45	11	5
ASYSGVSLFSNPVQYWEIQPSTFR		74	24	4
DLPPDTTLLDLQNNK		35	15	4
ITEIKDGDFK		38	10	4
VSPGAFTPLVK		27	11	4
GVISNSGGPVR		57	11	2
WSASFTVTK		45	9	2
EAGISDYLTIEELVK		42	15	5
FFNANQWADIFQASGAK		90	17	5
QLPAWFDQAK		31	10	5
WPTSGQLFLGHPK		73	13	5
YEDFGPLFTAK		62	11	5
EIVDSYLPVILDIIK		71	15	8
GCSFLPDPYQK		30	11	8

GSAVWCQNVK		33	10	8
KLVGYLDR		40	8	8
LGPGMADICK	1Met(ox)	40	10	8
PGEVCSALNLCESLQK		52	16	8
QEILAALEK		58	9	8
SDVYCEVCEFLVK		53	13	8
EKGYVTPVK		34	9	6
LYGMNEEGWR		47	10	6
MIELHNQEYR		75	10	6
NHCGIASAASYPTV		42	14	6
NSWGEEWGMGGYVK	1Met(ox)	38	14	6
VFQEPLFYEAPR		86	12	6
CSLQNAQEALIVTWQK		54	16	2
EVICQVLHLGTVTDFK		32	16	2
AALSMCK	1Met(ox)	56	7	12
AQGFTEDTIVFLPQTDK		87	17	12
AQGFTEDTIVFLPQTDKCMTEQ		36	22	12
EKFTAFCK		41	8	12
GPGEDFR		51	7	12
KAALSMCK		58	8	12
KNQCETR		30	7	12
MATLYSR		54	7	12
SPHWGSTYSVSVVETDYDQYALLYSQGSK		52	29	12
SVVAPATDGGLNLTSTFLR		95	19	12
TMLLQPAGSLGSYSYR	1Met(ox)	132	16	12
WFSAGLASNSSWLR		110	14	12
ILHCSCQACGK		51	11	4
LALFPDK		38	7	4
SAWCEAK		33	7	4
VDKLVEK		41	7	4
AANEVSSADVK		46	11	7
CEASAVPAPDFEWYR		34	15	7
EFEGEEEYLEILGITR		108	16	7
HSLEYSLR		46	8	7
QGDTAILR		83	8	7
SGIIFAGHDK		62	10	7
VTVNYPPTITESK		61	13	7
AFHFYLTR		28	8	8
ASHQQLDTVWENAK		85	14	8
EDYIYGFQFK		51	10	8
KTQIFLPMNFK		34	11	8
TCSHYECAFLGGLK		81	14	8
TENCIENLEYFQPIYVYNPGEK		96	22	8
TSQGTSFTFGGLNQAR		115	16	8
YVPQLLKEEK		40	10	8
LTVYTTLIDVTK		42	12	2
NMINTFVPSGK	1Met(ox)	37	11	2
ALLQAILQTEDMLK	1Met(ox)	29	14	5
GYFNEELSEILSDPSDDTK		35	19	5
SAIYQLEEEYENLLK		46	15	5
SVQNDSQAIAEVLNQLK		31	17	5
TLELQGLINDLQR		40	13	5
AMDLDQDVLSALAEVEQLSK	1Met(ox)	36	20	12
DILAQSPAAEPLK		42	13	12
EGFYDLSSEDPFGCK		45	15	12
ELAEQLEFIK		44	10	12

ELDSLQTEAESLDNTVK		91	17	12
GIETPQCDQSTGQCVCVEGVEGPR		28	24	12
IPSWTGAGFVR		45	11	12
NFLTQDSADLDSIEAVANEVLK		93	22	12
NIGNLFEEAEK		37	11	12
SLDIFTVGGSGDGVVTNSAWETFQR		81	25	12
VESLSQVEVILQHSAADIAR		46	20	12
YSDIEPSTEGEVIFR		38	15	12
CKPVNTFVHEPLVDVQNVCFQEK		51	23	7
HIIVACEGSPYVPVHFDASVEDST		64	24	7
PVNTFVHEPLVDVQNVCFQEK		33	21	7
QHMDSDSSPSSSSTYCNQMMR		90	21	7
RNMTQGR		32	7	7
SNSSMHITDCR		53	11	7
YPNCAYR		34	7	7
AAAAAAGEAR	1N-ac	56	10	6
AALDGTPGMIGYGMAK		62	16	6
EGGLLTLAGAK		66	11	6
MTDSFTEQADQVTAEVGK		111	18	6
NRPSSGSLIQVVTTEGR		31	17	6
NSGMPPGAAAIAVLPVTLDTPMNR		56	24	6
APLIPMEHCTTR		36	12	11
DIDKDLVI		28	8	11
FFETCDLDNDK		69	11	11
FFETCDLDNDKYIALDEWAGCFGIK		32	25	11
LEAGDHPVELLAR		77	13	11
LHLDYIGPCK		35	10	11
NVLVTLYER		51	9	11
NVLVTLYERDEDNNLLTEK		68	19	11
RLEAGDHPVELLAR		47	14	11
TFDSSCHFFATK		51	12	11
YIPPCLDSELTEFPLR		53	16	11
AGAFDDLTELTYLYLDHNK		39	19	5
FSDGAFLGVTTLK		98	13	5
LSHNPLK		30	7	5
NQLSSYPSAALSK		42	13	5
YLETLWLDNTNLEK		73	14	5
HFFSDYLMGFINSGILK		50	17	3
MCASSCDFVK		58	10	3
QCQPSDTVCASVR		35	13	3
CNSEFWSATSGSHAPASDDTPEFCAALR		52	28	5
GCPLNQQIDFQAFHTNAEGTGAR		84	23	5
VSGQHVEIQAK		54	11	5
VYQAEMDELPAAFVDGSK		31	18	5
YIGTTIVVR		41	9	5
ALFLETEQLK		54	10	6
ESLTLIVEGKPQIK		66	14	6
LDVPQNLMFGK		45	11	6
SVQYDDVPEYK		62	11	6
WKYEKPDGSPVFIAFR		35	16	6
YEKPDGSPVFIAFR		36	14	6
AFLFQDTPR		66	9	30
ALREDNAYCEDIDECAEGR		44	19	30
ASTATAEQFFQK		73	12	30
CLQGEVDCWPLPCPDVECEFSILPENECCPI		35	31	30
CSVCSCQNGFVMCR		67	14	30

CVTDPCQADTIR	61	12	30
CVTDPCQADTIRNDITK	39	17	30
DGYHDNGMFSPSGESCEDID 1Met(ox)	41	26	30
EDNAYCEDIDECAEGR	66	16	30
EFESWIDGCK	27	10	30
ENTMCVNTPGSFMCICK	35	17	30
FTGSSWIK	29	8	30
GYDFCSER	35	8	30
HGTECTLCQCK	47	11	30
HNGQIWVLENDR	39	12	30
IMELQDILAK	63	10	30
LDQCYCER	58	8	30
LSSQCLHQNGETLYNSGDTWVQNCQQCR	57	28	30
LVESSGCPALDCPESHQITLSHSCCK	29	26	30
MVCDCENPTVDLFCCPECDPR	52	21	30
NGHICCSVDPQCLQEL	64	16	30
NTVYSSSGVCVLYECK	129	16	30
PSTDLPLGTTFWLGQR	59	16	30
SALAYVDGK	66	9	30
SICQFQGR	40	8	30
TCLDEMNVVR	66	10	30
TCPTCNDFHGLVQK	73	14	30
VVEKPSTDLPLGTTFWLGQR	88	20	30
YLELESSGHR	61	10	30
YLELESSGHRNEVR	36	14	30
EIPSYTFSCIR	27	11	2
SYSSIVLKPQNIK	27	13	2
DAGTELTGHLVPHHDGLR	35	18	4
LAQAPGLR	31	8	4
LLGIGGHLSPQGK	45	13	4
SNQFFSLDPVTGAVTTAEELDR	138	22	4
DLTGELEYATK	48	11	3
FVESDADEELLFNIPFTGNVK	58	21	3
GLAYGLYLR	46	9	3
SLSGTAFGGFLMFK 1Met(ox)	92	14	4
TPVCTTGQGSGSTATVFAMAELQK	112	24	4
VTGPVPGALGAALWEAGSPVAFYASFSEGT	87	37	4
VWFELTQGSITK	84	12	4
DVQDFWISLPGTLCSEK	44	17	2
SFVQGLGVASDVVR	76	14	2
EPLDPNGIITQYEISYSSIR	41	20	2
GSGVSNFAQLIVR	61	13	2
EMLAVSVLAAVR	28	12	6
MNHQALVR	43	8	6
MTSGDVLSNR	59	10	6
VLALLDVPDK	73	10	6
VLALLDVPDKSQEK	62	14	6
WDICAGNAILK	51	11	6
DNTPNAIAIVQLQELSLR	74	18	2
TFYETPLQLLEK	45	12	2
ALTGGIAHLFK	48	11	1
ALGITEIFIK	71	10	16
AQLVEEWANSVK	80	12	16
ATGEDENILFSPLSIALAMGMM 3Met(ox)	47	32	16
DFDAATYLALINAVYFK	44	17	16
EFSNMVTAK	37	9	16

FTVEQEIDLK	33	10	16
FTVEQEIDLKDVLK	46	14	16
IANSLFVQNGFHVNEEFLQMM 2Met(ox)	28	22	16
NGEEFSFLK	52	9	16
QEVPLATLEPLVK	45	13	16
QKVEVYLPR	32	9	16
RTGTILFMGR	48	10	16
SQFRPENTR	40	9	16
TGTILFMGR	65	9	16
VEVYLPR	42	7	16
YFNAAVNHVDFSQNVAVANYINK	52	23	16
APEEPNIQVNPLGIPVNSK	69	19	11
EAEEETTNDNGVLVLEPAR	70	19	11
EETGQVLER	43	9	11
EVTVPVFYPTEK	33	12	11
GATLALTQVTPQDER	61	15	11
GPVLQLHDLKR	30	11	11
LPSGNHMK	28	8	11
TQLVNVAIFGPPWMAFK 1Met(ox)	66	17	11
VHIQSSQTVESSGLYTLQSILK	48	22	11
VSPAAPER	38	8	11
VWLEVEPVGMLK 1Met(ox)	58	12	11
DGNFGLQELGLK	55	12	2
LSSLSDLEQQYR	58	12	2
AFNNAGEGVPLYESATTR	66	18	15
DGIHLALGMDER	35	12	15
DVVPVLVSSR	33	10	15
EQNIEVDGLSYK	42	12	15
FLSEPSDAVTMR	35	12	15
GEEVIQLR	57	8	15
GNIQTFTVFFSR	75	12	15
GVGPLSDPILFR	40	12	15
GYIIGYGVGSPYAETVR	86	17	15
LSWRPPAEAK	42	10	15
NGDVVIPSDYFQIVGGSNLR	54	20	15
PAIPSSSVLPSAPR	63	14	15
QLYFLQR	28	7	15
VAVAGPLR	31	8	15
VVAYNEWGPGESSQPIK	95	17	15
AGAAAGGPGVSGVCVCK	71	17	14
AITQVSK	38	7	14
DNLAIQTR	35	8	14
EDAGEYECHASNSQGQASASAK	30	22	14
GEGEPCGGGGAGR	85	13	14
GTCEQGPSIVTPPK	82	14	14
GYCAPGMECVK	51	11	14
HEVTGWVLVSPLSK	79	14	14
ITVVDALHEIPVK	80	13	14
ITVVDALHEIPVKK	60	14	14
RGHYGVQR	47	8	14
SRYPVCGSDGTTYPSGCQLR	69	20	14
TELLPGDRDNLAIQTR	46	16	14
YPVCGSDGTTYPSGCQLR	115	18	14
APFTPTWPR	36	9	4
GYSAYDFSDQEDEMAK	101	16	4
MGSNVCGIADSVSSIFV	29	17	4

TGSTPYWIVR	48	10	4
ADDKVYPGEQYTYMLLATEEQSPGEGDGNC	47	33	56
AEEEHLGILGPQLHADVGDK	68	20	56
AETGDKVYVHLK	43	12	56
AGLQAFFQVQECNK	71	14	56
ALYLQYTDETFR	69	12	56
DDEEFIESNK	64	10	56
DIASGLIGPLIICK	63	14	56
DIFTGLIGPMK	67	11	56
DLYSGLIGPLIVCR	84	14	56
DSLDKEK	31	7	56
DVDKEFYLFPTVFDENESLLLEDNIR	53	26	56
EFYLFPTVFDENESLLLEDNIR	80	22	56
ERGPEEEHLGILGPVIWAEVGDTIR	47	25	56
EVGPTNADPVCLAK	77	14	56
EYTDASFTNR	60	10	56
EYTDASFTNRK	37	11	56
FNKNNEGTYYSPNYNPQSR	41	19	56
GAYPLSIEPIGVR	66	13	56
GEFYIGSK	44	8	56
GPEEEHLGILGPVIWAEVGDTIR	99	23	56
GSLHANGR	30	8	56
GVYSSDVFDIFPGTYQTLEMFPR	75	23	56
HYYIAAEEIIWNYAPSGIDIFTK	32	23	56
HYYIGIIETTWDYASDHGEK	51	20	56
IDTINLFPATLFDAYMVAQNPGEWMLSCQNL	31	35	56
IGGSYKK	48	7	56
IYHSHIDAPK	54	10	56
KAEEEHLGILGPQLHADVGDK	59	21	56
KALYLQYTDETFR	74	13	56
KDSLDKEK	36	8	56
KGSLHANGR	73	9	56
KLISVDTEHSNIYLQNGPDR	107	20	56
LISVDTEHSNIYLQNGPDR	75	19	56
MFTTAPDQVDKEDEDFQESN 1Met(ox)	78	21	56
MHAINGR	36	7	56
MYSVNGYTFGSLPGLSMCAE 1Met(ox)	71	22	56
MYYSAVDPTK	60	10	56
MYYSAVDPTKDIFTGLIGPMK	46	21	56
NLASRPYTFHSHGITYYK	30	18	56
NNEGTYYSPNYNPQSR	69	16	56
PVWLGFLGPIIK	59	12	56
PYTFHSHGITYYK	64	13	56
QKDVDKEFYLFPTVFDENESLLLEDNIR	66	28	56
QSEDSTFYLGER	93	12	56
QYTDSTFR	31	8	56
RQSEDSTFYLGER	66	13	56
SGAGTEDSACIPWAYYSTVDQVK	99	23	56
SVPPSASHVAPTETFTYEWTVPK	27	23	56
TTIEKPVWLGFLGPIIK	38	17	56
TYCSEPEK	30	8	56
TYCSEPEKVDKDNEDFQESNR	95	21	56
TYSDHPEK	38	8	56
TYYIAAVEVEWDYSPQR	40	17	56
VDKDNEDFQESNR	65	13	56
VNKDDEEFIESNK	74	13	56

VYPGEQYTYMLLATEEQSPGEGDGNCVTR	44	29	56
EDFDVEAADSAGNCLDSLVFVAGDR	74	25	21
EDTPNSVWEPAK	52	12	21
EPTMYVGSTSVQTSR 1Met(ox)	48	15	21
FYAAGLVSWGPQCGTYGLYTR	60	21	21
GDSGGAFAVQDPNDK	87	15	21
GDSGGAFAVQDPNDKTK	53	17	21
GFQVVVTLR	63	9	21
IIGGSDADIK	55	10	21
LLEVPEGR	33	8	21
LPVAPLR	29	7	21
LPVAPLRK	35	8	21
MGPTVSPICLPGTSSDYNLMDGDLGLISGWC	65	32	21
MLTPEHVFIHPGWK	50	14	21
NYVDWIMK	44	8	21
QFGPYCGHGFPGPLNIETK	32	19	21
REDFDVEAADSAGNCLDSLVFVAGDR	58	26	21
SNALDIIFQTDLTGQK	89	16	21
SSNNPHSPIVEEFQVPYNK	52	19	21
TMQENSTPRED	38	11	21
TNFDNDIALVR	81	11	21
VEDPESTLFGSVIR	83	14	21
AAYNLVR	32	7	5
EFVISDR	29	7	5
ELEEDFIK	27	8	5
FALITWIGENVSGLQR	76	16	5
TGTDKTLVK	36	9	5
EFVDIVQP	34	8	2
QYLLTGQVLSDGK	74	13	2
ELQVYISPK	31	9	10
GIQVELYSFPR	54	11	10
LEIDLLK	39	7	10
LEIELLK	34	7	10
NTVISVNPSTK	53	11	10
QSTQTLYVNVAPR	52	13	10
SIDGAYTIR	49	9	10
SLEVTFTPVIEDIGK	48	15	10
SQEFLEDADRK	66	11	10
VPSVYPLDR	49	9	10
ASAELALGENSEVLK	47	15	8
EYLPIGGLAEFCK	36	13	8
FVTVQTISGTGALR	84	14	8
IAAAILNTPDLR	52	12	8
IGASFLQR	52	8	8
KAEAQIAAK	34	9	8
NLDKEYLPIGGLAEFCK	29	17	8
TCGFDFTGAVEDISK	41	15	8
DILATNGVIHYIDELLIPDSAK	31	22	6
EGVYTVFAPTNEAFR	27	15	6
GDELADSALEIFK	71	13	6
TLFELAAESDVSTAIDLFR	109	19	6
VISTITNNIQQIIEIEDTFETLR	52	23	6
YGTLFTMDR	31	9	6
EVAGLWIK	33	8	7
IESVLSSSGK	60	10	7
IESVLSSSGKR	82	11	7

KPSQLSSFSWDNCDEGK	28	17	7
KPSQLSSFSWDNCDEGKDPAVIR	41	23	7
SEFVVPDLELPSWLTTGNYR	77	20	7
VDLVLEK	40	7	7
IPLENLQIIR	74	10	2
TIQEVAGYVLIALNTVER	48	18	2
EGDMLTLFDGDGPSAR 1Met(ox)	46	16	11
FEAFEEDR	35	8	11
GLISDAQSLYVELLSETPANPLLLSLR	48	27	11
ILLQVEILNVR	59	11	11
LLSSGPDLTLQFQAPPGPPNPGLGQGFVLHF	37	32	11
SGGSPLSPVIYDSDMDDVPER	98	21	11
TASDAGFPVGSHVQYR	29	16	11
TASHGDLIR	52	9	11
VAYEELLDNR	64	10	11
VSLDEDNDR	54	9	11
WVIEAAEGR	38	9	11
GSSSSPLGISVR	59	12	1
AYGYSGVSLK	36	10	6
GLLQNVHLVFENSVEDILSK	36	20	6
GTTQIDPNWVIR	37	12	6
LVFNPDQEDLDGDGR	81	15	6
QVMADSGPIYDQTYAGGR 1Met(ox)	45	18	6
SCDVTSNTCLGPSIQTR	101	17	6
ATNYNAGDR	44	9	7
GISLANWMCLAK	63	12	7
LGMDGYR	33	7	7
QYVQGCGV	34	8	7
RLGMDGYR	35	8	7
STDYGIFQINSR	77	12	7
TPGAVNACHLSCSALLQDNIADAVACAK	46	28	7
DLLFGSIVAVDEPTRPIYR	32	19	4
GSLLLGGLDAEASR	46	14	4
KGNLQLQGTR	84	10	4
LSDGQGFTQDDIQAGR	82	16	4
EAEEHQETQCLR	57	12	3
GLQVALEEFHK	48	11	3
HPPVQWAFQETSVESAVDTPFPAGIFVR	34	28	3
AYWDIMISNHQNSNR	78	15	6
EALQGVGDMGR	73	11	6
FRPDGLPK	38	8	6
GPGGVWAAK	53	9	6
SNEKAEEWGR	53	10	6
VYLQGLIDYYLFGNSSTVLEDSK	122	23	6
FLDDDITDDIMCAK	74	14	3
LEQWLCEKL	42	9	3
NICDISCDK	40	9	3
LLLTAGVSAGR	76	11	2
LVCYFTNWSQDR	54	12	2
ELLESYIDGR	68	10	26
ENLDRDIALMK	46	11	26
ETAASLLQAGYK	77	12	26
ETWTANVGK	36	9	26
GDACEGDSGGPFVMK	62	15	26
GQPSVLQVVNLPIVER	79	16	26
GQPSVLQVVNLPIVERPVCK	78	20	26

HQDFNSAVQLVENFCR	92	16	26
ITDNMFCAGYKPDEGK	33	16	26
IVEGSDAEIGMSPWQVMLFR	114	20	26
KPVAFSDYIHPVCLPDR	41	17	26
KSPQELLCGASLISDR	63	16	26
LAACLEGNCAEGLGTNYR	93	18	26
LAVTTHGLPCLAWASAQAK	77	19	26
LKKPVAFSDYIHPVCLPDR	32	19	26
NPDGDEEGVWCYVAGK	78	16	26
NPDSSTTGPWCYTTDPTVR	87	19	26
PVAFSDYIHPVCLPDR	40	16	26
SEGSSVNLSPPLEQCVPDR	86	19	26
SGIECQLWR	45	9	26
SPQELLCGASLISDR	84	15	26
TATSEYQTFFNPR	59	13	26
TFGSGEADCGLRPLFEK	50	17	26
VTGWGNLK	42	8	26
WYQMGIVSWGEGCDR	58	15	26
YGFYTHVFR	75	9	26
DWAESTLMTQK	50	11	4
EQANNILAR	31	9	4
ETYDFDIAVLR	42	11	4
NTEQEEGGEAVHEVEVVIK	49	19	4
ATLGPAVRPLPWQR	35	14	8
GDSGGPLVCGGVLEGVVTSGSR	68	22	8
KKPGIYTR	34	8	8
RPDSLQHVLLPVLDR	54	15	8
RTHHDGAITER	42	11	8
THHDGAITER	48	10	8
VASYAAWIDSVLA	50	13	8
VQVLLGAHSLSQPEPSKR	33	18	8
AFQYHSK	30	7	33
APWCHTTNSQVR	34	12	33
CQSWSSMTPHR	34	11	33
CTTPPPSSGPTYQCLK	59	16	33
DKYILQGVTSWGLGCAR	41	17	33
DVVLFEK	36	7	33
EAQLPVIENK	58	10	33
EQQCVIMAENR	75	11	33
EQQCVIMAENRK	44	12	33
FSPATHPSEGLEENYCR	42	17	33
FVTWIEGVMR	58	10	33
HSIFTPETNPR	44	11	33
KDIALLK	58	7	33
KSSIIIR	34	7	33
KVYLSECK	36	8	33
LFLEPTR	34	7	33
LFLEPTRK	30	8	33
LSSPAVITDK	51	10	33
LYDYCDVPQCAAPSFDCGKPQVEPK	33	25	33
MRDVVLFEK 1Met(ox)	32	9	33
NLDENYCR	50	8	33
NPDGDVGGPWCYTTNPR	111	17	33
NPDNDPQGPWCYTTDPEK	38	18	33
PNKPGVYVR	39	9	33
QLGAGSIEECAAK	67	13	33

RAPWCHTTNSQVR	43	13	33
TECFITGWGETQGTFGAGLLK	102	21	33
TPENFPCK	52	8	33
TPENYPNAGLTMNYCR	74	16	33
VILGAHQEVNLEPHVQEIEVSR	81	22	33
VIPACLPSPNYVVADR	58	16	33
VQSTELCAGHLAGGTDSCQGDSGGPLVCFE	66	31	33
WELCDIPR	39	8	33
CFEPQLLR	32	8	8
LASQACR	40	7	8
LHEAFSPVSYQHDLALLR	130	18	8
LQEDADGSCALLSPYVQPVCLPSGAAR	38	27	8
NPDNDIRPWCFVLNR	31	15	8
PAPEDLTVVLGQER	59	14	8
TTLSGAPCQPWASEATYR	75	18	8
VVGGLVALR	66	9	8
ALFVSEEEK	29	9	37
ALFVSEEEKK	66	10	37
CLVNLIEK	45	8	37
DAQYAPGYDK	38	10	37
DFHINLFQVLPWLK	37	14	37
DISEVVTPR	50	9	37
DLEIEVVLFHPNYNINGK	30	18	37
DLLYIGK	39	7	37
DLLYIGKDR	45	9	37
DNEQHVFK	38	8	37
EAGIPEFYDYDVALIK	55	16	37
EDYLDVYVFGVGPLVNQVNINALASK	39	26	37
EELLPAQDIK	47	10	37
EKLQDEDLGFL	60	11	37
FIQVGVISWGVVDVCK	42	16	37
FLCTGGVSPYADPNTCR	74	17	37
GDSGGPLIVHK	68	11	37
GDSGGPLIVHKR	42	12	37
ISVIRPSK	54	8	37
KCLVNLIEK	51	9	37
KDNEQHVFK	39	9	37
KEAGIPEFYDYDVALIK	37	17	37
LEDSVTYHCSR	78	11	37
LLQEGQALEYVCPSGFYPYPVQTR	53	24	37
LPPTTTCQQQK	57	11	37
LQDEDLGFL	35	9	37
PICLPCTEGTTR	33	12	37
PQGSCSLEGVEIK	80	13	37
QLNEINYEDHK	42	11	37
STGSWSTLK	39	9	37
VASYGVKPR	54	9	37
VKDISEVVTPR	73	11	37
VSEADSSNADWVTK	103	14	37
VSVGGEKR	57	8	37
WSGQTAICDNGAGYCSNPGIPIGTR	109	25	37
YGLVTYATYPK	76	11	37
YGQTIRPICLPCTEGTTR	58	18	37
AESPEVCFNEESPK	55	14	21
DADPDTFFAK	40	10	21
ELISLVEDVSSNYDGCCEGDVVQCIR	110	26	21

ESLLNHFLYEVAR	58	13	21
FLVNLVK	41	7	21
FTDSENVCQER	63	11	21
FTFEYSR	29	7	21
HELTDEELQSLFTNFANVVDK	38	21	21
HFQNLGK	33	7	21
HPDLSIPELLR	36	11	21
HVCGALLK	37	8	21
IAPQLSTEELVSLGEK	91	16	21
ICAMEGLPQK	50	10	21
KSDVGFLPPFPTLDPEEK	49	18	21
LCFFYNK	29	7	21
LKHELTDEELQSLFTNFANVVDK	32	23	21
LPNNVLQEK	43	9	21
NPFVFAPTLLTVAVHFEEVAK	32	21	21
RHPDLSIPELLR	50	12	21
TINPAVDHCCK	51	11	21
VMNHICSK	34	8	21
CDFEVLVVPWQNSSQLLK	35	18	2
DLSPDDPQVQK	51	11	2
AALEGFLAALQADPPQAER	86	19	21
AKMDLEER	47	8	21
CLPGEFVSEALLVPEGCR	62	18	21
DDTPMTLPK	45	9	21
EWAMADNQSK	41	10	21
FQVHTHLQVIEER	93	13	21
GGLQPPDSKDDTPMTLPK	31	18	21
GSTEQDAASPEKEK	53	14	21
HQEAQEACSSQGLILHGSGMLLPCGSDR	32	28	21
HYQHVAAVDPEK	76	12	21
LVETHATR	59	8	21
MDQCESSTR	65	9	21
MNPLEQYER	49	9	21
QALNEHFQSILQTLEEQVSGER	86	22	21
QINEVMR 1Met(ox)	36	7	21
QMYPELQIAR	67	10	21
RAALEGFLAALQADPPQAER	47	20	21
SGSCAHPHHQVVPFR	54	15	21
VEQATQAIPMER	69	12	21
VIALINDQR	70	9	21
VIALINDQRR	30	10	21
EHVAHLLFLR	48	10	7
EQLGEFYEALDCLCIPR	74	17	7
NWGLSFYADKPETTK	68	15	7
SDVMYTDWK	65	9	7
SDVMYTDWKK	50	10	7
TLMFGSYLDDEK	72	12	7
TLMFGSYLDDEKNWGLSFYAI 1Met(ox)	42	27	7
EVLTGNDEVIGQVLSTLK	94	18	6
LGASPLHVDLATLR	94	14	6
LPYTASSGLMAPR	81	13	6
LSIEDFTAYGGVFGNK	75	16	6
NVLLFLQDK	58	9	6
SEDVPYTAALTAVR	40	14	6
AALSGANVLTLIEK	85	14	36
ADGSGSVVLR	75	10	36

AIALDPR		32	7	36
ATALAIMGDK	1Met(ox)	73	10	36
AVTDEEPFLIFANR		59	14	36
CIPISWTCDLDDDCGDR		68	17	36
CPLNYFACPSGR		32	12	36
CSCYEGWVLEPDGESCR		41	17	36
DGILFWTDWDASLPR		66	15	36
EYAGYLLYSER		42	11	36
FCSEAQFECQNHR		32	13	36
GIALDPAMGK		39	10	36
GPVGLAIDFPESK		83	13	36
GVGGAPPTVTLLR		31	13	36
GWDTLYWTSYTTSTITR		67	17	36
IEAASMSGAGR		89	11	36
IFFSDIHFGNIQQINDDGSR		45	20	36
ILWIDAR		42	7	36
ITWPNGLTLDYVTER		38	15	36
IVFPHGITLDLVSR		42	14	36
LWWADQVSEK		34	10	36
LYWVDAFYDR		37	10	36
MYDAQQQQVGTNK		89	13	36
NAVVQGLEQPHGLVVHPLR		52	19	36
NEPVDRPPVLLIANSQNILATYLSGAQVSTITF		27	37	36
NGDTCVTLLDLELYNPK		63	17	36
NVIALAFDYR		56	10	36
QGLNNAVALDFDYR		49	14	36
SLDPFKPFIIFSNR		40	14	36
SVIVDTK		39	7	36
TGLSNPDGLAVDWVGGNLYWCDK		77	23	36
TNTQPFDLQVYHPSR		38	15	36
TTLLAGDIEHPR		40	12	36
TVLWPNGLSLDIPAGR		45	16	36
VFFTDYGQIPK		53	11	36
YVVISQGLDKPR		46	12	36
EQFLDGDGWTSR		56	12	4
FYALSASFEPFSNK		73	14	4
GKNVLINK		36	8	4
SGTIFDNFLITNDEAYAEEFGNETWGVTK		41	29	4
VSGPWEEADAEAVAR		89	15	1
CPAGVSLVLDGCGCCR		39	16	4
DGAPCIFGGTVYR		52	13	4
GLFCDFGSPANR		60	12	4
LPSPDCPFPR		28	10	4
FNALQYLR		57	8	11
ILGPLSYSK		46	9	11
ISETSLPPDMYECLR		29	15	11
ISNIPDEYFK		41	10	11
LKEDAVSAAFK		75	11	11
LPSGLPVSLLTLYLDNNK		76	18	11
NIPTVNENLENYYLEVNQLEK		91	21	11
NNQIDHIDEK		51	10	11
RFNALQYLR		45	9	11
SLEDLQLTHNK		80	11	11
SLEYLDLSFNQIAR		68	14	11
LENLLLLDLQHNR		32	13	3
NLMQLNLAHNILR		86	13	3

NQLEEVPSALPR		52	12	3
LLLGYNEISK		34	10	2
LQDIPYNIFNLPNIVELSVGHNK		42	23	2
DFALQNPSAVPR		45	12	9
DLHFLEELQLGHNR		48	14	9
LAELPADALGPLQR		66	14	9
LAYLQPALFSGLAELR		69	16	9
LEALPNSLLAPLGR		54	14	9
LEYLLLSR		51	8	9
LWLEGNPWDCGCPLK		45	15	9
NLIAAVAPGAFLGLK		91	15	9
VAGLLEDTFPGLLGLR		82	16	9
AIPVAQDLNAPSDWDSR		106	17	11
ANDESNEHSDVIDSQELSK		83	19	11
DSYETSQLDDQSAETHSHK		73	19	11
GDSVVYGLR		58	9	11
GKDSYETSQLDDQSAETHSHK		37	21	11
ISHELDSASSEVN		71	13	11
KANDESNEHSDVIDSQELSK		40	20	11
QLYNKYPDAVATWLNPDPSQK		28	21	11
QNLLAPQNAVSSEETNDFK		58	19	11
SKEEDKHLK		35	9	11
YPDAVATWLNPDPSQK		66	16	11
FTYTVLEDGCTK		31	12	2
INTDEIMTSLK		44	11	2
DICNDVLSLLEK		62	12	5
DSTLIMQLLR	1Met(ox)	46	10	5
EMQPTHPIR	1Met(ox)	46	9	5
GIVDQSQQAYQEAFEISK		108	18	5
TAFDEAIAELDTLSEESYK		86	19	5
AGFAGDDAPR		66	10	19
AVFPSIVGR		41	9	19
AVFPSIVGRPR		42	11	19
DDDIAALVVDNGSGMCK	1N-ac	82	17	19
DLTDYLMK		44	8	19
DLYANTVLSGGTTMYPGIADR	1Met(ox)	113	21	19
DSYVGDEAQSK		34	11	19
DSYVGDEAQSKR		67	12	19
EITALAPSTMK		52	11	19
GYSFTTTAER		57	10	19
HQGVMVGMGQK		60	11	19
IWHHTFYNELR		58	11	19
KDLYANTVLSGGTTMYPGIADR		35	22	19
LCYVALDFEQEMATAASSSSLEK		101	23	19
QEYDESGPSIVHR		46	13	19
RGILTLK		32	7	19
SYELPDGQVITIGNER		113	16	19
TTGIVMDSGDGVTHTVPIYEGYALPHAILR		41	30	19
VAPEEHPVLLTEAPLNPK		68	18	19
ITVETLSDKYK		34	11	1
AELQEGAR		50	8	36
AHVDALR		31	7	36
AKPALEDLR		57	9	36
AKVQPYLDDFQK		55	12	36
ATEHLSTLSEK		64	11	36
DLATVYVDVLK		77	11	36

DLATVYVDVLKDSGR		60	15	36
DLEEVKAK		43	8	36
DSGRDYVSQFEGSALGK		93	17	36
DYVSQFEGSALGK		102	13	36
EQLGPVTQEFWDNLEK		91	16	36
EQLGPVTQEFWDNLEKETEGLR		61	22	36
ETEGLRQEMSK		41	11	36
KWQEEMELYR		62	10	36
LAEYHAK		50	7	36
LEALKENGGAR		47	11	36
LHELQEK		49	7	36
LLDNWDSVTSTFSK		125	14	36
LREQLGPVTQEFWDNLEK		65	18	36
LREQLGPVTQEFWDNLEKETEGLR		38	24	36
LSPLGEEMR		52	9	36
PALEDLR		32	7	36
QEMSKDLEEVK	1Met(ox)	40	11	36
QGLLPVLESFK		54	11	36
QKLHELQEK		43	9	36
QKVEPLR		29	7	36
QKVEPLRAELQEGAR		29	15	36
THLAPYSDELR		55	11	36
VEPLRAELQEGAR		39	13	36
VKDLATVYVDVLK		86	13	36
VKDLATVYVDVLKDSGR		49	17	36
VQPYLDDFQK		47	10	36
VQPYLDDFQKK		30	11	36
VSFLSALEEYTK		85	12	36
VSFLSALEEYTKK		60	13	36
WQEEMELYR		49	9	36
AATVGSLAGQPLQER		99	15	34
AKLEEQAQQIR		97	11	34
ALMDETMK	1Met(ox)	52	8	34
ALMDETMKELK	1Met(ox)	53	11	34
AQAWGER		45	7	34
ARMEEMGSR		61	9	34
AYKSELEEQLTPVAEETR		108	18	34
DADDLQK		50	7	34
DADDLQKR		53	8	34
DRLDEVK		30	7	34
DRLDEVKEQVAEVR		70	14	34
ELQAAQAR		45	8	34
EQVAEVR		34	7	34
ERLGPLVEQGR		41	11	34
GEVQAMLGQSTEELR		126	15	34
LAVYQAGAR		59	9	34
LDEVKEQVAEVR		67	12	34
LEEQAQQIR		64	9	34
LGADMEDVCGR		82	11	34
LGPLVEQGR		69	9	34
LLRDADDLQK		33	10	34
LQAEAFQAR		61	9	34
LSKELQAAQAR		86	11	34
LVQYRGEVQAMLGQSTEELR		41	20	34
MEEMGSR		42	7	34
QQTEWQSGQR		59	10	34

QWAGLVEK	37	8	34
RLAVYQAGAR	53	10	34
SELEEQLTPVAEETR	103	15	34
SWFEPLVEDMQR	69	12	34
VEQAVETEPEPELR	77	14	34
VQAAVGTSAAPVPSDNH	111	17	34
WELALGR	27	7	34
WVQTLSEQVQEELLSSQVTQELR	134	23	34
AGTELVNFLSYFVELGTQPATQ	77	22	7
EPCVESLVSQYFQTVTDYGK	121	20	7
EQLTPLIK	29	8	7
SKEQLTPLIK	50	10	7
SKEQLTPLIKK	52	11	7
SPELQAEAK	69	9	7
VKSPELQAEAK	65	11	7
EFGNTLEDKAR	33	11	5
EWFSETFQK	28	9	5
IKQSELSAK	49	9	5
LKEFGNTLEDK	51	11	5
MREWFSETFQK	53	11	5
ESLSSYWESAK	34	11	3
STAAMSTYTGIFTDQVLSVLK	111	21	3
STAAMSTYTGIFTDQVLSVLK(1Met(ox)	35	24	3
DALSSVQESQVAQQAR	109	16	2
GWVTDGFSSLK	36	11	2
ALTDMPQMR	42	9	24
AQLVDMK	40	7	24
DSDWPFCSDEDWNYK	27	15	24
DSHSLTTNIMEILR	32	14	24
EVDLKDYEDQQK	71	12	24
EVVTSEDGSDCPEAMDLGTLSGIGTLDGFR	28	30	24
GDFSSANNR	39	9	24
GGSTSYGTGSETESPR	83	16	24
GLIDEVNQDFTNR	98	13	24
GSESGIFTNTK	71	11	24
HRHPDEAAFFDTASTGK	47	17	24
LEVDIDIK	43	8	24
MKPVPDLVPGNFK	31	13	24
NPGSSGTGGTATWKPGSSGPGSTGSWNS(	41	45	24
NPSSAGSWNSGSSGPGSTGNR	98	21	24
NSLFEYQK	42	8	24
PGSTGTWNPGSSER	44	14	24
PNNPDWGTFEEVSGNVSPGTR	124	21	24
QFTSSTSYNR	43	10	24
QLEQVIAK	44	8	24
RLEVDIDIK	34	9	24
TFPGFFSPMLGEFVSETESR	44	20	24
VQHIQLLQK	57	9	24
VTSGSTTTTR	27	10	24
AIQLTYNPDESSKPNMIDAATLK	32	23	20
ASTPNGYDNGIIWATWK	67	17	20
DNCCILDER	60	9	20
DTVQIHDITGK	42	11	20
EGFGHLSPTGTTEFWLGNEK	74	20	20
FGSYCPTTCGIADFLSTYQTK	113	21	20
IHLISTQSAIPYALR	80	15	20

KMLEEIMK	35	8	20
LDGSVDFK	39	8	20
LDGSVDFKK	42	9	20
LTIGEGQQHHLGGAK	45	15	20
LTYAYFAGGDAGDAFDGFDFGDDPSDK	70	27	20
QSGLYFIKPLK	55	11	20
RLDGSVDFK	32	9	20
RLDGSVDFKK	39	10	20
TSTADYAMFK	53	10	20
VELEDWNGR	49	9	20
VGPEADKYR	47	9	20
YEASILTHDSSIR	73	13	20
YLQEIYNSNNQK	81	12	20
AAPFLTYFGLFQVH	29	14	2
QIGEFIVTR	37	9	2
AIQDGTIVLMGTYDDGATK 1Met(ox)	109	19	13
DNWVFCGGK	51	9	13
GINVALANGK	66	10	13
ICLEDNVLMSGVK	61	13	13
LIADLGSTSITNLGFR	108	16	13
MASGAANVVGPK	84	12	13
MDASLGNLFAR	88	11	13
SALDTAAR	51	8	13
SPFEQHIK	77	8	13
TGEVLDTK	68	8	13
TKSPFEQHIK	33	10	13
YEGWPEVVEMEGCIPQKQD	29	19	13
YFDMWGGDVAPFIEFLK	82	17	13
AAPLQGMLPGLLAPLR	43	16	19
ALGSAIEYTIENVFESAPNPR	70	21	19
ALILVGLER	34	9	19
ALNLGYALDYAQR	53	13	19
DVVFLLDGSEGVR	46	13	19
EVYTFASEPNDVFFK	57	15	19
IIDELNVKPEGTR	28	13	19
ITEGVPQLLIVLTADR	58	16	19
LLTPITTLTSEQIQK	42	15	19
LLVLITGGK	27	9	19
LSDAGITPLFLTR	53	13	19
NADPAELEQIVLSPAFILAAESLPK	72	25	19
PGVISVMGT	36	9	19
QLGTVQQVISER	31	12	19
QLTLLGGPTPNTGAALEFVLR	43	21	19
VAVFFSNTPTR	30	11	19
VEFLLNAHSSKDEVQNAVQR	35	20	19
VPQIAFVITGGK	43	12	19
YPPPAVESDAADIVFLIDSSEGVRPDGFAHIR	27	32	19
AALTELSLGSAYQAMILGVDSK	63	22	66
AHLDIAGSLEGHLR	28	14	66
ALVEQGFTVPEIK	41	13	66
ATFQTPDFIVPLTDLR	43	16	66
AVSMPSFSILGSDVR	36	15	66
CSLLVLENELNAELGLSGASM 1Met(ox)	32	22	66
DKDQEVLLQTFLDDASPGDKR	27	21	66
DLKVEDIPLAR	39	11	66
EVGTVLSQVYSK	39	12	66

EYSGTIASEANTYLNSK		50	17	66
FDHTNSLNIAGLSLDFSSK		46	19	66
FQFPGKPGIYTR		27	12	66
FSVPAGIVIPSFQALTAR		28	18	66
FVTQAEGAK		40	9	66
GFEPTLEALFGK		55	12	66
GMTRPLSTLISSSQSCQYTLD	1Met(ox)	28	23	66
HINIDQFVR		27	9	66
HIQNIDIQHLAGK		32	13	66
IADFELPTIIVPEQTIEIPSIK		57	22	66
IAIANIIDEIIEK		76	13	66
IDDIWNLEVK		45	10	66
IEDGTLASK		45	9	66
IEGNLIFDPNNYLPK		30	15	66
IEIPLPFGGK		28	10	66
IGQDGISTSATTNLK		45	15	66
ILGEELGFASLHDLQLLGK		40	19	66
ITENDIQIALDDAK		45	14	66
LAAYLMLMR		72	9	66
LAPGELTIIL		36	10	66
LATALSLSNK		29	10	66
LGNNPVSK		41	8	66
LLLQMDSSATAYGSTVSK	1Met(ox)	49	18	66
LLSGGNTLHLVSTTK		37	15	66
LNTDIAGLASAIDMSTNYNSD	1Met(ox)	34	30	66
LPQQANDYLNSFNWER		60	16	66
LQDFSDQLSDYYEK		53	14	66
LSLESLTSYFSIESSTK		80	17	66
LSQLQTYMIQFDQYIK		54	16	66
NFVASHIANILNSEELDIQDLK		36	22	66
NFVASHIANILNSEELDIQDLKK		30	23	66
NIQEYLSILTDPDGK		61	15	66
NLTDFAEQYSIQDWAK		49	16	66
NSEEFAAAMSR	1Met(ox)	77	11	66
NSLFFSAQPFEITASTNNEGNLK		72	23	66
NTFTLSCDGSLR		28	12	66
NTLELSNGVIVK		59	12	66
QTVNLQLQPYSLVTTLNSDLK		60	21	66
QVFLYPEKDEPTYILNIK		31	18	66
SEILAHWSPAK		48	11	66
SGSSTASWIQNVDTK		35	15	66
SLWDFLK		31	7	66
SNTVASLHTEK		67	11	66
SPSQADINK		53	9	66
SVSLPSLDPASAK		28	13	66
TGISPLALIK		28	10	66
TILGTMPAFEVSLQALQK	1Met(ox)	37	18	66
TLADLTLLDSPIK		41	13	66
TQFNNNEYSQDLDAYNTK		54	18	66
TSSFALNLPTLPEVK		64	15	66
VELEVPQLCSFILK		33	14	66
VIGNMGQTMEQLTPELK		37	17	66
VNWEEEAASGLLTSLK		87	16	66
VPSYTLILPSLELPVLHVPR		29	20	66
YEDGTLSLTSTSDLQSGIIK		55	20	66
YGMVAQVTQTLK	1Met(ox)	27	12	66

YSQPEDSLIPFFEITVPESQLTVSQFTLPK	60	30	66
ESQAYYQR	35	8	10
GENFTETDVK	46	10	10
HMAGAAAAGAVVGGLGGYMl 3Met(ox)	42	26	10
PGGWNTGGSR	57	10	10
PIIHFGSDYEDR	64	12	10
QHTVTTTTK	51	9	10
QHTVTTTTKGENFTETDVK	27	19	10
VVEQMCITQYER	93	12	10
YPGQGSPGGNR	60	11	10
YPNQVYYR	47	8	10
DVSSSVYSTWK	52	11	3
LLCTDPGFVK	29	10	3
LVVQSSIDSSAFK	55	13	3
AIWNVINWENVTER	71	14	4
GDVTAQIALQPALK	68	14	4
GELLEAIKR	44	9	4
LTAASVGVQGSGWGWLGFNK	75	20	4
DLLAGLPAPGVEVYCLYGVGLPTPR	45	25	7
ITTTSPWMFPSR	40	12	7
LEPGQQEEYYR	33	11	7
SSGLVSNAPGVQIR	64	14	7
STELCGLWQGR	48	11	7
TYIYDHGFPYTDPVGVLYEDGDDTVATR	28	28	7
TYSVEYLDSSK	47	11	7
AGPPGPTDDFSVEYLVVGNR	39	20	6
DAVAGGPENCLTSLTQDR	87	18	6
LCDPSAPLAFLQASK	64	15	6
TCLPAPGVEGGGCEGVLEEGR	72	21	6
TYLGVESFDEVLR	73	13	6
WLDACLAGSR	61	10	6
ALAILTLR	40	8	5
FDSFADASVLGVLAPYVLR	38	19	5
IGESDFFFTVPVSR	41	14	5
QALVDSVFQVSVLPGNVGYLR	63	21	5
TAVDLESLASQLTADLQEVSGDHR	50	24	5
ADLFYDVEALDLESPK	108	16	15
ALDLINK	36	7	15
ALDLINKR	61	8	15
DGYLFQLLR	51	9	15
DSPVLIDFFEDTER	88	14	15
GEVLPLPEANFPSFPLPHHK	28	20	15
GGEGTGYFVDFSVR	87	14	15
HSHESQDLR	35	9	15
IADAHLDR	64	8	15
QIGSVYR	35	7	15
RDGYLFQLLR	41	10	15
RPSEIVIGQCK	52	11	15
SGFPQVSMFFTHTFPK	53	16	15
YKEENDDFASFR	64	12	15
YWNDCEPPDSR	46	11	15
APLTKPLK	35	8	6
ESDTSYVSLK	57	10	6
GYSIFSYATK	36	10	6
KAFVFPK	40	7	6
QDNEILIFWSK	38	11	6

RQDNEILIFWSK	28	12	6
AYSLFSYNTQGR	81	12	6
DNELLVYK	38	8	6
GYVIIKPLVWV	51	11	6
IVLGQEQDSYGGK	77	13	6
QGYFVEAQPK	39	10	6
VGEYSLYIGR	78	10	6
GHIYQGSEADSVFSGFLIFPSA	44	22	5
GLFQVVSGGMVLQLQQGDQ\ 1Met(ox)	29	24	5
KGHIYQGSEADSVFSGFLIFPSA	32	23	5
PAFSAIR	42	7	5
SLGFCDTTNK	38	10	5
FNAVLTNPQGDYDTSTGK	77	18	5
FQSVFTVTR	55	9	5
QTHQPPAPNSLIR	44	13	5
TNQVNSGGVLLR	62	12	5
VVTFCGHTSK	31	10	5
AEQCCEETASSISLHGK	104	17	14
AIEDYINEFSVR	86	12	14
ALPTTYEK	29	8	14
DRDGNTLTYYR	35	11	14
DVVLTTTFVDDIK	56	13	14
GTVIDVTDFVNWASSINDAPVLISQK	78	26	14
LSPIYNLVPVK	38	11	14
QCVPTEPCEDAEDDCGNDFQCSTGR	38	25	14
RPWNVASLIYETK	44	13	14
SIEVFGQFNGK	54	11	14
TAGYGINILGMDPLSTPFDNEFYNGLCNR	51	29	14
TEHYEEQIEAFK	59	12	14
TSNFNAAISLK	53	11	14
VVEESELAR	59	9	14
ALGHLDLSGNR	52	11	10
DGFDISGNPWICDQNLSDLYR	75	21	10
DLLLPQPDLR	42	10	10
ENQLEVLEVSWLHGLK	65	16	10
GPLQLER	52	7	10
LHLEGNK	46	7	10
NALTGLPPGLFQASATLDTLVLK	59	23	10
TLDLGENQLETLPPDLLR	80	18	10
VAAGAFQGLR	66	10	10
YLFLNGNK	28	8	10
AQITGYR	38	7	32
DLQFVEVTDVK	78	11	32
ESKPLTAQQTTK	52	12	32
GATYNIIVEALK	77	12	32
GATYNIIVEALKDQQR	95	16	32
GDSPASSKPISINYR	60	15	32
GLAFTDVDVDSIK	74	13	32
IAWESPQGQVSR	56	12	32
IGDQWDK	36	7	32
IGDTWSK	32	7	32
ISCTIANR	51	8	32
ITGYIIK	36	7	32
ITYGETGGNSPVQEFTVPGSK	80	21	32
IYLYTLNDNAR	66	11	32
LTVGLTR	38	7	32

MSESGFK		28	7	32
NTFAEVTGLSPGVTYYFK		92	18	32
QYNVGPSVSK		34	10	32
SDTVPSPR		38	8	32
SEPLIGR		31	7	32
SSPVVIDASTAIDAPSNLR		103	19	32
STTPDITGYR		46	10	32
SYTITGLQPGTDYK		73	14	32
TFYSCTTEGR		60	10	32
TYLGNALVCTCYGGSR		85	16	32
VEYELSEEGDEPQYLDLPSTATSVNIPDLLPC		88	34	32
VFAVSHGR		37	8	32
VGDTYERPK		50	9	32
VTIMWTPPESAVTGYR	1Met(ox)	61	16	32
VTWAPPPSIDLTNFLVR		66	17	32
WLPSSSPVTGYR		45	12	32
YEVSVYALK		51	9	32
DPEGLFLQDNIVAEFSVDETG	1Met(ox)	41	28	13
DPNGLPPEAQK		42	11	13
FSGTWYAMAK	1Met(ox)	66	10	13
GNDDHWIVDTDYDTYAVQYSCR		98	22	13
KDPEGLFLQDNIVAEFSVDET(	1Met(ox)	58	29	13
LIVHNGYCDGR		31	11	13
LLNLDGTCADSYSFVFSR		123	18	13
LLNNWDVCADMVGTFTDTEDPAK		116	23	13
MKYWGVASFLQK		41	12	13
QEELCLAR		37	8	13
VKENFDK		41	7	13
VKENFDKAR		64	9	13
YWGVASFLQK		50	10	13
AFIQLWAFDAVK		78	12	12
ECLQTCR		27	7	12
EDSCQLGYSAGPCMGMTSR		57	19	12
ETLLQDFR		72	8	12
EYCGVPGDGDEELLR		40	15	12
GECVPGEQEPEPILIPR		72	17	12
GVCEETSGAYEK		45	12	12
HHGPTITAK		34	9	12
MTVSTLVLGEGATEAEISMTS	2Met(ox)	108	23	12
TVAACNLPIVR		57	11	12
VVAQGVGIPEDSIFTMADR	1Met(ox)	62	19	12
WYNLAIGSTCPWLK		75	14	12
EQLGEFYEALDCLR		73	14	11
KDKCEPLEK		44	9	11
NWGLSVYADKPETTK		91	15	11
SDVVYTDWK		59	9	11
SDVVYTDWKK		51	10	11
SVQEIQATFFYFTPNK		43	16	11
TEDTIFLR		65	8	11
TYMLAFDVNDEK		62	12	11
TYMLAFDVNDEKNWGLSVYADKPETTK		61	27	11
WFYIASAFR		56	9	11
YVGGQEHFAHLLILR		87	15	11
AQLVPLPPSTYVEFTVSGTDCVAK		72	24	8
CNLLAEK		32	7	8
EHAVEGDCDFQLLK		77	14	8

FSVVYAK	36	7	8
HTFMGVVSLGSPSGEVSHPR	127	20	8
HTLNQIDEVK	62	10	8
QLKEHAVEGDCDFQLLK	46	17	8
QPNCDDPETEEAALVAIDYINQNLPWGYK	47	29	8
AADDTWEPFASGK	99	13	13
ALGISPFHEHAEVVFTANDSGPR	110	23	13
ALGISPFHEHAEVVFTANDSGPRR	44	24	13
CPLMVKVLDAVR	29	12	13
GSPAINVAVHVFR	97	13	13
GSPAINVAVHVFRK	52	14	13
KAADDTWEPFASGK	103	14	13
RYTIAALLSPYSYSTTAVVTNPK	72	23	13
RYTIAALLSPYSYSTTAVVTNPKE	78	24	13
TSESGELHGLTTEEEFVEGIYK	95	22	13
TSESGELHGLTTEEEFVEGIYKVEIDTK	67	28	13
YTIAALLSPYSYSTTAVVTNPK	106	22	13
YTIAALLSPYSYSTTAVVTNPKE	125	23	13
ADDKETCFAEEGKK	50	14	33
AEFAEVSKLVTDLTK	87	15	33
AFKAWAVAR	42	9	33
AVMDDFAAFVEK	81	12	33
CCTESLVNR	59	9	33
DAHKSEVAHR	61	10	33
DDNPNLPRLVR	33	11	33
DLGEENFK	44	8	33
EFNAETFTFHADICTLSEKER	54	21	33
ETYGEMADCCAK	61	12	33
FPKAEFAEVSK	58	11	33
HPYFYAPELLFFAKR	78	15	33
KVPQVSTPTLVEVSR	87	15	33
KYLYEIAR	49	8	33
LAKTYETTLEK	49	11	33
LDELRDEGKASSAK	47	14	33
LVRPEVDVMCTAFHDNEETFLK	70	22	33
LVRPEVDVMCTAFHDNEETFLKK	52	23	33
LVTDLTK	48	7	33
NECFLQHKDDNPNLPR	76	16	33
QEPERNECFLQHKDDNPNLPR	64	21	33
QTALVELVK	52	9	33
RHPDYSVVLLLR	73	12	33
RHPYFYAPELLFFAK	85	15	33
RHPYFYAPELLFFAKR	52	16	33
RPCFSALEVDETYVPK	70	16	33
SHCIAEVENDEMPADLPSLAADFVESK	139	27	33
TPVSDRVTK	46	9	33
TYETTLEK	40	8	33
VFDEFKPLVEEPQNLIK	81	17	33
VHTECCHGDLLECADDRADLAK	40	22	33
YICENQDSISSK	87	12	33
YLYEIAR	40	7	33
ADRDQYELLCLDNTR	74	15	80
AIAANEADAVTLDAGLVYDAYLAPNNLK	28	28	80
AIAANEADAVTLDAGLVYDAYLAPNNLKPVV/	67	38	80
APNHAVVTR	59	9	80
ASYLDCIR	47	8	80

AVANFFSGSCAPCADGTDFPQLCQLCPGCG	46	45	80
AVGNLRK	53	7	80
CDEWSVNSVGK	76	11	80
CGLVPVLAENYNK	39	13	80
CLKDGAGDVAFVK	78	13	80
CLVEKGDVAFVK	63	12	80
CSTSSLLEACTFR	100	13	80
DCHLAQVPSHTVVAR	54	15	80
DDTVCLAK	41	8	80
DGAGDVAFVK	92	10	80
DGAGDVAFVKHSTIFENLANK	49	21	80
DKEACVHK	33	8	80
DKSKEFQLFSSPHGK	31	15	80
DLLFKDSAHGFLK	38	13	80
DLLFRDDTVCLAK	60	13	80
DQYELLCLDNTR	77	12	80
DSAHGFLK	46	8	80
DSGFQMNQLR	78	10	80
DYELLCLDGTR	81	11	80
DYELLCLDGTRK	50	12	80
EDLIWELLNQAQEHFGK	89	17	80
EDLIWELLNQAQEHFGKDK	50	19	80
EDPQTFYYAVAVVK	98	14	80
EDPQTFYYAVAVVKK	54	15	80
EFQLFSSPHGK	74	11	80
EFQLFSSPHGKDLLFK	58	16	80
EGTCPEAPTDECK	40	13	80
EGTCPEAPTDECKPVK	58	16	80
EGYYGYTGAFR	60	11	80
FDEFFSEGCAPGSK	105	14	80
FDEFFSEGCAPGSKK	41	15	80
GDVAFVK	55	7	80
GDVAFVKHQTVPQNTGGK	48	18	80
HQTVPQNTGGK	54	11	80
HQTVPQNTGGKNPDPWAK	28	18	80
HSTIFENLANK	82	11	80
HSTIFENLANKADR	95	14	80
IECVSAETTEDCIAK	88	15	80
IMNGEADAMSLDGGFVYIAGK 1Met(ox)	130	21	80
KASYLDCIR	50	9	80
KCSTSSLLEACTFR	88	14	80
KDKEACVHK	43	9	80
KDSGFQMNQLR	81	11	80
KDSSLCK	35	7	80
KPVDEYK	38	7	80
KPVDEYKDCHLAQVPSHTVVAR	50	22	80
KPVEEYANCHLAR	79	13	80
KSASDLTWDNLK	105	12	80
KSCHTAVGR	60	9	80
LCMGSGLNLCEPNNK	84	15	80
LHDRNTYEK	31	9	80
LKCDEWSVNSVGK	90	13	80
MYLGYEYVTAIR	91	12	80
NLNEKDYELLCLDGTR	132	16	80
NLNEKDYELLCLDGTRK	84	17	80
NPDPWAK	28	7	80

NTYEKYLGEEYVK	61	13	80
PVVAEFYGSK	55	10	80
QQQHLFGSNVTDCSGNFCLFR	39	21	80
SAGWNIPIGLLYCDLPEPR	97	19	80
SASDLTWDNLK	84	11	80
SASDLTWDNLKGK	66	13	80
SCHTAVGR	44	8	80
SCHTGLGR	33	8	80
SDNCEDTPEAGYFAVAVVK	118	19	80
SETKDLLFR	38	9	80
SKEFQLFSSPHGK	72	13	80
SKEFQLFSSPHGKDLLFK	48	18	80
SMGGKEDLIWELLNQAQEHFGK	71	22	80
SVIPSDGPSVACVK	82	14	80
SVIPSDGPSVACVKK	36	15	80
TAGWNIPMGLLYNK	90	14	80
WCALSHHER	53	9	80
WCAVSEHEATK	67	11	80
YLGEEYVK	54	8	80
DGWHSWPIAHQWPQGPSAVDAAFSWEEK	36	28	21
DYFMPCPGR	29	9	21
EVGTPHGIILDSVDAAFICPGSSR	112	24	21
EWFWDLATGTMK 1Met(ox)	55	12	21
GDKVWVYPPEK	43	11	21
GDKVWVYPPEKK	52	12	21
GECQAEGVLFFQGDR	93	15	21
GGYTLVSGYPK	83	11	21
LHIMAGR	36	7	21
LLQDEFPGIPSPLDAAVECHR	57	21	21
LWWLDLK	38	7	21
LYLVQGTQVYVFLTK	77	15	21
NFPSPVDAAFR	50	11	21
QGHNSVFLIK	51	10	21
RLWWLDLK	46	8	21
SGAQATWTELPWPHEK	70	16	21
SLGPNSCSANGPGLYLIHGPNLYCYSDVEK	40	30	21
VDGALCMEK	55	9	21
VWVYPPEK	34	8	21
VWVYPPEKK	38	9	21
YYCFQGNQFLR	58	11	21
ASNLLLGFDR	70	10	16
DEQYLFLVR	56	9	16
ESAPGLIIATGSVGK	48	15	16
EYIVEYSR	33	8	16
GGTWEFLQAPAFTGYGEK	49	18	16
INCELSQGCSLHLAQR	27	16	16
ITTVSLSAPDALK	43	13	16
LEGELVPCPLAEENEFILYAVR	57	22	16
LSQLLNLQLR	65	10	16
NCPTTICDLDTQFR	65	14	16
NLLVNTLYTVR	53	11	16
SNVIVALAR	69	9	16
STVFTIFGSNK	51	11	16
VVVPYQGPSSDYVVVK	40	16	16
WESPYDSPDQDLLYAIAVK	43	19	16
YDLASGATEQLPLTGLR	35	17	16

AVEPQLQEEER	65	11	6
DPVASTSNLDMDFR	75	14	6
KVYDFLSTFITSGMR	88	15	6
SSVDELVGIDYSLMKDPVASTSNLDMDFR	122	29	6
TGLELSR	40	7	6
TMLQIGVMPMLNER 1Met(ox)	64	14	6
ADGEYWLGLQNMHLLTLK	65	18	3
DQDLFVQNCAALSSGAFWFR	61	20	3
FSTFDRDQDLFVQNCAALSSGAFWFR	44	26	3
AVGLAGTFR	36	9	10
DELLFPSWEALFSGSEGPLKPGAR	61	24	10
DFQPVLHLVALNSPLSGGMR	81	20	10
GADFQCFQQAR	40	11	10
IFSFDGK	40	7	10
IFSFDGKDVLR	27	11	10
LQDLYSIVR	75	9	10
LTESYCETWR	58	10	10
SVWHGSDPNGR	40	11	10
TEAPSATGQASSLLGGR	105	17	10
HVLFGTVGVPEHTYR	84	15	4
KHVLFGTVGVPEHTYR	78	16	4
VLYLSAFTSK	60	10	4
VTSLTACLVDQSLR	104	14	4
ATWSGAVLAGR	82	11	14
CEGPIPDVTFELLR	55	14	14
CLAPLEGAR	41	9	14
GVTFLLR	43	7	14
HQFLLTGDTQGR	65	12	14
LETPDFQLFK	41	10	14
LHDNQNGWSGDSAPVELILSDETLPAPEFSF	47	37	14
LLELTGPK	57	8	14
NGVAQEPVHLDSPAIK	71	16	14
SGLSTGWTQLSK	63	12	14
SLPAPWLSMAPVSWITPGLK	79	20	14
SWVPHTFESELSDPVELLVAES	42	22	14
TPGAAANLELIFVGPQHAGNYR	109	22	14
VTLTCVAPLSGVDFQLR	54	17	14
AADIEQQAVFAVFDENK	78	17	27
AEVDDVIQVR	54	10	27
ASEFLGYWEPR	32	11	27
AWAYYSAVNPEK	75	12	27
DIASGLIGLLLICK	97	14	27
DPDNIAAWYLR	56	11	27
DPPSDLLLLK	43	10	27
EDGILGPIIR	28	10	27
EFNPLVIVGLSK	55	12	27
EKPQSTISGLLGPTLYAEVGDIIK	38	24	27
ETDIEDSDDIPEDTTYK	64	17	27
ETDIEDSDDIPEDTTYKK	71	18	27
EVIITGIQTQGAK	90	13	27
GEYEEHLGILGPIIR	33	15	27
LAAALGIR	49	8	27
LEPEDEESDADYDYQNR	64	17	27
LLSLGAGEFK	53	10	27
LNNGGSYNAWSVEK	80	14	27
LSEGASYLDHTFPAEK	61	16	27

NFFNPPIISR	30	10	27
PGWWLLNTEVGENQR	81	15	27
QWLEIDLLK	42	9	27
SEAYNTFSER	57	10	27
SWYLEDNINK	30	10	27
TWNQSIALR	37	9	27
WIISSLTPK	29	9	27
WNILEFDEPTENDAQCLTR	99	19	27
DLFDPIIEDR	33	10	5
FCTGLTQIETLFK	63	13	5
GTGGVDTAAVGGVFDVSNADR	53	21	5
TDLNPDNLQGGDDLDPNYVLSSR	47	23	5
VLTPELYAELR	40	11	5
ECLCGALASYAAACAGR	97	17	6
GLWEQCQLLK	46	10	6
LLDLVFLLDGSSR	48	13	6
LPGLHNSLVK	36	10	6
YLFPGECQYVLVQDYCGSNPGTFR	32	24	6
YTLFQIFSK	36	9	6
ALALPPLGLAPLLNLWAK	27	18	7
DDWFMLGLR	40	9	7
IALGGLLFPASNLR	36	14	7
QAEISASAPTSLR	33	13	7
SCDVESNPGIFLPPGTQAEFNLR	59	23	7
TWDPEGVIFYGDTNPK	62	16	7
VVLSSGSGPGLDLPLVLGLPLQLK	47	24	7
ALPGTPVASSQPR	44	13	6
EVPLLQSLWLAHNEIR	29	16	6
LPGLPEGAFR	30	10	6
MDSNELTFIPR	39	11	6
SLQLNHNR	54	8	6
TVAAGALASLSHLK	78	14	6
AAFGQGSGPIMLDEVQCTGTEASLADCK	58	28	18
ASHEEVEGLVEK	45	12	18
AVDTWSWGER	71	10	18
ELSEALGQIFDSQR	93	14	18
GQWGTVCDNLWDLTDASVVCR	116	21	18
IDITLSSVK	66	9	18
IYTSPTWSAFVTDSSWSAR	136	19	18
KSQLVYQSR	37	9	18
LADGGATNQGR	89	11	18
LASAYGAR	40	8	18
RIDITLSSVK	65	10	18
SDLAVPSELALLK	49	13	18
SQLVYQSR	36	8	18
STHTLDLSR	50	9	18
TLQALEFHTVPFQLLAR	46	17	18
YSSDYFQAPSDYR	86	13	18
YYPYQSFQTPQHPSFLFQDK	33	20	18
YYPYQSFQTPQHPSFLFQDKR	28	21	18
DYEILFK	28	7	7
FFNVLTTNTDGK	58	12	7
IEFISTMEGYK	49	11	7
LDLTEKDYEILFK	38	13	7
NLDGISHAPNAVK	63	13	7
TAFYLAEFFVNEAR	66	14	7

YYIAASYVK	34	9	7
ALDENGHDLAVTFPGPGEDGLNPFLEVK	30	28	4
ELIDQYDVQR	38	10	4
IPAMVVDR	46	8	4
TVLQNWLK	35	8	4
AFNNVGEGIPLYESAVTR	56	18	17
DGTFLNLVSDDR	66	12	17
DGTFLNLVSDDRR	32	13	17
DVTVVSK	38	7	17
DVVASLVSTR	59	10	17
GMGPMSEAVQFR	69	12	17
GYAIGYGIGSPHAQTIK	50	17	17
HGPGVSTPDVAVR	94	13	17
HGSGESSAPLR	58	11	17
ITWADNSLPK	50	10	17
LIVAGLPR	29	8	17
QPLLLDDR	43	8	17
SDVTETLVSGTQLSQLIEGLDR	104	22	17
SGSAPQSPGASIR	39	13	17
TFTPFYFLVEPVDTLSVR	79	18	17
TIIVNWQPPSEANGK	60	15	17
TLSDVPSAAPQNLSLEVR	68	18	17
CSCASGFLLAADGK	76	14	12
CSCASGFLLAADGKR	35	15	12
EGETCGAEDNDSCGISLYK	73	19	12
FECPPNYVQVSK	36	12	12
GNEEGYFGTR	45	10	12
IGPAPAFTGDTIALNIIK	71	18	12
LNAYTGVVYLQR	55	12	12
QGSVTTFLAK	37	10	12
QGYQLAEDGHTCTDIDECAQGAGILCTFR	39	29	12
RVSEAEMAGR	35	10	12
SEFSQVASNTIPLPLPQPNTCK	43	22	12
TTCHDFLECQNSPAR	73	15	12
AVIHPDYDAASHDQDIMLLR	88	20	12
DSCQGDSGGPLVCGDHLR	83	18	12
EECEHAYPGQITQNMLCAGDI1Met(ox)	40	22	12
EKPGVYTNVCR	34	11	12
ESSQEQSSVVR	64	11	12
GLVSWGNIPCGSK	71	13	12
KPNLQVFLGK	58	10	12
LSELIQPLPLER	61	12	12
LVHGGPCDK	46	9	12
PNLQVFLGK	66	9	12
TADGDFPDTIQCAYIHLVSR	61	20	12
YTNWIQK	37	7	12
AGFSEDDYTALISQNILEGEK	113	21	4
ELQVPSEQVAFTVTAWDSQTAEK	33	23	4
LLVAQTSSPHSGHKPQK	35	17	4
VGADGTVFATR	98	11	4
FLYLGPFK	29	8	3
IVVEDVDEPPVFSK	41	14	3
LAYILQIR	47	8	3
DIQGSLQDIFK	77	11	8
GDIVTVVSPALLDR	41	14	8
INENTGSVSVTR	93	12	8

SIVVSPILIPENQR	54	14	8
TLFVHAR	37	7	8
TPHAEDMAELVIVGGK	43	16	8
VNSDGGLVALR	105	11	8
YEVSSPYFK	35	9	8
DYDPEDWLQVDAATGR	56	16	5
FSILQQGSPELFSIDELTGEIR	108	22	5
FTILEGDPDGQFTIR	78	15	5
TSLAEGAPPGTLVATFSAR	38	19	5
VSVQNEAPLQAAALR	66	15	5
ASLQFDLVK	34	9	3
NGESIASFFQFFGGWPK	29	17	3
VPPPSDAPLPFDR	28	13	3
IVIGLFGK	36	8	3
TPFVVEIADW	44	10	3
TVENFVALATGEK	77	13	3
EIVMTQSPPTLSLSPGER	30	18	2
LLIYGASTR	37	9	2
LFFLGQK	35	7	2
STLAPGLLWWDLAR	63	14	2
AIFSASIDDK	36	10	1
AELLVTEAPSKPITVTVEEQR	44	21	45
AHSSAGQQVAR	39	11	45
AQAGANTRPCPS	57	12	45
ATFSSVPLVASISAVSLEVAQPGPSNRPR	34	29	45
CAPGYYGNPSQGQPCQR	48	17	45
CEQCQPGYYGDAQR	63	14	45
CVASNAYGVAQSVVNLSVHGPPTVSVLPEG	56	35	45
DFISLGLQDGHLVFR	85	15	45
EVSEAVVDTLESEYLK	64	16	45
FDAGSGMATIR	58	11	45
FQGLDLNEELYLGGYPDYGAIPK	85	23	45
FSSGITGCVK	42	10	45
GMLEPVQRPDVVLVGAGYR 1Met(ox)	45	19	45
GMVFGIPDGVLELVPQR 1Met(ox)	34	17	45
GQTVTFTCVAIGVPTPIINWR	48	21	45
GSIQVDGEELVSGR	90	14	45
GSVYIGGAPDVATLTGGR	101	18	45
HLISTHFAPGDFQGFALVNPQR	27	22	45
HQTHGSLLR	30	9	45
IPGDQVVSVVFIK	66	13	45
IQVVVLSASDASPPPVK	41	17	45
LDVEFKPLAPDGVLLFSGGK	47	20	45
LGTVPQFPR	66	9	45
LLQVTPADSGEYVCR	31	15	45
LLSGPYFWSLPSR	53	13	45
LPAVEPTDQAQYLCR	60	15	45
LPQVSPADSGEYVCR	44	15	45
LVSEDPINDGEWHR	70	14	45
PGAPPPQPLDLQHR	71	14	45
RGSIQVDGEELVSGR	52	15	45
RPDGQPATR	33	9	45
SIEYSPQLEDAGSR	84	14	45
SLPEVPETIELEVR	58	14	45
SPAYTLVWTR	38	10	45
SPGPNVAVNAK	72	11	45

SPLPWQHR	30	8	45
SPVISIDPPSSTVQQGQDASFK	93	22	45
TSTASGLLLWQGVEVGEAGQGK	82	22	45
VGSSLPGR	43	8	45
VISSGSVASYVTSPQGFQFR	70	20	45
VTVTSEGGR	57	9	45
VVPYFTQTPYSFLPLPTIK	48	19	45
YELGSGLAVLR	74	11	45
YQLGSGEAR	53	9	45
YTLSYTAGPQGSPLSDPDVQITGNNIMLVAS(	59	40	45
VPAMDFYR 1Met(ox)	38	8	2
YEVSPVALQR	57	10	2
DYPDEVLQFAR	49	11	3
LFLGGLDALYSLR	65	13	3
VIALQAGGSAEPEEVVLEELQVFK	46	24	3
TNINFSLQGK	74	10	2
VSDSSNPFLNR	82	11	2
AALPYFPR	30	8	5
EVDYLETQNPALPCVEFDEK	54	20	5
FGGPAGLWTK	70	10	5
LDPQTLDTEQQWDTPCPR	82	18	5
QLYAWDDGYQIVYK	65	14	5
AIGPSQTHTIR	74	11	9
DQYYNIDVPSR	36	11	9
GFGGLTGQIVAALSTAK	112	17	9
GGSIQQYIYK	29	10	9
GLPNVVTSAISLPNIR	45	16	9
ITEVWGIPSPIDTVFTR	32	17	9
LVEVNPK	35	7	9
PALNYPVYGETTQVR	37	15	9
RPALNYPVYGETTQVR	38	16	9
DSDLFLLDTR	51	10	1
AQDAGVYQCLASNPVGTVVSR	112	21	26
AVVLWSK	37	7	26
DIGDTTIQLSWSR	76	13	26
EAAPSVAPSGLSGGGGAPGELIVNWTPMSR	41	30	26
ETIGDLTILNAQLR	90	14	26
EYQNGDGFGYLLSFR	75	15	26
FAQLNLAAEDTR	82	12	26
FGFLQEFSK	53	9	26
GFDNHSPIAK	39	10	26
GPPGPPGGVVVR	55	12	26
HFVSQTTGNLYIAR	39	14	26
IIVQAQPEWLK	60	11	26
LFAPSIK	33	7	26
LSLEDSGMYQCVAENK	91	16	26
NWIEIPVPEDIGHALVQIR	30	19	26
QGSTHWQTAR	44	10	26
RPPGNISWTFSSSSLSIK	38	18	26
TNPANIEGNAETAQVLGLTPWMDYEFR	47	27	26
TTGPGGDGIPAEVHIVR	62	17	26
VEVLAGDLR	57	9	26
VIASNILGTGEPSGPSSK	98	18	26
VISDTEADIGSNLR	113	14	26
VSGLHPNTK	45	9	26
VTVTPDGTLIIR	49	12	26

WLLNEFPNFIPTDGR	60	15	26
YTCMAQTVVDSASK	79	14	26
KDDPTLLSSGR	47	11	2
VIGIDHIK	41	8	2
AAAIQTMSLDAER	70	13	14
ATIVHQDQAYDDK	44	13	14
CISIYSSER	33	9	14
CLPDQQPIPTETFQVADR	58	18	14
DCENYITLLER	79	11	14
GESELYTSDTVMQNPQFIK	62	19	14
GYAPFSPDENSLVLFEGDEVYSTIR	96	25	14
IRGESELYTSDTVMQNPQFIK	42	21	14
LQDVFLLPDPSGQWR	48	15	14
MQASHGETFHVLYLTTDR	44	18	14
SVLQSINPAEPHK	39	13	14
TPLFHSK	37	7	14
VSLAPNSR	28	8	14
VYLFDFPEGK	64	10	14
AVPVAADQR	40	9	21
DRDGQPQVVR	39	10	21
ELPPHLGELTVAEETSSSLR	35	20	21
FDSFTVQYK	51	9	21
FLLYGLLGGK	35	10	21
FLLYGLSGR	56	9	21
GFEESEPLTGFLTTVPDGPTQLR	51	23	21
HGPLVAEAK	46	9	21
LGPISADSTTAPLEK	58	15	21
LGVLTVTDTTPDSMR 1Met(ox)	43	15	21
LNWEAPPGAFDSFLLR	47	16	21
LQVVPVAANQR	51	11	21
LSWTVAQGPFDSFVVQYR	36	18	21
LSWTVPEGQFDSFVVQFK	35	18	21
MHLYGLHEGR	31	10	21
MNLYGFHGGQR	41	11	21
SPVSVEAK	36	8	21
VGGEESEVTVGGLEPGR	36	17	21
VGPISAVAITAGR	32	13	21
VGPVSAVGVTAPEEESPDAPLAK	76	23	21
YEVTVVSVR	32	9	21
ENNAVYAFLGLTAPPGSK	46	18	5
ENVPENSRPATGYPLPPQIFNESQYR	28	26	5
GDYDAFFEAR	53	10	5
QQDVLGFLEANK	45	12	5
VYIASSSGSTAIK	64	13	5
GSFALSFPVESDVAPIAR	54	18	5
LEAGINQLSFPLSSEPIQGSYR	95	22	5
MFIFAILPDGEVVGDSEK 1Met(ox)	30	18	5
NELIPLIYLENPR	28	13	5
SLFTDLVAEK	41	10	5
DFADIPNLR	44	9	12
DRIEEIR	36	7	12
EKETVIIPNEK	62	11	12
ESAYLYAR	39	8	12
LDFTGNLIEDIEDGTFSK	94	18	12
LEGNPIVLGK	68	10	12
LPVLPPK	34	7	12

LSLLEELSLAENQLLK		112	16	12
LTLFNAK		27	7	12
RLDFTGNLIEDIEDGTFSK		89	19	12
RLPIGSYF		41	8	12
VIHLQFNNIASITDDTFCK		56	19	12
AEAAAPYTVLAQSAPR		73	16	8
AGPDLASCLDVDECR		90	15	8
DGGCSLPILR		28	10	8
GCQLCPPFGSEGFR		45	14	8
GGYTCVCPDGFLLDSSR		72	17	8
GSACEEDVDECAQEPPPCGPGR		96	22	8
GYLAPSGDLSLR		38	12	8
VSAPDGPCPTGFER		69	14	8
LLSQNPPSQIFQSLSGNSR		90	19	2
VLYDPFLPPLR		28	11	2
LAADDPEVR		42	9	1
IFVFLEHQTK		42	10	2
SFTILLSNTENQEK		68	14	2
DTANWLEINPDTGAISTR		81	18	2
GQVPENEANVVITTLK		29	16	2
AILQSGSFNAPWAVTSLYEAR		117	21	2
IFFPGVSEFGK		41	11	2
CPVPSFHVELCR		50	12	8
DLEEDPYLPGNPR		29	13	8
ELIAYSQYPR		52	10	8
LFPYLDPFDSASQLMEPGR	1Met(ox)	52	19	8
LYDFNSYWR		35	9	8
QSTEQAIQLLEK		71	12	8
QSYFASVSYLDTQVGR		90	16	8
VSFLTGR		38	7	8
EQVMDTLVR		63	9	1
AQQEQELAADAFK		79	13	3
LWEEQLAAAK		46	10	3
SLEDQVEMLR		60	10	3
FLVGPDGIPIMR	1Met(ox)	48	12	7
LFWEPMK		35	7	7
MDILSYMR		52	8	7
NSCPPTSELLGTSDR		72	15	7
PGGGFVPNFQLFEK		46	14	7
QEPGENSEILPTLK		90	14	7
YVRPGGGFVPNFQLFEK		50	17	7
ETALLIDPK		39	9	8
FALEVAAK		33	8	8
LGLQNDLFSLAR		83	12	8
LSVEGFAVDK		60	10	8
VALSNMNVIDR		73	11	8
VLGATLLPDLIQK		63	13	8
VTLSFPSTLQTGTGTLK		38	17	8
YAAVTQFEATDAR		62	13	8
ALASGGSALDAVESGCAMCER		123	21	5
FLPSYQAVEYMR	1Met(ox)	63	12	5
RGEDPTIACQK		52	11	5
TGHIAAGTSTNGIK		53	14	5
VGDSPIPGAGAYADDTAGAAAATGNGDILMF		87	31	5
NIIHGSDSVK		32	10	1
AGALNSNDAFVLK		75	13	48

AGKEPGLQIWR		58	11	48
AMAELAA	1Met(ox)	40	7	48
AQPVQVAEGSEPDGFWEALGGK		96	22	48
ATEVPVSWESFNNGDCFILDLGNNIHQWCG		32	35	48
AVEVLPK		43	7	48
DPDQTDGLGLSYLSSHIANVER		107	22	48
DSQEEEKTEALTSAK		80	15	48
EGGQTAPASTR		51	11	48
EPAHLMSLFGGK		31	12	48
EPAHLMSLFGGKPMIIYK	1Met(ox)	34	18	48
EPGLQIWR		55	8	48
EVQGFESATFLGYFK		102	15	48
FDLVPVPTNLYGDFFTGDAYVILK		48	24	48
GASQAGAPQGR		68	11	48
GGVASGFK		45	8	48
GIRDNER		32	7	48
HVVPNEVVVQR		73	11	48
IEGSNKVPVDPATYGQFYGGDSYIILYNYR		49	30	48
KGGVASGFK		76	9	48
KMDAHPPR		60	8	48
LFACSNK		36	7	48
LKATQVSK		39	8	48
MDAHPPR		41	7	48
NWRDPDQTDGLGLSYLSSHIANVER		75	25	48
PALPAGTEDTAK		64	12	48
PALPAGTEDTAKEDAANR		98	18	48
PALPAGTEDTAKEDAANRK		62	19	48
PNSMVVEHPEFLK		73	13	48
QGQIIYNWQGAQSTQDEVAASAILTAQLDEE		75	40	48
QTQVSVLPEGGETPLFK		81	17	48
RTPITVVK		29	8	48
RYIETDPANR		30	10	48
SEDCFILDHGK		68	11	48
SEDCFILDHGKDGK		59	14	48
TASDFITK		48	8	48
TEALTSAK		30	8	48
TGAQELLR		77	8	48
TPITVVK		49	7	48
TPSAAYLWVGTGASEAEK		117	18	48
VEKFDLVPVPTNLYGDFFTGDAYVILK		41	27	48
VHVSEEGTEPEAMLQVLGPK		91	20	48
VHVSEEGTEPEAMLQVLGPKPALPAGTEDT/		55	32	48
VPEARPNSMVVEHPEFLK		27	18	48
VPFDAATLHTSTAMAAQHGMDDDGTGQK		54	28	48
VPVDPATYGQFYGGDSYIILYNYR		79	24	48
VSNGAGTMSVSLVADENPFAQGALK		108	25	48
YIETDPANR		48	9	48
VSGSQIVDIDK		68	11	1
DHNCHNLPEGVADLTQIDVNVQDHFWDGK		27	29	4
DVFFGPK		32	7	4
GCEMICYCNFSELLCCPK		98	18	4
ISFVIPCNNQ		29	10	4
ALEGLQYPFAVTSYGK		65	16	12
EDLSPSITQR		35	10	12
ESHPGLFPPTFGAVAPFLADLDTTDGLGK		37	29	12
GNLYWTDWNR		45	10	12

MVYWTDITEPSIGR	1Met(ox)	38	14	12
NIFWTDSNLDR		38	11	12
QAEVTFVGHPGNLVIK		29	16	12
QDLGSPEGIAVDHLGR		54	16	12
QELFPFGPGQGDLELEDGDDFVSPALELSG/		33	33	12
SNGAYNIFANDR		69	12	12
VIIGLAFDCVDK		44	12	12
VLFETDLVNPR		50	11	12
DGMWEAFQDGEK		46	12	4
FDATQAFVGELSQFNIWDR		76	19	4
TESTLNALLQR		42	11	4
VAQLPLFVSDGK		35	12	4
SPTFAGGLFSISK		66	13	3
TVVVSPDIVTIDLNTFEFAK		88	20	3
TVYSVLHTTPAILLK		50	15	3
GSDPVTIFLR		56	10	3
VATLSTLLFK		59	10	3
VPGDVSLQLSTLEMDDR	1Met(ox)	54	17	3
AFGAPVPSVQWLDEDGTTVLQDER		105	24	8
AQLLVVGSPGPVPR		78	14	8
DATQITQGPR		64	10	8
DLQELGDSDKYFIEDGR		36	17	8
LVLSDLHLLTQSQVR		61	15	8
VKDATQITQGPR		90	12	8
VQAVNSQGK		34	9	8
YGPGEPSPVSETVVTPEAAPEK		27	22	8
AVSEKEVDSGNDIYGNPIKR		28	20	14
CPMIPCYISSPDECLWMDWV	1Met(ox)	62	23	14
DIEFIYTAPSSAVCGVSLDVGGK		83	23	14
DIEFIYTAPSSAVCGVSLDVGGKK		68	24	14
EVDSGNDIYGNPIK		86	14	14
EVDSGNDIYGNPIKR		83	15	14
EYLIAGK		45	7	14
GAAPPKQEFLDIEDP		42	15	14
GPEKDIEFIYTAPSSAVCGVSLDVGGK		44	27	14
KEYLIAGK		56	8	14
MHITLCDFIVPWDTLSTTQK		50	20	14
NINGHQAK		28	8	14
QEFLDIEDP		49	9	14
SDGSCAWYR		62	9	14
FLLEYIAPMTEK		54	12	4
IFQNLDGALDEVVLK		64	15	4
LVAEWEGQDSDSDQLFYTK		59	19	4
SEDYVDIVQGR		39	11	4
ELISNASDALDK		60	12	6
FAFQAEVNR		63	9	6
GVVDSDDLPLNVSR		88	14	6
IKEDEDDKTVLDLAVVLFETATLR		30	24	6
IYFMAGSSR		54	9	6
SILFVPTSAPR		41	11	6
CNPGTGQCVCPAGWVGEQCQHCGGR		37	25	13
DLDMFINASK	1Met(ox)	57	10	13
IMQSSQSMSK	1Met(ox)	44	10	13
IYMYGGK		31	7	13
KVEFVLK		41	7	13
LADDLYR		39	7	13

LTGSSGFVTDGPGNYK	109	16	13
LTLTPWVGLR	52	10	13
NHNALLASLTTQK	77	13	13
SCALDQNCQWEPR	77	13	13
YDVDTQMWTILK	95	12	13
YGHSLALYK	34	9	13
YYTAINFVATPDEQNR	33	16	13
APQTVELPAVAGHTLTAR	65	18	32
CESCLQGYFLLDGK	77	14	32
CGSGGPGSCPVPQECVPQDGAAGAGLCR	55	28	32
CMEGGLSGPR 1Met(ox)	36	10	32
FHVELAAPSPELYSLHCPDR	38	20	32
FLDTGVVQSDR	75	11	32
GAMYLLGGLTAGGVTR 1Met(ox)	83	16	32
GFIYPMLPGGPGGPGAEDVAVWTR	70	24	32
GNSHICISR	35	9	32
GPDTENMEEVGR 1Met(ox)	73	12	32
GPESCSLGCAQATQCALCLR	120	20	32
HSECAGVGAR	39	10	32
KYSLDPEEIENWVTEGPSEDEAVCVNCQNN	33	35	32
LDGGQLVWETLMDSR	45	15	32
LFHASALLGDTMVVLGGR	84	18	32
LFPLPGR	33	7	32
LGCGGSPCSPMPR	32	13	32
LLGDCQACLAFSSPTAPPR	46	19	32
LSADTASR	42	8	32
LYISGGFGGVALGR	98	14	32
SASVGPPMEESVAHAVAAVG 1Met(ox)	48	22	32
SDPDEFSSDVLLYQVNCNAWLLPDLTR	53	27	32
SFHAAAYVPAGR	35	12	32
SLIAAFCGQR	55	10	32
SSSCTSYSSCLGCLADQGCGWCLTSATCHL	38	31	32
STTITLTPSAETDVSLVYR	108	19	32
TGVPGGSEISFFFLEPYR	59	18	32
TLQPGDGEASTPR	67	13	32
TWSLLAPSQGAK	51	12	32
WCTNCPEGACIGR	51	13	32
WTQMLAGAEDGGPGPSPR	97	18	32
WVAHQEK	33	7	32
ATAVVDGAFK	58	10	5
EGGLGPLNIPLLADVTR	57	17	5
GLFIIDGK	30	8	5
IGKPAPDFK	27	9	5
KEGGLGPLNIPLLADVTR	33	18	5
FAYGYIEDLK	29	10	4
IEEQLTLEK	49	9	4
LGVQDLFNSSK	56	11	4
TYNFLPEFLVSTQK	35	14	4
ASISGGGLPAPYQAK	71	15	6
EQILAFSQK	32	9	6
FFFSGYDK	29	8	6
QSPINIVTTK	49	10	6
VVWTVFR	35	7	6
YLGSLTTPTCDEK	46	13	6
EENFYVDETTVVK	41	13	7
GLASANVDFAFSLYK	91	15	7

GTWTQPFDLASTR	55	13	7
HLVALSPK	43	8	7
MNTVIAALSR	73	10	7
SETEIHQGFQHLHQLFAK	33	18	7
WSAGLTSSQVDLYIPK	79	16	7
GENSWFSTQVDTVATK	69	16	2
GQSEDPGSLLSLFR	46	14	2
DAVEDLESVGK	75	11	2
ENAGEDPGLAR	57	11	2
DGQLEVILDR	60	10	10
EAVVVDYGVR	70	10	10
FVVLFNPLEQER	33	12	10
GAEVLYSLAAAHAR	64	14	10
GQKPELQMLTVSEELPFDNVDGGVWR	36	26	10
SQISVLQNR	47	9	10
TLQAEEDTLPSAETALILHR	53	20	10
TVIQLDSSPR	36	10	10
VIDSGTSDFALSNR	97	14	10
YMQVWFSGLTGLLK	37	14	10
AFQVWSDVTPLR	66	12	10
CGNPDVANYNFFPR	54	14	10
ELAVQYLNTFYGCPK	65	15	10
ESCNLFVLK	34	9	10
FFGLPQTGDLDQNTIETMR	90	19	10
FPFLFNGK	41	8	10
GEIFFFK	30	7	10
IIGYTPDLDPETVDDAFAR	85	19	10
QDIVFDGIAQIR	83	12	10
TDKELAVQYLNTFYGCPK	39	18	10
AGLAASLAGPHSIVGR	81	16	6
AVVVHAGEDDLGR	55	13	6
GGNQASVENGNAGR	73	14	6
LACCVVGVCGPGLWER	87	16	6
VTEIWQEVMQR	100	11	6
VTGVVLFR	45	8	6
ALGGDLASINNK	75	12	12
DSTFSAWTGLNDVNSEHTFLWTDGR	35	25	12
DYQYYFSK	32	8	12
GEDLFFNYGNR	45	11	12
GTFQWTIEEEVR	44	12	12
NFGDLVSIQSESEK	71	14	12
SQGPEIVEVEK	49	11	12
TGIAGGLWDVLK	73	12	12
TNFWIGLFR	46	9	12
WVSESQIMSVAFK	31	13	12
YFWTGLSDIQTK	74	12	12
YTNWAADEPK	47	10	12
DFYVVEPLAFEGTPEQK	61	17	6
GLETFSQLVWK	50	11	6
GVQAQPLNVGFCEQEFEQT	42	19	6
GYVVWQEVFDNK	32	12	6
QLESFYIQTLLDIVSSYGK	51	19	6
YVLYPNNFQFQYDVSSAAQPGCSVLDEAFQ	40	31	6
SSTGPGEQLR	53	10	1
ALDLSLK	34	7	5
DYIFGNYIER	42	10	5

EHLVQATPENLQEAR		50	15	5
EVSFDVELPK		30	10	5
LWAYLTIEQLLEK		38	13	5
LVQTAELTK		36	9	1
NAFVFLQYDK		51	10	2
SFFEFLVLNK		48	10	2
NANTFISPQQR		61	11	1
EDTSPAVLGLAAR		73	13	10
ESYNVQLQLPAR		36	12	10
ETQYVDYDFPTDFPAIAPFLADIDTSHGR		44	29	10
FSNLYVGTNGIISTQDFPR		76	19	10
GEADDLKSEGPYFSLTSTEQSVK		32	23	10
HSGQFTDEYLPEQR		53	14	10
ILINTDIGLPNGLTFDPFSK		63	20	10
ITQTAEGLDPENYLSIK		48	17	10
LANPLHFYEAR		49	11	10
VLFYTDLVNPR		49	11	10
EGAMSAQLGYPVVGWHIANK		27	20	9
GVHYISVSATR		58	11	9
KVVENGALLSWK		37	12	9
LGCSLNQNSVPDIHGVEAPAR		88	21	9
LVPVVNNR		42	8	9
SFSEVELHNMK		52	11	9
TASPDPGEVVSSACAADEPVTVLTVILDADL		74	33	9
VTIPTDLIASSGDIIK		112	16	9
VVENGALLSWK		65	11	9
GITEPPFGIFVFNK		50	14	3
ILDVNDNIPVVENK		34	14	3
IVSLEPAYPPVFYLNK		32	16	3
DLDIFDR		32	7	2
LSVQGEVSTFTGK		32	13	2
DDFLIYDR		48	8	4
DMPASEDLQDLQK		42	13	4
LPTDSELAPR		71	10	4
QPPAWSIR		28	8	4
EALVPLVADHK		47	11	4
TTDVTQTFGIEK		83	12	4
VANYVDWINDR		69	11	4
VQLSPDLLATLPEPASPGR		92	19	4
APAVAEENPK		72	10	8
CPGCGQGVQAGCPGGCVEEEDGGSPAEG		59	37	8
CPGCGQGVQAGCPGGCVEEEDGGSPAEG		28	38	8
GAQTLYVPNCDHR		55	13	8
GPCWCVDR		49	8	8
HLDSVLQQLQTEVYR		112	15	8
RGPCWCVDR		29	9	8
SLPGSPDGNGSSSCPTGSSG		37	20	8
ALSMCPPSPLGCELVK	1Met(ox)	39	16	2
HMEASLQELK	1Met(ox)	28	10	2
AFPALTSLDLSDNPGLGER		111	19	14
ATVNPSAPR		48	9	14
ELTLEDLK		39	8	14
FPAIQNLALR		67	10	14
GLMAALCPHK		48	10	14
ITGTMPPLPLEATGLALSSLR	1Met(ox)	84	21	14
LKELTLEDLK		44	10	14

LTVGAAQVPAQLLVGALR	72	18	14
RVDADADPR	36	9	14
STLSVGVSGTLVLLQGAR	59	18	14
SWLAELQQWLK	50	11	14
SWLAELQQWLKPGLK	65	15	14
VLDLSCNR	47	8	14
VLSIAQAHSPAFSCEQVR	61	18	14
ENFAILTIDGDEASAVR	68	17	10
HELQHPIIAR	42	10	10
LLNTPDGSPYTWWVGK	51	16	10
LNQDLFSVSFQFR	68	13	10
MSQIDISSGSGLNDGQWHEVR	41	21	10
TWNPNGLLVFSHFADNLGNVEIDLTESK	31	28	10
VDNAPDQQNSHPDLAQEEIR	29	20	10
VQFNQIAPLK	46	10	10
YNTPGFTGCLSR	36	12	10
YSSSDWVTQYR	94	11	10
ADQVCINLR	59	9	17
CVNHYGGYLCLPK	53	13	17
DIDECDIVPDACK	72	13	17
EHIVDLEMLTVSSIGTFR 1Met(ox)	45	18	17
FSCMCPQGYQVVR	46	13	17
GEQCVDIDECTIPPYCHQR	35	19	17
GSFACQCPPGYQK	84	13	17
IPSNPSHR	54	8	17
IQCAAGYEQSEHNVCQDIDECTAGTHNCR	50	29	17
LNCEDIDECR	70	10	17
NPADPQR	31	7	17
NPCQDPYILTPENR	48	14	17
QTSPVSAMLVLVK	47	13	17
RNPADPQR	37	8	17
SGNENGEFYLR	57	11	17
SVPSDIFQIQATTIYANTINTFR	92	23	17
TCQDINECETTNECR	114	15	17
DWEYIQDPR	40	9	4
SPVCMEFQYQATGGR	58	15	4
VFQANNDATEVVLNK	89	15	4
YDWLDIWDGIPHVGPLIGK	27	19	4
FNTFIHEDIWNIR	41	13	4
KMTLYHCK	27	8	4
QHVHPEETGGSDR	33	13	4
RVVIACEGNPQVPVHFDG	32	18	4
CALEDETYADGAETEVDCNR	88	20	5
LDSSEFLK	28	8	5
LSFQEFLK	47	8	5
SVSPSASPVVCYQSNR	75	16	5
YVQELQK	29	7	5
AGEQVTYTCATYYK	85	14	47
AQTTVTCMENGWSPTPR	89	17	47
AVYTCNEGYQLLGEINYR	93	18	47
CFEGFGIDGPAIAK	51	14	47
CTSTGWIPAPR	63	11	47
DGWSAQPTCIK	42	11	47
DTSCVNPPTVQNAYIVSR	62	18	47
ECDTDGWTNDIPICEVVK	69	18	47
EFDHNSNIR	52	9	47

EIMENYNIALR		66	11	47
EQVQSCGPPPELLNGNVK		72	18	47
EYHFGQAVR		42	9	47
FVCNSGYK		29	8	47
GDAVCTESGWRPLPSCEEK		65	19	47
GEWVALNPLR		51	10	47
HGGLYHENMR		71	10	47
IDVHLVPDR		51	9	47
IDVHLVPDRK		44	10	47
IEGDEEMHCSDDGFWSK		45	17	47
IIYKENER		42	8	47
IVSSAMEPDR		79	10	47
IVSSAMEPDREYHFGQAVR		53	19	47
KGEWVALNPLR		56	11	47
KGEWVALNPLRK		44	12	47
LGYVTADGETSGSITCGK		113	18	47
LSYTCEGGFR		34	10	47
NDFTWFK		31	7	47
NGQWSEPPK		38	9	47
NTEILTGSWSDQTYPEGTQAIYK		97	23	47
RPYFPVAVGK		41	10	47
SCDNPYIPNGDYSPLR		31	16	47
SIDVACHPGYALPK		62	14	47
SITCIHGVWTQLPQCVAIDK		43	20	47
SLGNVIMVCR	1Met(ox)	61	10	47
SPDVINGSPISQK		65	13	47
SPPEISHGVVAHMSDSYQYGI	1Met(ox)	28	26	47
SSIDIENGFISESQYTYALK		82	20	47
SSNLIILEEHLK		71	12	47
SSQESYAHGTK		48	11	47
TDCLSLPSFENAIPMGEK	1Met(ox)	28	18	47
TGDEITYQCR		67	10	47
TGESVEFVCK		67	10	47
TGESVEFVCKR		64	11	47
TKEEYGHSEVVEYYCNPR		53	18	47
TKNDFTWFK		37	9	47
VSVLCQENYLIQEGEEITCK		81	20	47
WSSPPQCEGLPCK		35	13	47
AHSDGGDGVVSQVK		51	14	43
APPSQPPR		65	8	43
ATSVALTWSR		62	10	43
AVDLIPWMEYEFR		58	13	43
DAGIYYCLASNNYGMVR	1Met(ox)	83	17	43
DETMSPSTAFQVK		58	13	43
DGEYVVEVR		67	9	43
ELTITWAPLSR		44	11	43
FIPLIPIPER		33	10	43
FVSQTNGNLYIANVEASDK		122	19	43
GDGPYSLVAVINSAQDAPSEAPTEVGVK		81	28	43
GEPSIPSNR		49	9	43
GFGPIFEEQPINTIYPEESLEGK		80	23	43
GMVLLCDPPYHFPDDLSYR		36	19	43
GPPGPPGGLR		28	10	43
HSIEVPIPR		43	9	43
IFNIQLEDEGIYECEAENIR		99	20	43
IKTDGAAPNVAPSDVGGGGGR		87	21	43

ILALAPTFEMNPMK		63	14	43
IVESYQIR		60	8	43
KAPPSQPPR		53	9	43
KVLEPMPSTAEISTSGAVLK		48	20	43
KVTVTNPDTGR		66	11	43
MNNGDVDLTSDR		94	12	43
NDGGIYTCFAENNR		85	14	43
NFMLDSNGELLIR	1Met(ox)	51	13	43
STEATLSFGYLDPFPPEER		37	19	43
STEATLSFGYLDPFPPEERPEVR		51	23	43
TDGAAPNVAPSDVGGGGGR		65	19	43
TDPPIIEGNMEAAR		60	14	43
TILSDDWK		44	8	43
TTKPYPADIVVQFK		56	14	43
VIIECKPK		29	8	43
VLEPMPSTAEISTSGAVLK		100	19	43
VLYRPDGQHDGK		39	12	43
VQVTSQEYSAR		66	11	43
VTVTNPDTGR		52	10	43
VVATNTLGR		49	9	43
WLLNEFPVFITMDK	1Met(ox)	63	14	43
YGHGVSEEDKGFGPIFEEQPINTIYPEESLEC		28	33	43
YSMVGGNLVINNPDK		82	15	43
YTCTAQTIVDNSSASADLVVR		100	21	43
YWAAHDKEEAANR		87	13	43
DFLQSLK		28	7	15
EDFLEQSEQLFGAK		35	14	15
EQQDSPGNKDFLQSLK		51	16	15
GDKLFGPDLK		29	10	15
GFPIKEDFLEQSEQLFGAK		74	19	15
GFPIKEDFLEQSEQLFGAKPVSLTGK		49	26	15
GISEQSLVVSGVQHQSTLELS	1Met(ox)	34	37	15
HQMDLVATLSQLGLQELFQAPDLR		57	24	15
LCQDLGPGAFR		43	11	15
LGNQEPGGQTALK		47	13	15
NKFDPSLTQR		33	10	15
NPNPSAPR		41	8	15
QEDDLANINQWVK		69	13	15
QLTSGPNQEQVSPLTLLK		67	18	15
WFLLEQPEIQVAHFPFK		28	17	15
IPGMVVDR		50	8	3
MSPINMLYFNDK		63	12	3
YPHTHLVQQANPR		53	13	3
LYLDPGLR		34	8	2
QVGYEDQWLQLLR		53	13	2
DGLIPLEIR		29	9	6
FLHNPDAAQGFVGCALSSTIQR		76	22	6
GGFVLLDGETFEVK		69	14	6
HEIVQTLSLK		46	10	6
HNVMISTEWAAPNVLR	1Met(ox)	51	16	6
LVLPSLISSR		64	10	6
AFLLTPR		32	7	3
EFPEVHLGQWYFIAGAAPTK		32	20	3
SLTSCLDSK		37	9	3
LLDDNGNIAEELSILK		115	16	2
WNTDSVEEFLSEK		57	13	2

ADMDQFTASISETPVDVR	1Met(ox)	60	18	9
CLVGEFVSDVLLVPEK		78	16	9
EMIFNAER		29	8	9
EWEEAELQAK		46	10	9
GSGVGEQDGGLIGAEEK		124	17	9
HYQHVLAVDPEK		51	12	9
QQLVETHLAR		31	10	9
VPYVAQEIQEEIDELLQEQR		138	20	9
VSIDNWCR		50	8	9
AASQPGELKDWFVGR		63	15	13
EALVSVWLQCTAISR		50	15	13
EGGPNNHLLK		34	10	13
ELLVIELSDNPGVHEPGEPEFK		35	22	13
IHIMPSLNPDGFEK		35	14	13
LLIPGNYK		36	8	13
LQQEDGISFEYHR		87	13	13
LTASAPGYLAITK		66	13	13
NSLISYLEQIHR		63	12	13
SGSAHEYSSSPDDAIFQSLAR		108	21	13
SNAQGIDLNR		67	10	13
VAVPYSPAAGVDFELESFSER		97	21	13
YIGNMHGNEAVGR		58	13	13
EELLLLQSTAEQLR		88	14	3
VAELEHGSSAYSPPDAFK		81	18	3
VAQLPLSLK		37	9	3
DGSGVGVFK		34	9	1
FVSISDLLVPK		30	11	4
SPYLYPLYGLGELPQGFAR		56	19	4
TDDYLDQPCYETINR		59	15	4
VPSTEAEALASSLMGLFEK		30	19	4
SFTIWLSDK		43	9	3
VLNDGSVYTAR		85	11	3
VNLEECFR		44	8	3
LDFGNSQGK		59	9	3
NPSAAFFCVAR		40	11	3
VSEEIEDIIK		42	10	3
IENIDHLGFFIYR		42	13	3
LTAYLDLNLDK		46	11	3
NLLELLINIK		40	10	3
ADGESCSASMMYQEGK	1Met(ox)	38	16	21
AFLEVNEEGSEAAASTAVVIAGR		129	23	21
DDLYVSDAFHK		54	11	21
EQLQDMGLVDLFSPEK		90	16	21
EVPLNTIIFMGR		67	12	21
FATTFYQHLADSK		79	13	21
FDTISEK		32	7	21
FRIEDGFSLK		55	10	21
GDDITMVLILPKPEK	1Met(ox)	36	15	21
IEDGFSLK		40	8	21
ITDVIPSEAINELTVLVLVNTIYFK		84	25	21
LPGIVAEGR		53	9	21
LQPLDFKENAEQSR		52	14	21
NDNDNIFLSPLSISTAFAMTK		91	21	21
RVAEGTQVLELPFK		27	14	21
RVWELSK		29	7	21
SKLPGIVAEGR		41	11	21

SKLPGIVAEGRDDLYVSDAFHK		33	22	21
TSDQIHFFFAK		39	11	21
VAEGTQVLELPFK		74	13	21
VANPCVK		32	7	21
AAMVGMLANFLGFR		71	14	18
ADSQAQLLLSTVVGVFTAPGLHLK		96	24	18
ALQDQLVLVAAK		92	12	18
ANAGKPKDPTFIPAPIQAK		43	19	18
DPTFIPAPIQAK		69	12	18
FMQAVTGWK		62	9	18
LDAHKVLSALQAVQGLLVAQGR		60	22	18
LDTEDKLR		71	8	18
LQAILGVPWK		73	10	18
PKDPTFIPAPIQAK		79	14	18
QPFVQGLALYTPVVLPR		105	17	18
SLDFTELDVAAEK		98	13	18
SLDFTELDVAAEKIDR		47	16	18
TIHLTMPQLVLQGSYDLQDLLAQAELPAILHT		47	38	18
VANPLSTA		28	8	18
VEGLTFQQNSLNWMK		71	15	18
VGEVLNSIFFELEADER		125	17	18
VLSALQAVQGLLVAQGR		91	17	18
AACAQLNDFLQEYGTQGCQV		150	20	2
DDPDAPLQPVTPLQLFEGR		52	19	2
IITDFPSLTR		29	10	2
SLEDVVIDIQSSLSK		55	15	2
ADNFLLENTLPAQSTFTLAISAYALSLGDK		39	30	33
ALVEGVDQLFTDYQIK		58	16	33
ATLLDIYK		40	8	33
DINYVNPVIK		46	10	33
DSEITFIK		46	8	33
DSSVPNTGTAR		38	11	33
DVFLEMNIPYSVVR		39	14	33
ELSYYSLEDLNNK		72	13	33
ENSQYQPIK		28	9	33
ESYSGVTLDPR		46	11	33
FQNSAILTIQPK		66	12	33
GEQIQLK		40	7	33
GGSASTWLTAFALR		81	14	33
GIYGTISR		29	8	33
IDTALIK		39	7	33
IDTQDIEASHYR		74	12	33
IPLDLVPK		39	8	33
KIEEIAAK		36	8	33
LNLVATPLFLKPGIPYPIK		41	19	33
LQGTLPVEAR		47	10	33
LSMDIDVSYK	1Met(ox)	55	10	33
MVETTAYALLTSLNLK	1Met(ox)	60	16	33
NADYSYSVWK		52	10	33
QCTMFYSTSNIK		41	12	33
QLPGGQNPVSYVYLEVVSK		51	19	33
SDLGCGAGGGLNNANVFHLAGLTFLTNANAI		46	41	33
SIVSALKR		38	8	33
TSTSEEVCSFYLK		81	13	33
VFQFLEK		39	7	33
VLGQVNK		31	7	33

VSITSITVENVFVK	67	14	33
YGGGFYSTQDTINAIEGLTEYSLLVK	64	26	33
YVEQNQNSICNSLLWLVENYQLDNGSFK	41	28	33
EPGLCTWQSLR	36	11	8
FVYTPAMESVCGYFHR	56	16	8
GFQALGDAADIR	66	12	8
HLACLPR	33	7	8
LQDGLLHITTCSFVAPWNSLSLAQR	67	25	8
LQSGTHCLWTDQLLQGSEK	77	19	8
SEEFLIAGK	49	9	8
TYTVGCEECTVFPCLSIPCK	67	20	8
AFCSFQIYAVPWQGTMTLSK	64	20	14
AFCSFQIYAVPWQGTMTLSKSTCQDA	50	26	14
ALDFAVGEYNK	75	11	14
ARKQIVAGVNYFLDVELGR	37	19	14
ASNDMYHSR	69	9	14
KAFCSFQIYAVPWQGTMTLSK 1Met(ox)	49	21	14
KAFCSFQIYAVPWQGTMTLSKSTCQDA	29	27	14
KQIVAGVNYFLDVELGR	133	17	14
LVGGPMDASVEEEGVR	94	16	14
LVGGPMDASVEEEGVRR	45	17	14
QIVAGVNYFLDVELGR	106	16	14
RALDFAVGEYNK	48	12	14
TQPNLDNCPFHDQPHLK	60	17	14
TQPNLDNCPFHDQPHLKR	29	18	14
GLQYAAQEGLLALQSELLR	87	19	3
LAEGFPLPLLK	31	11	3
SPVTLLAAVMSLPEEHNK	44	18	3
AATGECTATVGK	45	12	6
ENFLFLTPDCK	55	11	6
FSVATQTCQITPAEGPVVTAQYDCLGCVHPI	44	43	6
LGQSLDCNAEVYVVPWEK	83	18	6
QVVAGLNFR	53	9	6
TVGSDTFYSFK	57	11	6
DFNLLQVSEPSEPCVR	47	16	2
VAQVDSLKDK	46	10	2
DAVQALQEAQGR	27	12	3
DFEAQAAAQAR	33	11	3
LAGLDAGLHQLHVR	34	14	3
IEEDSEVLMMIK 2Met(ox)	73	12	4
LAACVNLIPQITSIYEWK	60	18	4
SVHPYEVAEVIALPVEQGNFPYLQWVR	32	27	4
TQSSLVPALTDFVR	55	14	4
VEVVLLHGK	37	9	1
TWASPVVTPGAR	27	12	1
ASVGQDSPEPR	53	11	8
DLWVNIDQMEK	45	11	8
DLWVNIDQMEKDK	43	13	8
EIPVLVTQISSTNHPVK	54	17	8
IHGILSNTHR	31	10	8
IYGPSDSASR	37	10	8
MLTATQYIAPLMANFDPSVSR	52	21	8
VGLSDAFVVVHR	97	12	8
ESDVPLKTEEFEVTK	51	15	4
GAEIEYAMAYSK	68	12	4
SYLYFTQFK	56	9	4

TAVAHRPGAFK	40	11	4
DTQFSYAVFK	56	10	6
LQAQDAGIYECHTPSTDTR	51	19	6
LQGDAVVLK	49	9	6
LVAQLDTEGVGSLGPGYEGR	95	20	6
VLPDVLQVSAAPPGPR	86	16	6
VVAGEVQVQR	71	10	6
FWVVDGR	32	7	4
TFQTYWVLGVPK	53	12	4
TSTDLQVLAAR	67	11	4
YAVGLDGQAPGQDCVWQG	51	18	4
DLQLVLPDYFPER	40	13	3
FNFLEQAFDK	37	10	3
SPSGGAAGPLLTPSQSLDGSR	53	21	3
EALPGSGQAR	38	10	2
FSVLLLHGIR	30	10	2
CDQAGVIIVGNLNSLSR	102	17	12
CELPCQDGTYGLNCAER	77	17	12
CLPGWSGVHCDSVCAEGR	45	18	12
CPLGFYGK	29	8	12
CPPGYTGAFCEDLCPPGK	75	18	12
CPSGTYGYGCR	60	11	12
CQDECPVGTYGVLCAETCQCVNGGK	32	25	12
CTPGWTGLYCTQR	45	13	12
CYHVSGACLCEAGFAGER	75	18	12
DCALICQCQNGADCDHISGQCTCR	27	24	12
ICSPGFYGHR	33	10	12
NGASCSPDDGICECAPGFR	79	19	12
ALEQDLPVNIK	58	11	15
DGSTIPIAK	28	9	15
EWVAIESDSVQPVPR	106	15	15
FIIEGMEEAGSVALEELVEK 1Met(ox)	141	20	15
FRQELFR	32	7	15
GATDNKGPVLAWINAVSAFR	43	20	15
GTVCFYGHLDVQPADR	89	16	15
HLEDVFSK	47	8	15
MMAVAADTLQR	78	11	15
MVVSMTLGLHPWIANIDDTQYLAAK	36	25	15
SVVLIPLGAVDDGEHSQNEK	72	20	15
VASVDMGPQQLPDGQSLPIPI 1Met(ox)	33	33	15
VFQYIDLHQDEFVQTLK	103	17	15
WNYIEGTK	52	8	15
YPSLSIHGIEGAFDEPGTK	72	19	15
ALAAVLLQALDR	68	12	12
ASWGEFQAR	29	9	12
AYQGVAAPFPK	70	11	12
FGEGVSSPK	59	9	12
GLQEAAEER	46	9	12
LADLASDLLLQYLLQGGAR	105	19	12
LLQQGLAQVEAGR	91	13	12
NSEPQDEGELFQGVDPR	76	17	12
RPESALLGGSEAGER	44	15	12
THLGEALAPLSK	53	12	12
VGEEDEEAAEAEAEAEEAER	143	20	12
VNLESPGPER	63	10	12
DQNVFVAQK	74	9	5

LQPVLQPLPSPGVGGK	46	16	5
LVDFLSR	41	7	5
SLDEISQPAQELK	59	13	5
TLSGTPEESKR	43	11	5
SHYAAFSVGR	33	10	1
ADQLTEEQIAEFK	75	13	4
EAFSLFDKDGDGTITTK	62	17	4
SLGQNPTEAELQDMINEVDAD 3Met(ox)	51	37	4
VFDKDGNGYISAAELR	90	16	4
AISQSGVALSPWVIQK	76	16	6
LGLLGDSVDIFK	58	12	6
NPLFWAK	35	7	6
TTFDVYTESWAQDPSQENK	74	19	6
TVVDFETDVLFLVPTEIALAQHR	32	23	6
VGPLGFLSTGDANLPGNYGLR	30	21	6
ATPEQYQILK	36	10	1
DIDPGAFQDLNK	57	12	9
EWLENIPK	36	8	9
ILILNNNLLR	74	10	9
KGFTSLQR	68	8	9
LGSEVLMSDLK	30	11	9
LSNVQELFLR	33	10	9
RLHINNNK	29	8	9
SLPVDVFAGVSLSK	86	14	9
WLYMDSNYLDTLSR	54	14	9
AEWLAVKDER	34	10	8
AQEENTWFSYLK	50	12	8
AVPWVILSDGDGTVEK	83	16	8
EWTTTTGDVVNENPEWVK	109	18	8
FIPNTDDQIIVALK	60	14	8
GSVDHENWVSNYNALR	86	16	8
LYVGGLGK	34	8	8
YEGIEFI	45	7	8
EEINALVQELGFYR	74	14	2
IPLSEMTR	37	8	2
LLQAPDTDLR	34	10	6
LSFSPNLDDYNPDIPEWR	40	18	6
LYDYNLGSITESSLWR	81	16	6
TILSYLYVCPTNK	46	13	6
VSENPYTSGIIASK	70	14	6
WQLIQEGVVPNR	49	12	6
EGSCPQVNINFPQLGLCR	32	18	2
VSCVTPNF	30	8	2
CAGTVEVEIQR	80	11	13
EAEFGQGTGPIWLNEVK	75	17	13
FQGEWGTICDDGWDSYDAAVACK	94	23	13
GPDTLWQCPSSPWEK	66	15	13
HYCNHNEDAGVTCSDGSDLELR	43	22	13
LASPSEETWITCDNK	100	15	13
LEVFYNGAWGTVGK	60	14	13
LQEGPTSCSGR	65	11	13
LVDGVTECSGR	49	11	13
LVGGDIPCSGR	51	11	13
QLGCPTAVTAIGR	74	13	13
SSMSETTVGVVCR	63	13	13
WGTVCDDNFNIDHASVICR	27	19	13

SQGGEPTYNVAVGR		39	14	1
AAGTEGPFQEVDGVATTR		88	18	9
FEVIEFDDGAGSVLR		42	15	9
NVLELSNVVR		45	10	9
SDMGVGVFTPTIEAR	1Met(ox)	27	15	9
TGEQAPSSPPR		41	11	9
TQQGVPAQPADFQAEVESDTR		56	21	9
VLAFTAVGDGPPSPTIQVK		80	19	9
VLAVNSIGR		30	9	9
YSIGGLSPFSEYAFR		85	15	9
FSVLGSGLNR		53	10	1
IEFPILEDSSELQLK		81	15	8
ISVTQPDSIVGIVAVDK		40	17	8
NNVVITVTQR		32	10	8
NSLGGFASTQDTTVALK		41	17	8
SNLIQQWLSQQSDLGVISK		45	19	8
SSMAVHSLFK		46	10	8
TLTLPSLPLNSADEIYELR		42	19	8
TNIQVTVTGPSSPSPVK		69	17	8
SSHQLQALAAGR		60	12	1
ADFPLSVVR		48	9	9
ALVTVDEVLKDEK		60	13	9
EPSFTCASDGLTYYNR		69	16	9
GITLAVVTCR		48	10	9
LVTSFCR		27	7	9
PRPEITWEK		39	9	9
SDFVILGR		46	8	9
VSELTEEPDSGR		59	12	9
WAYNSQTGQCQSFVYGGCEGNGNNFESR		44	28	9
IQDEQTTYVFFDNK		65	14	2
IQGDLAGR		36	8	2
AIFATAVAQELLQR		59	14	3
DLIGVVPLAMEAEILNTAILTGK		34	23	3
KGDVLTFPVSISR		41	13	3
AYGPLFLR		36	8	2
LLDLYASGER		51	10	2
IGPGDVLTFYDGDDLTAR		69	18	3
NGDNVEAPPVYDSYEVEYLPIEGLLSSGK		65	29	3
RPAYGDVTVTSLHPGGSAR		31	19	3
ADSSPVK		27	7	8
ATLVCLISDFYPGAVTVAWK		77	20	8
FSGSNSGNTATLTISR		109	16	8
LTVLGQPK		35	8	8
PGQAPVLVVYDDSDRPSGIPER		62	22	8
TVAPTECS		37	8	8
YAASSYLSLTPEQWK		95	15	8
YAASSYLSLTPEQWKSHR		27	18	8
AAECPAGFVRPPLIIFSVDGFR		28	22	31
AGTFFWSVVIPHER		52	14	31
CFELQEAGPPDCR		81	13	31
CVNVIFVGDHGMEDVTCDR		91	19	31
DIEHLTSLDFFR		57	12	31
GDCCTNYQVVCK		49	12	31
GESHWVDDDCEEIK		67	14	31
IEDIHLLVER		63	10	31
IVGQLMDGLK	1Met(ox)	67	10	31

KPDQHFKPYLK	31	11	31
KPLDVYK	35	7	31
LDELNKR	32	7	31
LHYANNR	32	7	31
NGVNVISGPIFDYDYDGLHDTEDK	40	24	31
NKLDELNKR	43	9	31
QMSYGFLFPPYLSSSPEAK	48	19	31
QYVEGSSIPVPTHYYSIITSCLDFTQPADK	33	30	31
RIEDIHLLVER	44	11	31
RLHYANNR	41	8	31
RVWNYFQR	33	8	31
SYPEILTLK	52	9	31
SYTSCCHDFDELCLK	73	15	31
TEFLSNYLTNVDDITLVPGTLGR	87	23	31
TYLHTYESEI	42	10	31
VNSMQTVFVGYGPTFK	96	16	31
VSPSFSQNCLAYK	71	13	31
VWNYFQR	39	7	31
WVEELMK	29	7	31
WWGGQPLWITATK	77	13	31
YDAFLVTNMVPMYPAFK 2Met(ox)	65	17	31
YGPFGPEMTNPLR	41	13	31
AASTFNIR	45	8	11
ETGTPIWTSR	42	10	11
GSPGKEHPEER	40	11	11
IEFTSDQAR	48	9	11
IHVGEEK	31	7	11
IHVGEEKR	37	8	11
LLLHDKDR	41	8	11
LYSSTPDLTIQFHSDPAGLIFGK	33	23	11
SALLYDSLQTESVPFEGLLSEGNTIR	59	26	11
SPTNTISVYFR	82	11	11
TTSHTELVR	29	9	11
APVLELEK	41	8	16
DFQWTDNTGLQFENWR	99	16	16
ELGGEVFYVGPAR	62	13	16
GIEDEQDLVPLEVTGVVFHYR	48	21	16
GTVLCGPPPAVENASLIGAR	75	20	16
HLQAAFEDGFDNCDAGWLSDR	39	21	16
LGSGSVQAALAELVALPCLFTLQPR	32	25	16
LSSAIIAAPR	83	10	16
LTLAGAR	30	7	16
QDLPILVAK	61	9	16
RLTLAGAR	31	8	16
RNPQELYDVYCFAR	31	14	16
SSLPGVR	32	7	16
VSLPSYPR	45	8	16
YALTFAEAQEACR	86	13	16
YPIQTPR	37	7	16
AEIENYVLTYK	58	11	10
GLCVDGECVCEEPYTGEDCR	125	20	10
ITFTPSSGIASEVTVPK	74	17	10
LILNYSPR	40	8	10
LNPATEYEISLNSVR	78	15	10
SPPTSASVSTVIDGPTQILVR	56	21	10
TATSLDLEWDNSEAEVQEYK	87	20	10

TSYTLTDLEPGAEYIISVTAER	86	22	10
VGFGNVEDEFWLGLDNIHR	53	19	10
YEVSVSAVR	59	9	10
IVISQLNPYTLR	67	12	2
VFLCPGLLK	29	9	2
AGLLRPDYALLGHR	38	14	9
DGSPDVTTADIGANTPDATK	92	20	9
EFTEAFLGCPAIHPR	31	15	9
EYGVVLAPDGSTVAVEPLLAGLEAGLQGR	48	29	9
GSQTQSHPDLGTEGCWDQLSAPR	27	23	9
PSLSHLLSQYYGAGVAR	100	17	9
QNGAALTSASILAQQVWGTLVLLQR	49	25	9
TDCPGDALFDLLR	46	13	9
VINLPLDSMAAPWETGDTFPDVVAIAPDVR	51	30	9
DSYYMTSSQLSTPLQQWR	75	18	3
TPECPSHTQPLGVYLLTPAVQDLWLR	47	26	3
VPTGGVEEGLLER	71	13	3
FHLGEPEASTQFMTQNYQDS 1Met(ox)	42	27	7
FPGAVDGATYILVMVDPDAPS 1Met(ox)	92	22	7
HWLVTDIK	46	8	7
IQGQELSAYQAPSPPAHSGFHR	43	22	7
ITSWMEPIVK	47	10	7
VISLLPK	46	7	7
YQFFVYLQEGK	47	11	7
AAVYHHFISDGVR	62	13	123
ACEPGVDYVYK	72	11	123
ADIGCTPGSGK	47	11	123
ADIGCTPGSGKDYAGVFSDAGLTFTSSSGQ(	62	35	123
AEDLVGK	43	7	123
AGDFLEANYMNLQR 1Met(ox)	84	14	123
AKDQLTCNK	42	9	123
AKDQLTCNKFDLK	59	13	123
APSTWLTAYVVK	83	12	123
ASHLGLAR	60	8	123
AVLYNYR	38	7	123
AYYENSPQQVFSTEFEVK	77	18	123
CAEENCFIQK	58	10	123
DAPDHQELNLDVSLQLPSR	83	19	123
DFDFVPPVVR	52	10	123
DICEEQVNSLPGSITK	85	16	123
DMALTAFVLISLQEAK	50	16	123
DQLTCNKFDLK	62	11	123
DSCVGSLVVK	49	10	123
DSITTWEILAVSMSDK	80	16	123
DSITTWEILAVSMSDKK	87	17	123
DTWVEHWPEEDECQDEENQK	63	20	123
DYAGVFSDAGLTFTSSSGQQTAQR	164	24	123
EALKLEEK	44	8	123
EDIPPADLSDQVPDTESETR	112	20	123
EGVQKEDIPPADLSDQVPDTESETR	45	25	123
ENEGFTVTAEGK	81	12	123
EPGQDLVVLPLSITTDFIPSFR	93	22	123
EVVADSVWVDVK	87	12	123
EYVLPSFEVIVEPTEK	87	16	123
FISLGEACK	49	9	123
FISLGEACKK	39	10	123

FVTVQATFGTQVVEK		102	15	123
FYHPEKEDGK		56	10	123
FYYIYNEK		34	8	123
GLEVTITAR		61	9	123
GQGTLSVVTMYHAK		83	14	123
GVFVLNK		41	7	123
GYTQQLAFR		67	9	123
HQQTVTIPPK		48	10	123
IEGDHGAR		50	8	123
IFTVNHK		35	7	123
IHWESASLLR		47	10	123
ILLQGTPVAQMTEDAVDAER		123	20	123
IPIEDGSGEVVLSR		103	14	123
ISLPESLK		37	8	123
ISLPESLKR		52	9	123
IWDVVEK		38	7	123
KGYTQQLAFR		99	10	123
KHYLMWGLSSDFWGEKPNLSYIIGK		27	25	123
KLVLSSEK		29	8	123
KQELSEAEQATR		75	12	123
KVEGTAFVIFGIQDGEQR		65	18	123
KVFLDCCNYITELR		93	14	123
KVFLDCCNYITELRR		27	15	123
KVLLDGVQNPR		82	11	123
LESEETMVLEAHDAQGDVPVTVTVHDFPGK		48	30	123
LKGPLLNK		35	8	123
LMNIFLK	1Met(ox)	50	7	123
LPYSVVR		49	7	123
LSINTHPSQKPLSITVR		70	17	123
LVAYYTLIGASGQR		100	14	123
LVLSSEK		38	7	123
NEQVEIR		52	7	123
NNNEKDMALTAFVLISLQEAK	1Met(ox)	27	21	123
NRWEDPGK		27	8	123
NTLIIYLDK		51	9	123
NTMILEICTR	1Met(ox)	84	10	123
PGMPFDLMVFVTNPDGSPAYR		81	21	123
PNLSYIIGK		33	9	123
QCQDLGAFTESMVVFGCPN	1Met(ox)	81	19	123
QDSLSSQNQLGVLPLSWDIPELVNMGQWK		55	29	123
QELSEAEQATR		81	11	123
QGALELIK		49	8	123
QGALELIKK		62	9	123
QKPDGVFQEDAPVIHQEMIGGLR		62	23	123
QLANGVDR		56	8	123
QLYNVEATSYALLALLQLK		142	19	123
QLYNVEATSYALLALLQLKDFDFVPPVVR		42	29	123
QPSSAFAAFVK		46	11	123
QPVPGQQMTLK	1Met(ox)	57	11	123
RAPSTWLTAYVVK		52	13	123
RIPIEDGSGEVVLSR		85	15	123
RQGALELIK		35	9	123
SDDKVTLEER		63	10	123
SEETKENEGFTVTAEGK		86	17	123
SEFPESWLWNVEDLK		69	15	123
SEFPESWLWNVEDLKEPPK		71	19	123

SGIPIVTSPYQIHFTK		60	16	123
SGQSEDRQPVPGQQMTLK		57	18	123
SGSDEVQVGQQR		84	12	123
SLKVVPEGIR		48	10	123
SNLDEDIIAEENIVSR		99	16	123
SSLSVPYVIVPLK		54	13	123
SVQLTEK		54	7	123
SVQLTEKR		42	8	123
SYTVAIAGYALAQMGR		60	16	123
TELRPGETLNVNFLLR		64	16	123
TFISPIK		38	7	123
TGLQEVEVK		70	9	123
TIYTPGSTVLYR		53	12	123
TKKQELSEAEQATR		90	14	123
TMQALPYSTVGNSNNYLHLSVLR		86	23	123
TVMVNIENPEGIPVK		73	15	123
VEGTAFVIFGIQDGEQR		113	17	123
VELLHNPAFCSLATTK		68	16	123
VFLDCCNYITELR		88	13	123
VFSLAVNLIAIDSQVLCGAVK		60	21	123
VHQYFNVELIQPGAVK		85	16	123
VLLDGVQNPR		82	10	123
VPVAVQGEDTVQSLTQGDGVAK		114	22	123
VQLSNDFDEYIMAIEQTIK	1Met(ox)	132	19	123
VRVELLHNPAFCSLATTK		62	18	123
VSHSEDDCLAFK		80	12	123
VTIKPAPETEK		52	11	123
VVLVAVDK		53	8	123
VVLVAVDKGVFVLNK		47	15	123
VVLVSLQSGYLFIQTDK		71	17	123
VVPEGIR		27	7	123
VYAYYNLEESCTR		89	13	123
YELDKAFSDR		47	10	123
YISKYELDK		45	9	123
YYTYLIMNK	1Met(ox)	58	9	123
AVILQGSNDVELVAEGNSR		57	19	7
EMATQLAFMR	1Met(ox)	34	10	7
GETGPSGPVGPAGAVGPR		69	18	7
GPAGPSGPAGK		31	11	7
GVGLGPGPMGLMGPR	1Met(ox)	32	15	7
NSIAYMDEETGNLK		77	14	7
SLNNQIETLLTPEGSR		65	16	7
ANRPFLVFIR		40	10	3
ATEDEGSEQKIPEATNR		68	17	3
IPEATNR		32	7	3
EWIQVDLGLLR		61	11	5
FVTAVGTQGAISK		55	13	5
LEIWDGFPDVGPHIGR		30	16	5
LNYPENGWTPGEDSYR		48	16	5
SFEGNNNYDTPELR		51	14	5
ASLINNAFQLVSIGK		47	15	6
ILASTQFEPTAAR		67	13	6
NPVGYPLAWQFLR		33	13	6
TDVLILPEEVEWIK		45	14	6
TQEFPQILTLIGR		72	13	6
YQFSLSSTEK		56	10	6

ETFNVYYLETEADLGR	62	16	3
GLATFPATAAESAFSTLVEVAGTCVAHSEGE	33	37	3
KIDTIAADESFTQGDLGER	43	19	3
LPLQIEVSDTELSQIK	61	16	3
SILAVGVGNVR	53	11	3
WPVVVAEGEGQGPLIR	67	16	3
AGEVQEPELR	56	10	20
AREDIFMETLK	67	11	20
AYLEEECPATLR	64	12	20
CLAYDFYPGK	47	10	20
DYIEFNK	39	7	20
EDIFMETLK	33	9	20
EIPAWVPFDPAAQITK	47	16	20
HVEDVPAFQALGSLNDLQFFR	89	21	20
IDVHWTR	46	7	20
KSQPMGLWR	30	9	20
NILDRQDPPSVVVTSHQAPGEK	39	22	20
QDPPSVVVTSHQAPGEK	50	17	20
QDSQLQK	33	7	20
QKWEAEPVYVQR	45	12	20
QVEGMEDWK	50	9	20
QVEGMEDWKQDSQLQK	39	16	20
SSGAFWK	35	7	20
WEAEPVYVQR	63	10	20
YSLTYIYTGLSK	88	12	20
YYYDGKDYIEFNK	76	13	20
AEAIGYAYPTR	38	11	12
AIQYQQHFSR	74	10	12
ATIADLILSALER	69	13	12
EFQLTLQPGFWK	47	12	12
EKWAQEPLLQPLSLR	27	15	12
LEAAIQR	55	7	12
LPEINLDGMVGVR	76	13	12
RAEAIGYAYPTR	45	12	12
SDVCLVQLLGTGTDSSEPCGLSDLCR	119	26	12
VLEEQLK	36	7	12
WAQEPLLQPLSLR	47	13	12
WLEAILSWQK	40	10	12
AVFVDLEPTVIDEVR	69	15	1
GLTVPIASIDIPSGWDVEK	44	19	2
LFGYEPTIYYPK	32	12	2
SLGEENFEVVK	44	11	1
VVLAPQDVVVAR	28	12	1
AYSPEFYFDTPNPTR	78	15	3
ESENIQLGQDLK	50	12	3
GQLLEYILTDLR	87	12	3
ANSVFEDLSVTLR	105	13	9
DHSPDLYSLELAGLDEIGK	29	19	9
DHSPDLYSLELAGLDEIGKR	41	20	9
FADDMYSLYGGNAVVELVTVI 1Met(ox)	57	21	9
ILVDALQK	32	8	9
IPDVAALSMGFSVK 1Met(ox)	66	14	9
LFQENSVLSSLPLNSLSR	63	18	9
SFDTSLIR	47	8	9
SPGSVVFR	51	8	9
ACANPAAGSVILLENLR	50	17	10

AGGFLMK	42	7	10
GCITIIGGGDTATCCAK	100	17	10
IQLINNMLDK	37	10	10
ITLPVDFVTADKFDENAK	43	18	10
LGDVYVNDAFGTAHR	43	15	10
QIVWNGPVGVFEWEAFAR	40	18	10
VLNNMEIGTSLFDEEGAK	87	18	10
VLPGVDALSNI	39	11	10
YSLEPVAVELK	42	11	10
DQVYTVNLNEMPK	78	13	7
LDFQLMLK	31	8	7
LSTLEYDGEEISGLAR	88	16	7
LYSATVADFLASDAVIYR	80	18	7
QTNVALFADGK	58	11	7
TPDSVWTAVPEDK	71	13	7
TSIDFPDETLSFIK	42	14	7
DQLVIPDGQEEEQEAAGEGR	98	20	4
DTINLLDQR	47	9	4
EETNEIQVVNEEPQR	73	15	4
LQQDVLQFQK	32	10	4
AAPSAEFSVDR	62	11	9
AQWPAWQPLNVR	35	12	9
EGYTEFSLR	43	9	9
IIYFQDEGSLTK	33	12	9
LCGGGIQER	36	9	9
QQSDEVLTVIK	58	11	9
SEQLKEESEGEQFPGCR	74	17	9
SLAELGDCNEDLEQVEK	63	17	9
VEGDPDFYKPGTSYR	50	15	9
FYFENLLSK	28	9	2
ITEQLIEAINNGDFEAYTK	41	19	2
FYFENALSK	30	9	2
VTEQLIEAINNGDFEAYTK	92	19	2
ILPDTPQEPFALWEILNK	30	18	2
VTVTPIYTDGEGVSVSAPGK	45	20	2
DQAGEYECSAENDVSFPDVR	77	20	11
DYSLQIQNVDVTDDGPYTCSVQTQHTPR	41	28	11
KGDTAVLR	68	8	11
LFNGQQGIIIQNFSTR	73	16	11
PFENGQYLDIYGITR	89	15	11
SGTVTPGR	30	8	11
SSIIFAGGDK	69	10	11
TMQVHLTVQVPPK	55	13	11
VSISTLNK	36	8	11
VSISTLNKR	46	9	11
VVVNFAPTIQEIK	56	13	11
FPVVNTAYGR	38	10	3
GGGGPGGGAPGGPGLGLGSLGEER	61	24	3
GNYGLLDQIQALR	48	13	3
DDGGIIICEAQNQALPSGHSK	47	21	9
EGDTLVLTCAVTGNPR	67	16	9
EGDTLVLTCAVTGNPRPNQIR	28	21	9
EQAVEGGEVELSCLVPR	77	17	9
FQLEEFSPR	46	9	9
IHASQAVVR	39	9	9
LHQYDGSIVVIQNPAR	81	16	9

QTQYVLDVQYSPTAR	83	15	9
VWSVASTVR	46	9	9
DAGPNGVASYELQAGPEAQELFGLQVAEDQ	70	33	6
GLFTISPETGEIQVK	52	15	6
NTGLITVQGPVDREDLSTLR	36	20	6
TGDIFTTETSIDR	58	13	6
VPEEQPPNTLIGSLAADYGFPDVGHLYK	27	28	6
YFLQTTTPLDYEK	48	13	6
DFQTAEVAYYSPTTR	40	15	2
VPNLLSTSWTFPR	43	13	2
AESESDLVAEIANVVQK	45	17	27
AGIEVQEIK	67	9	27
ANILYAWAR	51	9	27
DCSGVSLHLTR	70	11	27
DGNYWVTDVALHQVFK	53	16	27
EEEEVLDQGDFYSLLSK	79	17	27
EGPVLILGR	46	9	27
FITQWGEESSGSSPLPGQFTVPHSLALVPLL	27	39	27
GDHVWDGNSFDSK	52	13	27
GNAILVR	48	7	27
GSGGLNLGNFFASR	72	14	27
HFDMPHDIVASEDGTVYIGDA 1Met(ox)	30	28	27
IPLLQQPK	43	8	27
IPVDEEAFVIDFKPR	53	15	27
IVQFSPSGK	39	9	27
MPGVTPK	28	7	27
NGQWTLIGR	57	9	27
NLFYLPHGLSIDK	28	13	27
NNLVIFHR	41	8	27
NVFAISYIPGLLFAVNGK	66	18	27
NYPMHVFAYR	33	10	27
PVVPIDSSDFALDIR	55	15	27
REEEEVLDQGDFYSLLSK	42	18	27
SMQPGSDQNHFCQPTDVAVDPGTGAIYVSD	36	36	27
TIPPEANIPIPVK	29	13	27
VHTHHLGK	31	8	27
VVNSDISCHYK	67	11	27
GQLTFYAQPNWIQK	42	14	13
LECFALGNPVPTIIWR	45	16	13
LQFAYLDNFK	63	10	13
LSAAADLIVR	57	10	13
SDAGSYTCIATNHFGTASSTGNLVVK	46	26	13
SELVITWETVPEELQNGR	46	18	13
SLSATDIEVFWASPLEK	44	17	13
TEEALPEVTPANVSGGGGSK	41	20	13
TFTATVVGLNPWVEYEFR	76	18	13
TPFSVGWQAVSTVPELIDGK	36	20	13
TVAANVIGIGEPSRPSEK	60	18	13
VGGQDSAGDLMIR	73	13	13
VLGPPTPLILR	38	11	13
CYTAVVPLVYGGETK	62	15	3
IVLVDNK	37	7	3
SSEDPNEDIVER	65	12	3
ESTLHLVLR	42	9	3
IQDKEGIPPDQQR	65	13	3
TLSDYNIQK	45	9	3

AGLPFGLMR		59	9	9
CPGSDVIMVENANYGR		87	16	9
EEPVSLTFPNPYQFISSVDYNPR		43	23	9
ELACEGYPIELR		54	12	9
LVVSQLNPYTLR		70	12	9
SVYVDDDSEAAGNRVDYAFNTNANR		33	25	9
TDTLTEYASWEDYVAAR		56	17	9
TQCVVVAGSDAFPDPCPGTYK		70	21	9
YLEVQYDCVPYK		42	12	9
FLVAPLTFSNR		30	11	2
TNDQLVAFLSR		64	11	2
ETSLQVDNLPR		33	11	2
VEITAVATQGR		74	11	2
AAHFVFR		36	7	18
AGEQDATIHLK		74	11	18
ASWTRPEK		28	8	18
AVGEEVWHSK		52	10	18
CVVTGEDGSESEATVNVK		128	18	18
DGEQIEQEEDDEKYIFSDDSSQLTIK		63	26	18
DIQVIVNVPPTIQAR		101	15	18
FFLCQVAGDAK		74	11	18
FIVLSNNYLQIR		81	12	18
GLGEISAASEFK		85	12	18
LAALNGK		35	7	18
LEGQMGEDGNSIK		96	13	18
LPSGSDHVMLK		70	11	18
NDEAEYICIAENK		77	13	18
QDDGGSPIR		46	9	18
QETLDGHMVVR		43	11	18
TQPVQGEPSAPK		49	12	18
YIFSDDSSQLTIK		83	13	18
ANLQSVPHASASRPR		42	15	6
EFLGENISNFLSLAGNTYQLTR		100	22	6
ILQADQEL		38	8	6
LNILNNNYK		33	9	6
VNEPSILEMSR		46	11	6
VTEPISAESGEQVER		92	15	6
FYFENLWSR		56	9	3
GAILTTMLATR	1Met(ox)	60	11	3
ITQYLDAGGIPR		56	12	3
AVDAALKK		36	8	8
DIPTNSPELEETLTHTITK		67	19	8
IASFSQNCDIYPGK		59	14	8
IGEIKEETTSHLR		71	13	8
KYFIDFVAR		40	9	8
VQVVAGK		39	7	8
YFIDFVAR		54	8	8
YNSQNQSNNQFVLYR		63	15	8
DTADGILTDVILK		49	13	2
IPLYSFSMVLVDK	1Met(ox)	32	13	2
ADGLAVIGVLMK		56	12	4
ESISVSSEQLAQFR		41	14	4
SLLSNVEGDNAVPMQHNNRPTQPLK		49	25	4
VLDALQAIK		68	9	4
AAEDYGVIK		56	9	15
ANKGPSYGMSR	1N-ac	38	11	15

DMAAVQR	33	7	15
EFTESQLQEGK	79	11	15
GASQAGMTGYGR	70	12	15
GDPNWFMK	30	8	15
GPSYGMSR	32	8	15
LGFQVWLK	56	8	15
LVEWIIVQCGPDVGRPDR	61	18	15
LVNSLYPDGSKPVK	56	14	15
NGVILSK	27	7	15
QMEQVAQFLK	39	10	15
TDMFQTVDLFEGK	53	13	15
TLMALGSLAVTK	83	12	15
VPENPPSMVFK	40	11	15
AVDHINSTIAPALISSGLSVVEQEK	34	25	11
DATNVGDEGGFAPNILENSEALELVK	73	26	11
DGKYDLDFK	38	9	11
DYPVVSIEDPFDQDDWAAWSK	69	21	11
IEEELGDEAR	71	10	11
LAMQEFMILPVGAESFR	30	17	11
LDNLMLELDGTENK	60	14	11
LGAEVYHTLK	81	10	11
SGETEDTFIADLVVGLCTGQIK	49	22	11
VNQIGSVTEAIQACK	86	15	11
YITGDQLGALYQDFVR	107	16	11
AQVTPEVQPGCR	58	12	19
DGAGGWSPLVSNK	49	13	19
EASLQVDQLTPK	54	12	19
EYLSVIIAK	35	9	19
FIPLEWNPK	39	9	19
GEFNLMNLDYEISFGGIPAPGI 1Met(ox)	51	22	19
LFLLINSGEAK	66	11	19
LISISGK	27	7	19
LPSDITAGVELNDGQWHSVSLSAK	32	24	19
QDGTPLSWWVGR	69	12	19
QVHLSSGTEFSAVK	31	14	19
SLQLNGMTLDLEER	60	14	19
SLSPIKDIISLK	48	12	19
SPLGGFQGCMR	54	11	19
TMQSDGILLHR	61	11	19
TPSLLLFVSSFYK	76	13	19
TTASSGVFLENLGIADFIR	84	19	19
YQEPDVVNFDFK	38	12	19
YQWLQIDLGER	33	11	19
FSSFEEAVK	30	9	4
LVEEIGWSYTGALNQK	87	16	4
QSPINIDEDLTQVNVNLK	81	18	4
QVFSSYTGK	35	9	4
EKLEATINELV	35	11	5
LEATINELV	37	9	5
TAFQEALDAAGDK	45	13	5
VGEFSGANK	47	9	5
VGEFSGANKEK	35	11	5
IEAELQDICNDVLELLDK	75	18	4
NLLSVAYK	37	8	4
QTTVSNSQQAYQEAFEISK	104	19	4
TAFDEAIAELDTLNEESYK	81	19	4

GTVTGQVQGPEDK		84	13	3
IGVVGWVK		32	8	3
SVDYEVFGR		34	9	3
AGWNAYIDNLMADGTCQDAA	1N-ac	101	25	5
DSLLQDGEFSMDLR	1Met(ox)	48	14	5
DSPSVWAAVPGK		47	12	5
SSFYVNGLTLGGQK		69	14	5
TFVNITPAEVGVLVGK		54	16	5
GGHAGATYIFGK		66	12	3
IDSAPGLGDFLQLHIEQGK		60	19	3
LFQGQLSGLYYDGLK		74	15	3
AREENVATFR		64	10	21
CENVATLDPINFETPEAYISLPK		92	23	21
DGAVSLVINLGSGAFEAIVEPVNGK		53	25	21
DLFIDGR		30	7	21
EASILSYDGSMYMK	2Met(ox)	51	14	21
EENVATFR		40	8	21
FNDNAWHDVK		70	10	21
FSMDCAETAVLSNK		106	14	21
GLILDLK		39	7	21
GPETLYAGQK		39	10	21
LAVGFSTTVK		44	10	21
LMVNLGK		31	7	21
NGLILHTGK		48	9	21
NIIADPVTFK		41	10	21
PSALTLDGVQAMPGFK		34	16	21
QLAEMQNAAGVK	1Met(ox)	64	12	21
SDLSFQFK		50	8	21
SGGLILYTWPANDRPSTR		34	18	21
SGTISVNSR		50	9	21
VLNMAAENNPNIK		55	13	21
YNRPVEEWLQEK		76	12	21
TGVITSPDFPNPYPK		51	15	1
ADTFLGIENLEYLQADYNLIK		65	21	7
ETNKQELCPMGTGSDFDVR		27	19	7
FASLTHLDIR		42	10	7
LPYIGVLEHIGR		42	12	7
LQNIEGGAFLGLSALK		111	16	7
QLHLNNNELK		28	10	7
VLILNDNLISFLPDNIFR		93	18	7
LTLEEAR		28	7	2
GAEIATTGQLYAAWDGGLDHCSPGWLADG		40	32	2
VEADIPGHGQEVLIR		28	15	2
YLEFISECIIQVLQSK		53	16	2
LSEGVTISYK		36	10	1
AGALQLLLVGDK		113	12	2
VYDFLSTFITSGMR	1Met(ox)	56	14	2
ADLGALELWR		58	10	4
DPIYFTGLASEPGAR		67	15	4
FSTDEGQCWQTYTFTR		70	16	4
TEFGMAIGPENSGK		55	14	4
DLADELALVDVIEDK		76	15	8
DLADELALVDVIEDKLK		47	17	8
DQLIYNLLK		40	9	8
LLIVSNPVDILTYVAWK		63	17	8
LVIITAGAR		55	9	8

QVVESAYEVIK	43	11	8
SADTLWGIQK	58	10	8
VTLTSEEEAR	55	10	8
ETWLFSR	37	7	2
SPTPDELPTCYPGDDWSGVSLR	88	22	2
AEAQAQYSAAVAK	97	13	35
AGFSWIEVTFK	67	11	35
AISGGSIQIENGYFVHYFAPEGLTTMPK	30	28	35
ANTVQEATFQMELPK	104	15	35
DQFNLIVFSTEATQWRPSLVPASAENVNK	35	29	35
EKAEAQAQYSAAVAK	79	15	35
ETLFSVMPGLK	50	11	35
FAHTVVTSR	31	9	35
FKPTLSQQQK	30	10	35
FSSHVGGTLGQFYQEVLWGSPAASDDGRR	36	29	35
GPDVLTATVSGK	62	12	35
GSEMVVAGK	74	9	35
IHEDSDSALQLQDFYQEVANPLLTAVTFEYPS	36	44	35
ILDDLSPR	53	8	35
ITFELVYEELLK	81	12	35
LALDNGGLAR	68	10	35
LGVYELLLK	50	9	35
LPEGSVSLIILLTDGDPTVGETNPR	107	25	35
LWAYLTIQQLLEQTVSASDADQQALR	82	26	35
NGIDIYSLTVDSR	76	13	35
NMEQFQVSVSVAPNAK	83	16	35
NPLVWVHASPEHVVVTR	58	17	35
NVHSAGAAGSR	39	11	35
NVVFVIDK	61	8	35
PSLVPASAENVNK	33	13	35
QGPVNLLSDPEQGVEVTGQYER	96	22	35
RLDYQEGPPGVEISCWSVEL	74	20	35
SFAAGIQALGGTNINDAMLMA 1Met(ox)	54	33	35
SIQNNVR	32	7	35
SPEQQETVLDGNLIIR	99	16	35
TGLLLLSDPDK	44	11	35
VQGNDHSATR	79	10	35
VRPQQLVK	32	8	35
VTIGLLFWDGR	60	11	35
WKETLFSVMPGLK	28	13	35
GDMFLVANLGTK 1Met(ox)	71	12	6
LSSGLVTAALYGR	101	13	6
QEALELMNQNLDIYEQQVMTAAQK	53	24	6
SHLIIAQVAK	54	10	6
TSIYPFLDFMPSPQVVR	73	17	6
VDLITFDTPFAGR	73	13	6
FFADLLDYIK	55	10	4
NVLIVEDIIDTGK	63	13	4
TMQTLLSLVR	54	10	4
VIGGDDLSTLTGK	51	13	4
DLAHEATK	30	8	25
EETGFSTYNPQVIIR	61	15	25
EIAEDELVAAEALLK	80	15	25
FIDFLAIEMR	52	10	25
GSYNNIVVNVK	58	11	25
HMAAPLIGQLTR	50	12	25

IGPVTYSIDGCVR		41	13	25
IIADADATVK		43	10	25
IQDMSGWYLTDLPGR		61	15	25
ISEISMEVAEQGR	1Met(ox)	35	13	25
IYFGGLPTLR		34	10	25
LADEINSIIDYVEDIQTK		92	18	25
LIQLAEGNLNTLVTEMNELLTR		55	22	25
LTIELEVR		38	8	25
SGFFNLQEDNWK		48	12	25
SLGLICDGCPVGYTGPR		58	17	25
TAVADNLLFYLGSAK		72	15	25
TFSSSALLMYLATR	1Met(ox)	32	14	25
TNAVVKDPSK		38	10	25
TPYNILSSPDYVGVTK		41	16	25
TYKPEIK		57	7	25
VAPQQDDLDSPQQISISNAEAR		51	22	25
VEYPDLTIDDSYWYR		44	15	25
VSFLWDVGSGVGR		67	13	25
VTADGEQTGQDAER		81	14	25
EVQPVELPNCNLVK		35	14	9
GIETGSEDMEILPNGLAFISSGLK		64	24	9
IFFYDSENPPASEVLR		93	16	9
ILLMDLNEEDPTVLELGITGSK	1Met(ox)	99	22	9
IQNILTEEPK		68	10	9
LLIGTVFHK		63	9	9
STVELFK		35	7	9
VVAEGFDFANGINISPDGK		62	19	9
YVYIAELLAHK		36	11	9
ADDLGKGGNEESTK		61	14	9
DGVADVSIEDSVISLSGDHCIIGR		68	24	9
GDGPVQGIINFEQK		51	14	9
HGGPKDEER		57	9	9
HVGDLGNVTADK		90	12	9
HVGDLGNVTADKDGVADVSIEDSVISLSGDH		88	36	9
KHGGPKDEER		57	10	9
LACGVIGIAQ		46	10	9
TLVVHEK		29	7	9
GNLCVNLMR		44	9	3
IAFSATR		55	7	3
LEQGENVFLQATDK		90	14	3
DSFDIIK		28	7	5
DSFDIIKR		46	8	5
EFTRPEEIIFLR		41	12	5
LEMNYVVGGVVSHR		76	14	5
YMDGMTVGVVR		79	11	5
AGAHLQGGAK		41	10	11
GALQNIIPASTGAAK		40	15	11
IISNASCTTNCLAPLAK		85	17	11
LISWYDNEFGYSNR		62	14	11
LTGMAFR		28	7	11
LVINGNPITIFQER		81	14	11
VGVNGFGR		36	8	11
VIHDNFGIVEGLMTTVHAITATQK		41	24	11
VPTANVSVVDLTCR		106	14	11
VVDLMAHMASKE	2Met(ox)	44	12	11
WGDAGAEYVVESTGVFTTME	1Met(ox)	73	21	11

APPSVFAEVPQAQPVLVFK		102	19	20
ARLEALK		30	7	20
EPESILQVLSQMEK		70	14	20
FLFPFFDSAYQGFASGNLER		99	20	20
IANDNSLNHEYLPILGLAEFR		38	21	20
IGADFLAR		59	8	20
INVSGLTTK		46	9	20
ITWSNPPAQGAR		53	12	20
IVASTLSNPELFEEWTGNVK		79	20	20
LALGDDSPALK		58	11	20
NFGLYNER		36	8	20
NLDYVATSIHEAVTK		72	15	20
NTPVYVSSPTWENHNAVFSAAGFK		93	24	20
QVEYLVNEK		44	9	20
RVGGVQSLGGTGALR		59	15	20
TDDCHPWVLPVVK		29	13	20
TPGTWNHITDQIGMFSFTGLNPK		38	23	20
VGGVQSLGGTGALR		48	14	20
VGNLTVVGK		72	9	20
VNLGVGAYR		45	9	20
FDLNDVYVYWQTSESK		37	16	2
GLYDVVSVLR		71	10	2
FIIPQIVK		29	8	15
GEMMDLQHGSLFLQTPK	1Met(ox)	33	17	15
GLTSVINQK		49	9	15
GMYGIENEVFLSLPCILNAR	1Met(ox)	36	20	15
GYTNWAIGLSVADLIESMLK	1Met(ox)	122	20	15
ITVVGVGQVGMACAISILGK		55	20	15
IVADKDYSVTANSK		89	14	15
IVVVTAGVR		75	9	15
LIAPVAEEEATVPNNK		55	16	15
LKDDEVAQLK		66	10	15
LKDDEVAQLKK		69	11	15
MVVESAYEVIK		67	11	15
SADTLWDIQK		71	10	15
SLADELALVDVLEDK		88	15	15
VIGSGCNLDSAR		74	12	15
ACGLVASNLNLKPGECLR		70	18	7
DGGAWGTEQR		39	10	7
DSNNLCLHFNPR		52	12	7
LNLEAINYMAADGDFK		107	16	7
LPDGYEFK		31	8	7
SFVLNLGK		31	8	7
VRGEVAPDAK		43	10	7
FYPEDVSEELIQDITQR		44	17	3
IGFPWSEIR		58	9	3
LFFLQVK		28	7	3
APVAGTCYQAEWDDYVPK		86	18	12
CDEPILSNR		52	9	12
GNDISSGTVLSDYVGSGPPK		122	20	12
LYEQLSGK		53	8	12
LYTLVLTDPDAPSR		75	14	12
LYTLVLTDPDAPSRK		37	15	12
NRPTSISWDGLDSGK		65	15	12
VLTPTQVK		30	8	12
WSGPLSLQEVDEQPQHPLHVTYAGAAVDEL		42	32	12

YREWHHFLVVNMK		49	13	12
YVWLVYEQDR		31	10	12
YVWLVYEQDRPLK		57	13	12
DPNTPLLQGIADYRPK		44	16	5
LGQHLLPWMDR		38	11	5
LSSLQAGTKEDLYLHSLK		40	18	5
TYIDLIFPDCLAPR		42	14	5
WFLEDEKR		33	8	5
IGPITPLEFYR		51	11	1
ALVGDEVELPCR		46	12	3
DHSYQEEAAMELK		44	13	3
FSDEGGFTCFFR		61	12	3
FFQMVGLK		36	8	5
IGVDEFSTLVAES		59	13	5
SGFIEEDELGFILK		71	14	5
SMTDLLNAEDIKK	1Met(ox) 1N-a	36	13	5
VFHMLDKDK		35	9	5
AFLASPEYVNLPINGNGK		86	18	8
ALPGQLKPFETLLSQNQGGK		33	20	8
ASCLYGQLPK		50	10	8
DQQEAALVDMVNDGVEDLR	1Met(ox)	133	19	8
FQDGDLTLYQSNTILR		113	16	8
MLLADQGQSWK		51	11	8
PPYTVVYFPVR		61	11	8
YISLIYTNYEAGKDDYVK		67	18	8
AANAYGISDPSQISDPVK		75	18	14
ARPFFNEFQGADSEIK		27	16	14
DGSPLDDKDER		37	11	14
ESEVAELTVLER		54	12	14
FSVSQTGDLTITNVQR		85	16	14
GLKPNAIYLFLVR		43	13	14
IVEHPSDLIVSK		47	12	14
NSVVIPDLR		30	9	14
NSVVIPDLRK		46	10	14
RPSNLAVTVDDSAEFK		51	16	14
SRPDEGVYVCVAR		41	13	14
TLEEAPSAPPQGVTVSK		59	17	14
TVDGSTFSVVIPFLVPGIR		43	19	14
YSVEVAASTGAGSGVK		77	16	14
ALGVNAMLR		40	9	5
QDGDQFYIK		43	9	5
SLATWENENK		36	10	5
SSENFDELLK		34	10	5
VAVAAASKPHVEIR		32	14	5
DALVDFSEQYTPEADPYFIQDR		80	22	2
EAQPGQSQVSYQGLPVQK		60	18	2
AALLLDQYR		48	9	10
ICCQFDFK		31	8	10
IEQLEQLLEENHEIISHIK		41	19	10
LYNLPGVEGLSLDISSLVDIR		93	21	10
PSFFSISPQDCQFALGGR		63	18	10
SNVLLVPLGDDFR		39	13	10
TGVEPGARPPGFPVLSGDFFSYADR		41	25	10
WWDNINVQK		34	9	10
YDKPQEWDAQFFNYQR		35	16	10
YPLSDFTLLTEAR		88	13	10

AGLGHPAAFGR	1N-ac	56	11	7
DENATLDGGDVLFTGR		73	16	7
GAEILADTFK		63	10	7
GHVLLHR		35	7	7
LTVPDDIAANCIYLNIPNK		62	19	7
SFCSMAGPNLIAIGSSESAQK		41	21	7
VDGLLTCCSVLINK		60	14	7
AFELYDQDGNGYIDENELDALLK		114	23	4
ELQNLIQELQQAR		65	13	4
LAEYTDLMLK		49	10	4
LLPVQENFLLK		34	11	4
ALSGNVIAWAESHIEIYGGATK		47	22	12
FQLTFPLR		44	8	12
IETALTSLHQR		85	11	12
KLTPGEVYNLATCSTK		66	16	12
LENLEQYSR		57	9	12
LPFVINDGK		52	9	12
LTPGEVYNLATCSTK		84	15	12
SSVLQLR		51	7	12
TPAAETLSQLGQTLQSLK		126	18	12
TRLENLEQYSR		56	11	12
VKIETALTSLHQR		75	13	12
WTFEACR		39	7	12
ELEAVCQDVLSLLDNYLIK		69	19	3
NVTELNEPLSNEER		73	14	3
TAFDDAIAELDTLNEDSYK		52	19	3
CDENILWLDYK		52	11	16
DPVQEAWAEDVDLR		78	14	16
EAEAAMFHR		45	9	16
FGVEQDVDMVFASFIR	1Met(ox)	54	16	16
GADFLVTEVENGGSLGSK		91	18	16
GDLGIEIPAEK		49	11	16
GDYPLEAVR		41	9	16
GIFPVLCKDPVQEAWAEDVDLR		35	22	16
GVNLPGAAVDLPAVSEK		71	17	16
IENHEGVR		33	8	16
IYVDDGLISLQVK		84	13	16
LDIDSPPITAR		69	11	16
LFEELVR		43	7	16
NTGIICTIGPASR		62	13	16
RFDEILEASDGIMVAR		40	16	16
TATESFASDPILYRPVAVALDTK		33	23	16
GADQAELEEIAFDSSLVFIPAEFR		29	24	7
IEEGVPQFLVLISSGK		29	16	7
ISLSPEYVFSVSTFR		61	15	7
LLPSFVSSENAFYLSPDIR		64	19	7
LNLLDLDYELAEQLDNIAEK		125	20	7
QINVGNALEYVSR		68	13	7
VGVVQFSNDVFPEFYLK		73	17	7
ALSSEWKPEIR		51	11	9
DISWFSPNGEK		30	11	9
DKDISWFSPNGEK		64	13	9
EASMEGIVTIVGLKPETTYAVF	1Met(ox)	31	22	9
EGEDAVIVCDVVSSLPPTIIWK		78	22	9
KTDEGTYR		38	8	9
KVDKNDEAEYICIAENK		34	17	9

NAPTPQEFR	29	9	9
VSSLTLK	29	7	9
APEVSQYIYQVYDSILK	33	17	4
DIPNENELQFQIK	30	13	4
LQNNNVYTIAK	43	11	4
QVFLATWK	37	8	4
ENSLLFDPLSSSSSNK	41	16	2
QQQDYWLIDVR	32	11	2
EVSQFTPVAFPIAK	29	14	2
VEESGVSISWNPPNGPAAR	42	19	2
APSSQNWLTVNGGK	76	14	37
CGILSSGNNLFFNEDGLR	55	18	37
CIQGNCVCDEQWGGLYCDDPETSLPTQLK	45	29	37
CSGSVSQPSVFFPTK	37	15	37
DKTHNALSSR	36	10	37
DLTLKPGYVLQFK	35	13	37
EDNWFFYPGGNIGLYCPYSSK	66	21	37
EHITLDTLSYSSYK	55	14	37
ELDFMSFLEPQIISIDLPQDAK 1Met(ox)	49	22	37
FLQFTLR	40	7	37
FLQYWGR	28	7	37
FSYSDPSIIVLYAK	63	14	37
FVYLELPAAAK	71	11	37
GAEVSFGCGVLASGK	69	15	37
GENVQFQWK	54	9	37
HDGLDQNDWAIDNVLISGSADQR	33	23	37
IISVELPGDAK	63	11	37
ITGAQVGTGCGTLNDGK	95	17	37
ITIPLPNAALTR	49	12	37
ITVYLPLSTISPR	44	13	37
ITYPLPESLVGNPVR	37	15	37
KQNYMMNFSR	35	10	37
LSSYHNFYSIR	29	11	37
NPDFLKDDFEGQLESDR	31	17	37
QAATKPLDLTR	39	11	37
QAVTQDLDLR	79	10	37
QLITSFLDSSQSR	89	13	37
QTGPILGNMWAIDNVYIGPSCLK	39	23	37
QVVLEDSLDPVDTGNWLFFPGATVK	84	25	37
RVIVLLPQK	30	9	37
SGTSLIFK	45	8	37
TSGITCIKPR	29	10	37
VIVLLPQK	53	8	37
VPSLVSVVINPELQTPATK	94	19	37
VSYNVPLEAR	45	10	37
WWQPFVISNGIVVSGVER	53	18	37
YIALEIPLK	35	9	37
AGVQVWLGANGK	34	12	44
AISGLTIDGHAVGAK	83	15	44
ALASYVAACQAAGVVIEDWR	86	20	44
APGWDPLCWDECR	55	13	44
CSVQNGLLGCYPDR	85	14	44
EEFLTAFLQNYQLAYSK	72	17	44
EYPGQVLVDDVLQYLPFQAADGQVQVFR	35	28	44
FAVLQENVAWGNGR	60	14	44
FLLSQGVCIPVQDCGCTHNGR	27	21	44

FQDQVCGLCGNYNGDPADDFLTPDGALAPD	38	39	44
GATTSPGVYELSSR	74	14	44
GCGEGCGPQGCPVCLAEETAPYESNEACG(	35	32	44
GEVGFVLVDNQR	70	12	44
GGGQAANALAFGNSWQEETRPGCGATEPG	35	33	44
GNPAVSYVR	46	9	44
GVWVNGLR	36	8	44
ISVAQGASK	53	9	44
LASVSVSR	37	8	44
LDDGDYLCEDGCQNNCPACTPGQAQHYEG	48	31	44
LDSLVAQQLQSK	67	12	44
LLISSLSESPASVSILSQADNTSK	96	24	44
LPVSLSEGR	58	9	44
LPVVLANGQIR	46	11	44
LTYNHGGITGSR	50	12	44
NAAGHLQR	37	8	44
NECGILADPK	27	10	44
NEVTYDPYLVLIPDVAAYCPAYVVK	31	25	44
NPQGPFATCQAVLSPSEYFR	72	20	44
PAGWQVGGAQGCGECVSK	80	18	44
PFLEQCVYDLCVVGGER	71	17	44
SLAAYTAACQAAGVAVKPWR	58	20	44
SPANCPLSCPANSR	78	14	44
SRLPVSLSEGR	37	11	44
TCQGSCAALSGLTGCTTR	109	18	44
TPDGSLLVR	49	9	44
VAVIVSNDHAGK	72	12	44
VAYDLVYYVR	72	10	44
VLVENEHR	52	8	44
VNGVLTALPVSVADGR	82	16	44
VPSSYAEALCGLCGNFNGDPADDLALR	89	27	44
VSYVGLVTVR	73	10	44
VTLQPYNVAQLQSSVDLSGSK	118	21	44
VVAEVQICHGK	37	11	44
YDLAFVVASQATK	73	13	44
EGDDIEMPCAFR	35	12	3
ELLHELALSVPGAR	52	14	3
LQDEGVYECR	47	10	3
DRETVLSSAL	33	10	3
SSHLVFINTR	33	10	3
WLDGTSPDYK	38	10	3
DYGGYLSTYILPAK	59	14	5
IYFLSTEDLPR	76	11	5
MFDLETNEHVK	52	11	5
VSALEEQQFLIIHPTADEK	27	19	5
VTVEDLFSEDFK	73	12	5
VPLALFALNR	46	10	1
ALLSYDGLNQR	65	11	4
DGVMFQIDQATK	54	12	4
FFDIQLGIK	45	9	4
LFEYILLYK	49	9	4
CFETVYDGYSK	55	11	2
CPAMVAYCMTTR	47	12	2
IFNSFVYTEK	58	10	2
ISVLTVADTVR	59	11	2
CGLQELGPGLFR	47	12	5

GLAALQYLYLQDNALQALPDDTFR	63	24	5
IDAAAFTGLALLEQLDLSDNAQLR	54	24	5
SVDPATFHGLGR	44	12	5
VAHVHPHAFR	37	10	5
ATLITFLCDR	48	10	3
WYTSYACPEEPLECVVTDPSTLEQYDLSSLA	29	32	3
YVDQVLQLVYK	27	11	3
EDQLPSGFPNIDMGPQLK	53	18	10
GGQFLTPLGSPEDMDLEELIQDISR	60	25	10
HNVDDSLLTTVGSLLEDETYTVR	59	23	10
NVLELTDVK	27	9	10
SPQGLGAFTPVVR	63	13	10
TGEQAPASAPR	96	11	10
TSVLLSWEFPDNYNSPTPYK	58	20	10
VLAFTSVGDGPLSDPIQVK	76	19	10
WEPPAGTAEDQVLGYR	43	16	10
YSIGGLSPNSEYEIWVSAVNSIGQGGPPSESV'	52	34	10
TDVVTVNPSIITER	47	14	1
PTLSALPSPLVTSGK	27	15	3
SPMDTFLLIK	33	10	3
SVTLLCQSR	47	9	3
DVHEGQPLLNVK	35	12	13
DWVIPPINLPENSR	44	14	13
ESAEVEEIVFPR	51	12	13
FLEAGIYEVPIIITDSGNPPK	73	21	13
FLIYAQDK	29	8	13
FLIYAQDKETQEK	55	13	13
IDPVNGQITTIAVLDR	56	16	13
LNGDFAQLNLK	48	11	13
LSLKPTLTEESVK	42	13	13
SFPLSSEHAK	28	10	13
TGFPEDVYSAVLSK	79	14	13
VQYESSEPADFK	54	12	13
VQYESSEPADFKVDEDGMVY 1Met(ox)	32	23	13
DTCLGDSGGAFVIFDDLSQR	89	20	3
NAEPGLFPWQALIVVEDTSR	29	20	3
TLSDVLQYVK	52	10	3
ELQDLALQGAK	55	11	5
GLSAEPGWQAK	54	11	5
SGEATDGARPQALPEPMQESK	43	21	5
SGELEQEEER	80	10	5
YPGPQAEGDSEGLSQGLVDR	95	20	5
KKDEGLYECR	55	10	4
VNANSHAR	53	8	4
VQGNDISHK	31	9	4
VTDANYGELQEHK	77	13	4
ASLTLTLLR	58	9	6
GAVLTATVLAR	40	11	6
NVAVTVEYGPR	62	11	6
SDGGAVLALGLLGPVTR	44	17	6
SGELGAVIEGLLR	87	13	6
TFSLSPDAPR	46	10	6
IMGITLVSK	37	9	2
LLGLSLAGK	60	9	2
AICDHVR	30	7	12
DLDVAILVGSMPR	50	13	12

DVIATDKEDVAFK	56	13	12
ELTEEKESAFEFLSSA	47	16	12
ESAFEFLSSA	38	10	12
EVGVYEALKDDSWLK	55	15	12
FVEGLPINDFSR	47	12	12
GEFVTTVQQR	63	10	12
KLSSAMSAAK	47	10	12
LGVTANDVK	50	9	12
VIVVGNPANTNCLTASK	92	17	12
VLVTGAAGQIAYSLLYSIGNGSVFGK	41	26	12
ANTFVAELK	37	9	4
IFGVTTLDIVR	51	11	4
SQETECTYFSTPLLLGK	49	17	4
VAVLGASGGIGQPLSLLLK	62	19	4
AIIDGVESVSR	73	11	8
EENDIQTGSALLPLSPESK	39	19	8
FAVLYQQLDGEDQTK	96	15	8
FPLEMQIYCFDADR	45	14	8
TSLENTFIHNTGK	78	13	8
TVEINLTNDYR	42	11	8
VSGGVSEMVFK	92	11	8
VTSVSSDSQTGMDR	72	14	8
GDPGEAGPQGDQGR	54	14	10
GDPGFEGER	56	9	10
GLEQLLVGGSHLK	58	13	10
GTYTDCAIKK	38	10	10
GVFHQTVSR	42	9	10
IALVITDGR	77	9	10
LKPYGALVDK	33	10	10
LLLFSDGNSQGATPAAIEK	108	19	10
TDPAHDVR	34	8	10
VPSYQALLR	58	9	10
ADEGIQPDPYYGLK	31	14	2
ILSVAVNDEGSR	40	12	2
ASSIIDELFQDR	87	12	21
ASSIIDELFQDRFFTR	42	16	21
EILSVDCSTNNPSQAK	87	16	21
EIQNAVNGVK	39	10	21
ELDESLQVAER	80	11	21
EPQDTYHYLPFSLPHR	51	16	21
FMETVAEK	47	8	21
IDSLLENDR	66	9	21
KTLLSNLEEAK	70	11	21
KTLLSNLEEAKK	87	12	21
KYNELLK	58	7	21
LANLTQGEDQYYLR	84	14	21
LFDSDPITVTVPVEVSR	114	17	21
QQTHMLDVMQDHFSR	43	15	21
RELDESLQVAER	80	12	21
RPHFFFPK	39	8	21
SGSGLVGR	66	8	21
TLLSNLEEAK	74	10	21
TLLSNLEEAKK	44	11	21
VTTVASHTSDSDVPSGVTEVVVK	94	23	21
YVNKEIQNAVNGVK	59	14	21
FDGGVEAIATR	68	11	2

SSSGLTYIAEWK		28	12	2
DFTCVHQALK		51	10	18
FPVFMGR		37	7	18
FQPTLLTLPR		59	10	18
GVTSVSQIFHSPDLAIR		81	17	18
HRLEDMEQALSPSVFK		51	16	18
IKVTTSQDMLSIMEK		70	15	18
KVETNMAFSPFSIASLLTQVLL	1Met(ox)	32	29	18
KYPVAHFIDQTLK		54	13	18
LEDMEQALSPSVFK	1Met(ox)	90	14	18
LLDSLPSDTR		64	10	18
LVLLNAIYLSAK		90	12	18
LYHAFSAMK		53	9	18
TLYSSSPR		33	8	18
TNLESILSYPK		94	11	18
VETNMAFSPFSIASLLTQVLLC	1Met(ox)	76	28	18
VPMMNSK	2Met(ox)	31	7	18
VTTSQDMLSIMEK		91	13	18
YPVAHFIDQTLK		53	12	18
ACDGINDCGDQSDELCCK		127	18	17
ADSPMDDFFQCVNGK		43	15	17
AQLGDLPWQVAIK		71	13	17
EANVACLDLGFQQGADTQR		110	19	17
EMECAGTYDGSIDACK		97	16	17
GLETSLAECTFTK		77	13	17
HGNTDSEGIVEVK		79	13	17
IIFHENYNAGTYQNDIALIEMK		54	22	17
IVIEYVDR		58	8	17
KYTHLSCDK		31	9	17
PFISQYNV		42	8	17
RAQLGDLPWQVAIK		29	14	17
TMGYQDFADVVCYTQK	1Met(ox)	87	16	17
VANYFDWISYHVGR		56	14	17
VFSLQWGEVK		61	10	17
VTYTSQEDLVEK		70	12	17
YQIWTTVVDWIHPDLK		46	16	17
EANNYEEDPNKPTSWTENQAGK		46	22	15
EKETLITIMK		47	10	15
ELSAERPLNEQIAEAEEDK		69	19	15
FQDDPDGLHQLDGTPLTAEDIVHK		44	24	15
GENDETVSNTLTLTNGLER		94	19	15
GILDKEEAEAIKR		69	13	15
GKTEAYLEAIRK		35	12	15
LLNLGLITESQAHTLEDEVAEVLQK		75	25	15
LNVEDVDSTK		44	10	15
TEAYLEAIR		49	9	15
TEAYLEAIRK		33	10	15
TLIDFVK		34	7	15
TYSEDNFEELQYFPNFYALLK		66	21	15
VTPMAAIQDGLAK		45	13	15
YGTISPEEGVSYLENLDEMIAL	1Met(ox)	66	25	15
QGFSYQCPQGQVIVAVR		48	17	2
YFESVLDR		37	8	2
ALLEAPLK		41	8	6
EICPAGHGYTYASSDIR		72	17	6
EQDAPVAGLQPVER		64	14	6

IYFCQIPCLNGGR	41	13	6
QSTFTLPLSNQLASVNPSLVK	77	21	6
STPLGQQQPAPR	86	12	6
ELTIQMDQQLR	40	11	2
LALHLNLGDSK	29	11	2
AAISGENAGLVR	96	12	22
ADVQAHGEGQEFSITCLVDEEEMK	30	24	22
ADVQAHGEGQEFSITCLVDEEEMKK	30	25	22
DKVTAWK	30	7	22
ELAAQTIKK	31	9	22
EVAFDLEIPK	56	10	22
FAHYVVTSQVVNTANEAR	120	18	22
GFSLDEATNLNGGLLR	98	16	22
GHMLENHVER	53	10	22
GSLVQASEANLQAAQDFVR	117	19	22
IADNKQSSFK	40	10	22
ILGDMQPGDYFDLVLFGTR 1Met(ox)	70	19	22
IYEDHDATQQLQGFYSQVAK	40	20	22
LDAQASFLPK	71	10	22
LWAYLTIQELLAK	96	13	22
NHMQYEIVIK	43	10	22
QAVDTAVDGVFIR	87	13	22
QYYEGSEIVVAGR	80	13	22
TAFISDFAVTADGNAFIGDIK	99	21	22
TMEQFTIHLTVNPQSK 1Met(ox)	44	16	22
VTFQLTYEEVLK	59	12	22
VTYDVSR	28	7	22
TPFSVGWQTVTTVPEVIDGK	63	20	2
VLLNWEQVK	39	9	2
AQWANPFDPSK	48	11	6
FSISATYDLGATLLK	81	15	6
GTEAAAVPEVELSDQPENTFLHPIIQIDR	34	29	6
GWVDLFVPK	54	9	6
NALALFVLPK	59	10	6
SILFLGK	29	7	6
EYYFAEAQIADFSDPAFISK	110	20	14
FAFNLYR	39	7	14
GGETAQSADPQWEQLNNK	95	18	14
GNFLAANDQELDCDILQLEYV 1Met(ox)	49	33	14
GPLDQLEK	39	8	14
IAIDLFK	36	7	14
LNILNAK	45	7	14
NYNLVESLK	44	9	14
QFPILLDFK	48	9	14
SVNDLYIQK	30	9	14
TLEAQLTPR	43	9	14
TNNHIMK	29	7	14
TSCLLFMGR	50	9	14
YEITTIHNLFR	44	11	14
AGYIIPLQGPGLTTTESR	47	18	3
GAYTQVIFLAR	30	11	3
VTSEGAGLQLQK	43	12	3
LLCGLLAER	46	9	3
PMFIVNTNVPR	77	11	3
VYINYYDMNAANVGWNNSTFA	62	21	3
DILQSCQTSEECELAR	91	16	3

DPVASSLSPYFGTK	65	14	3
LASLFPALFSR	45	11	3
NEDIIDPVEDR	51	11	1
TFTVSSTLDFR	45	11	3
TSGSKPAADIR	28	11	3
VDQNDNTSLQWSNPAQQTLYFDDKK	29	25	3
GFGPPATNQFTTK	57	13	1
DVDECSLKPSICGTAVCK	36	18	16
FRLPEISR	32	8	16
FSAEFDFR	49	8	16
GGKIEVQLK	39	9	16
HCLVTVEK	32	8	16
IETISHEDLQR	65	11	16
IQALSLCSDQQSHLEFR	51	17	16
ITTGGDVINNGLWNMVSVEEL 1Met(ox)	29	28	16
KVESELIKPINPR	92	13	16
NNLELSTPLK	57	10	16
QLAVLDK	27	7	16
QSTNAYPDLR	68	10	16
SCEVVSVCLPLNLDTK	97	16	16
SFQTGLFTAAR	79	11	16
SQDILLSVENTVIYR	103	15	16
VYFAGFPR	35	8	16
IPGIFELGISSQSDR	40	15	6
KPYNVESYTPQTQGK	73	15	6
LLCNGDNDCGDQSDEANCR	136	19	6
SGFSFGFK	63	8	6
VEPLYELVTATDFAYSSTVR	88	20	6
VKVEPLYELVTATDFAYSSTVR	33	22	6
DQPFTILYR	36	9	4
GPYSNPYSTPYSGPYPAAAPPLSAPNYPTISI	46	32	4
IYVSQYPF	42	8	4
QSGQCLDIDECR	51	12	4
DTVNLYTSSGCLCPPLNVNEE 1Met(ox)	27	32	3
LLLVEGSIAEK	57	11	3
NNYNYVIR	45	8	3
AEQCAEFDGAEFQGR	29	15	2
LSCSDDTAPESQAAWQK	47	17	2
EYGGLDVLVNNAGIAFK	93	17	3
GIGLAIVR	43	8	3
VVNVSSIMSVR	32	11	3
AAGLLSTYR	52	9	5
AFLSSHLQDLSTIVR	77	15	5
GGVLFAITDAFQK	54	13	5
LVDNYCEAWR	64	10	5
TADTAVTGLASPLSTGK	87	17	5
DLELLIQTATR	67	11	3
LVTLEEFLASTQR	65	13	3
YLQEVIDVLETDGHFR	31	16	3
GQDHCGIESEVVAGIPR	61	17	5
LCGTFLGGPKPPQR	53	14	5
NGPVEGAFSVYSDFLLYK	82	18	5
SGVYQHVTGEMMGGHAIR	63	18	5
VMFTEDLK	39	8	5
LESLPAHLPR	78	10	3
SDTAYQWNLK	62	10	3

SLEVLNLSSNK	83	11	3
AGNSQGDFYIR	48	11	2
EQPSSIVHR	31	9	2
AASNIIFSNGNLDPWAGGGIR	66	21	2
DLFLQGAYDTVR	64	12	2
DYFIATCK	49	8	22
ESEQGVYTCTAQGIWK	57	16	22
FCGQLGSPLGNPPGK	65	15	22
GGGALLGDR	46	9	22
IQYYCHEPYYK	43	11	22
LFGEVTSPLFPK	69	12	22
LFGEVTSPLFPKPYPNNFETTTVITVPTGYR	53	31	22
LGNHPIR	39	7	22
LPVANPQACENWLR	80	14	22
LVFQQFDLEPSEGCFYDYVK	84	20	22
MDVFSQNMFCAGHPSLK	52	17	22
MGNFPWQVFTNIHGR	67	15	22
PVNPVEQR	55	8	22
PYPNNFETTTVITVPTGYR	105	19	22
QDACQGDSGGVFAVR	135	15	22
QRPPDLDTSSNAVDLLFFTDESGDSR	70	26	22
TLDEFTIIQNLQPQYQFR	83	18	22
VLNYVDWIK	50	9	22
VLNYVDWIKK	41	10	22
VSVHPDYR	44	8	22
WILTAAHTLYPK	57	12	22
YTTTMGVNTYK	50	11	22
DKLTELQLR	28	9	2
MALFAGGK 1Met(ox)	28	8	2
AITPPHPASQANIIFDITEGNLR	38	23	2
SAATLQQEK	55	9	2
AAQAQGQSCEYSLMVGYQC(1Met(ox)	72	24	27
CCHCCLLGR	35	9	27
CENTLGSYLCSCSVGFR	47	17	27
CLAFECPENYR	36	11	27
CVDVDECAPPAEPCGK	79	16	27
DCSLPYATESK	51	11	27
DIDECESGIHNCLPDFICQNTLGSFR	42	26	27
DLLLTVK	39	7	27
DSSCGTGYELTEDNSCK	92	17	27
GYHLNEEGTR	49	10	27
GYQLSDVDGVTCEDIDECALPTGGHICSYR	80	30	27
HGTVSSFVAK	65	10	27
IIEVEEEQEDPYLNDR	102	16	27
ITYYHLSFPTNIQAPAVVFR	88	20	27
KVSPHSGVVALTK	42	13	27
KVSPHSGVVALTKPVPEPR	77	19	27
LGESCINTVGSFR	89	13	27
LPCHENR	46	7	27
MCVDVNECQR	69	10	27
MGPSSAVPGDSMQLAITGGN 1Met(ox)	44	28	27
MVQEQCCHSQLEELHCATGISLANEQDR	41	28	27
RGYQLSDVDGVTCEDIDECALPTGGHICSYR	43	31	27
SQETGDLDVGGLQETDK	130	17	27
SQETGDLDVGGLQETDKIIEVEEEQEDPYLNI	68	33	27
TGYYFDGISR	77	10	27

VSPHSGVVALTK		29	12	27
VSPHSGVVALTKPVPEPR		81	18	27
AASGTQNNVLR		76	11	33
ACGACPLWGK		39	10	33
CFSGQCISK		32	9	33
DGFVQDEGTMFPVGK	1Met(ox)	73	15	33
DSCTLPASAEK		36	11	33
ELENALK		31	7	33
ELSHLPSLYDYSAYR		47	15	33
EQTMSECEAGALR		84	13	33
GCPTEEGCGER		41	11	33
GGGAGFISGLSYLELDNPAGNK		105	22	33
GQSISVTSIRPCAAETQ		60	17	33
ILPLTVCK		31	8	33
KVFSGDGK		31	8	33
KVFSGDGKDFYR		98	12	33
LIDQYGTHYLQSGSLGGEYR		71	20	33
LKQNDFNSVEEK		74	12	33
LSGNVLSYTFQVK		80	13	33
LTPLYELVK		27	9	33
MHVLHCQGR		33	9	33
MPYECGPSLDVCAQDER	1Met(ox)	81	17	33
NVVYTCNEGYSLIGNPVAR		84	19	33
QNDFNSVEEKK		50	11	33
SLVCNGDSDCDEDSADEDR		128	19	33
SLVCNGDSDCDEDSADEDRCEDSER		44	25	33
SRECNNPPPSGGGR		47	14	33
SSGWHFVVK		42	9	33
SVAVYGQYGGQPCVGNAFETQSCEPTR		79	27	33
SYTSHTNEIHK		53	11	33
VFSGDGKDFYR		36	11	33
VLFYVDSEK		47	9	33
VTVSCSGGMSLEGPSAFLCGSSLK		82	24	33
WLVGEMHCQK		45	10	33
YSAWAESVTNLPQVIK		53	16	33
AEDEENEKETAVSTEDDSHHK		43	21	18
EEDIDENLLF		29	10	18
GHQLQLDYFGACK		73	13	18
HSASDDYFIPSQAFLEAER		75	19	18
KGHQLQLDYFGACK		30	14	18
LSENTDFLAPGVSSFTDSNQQESITK		51	26	18
LSENTDFLAPGVSSFTDSNQQESITKR		40	27	18
MQEDEFDQGNQEQEDNSNAI	1Met(ox)	61	31	18
NHGVDDDGDDDGDDGGTDGPR		57	21	18
NILMQLYEANSEHAGYLNEK		72	20	18
NYHMYVYPVHWQFSELDQHF	1Met(ox)	33	23	18
QEEDNTQSDDILEESDQPTQVSK		69	23	18
SIPTCTDFEVIQFPLR		42	16	18
SSSQELGLK		53	9	18
TVSEALLMEPTDDGNTTPR	1Met(ox)	102	19	18
VHAVDSCMSFQCK		65	13	18
VHENENIGTTEPGEHQEAK		88	19	18
VLTHSELAPLR		58	11	18
GQVEQANQELQELIQSVK		104	18	6
LQEGQTLEFLVASVPK		62	16	6
LVTPGETPSWTGSGFVR		57	17	6

QLDALLEALK	43	10	6
SLADVDAILAR	68	11	6
YSEIEPSTEGEVIYR	61	15	6
APLQGTLLGYR	68	11	3
LAYQGQDTPEVLMDIGLR	77	18	3
TATITVLPQQPR	40	12	3
APSFWYK	33	7	12
CYLITVTPVYADGPGSPESIK	46	21	12
DASTWSQIPPEDTASTR	49	17	12
DNMLWVEWTTPR	31	12	12
IDPSHTQGYR	45	10	12
ILDYEVTLTR	75	10	12
LTWTNPSIK	34	9	12
NEAVLEWDQLPVDVQNGFIR	75	20	12
SSFTVQDLKPFTEYVFR	30	17	12
TVQLVWK	32	7	12
YILEWCVLSDK	38	11	12
YLATLTVR	50	8	12
ALSIGFETCR	43	10	2
YGFIEGHVVIPR	32	12	2
LLAPAALAFR	53	10	2
NFVQLCLSAEPSEAPR	91	16	2
GAHLASADELRR	36	12	2
LVSISVGR	53	8	2
ALSLAPLAGAGLELQLER	67	18	2
SYSESSSTSSSESLNSSAPR	134	20	2
CYPHPGSELPLQALVMGEGTCEK	37	23	12
EKVTEQHR	35	8	12
EPGCGCCSVCAR	44	12	12
GDPECHLFYNEQQEAR	66	16	12
GECWCVNPNTGK	61	12	12
HGLYNLK	40	7	12
HHLGLEEPK	45	9	12
HHLGLEEPKK	59	10	12
LAACGPPPVAPPAAVAAVAGGAR	62	23	12
LEGEACGVYTPR	80	12	12
LIQGAPTIR	46	9	12
TPCQQELDQVLER	103	13	12
GIMGEDTYPYQGK	58	13	3
MALNQFSDMSFAEIK	79	15	3
NMCGLAACASYPIPLV	42	16	3
DRDLEVDTTLK	61	11	12
FTYSVTVDGCTSHTGAWGK	97	19	12
GEAGPQGPR	41	9	12
GFSGLDGAK	39	9	12
GPAGPQGPR	36	9	12
NCPGAEVPEGECCPVCPDGSESPTDQETTC	42	35	12
NSVAYMDQQTGNLKK	72	15	12
SGEYWIDPNQGCNLDAIK	74	18	12
SLSQQIENIR	83	10	12
STGGISVPGPMGPSGPR	30	17	12
VFCNMETGETCVYPTQPSVAQK	67	22	12
VLCDDVICDETK	67	12	12
ADPEEELSLTFALR	55	14	5
AYPDVAALSDGYWVVSNR	58	18	5
LSELVQAVSDPSSPQYGK	72	18	5

VPIPWVSGTSASTPVFGGILSLINEHR	36	27	5
YLTLENVADLVR	65	12	5
AFDITYVR	27	8	13
DGSEASLEWSSER	76	13	13
DTLQEANDILNNLK	83	14	13
EAQQALGSAAADATEAK	100	17	13
HKQEADDIVR	27	10	13
LNTFGDEVFNDPK	63	13	13
LQELESLIANLGTGDEMVTDQAFEDR	30	26	13
LSAEDLVLEGAGLR	80	14	13
NTIEETGNLAEQAR	113	14	13
PSAYNFDNSPVLQEWVTATDIR	41	22	13
SAGYLDDVTLASAR	73	14	13
TFAEVTDLDNEVNNMLK	91	17	13
VSVPLIAQGNSYPSETTVK	48	19	13
EDQTSPAPGLR	44	11	3
GPWCYVSGEAGVPEK	90	15	3
SEAAAVQPVIGISQR	74	15	3
AHYGGFTVQNEANK	79	14	10
DNDGWLTSDPR	50	11	10
DNENVVNEYSSELEK	70	15	10
EDGGGWWYNR	37	10	10
HQLYIDETVNSNIPTNLR	71	18	10
MGPTELLIEMEDWK	48	14	10
NSVDELNNNVEAVSQTSSSSFQYMYLLK	82	28	10
NYCGLPGEYWLGNDK	37	15	10
QGFGNVATNTDGK	70	13	10
YQISVNK	37	7	10
ALVILAK	37	7	9
APLVLKD	29	7	9
DGLILTSR	33	8	9
EGPYDVVVLPGGNLGAQNLSESAAVK	59	26	9
EILKEQENRK	36	10	9
EVAAQVK	29	7	9
GAEEMETVIPVDVMR 1Met(ox)	31	15	9
GPGTSFEFALAIVEALNGK	88	19	9
VTTHPLAK	36	8	9
FASEYGYQSWPSFSTLEK	62	18	11
FQSAVLYAAQQSK	79	13	11
GGEAVCLYEEPVSELLR	67	17	11
GSPGLSFYFK	39	10	11
LLLQSVVDANMNTLR	57	15	11
LQTQQTYSIELQPGK	32	15	11
NELEQSFHVTSLTDIY	27	16	11
TILFYPWEPTSK	28	12	11
VNLILEGVDTVSK	55	13	11
VSSTEDWSFNSK	107	12	11
YSFDITNVVR	32	10	11
ATFGCHDGYSLDGPEEIECTK	62	21	14
ATVVYQGER	58	9	14
CSYTEDAQCIDGTIEVPK	93	18	14
EHSSLAFWK	30	9	14
FICPLTGLWPINTLK	44	15	14
FKNGMLHGDK	44	10	14
KATVVYQGER	74	10	14
KFICPLTGLWPINTLK	55	16	14

NGMLHGDK	1Met(ox)	48	8	14
TCPKPDDLPFSTVVPLK		40	17	14
TFYEPGEEITYSCK		39	14	14
TFYEPGEEITYSCKPGYVSR		94	20	14
VCPFAGILENGAVR		76	14	14
WSPELPVCAPIICPPPSIPTFATLR		54	25	14
DVTVLQNTDGNNNEAWAK		76	18	12
EDAIWNLLR		46	9	12
FQLFGSPSGQK		28	11	12
GGSFQLNELQGLK		32	13	12
IDSGLYLGSGYFTAIQNLR		91	19	12
KPVTEAR		49	7	12
KSEEEVAAR		35	9	12
LADFALLCLDGK		60	12	12
LCAGTGENK		31	9	12
LRPVAAEVYGTERQPR		42	16	12
RKPVTEAR		30	8	12
YLGPQYVAGITNLK		54	14	12
FDDYNSDSSLTLR		89	13	5
GPDVGVGESQAEEPR		115	15	5
VLQSIGVDPLPAK		50	13	5
VLVVDIQAQK		64	10	5
VYPESQAQEPGVAASLR		88	17	5
AVRPGYPK		27	8	12
CQCDELCSYYQSCCTDYTAECKPQVTR		55	27	12
DVWGIEGPIDAAFTR		77	15	12
DWHGVPGQVDAAMAGR		47	16	12
FEDGVLDPDYPR		81	12	12
GQYCYELDEK		39	10	12
IYISGMAPR		73	9	12
MDWLVPATCEPIQSVFFFSGDK		37	22	12
QPQFISR		33	7	12
RVDTVDPPYPR		37	11	12
SIAQYWLGCPAPGHL		75	15	12
VDTVDPPYPR		48	10	12
GVAALTSDPAVQAIVLDTASDVLDK		29	25	2
VLVQNAAGSQEK		38	12	2
DLVGELGTALR		84	11	1
ACTSQGCGKPITEESSTLGEGSK		102	23	38
CTASNFLGTATHDFHVIVEEPPR		57	23	38
DGENYATVVGYSAFLHCEFFASPEAVVSWQ		31	31	38
DGNPFYFTDHR		29	11	38
DTATLSWGLPK		49	11	38
EKIDPLEVEEGDPIVLPCNPPK		52	22	38
GAGPESEPYIFQTPEGVPEQPTFLK		71	25	38
GDLYFANVEEK		65	11	38
GEILLLECFAEGLPTPQVDWNK		41	22	38
GNPEPTFSWTK		44	11	38
GYQINWWK		40	8	38
IDPLEVEEGDPIVLPCNPPK		55	20	38
IENVSYQDK		43	9	38
IENVSYQDKGNYR		69	13	38
IGGDLPK		44	7	38
IPNEGHISHFQGK		53	13	38
KPQSAVYSTGSNGILLCEAEGEPQPTIK		29	28	38
KTTVILPLAPFVR		57	13	38

LGIAMSEEIEFIVPSVPK		53	18	38
LHMLELHCESK		34	11	38
LLLPPTESGSESSITILK		74	18	38
NDYCCFAAFPR		29	11	38
PITEESSTLGEGSK		106	14	38
SMEQNGPGLEYR	1Met(ox)	67	12	38
SQPSQPSDHHETPPAAPDR		62	19	38
SQPSQPSDHHETPPAAPDRNPQNIR		32	25	38
TAVTANLDIR		47	10	38
THPVEVFEPGAEHIVR		28	16	38
TLKIENVSYQDK		76	12	38
TTEEDAGSYSCWVENAIGK		78	19	38
TTVILPLAPFVR		51	12	38
VDKDTATLSWGLPK		72	14	38
VEEVKPLEGR		45	10	38
VIAVNEVGR		60	9	38
VIKVDKDTATLSWGLPK		67	17	38
VMTPAVYAPYDVK	1Met(ox)	62	13	38
VNGSPVDNHPFAGDVVFPR		65	19	38
VQVAFPFDEYFQIECEAK		83	18	38
DQQVVTAVEYQEAILACK		55	18	2
TGTLQFNTVSK		61	11	2
GPVSVGVDAR		45	10	1
AQYVLISPEASSR		44	13	8
GAVYVYFGSK		41	10	8
IADVTSGLIGGEDGR		111	15	8
ILEGFQPSGR		40	10	8
LSGALHVYSLGSD		27	13	8
QVLLVGAPTYDDVSK		52	15	8
TMFIGGSQLSQK		68	12	8
VAFLTVTLHQGGATR		59	15	8
SLGLPENHIVFPVPIDQCIDG		34	21	2
VPLQQNFQDNQFQGK		48	15	2
LGPEAFWFNSGR		61	12	1
CPPGAHACGPCLQPFQEDQQGLCVPR		43	26	2
LEDEIDFLAQELAR		106	14	2
EPGEGLAVTVSLIGAYK		48	17	15
FCGDAVPGSISSEGNELLVQFVSDLSVTADG		57	37	15
FCGTFRPAPLVAPGNQVTLR		42	20	15
FDLEPDTYCR		30	10	15
GESGYVASEGFPNLYPPNK		35	19	15
GFLLWYSGR		48	9	15
GVSYLLMGQVEENR		48	14	15
MTTDEGTGGR		37	10	15
PAPLVAPGNQVTLR		47	14	15
TEESPSAPDAPTCPK		61	15	15
TGGLDLPSPPTGASLK		46	16	15
TGTLQSNFCASSLVVTATVK		98	20	15
VFDLELHPACR		48	11	15
YDALEVFAGSGTSGQR		81	16	15
YDSVSVFNGAVSDDSR		101	16	15
DIPNLTALVR		55	10	16
DLAEVPASIPVNTR		69	14	16
HLEILQLSK		50	9	16
IEVGAFNGLPSLNTLELFDNR		29	21	16
LDLGELK		36	7	16

LDLIRPGSFQGLTSLR		36	16	16
LEELELSGNR		69	10	16
LEYISEAAFEGLVNLR		49	16	16
LTTVPTQAFEYLSK		50	14	16
NAFDDLK		29	7	16
NNPIESIPSYAFNR		57	14	16
PGSFQGLTSLR		48	11	16
RLDLGELK		32	8	16
RLEYISEAAFEGLVNLR		30	17	16
YLNLGMCNLK		48	10	16
YLNLQENGIQVIR		87	13	16
NSVNSHTIGR		45	10	1
LMVELHNLYR		63	10	2
WDEELAAFAK		53	10	2
LLDGSAEFDGK		60	11	2
TPSSDGTVEVR		84	11	2
AFVSLGAPWGGVAK		38	14	2
APNENGPYFLALR		42	13	2
AVVHGILMGVPVPFPIPEPDGCK		75	23	9
DCGSVDGVIK		53	10	9
DKTYSYLNK		39	9	9
EVNVSPCPTQPCQLSK		42	16	9
LVVEWQLQDDK		77	11	9
NQSLFCWEIPVQIVSHL		78	17	9
SEYPSIK		29	7	9
SGINCPIQK		45	9	9
TYSYLNKLPVK		62	11	9
AEELVNTAPLTGVPQHVPVR		35	20	12
ALIPLAK		30	7	12
GTAHPGEVTATCWAQSALPAPK		28	22	12
IELTDTTLEQVR		87	12	12
LTVWAPLLPLR		42	11	12
PADPPQYQEVPLDEAVTLR		38	19	12
PVTWQLEYPGQAPEAEK		64	17	12
SETFLLLQPWPR		52	12	12
SPLSDSILGEQALAVTDDK		53	19	12
VASLEGGR		28	8	12
VPGPAEGPAEPAAEASDEAER		115	21	12
VQPVMGISLTLSR	1Met(ox)	51	13	12
DVAIEEIDGLQLVK		39	14	6
GILLGVVGTDVPVK		33	14	6
INLFVGAEQLTNQDFLK		64	17	6
LWASAFGGEIK		56	11	6
SNVVTASTSIQLLDER		73	16	6
VLVMTNDYYYTDIK	1Met(ox)	52	14	6
AGQSAAGAAPGGGVDTR		57	17	7
ALTQTGGPHVK		33	11	7
VANPSGNLTETYVQDR		86	16	7
VKVEPSHDASK		40	11	7
VNVGAGSHPNK		49	11	7
VPVHDVTDASK		41	11	7
YTPVQQGPVGVNVTYGGDPIPK		58	22	7
AGTGLAAFSINQDTGMIQTLAPLDR		81	25	66
ALPFSIDPYLGIISTSK		49	17	66
ATDSGQPPLSASVR		86	14	66
AVAAQDPVIYSLVR		73	14	66

DGGTPSLQSEEEVLVTVR		78	18	66
DGVFSMNSYSGLISTQK		54	17	66
DPDQGANAQVVYSLPDSAEGHFSIDATTGVI		27	32	66
DSPVIQVLAYDADEGQNADVTYSVNPEDLVK		39	31	66
DVFNLVAK		29	8	66
DVIEINPVTGVVK		72	13	66
DVNDSPPR		41	8	66
DVSYQIVEDGSDVSK		87	15	66
EVSIFLQLR		47	9	66
FSEPLYTFSAPEDLPEGSEIGIVK		52	24	66
FSGQSYVR		55	8	66
FSLHPLTGELVVTGHLDR		29	18	66
FTQLHYEASVPDTIAPGTELLQVR		54	24	66
GKHELQVLAVDR		38	12	66
GNSEGFFNINALLGIITLAQK		82	21	66
GSVVENSEPGELVATLK		53	17	66
GVNAEVHYSLLK		58	12	66
IDPYLGDISLK		44	11	66
IHATDRDPQDTLTYSLAEEETLGR		32	24	66
IIAAQGLPR		63	9	66
IILTDENDNPPQFK		62	14	66
IISGDVANVFK		74	11	66
ILEPEALK		35	8	66
ISEGNGFGSLAALVVHVEPALR		35	22	66
ISLTQVLDK		36	9	66
KPFDYQALNK		33	10	66
LEYLILSGNQDR		80	12	66
LIYNIVEEEPLMLFTTDFK	1Met(ox)	114	19	66
LKVPEDLPPGTVLTFLDASDPDLGPAGEVR		33	30	66
LMDFHDR	1Met(ox)	31	7	66
LSILTPR		41	7	66
LSPVSPGPVYR		58	11	66
LVASDLDEGLNGR		81	13	66
PAQVIIHVR		50	9	66
QVTCYITEGDPLGQFGISQVGDEWR		97	25	66
SCQADITLHVEDVNDNAPR		45	19	66
SEYFVEIPESIPVGSPILLVSAM	1Met(ox)	27	33	66
SGSGPYFYSQIR		51	12	66
SLQFDQDVYWAAVK		77	14	66
SNEFSLVSVK		41	10	66
STFVGQISEAAPLYSMIMDK	1Met(ox)	63	20	66
TEEYVVGNFCFLR		53	13	66
TGNADEAVTIHPVTGSISVLNPAFLGLSR		43	29	66
TGVLTQFTK		60	9	66
TGVLTVTGPLDYESK		53	15	66
TLDADISEQNR		63	11	66
TLELEALTR		43	9	66
TPVAVVFAR		52	9	66
VDLMTGALILER		51	12	66
VIDLLAIDK		48	9	66
VIDLLAIDKDSGPYGTIDYTIINK		48	24	66
VLQLILSDPDSPENGPPYSFR		57	21	66
VPQDTVPGVELLR		44	13	66
VQAIDPDSR		37	9	66
VQAIDQDKGK		54	10	66
VQLSEFSPPGSR		52	12	66

VSIEDVNDNPPK		91	12	66
VTISEDMPLK		44	10	66
VTPDGWLVTAEGLSR		56	15	66
VVVGILDVNDNPPIFSHK		59	18	66
YFLSIECSR		51	9	66
YVLMDGAHGTFR	1Met(ox)	32	12	66
LNIPVSQVNPR		38	11	1
QLPSFQTFFAPALDVIR		29	17	2
SKPAVQYQWDR		62	11	2
DAEGILEDLQSYR		74	13	2
DQPPNSVEGLLNALR		58	15	2
AVISPGFDVFAK		55	12	23
DFHINLFR		35	8	23
DHENELLNK		47	9	23
DMTEVISSLENANYK		68	15	23
EILNINQK		31	8	23
ELNELGSK		33	8	23
ESASLMVDR		62	9	23
EVVTDQFLCSGTQEDESPCK		98	20	23
EVVTDQFLCSGTQEDESPCKGESGGAVFLE		54	31	23
FFQVGLVSWGLYNPCLGSADK		67	21	23
FGHGDKVR		45	8	23
GESGGAVFLER		71	11	23
HAFILQDTK		34	9	23
HAIILLTDGK		84	10	23
KNQGILEFYGDDIALLK		53	17	23
NDYLDIYAIGVGK		52	13	23
NQGILEFYGDDIALLK		90	16	23
PICLPCTMEANLALR		48	15	23
QHLGDVLNFLPL		56	12	23
RNDYLDIYAIGVGK		45	14	23
TAVDHIR		54	7	23
VLMSVLNDNSR		39	11	23
VNVGDPK		29	7	23
AEVSADQVATVMWDYFSQLSNNAK		82	24	18
GNTEGLQK		29	8	18
IDQNVEELK		59	9	18
IDQTVEELR		44	9	18
ISASAEELR		71	9	18
LAPLAEDVR		49	9	18
LEPYADQLR		32	9	18
LGPHAGDVEGHLSFLEK		92	17	18
LKEEIGK		50	7	18
LLPHANEVSQK		29	11	18
LTPYADEFK		38	9	18
SELTQQLNALFQDK		96	14	18
SLAELGGHLDQQVEEFR		92	17	18
SLAELGGHLDQQVEEFRR		40	18	18
SLAPYAQDTQEK		67	12	18
TLSLPELEQQQEQQQEQQQEQVQMLAPLES		45	30	18
TQVNTQAEQLR		76	11	18
VNSFFSTFK		54	9	18
GTTGLPQEVHVLNLR		67	15	5
NFLSLNYLAEYLQPK		39	15	5
NLILILK		27	7	5
SIRDDIPSTQGNLVK		79	15	5

SVNWVIK		44	7	5
CRPPVGCEELVR		46	12	3
GELDCHQLADSFRE		28	14	3
THEDLYIIPIPNCDR		36	15	3
AEDHFSVIDFNQNIR		44	15	30
AGELEVFNGYFVHFFAPDNLDPIPK		33	25	30
AHVSFKPTVAQQR		63	13	30
ALYAQAR		40	7	30
DKHADPDFTR		36	10	30
ETAVDGELVVLYDVK		64	15	30
FLHVPDTFEGHFDGVPVISK		34	20	30
FYNQVSTPLLR		46	11	30
HADPDFTR		40	8	30
HLEVDVWVIEPQGLR		35	15	30
IQPSGGTNINEALLR		97	15	30
IYGNQDTSSQLK		55	12	30
IYGNQDTSSQLKK		49	13	30
IYLQPGR		39	7	30
KFYNQVSTPLLR		39	12	30
KLGSYEHR		66	8	30
KLWAYLTINQLLAER		88	15	30
LDQIESVITATSANTQLVLETL/	1Met(ox)	33	34	30
LGSYEHR		46	7	30
LSNENHGIAQR		74	11	30
MATTMIQSK	1Met(ox)	58	9	30
MKQTVEAMK	1Met(ox)	39	9	30
MLADAPPQDPSCCSGALYYG	1Met(ox)	42	22	30
NDLISATK		45	8	30
QTVEAMK	1Met(ox)	32	7	30
SSALDMENFR	1Met(ox)	70	10	30
TEVNVLPGAK		52	10	30
TILDDLRAEDHFSVIDFNQNIR		42	22	30
VQFELHYQEVK		55	11	30
VVNNSPQPQNVVFDVQIPK		69	19	30
IYVVDVGSEPR		103	11	6
LNPNFLVDFGK		29	11	6
LYITTSLYSAWDK		67	13	6
NEGGTWSVEK		47	10	6
NTGTEAPDYLATVDVDPK		39	18	6
VAGGPQMIQLSLDGK	1Met(ox)	48	15	6
APLGSPSPRPR		39	11	1
CACTYGFTGPQCER		42	14	2
GYILQEDGR		34	9	2
AFLLSLAALR		51	10	3
IAVSKPSGPQPQADLQALLQSGAQVR		35	26	3
SQLEAIFLR		52	9	3
AKPLVLGPEFK		37	11	1
KPQDFLEELK		36	10	4
LNPGALLPVLTDPALNDLYVISTFK		44	25	4
QTEQTYWQATPFR		62	13	4
VHLVVFNNLQLADGR		32	15	4
ATLDVDEAGTEAAAATSFAIK		84	21	10
EIEEVLTPEMLMR		35	13	10
FSISGSYVLDQILPR		52	15	10
FYYLIASETPGK		57	12	10
GDATVFFILPNQGK		51	14	10

GFQHLLHTLNLPGHGLETR	44	19	10
IAPANADFAFR	70	11	10
LFHTNFYDTVGTIQLINDHVK	40	21	10
LGFTDLFSK	45	9	10
VGSALFLSHNLK	49	12	10
ALILGELEK	34	9	8
ASEQAELPR	61	9	8
FWLEQGVDSSVFEALPK	64	17	8
GQSQFQALCFVTQLQHNEIIPSEAMAK	28	27	8
LPDGQVTEESLQADSDADSISLELR	33	25	8
QLCLWDEDPYPG	33	12	8
SYSFDFYVPQR	38	11	8
VRLPDGQVTEESLQADSDADSISLELR	33	27	8
CQLSSLPGNIFR	40	12	3
LFLQNNLIR	58	9	3
SLEPDTFQGLER	33	12	3
AAFFLSYEELLQR	50	13	15
AQALAVSYR	57	9	15
DALFTILHDLRPQDR	34	15	15
DSGVTVNGELIGAPAPPNGHK	32	21	15
ELLSSWLQSDDEPEK	67	15	15
FSIIGFSNR	40	9	15
GHQVPVVWK	40	9	15
IAQNGILGDFIIR	66	13	15
IYNGEEQIDCWFAR	90	14	15
LSLENCGLTR	61	10	15
LWSYLTTK	28	8	15
SYLEITPSR	51	9	15
TITILINKPER	40	11	15
VHEEEDAGSQLIGFYDEIR	34	19	15
YAFTTVSCR	34	9	15
DFSSIIQTCSGNIQR	53	15	2
QLEADILDVNQIFK	81	14	2
ELVDEEADEAYELLSQAESWQR	34	22	3
SLLSDVEELVEK	83	12	3
TLFPVVLEQLDDYNAK	59	16	3
DTGTYGFLLPER	42	12	2
YPLYVLK	27	7	2
DLLDDLKSELTGK	30	13	2
GTVTDFPGFDER	38	12	2
AKPSAPVVSGPAAR	61	14	11
EITQDTNDITYADLNLPK	87	18	11
GTANLSETIR	65	10	11
KGSPDDVEFK	71	10	11
LTCQVEHDGQPAVSK	33	15	11
NGNELSDFQTNVDPVGESVSYSIHSTAK	63	28	11
SGAGTELSVR	68	10	11
SVLVAAGETATLR	101	13	11
TETASTVTENK	63	11	11
VPPTLEVTQQPVR	69	13	11
VTTVSDLTK	40	9	11
QDATASLILLWK	55	12	2
VTATGFQQCSLIDGR	93	15	2
AAPYWITAPQNLVLSPGEDGTLICR	68	25	42
AETYEGVYQCTAR	74	13	42
ASEPDKNPTAVEGLGSEPDNLVITWK	29	26	42

ASEPDKNPTAVEGLGSEPDNLVITWKPLNGF	30	41	42
DSTGTYTCVAR	40	11	42
EDGMLPK	31	7	42
ENIVIQCEAK	39	10	42
ERPPTFLTPEGNASNK	72	16	42
ERPPTFLTPEGNASNKEELR	31	20	42
FYFYAQTSAGSGSQITEEAVTTVDEAGILPPD	31	37	42
GAAVSNNIVVR	77	11	42
GAAVSNNIVVRPSR	43	14	42
GEGPASPDR	34	9	42
GKPPPSFSWTR	29	11	42
GNVLSLECIAEGLPTPIIYWAK	53	22	42
GSMVSFECK	50	9	42
IDGDTIIFSNVQER	82	14	42
ILTFQGSK	31	8	42
ILTPANTLYQVIANR	98	15	42
ISWLTNGVPIEIAPDDPSR	61	19	42
IVNPTLDSLTLEWDPPSHPNGILTEYTLK	32	29	42
KIDGDTIIFSNVQER	96	15	42
KILTFQGSK	49	9	42
LLEDLVQPPTITQQSPK	46	17	42
LSPYVNYSFR	58	10	42
NALGAIHHTISVR	70	13	42
NEVHLEIK	55	8	42
NEVHLEIKDPTWIVK	39	15	42
NPTAVEGLGSEPDNLVITWKPLNGFESNGPC	32	35	42
PALLDCAFFGSPLPTIEWFK	58	20	42
QPEYAVVQR	46	9	42
SLPSEASEQYLTK	84	13	42
SVQLSWTPGDDNNSPITK	87	18	42
THGMLPGLEPFSHYTLNVR	36	19	42
TLQIIHVSEADSGNYQCIAK	68	20	42
VFNTPEGVPSAPSSLK	68	16	42
VMAVNSIGK 1Met(ox)	56	9	42
VQALNDMGFAPEPAVVMGHSGEDLPMVAPC	62	33	42
VSQGLNGDLYFSNVLPEDTR	88	20	42
VSQGLNGDLYFSNVLPEDTREDYICYAR	32	28	42
VVNGKGEGPASPDR	59	14	42
YIVSGTPTFVPYLIK	77	15	42
DTMADGPWDSPALILELEDAV 1Met(ox)	58	22	8
ELDVLQGR	33	8	8
IDRLEQELPAR	39	11	8
IRADQDTIR	46	9	8
LVEAFGGATK	65	10	8
MDQLEGQLLAQVLALEK	99	17	8
QTALQQEAR	57	9	8
VALSHSSR	51	8	8
QEDPTNVGPEVK	60	12	1
GVEIYEPFFTQGETK	47	15	2
TGQFDSQEYTEYAVK	81	15	2
GEAGGQAEAEGDAPGPR	129	17	3
NTQNDFEVHIVQVENDEI	39	18	3
VLETQDLNGDGLMTPAELINF 1Met(ox)	40	27	3
ATITGYR	34	7	26
DDKESVPISDTIIPAVPPPTDLR	80	23	26
DLEVVAATPTSLLISWDAPAVTVR	111	24	26

EINLAPDSSSVVVSGLMVATK 1Met(ox)	107	21	26
EVTSDSGSIVVSGLTPGVEYVYTIQVLR	51	28	26
FLATTPNSLLVSWQPPR	84	17	26
FTNIGPDTMR	35	10	26
FTQVTPTSLSAQWTPPNVQLTGYR	85	24	26
GEWTCIAYSQLR	53	12	26
GEWTCKPIAEK	37	11	26
GLKPGVVYEGQLISIQQYGHQEVTR	49	25	26
GNLLQCICTGNGR	71	13	26
HTSVQTTSSGSGPFTDVR	93	18	26
HYQINQQWER	58	10	26
LGVRPSQGGEAPR	43	13	26
NEEDVAELSISPSDNAVVLTNLLPGTEYVVS\	53	45	26
NSITLTNLTPGTEYVVSIVALNGR	117	24	26
PAQGVVTTLENVSPPR	98	16	26
RPGGEPSPEGTTGQSYNQYSQR	50	22	26
TAGPDQTEMTIEGLQPTVEYV 1Met(ox)	45	48	26
TGLDSPTGIDFSDITANSFTVHWIAPR	40	27	26
TGQEALSQTTISWAPFQDTSEYIISCHPVGTI	38	39	26
TKTETITGFQVDAVPANGQTPIQR	100	24	26
TYHVGEQWQK	42	10	26
VDVIPVNLPGEHGQR	63	15	26
VPGTSTSATLTGLTR	102	15	26
AAAVSSGFDGAIQLVSLGGR	60	20	14
ALQSNHFELSLR	64	12	14
AYGTGFVGCLR	58	11	14
EGSLQVGNEAPVTGSSPLGATQLDTDGALW	42	45	14
GKDFLALALLDGR	53	13	14
SADGLTASCLCPATCR	52	16	14
SAGDVDTLAFDGR	90	13	14
SFLAFPTLR	49	9	14
SIESTLDDLFR	67	11	14
STVPVNTNR	31	9	14
TEATQGLVLWSGK	67	13	14
TFVEYLNAVTESEK	93	14	14
VCGSDGVTYGNECQLK	65	16	14
VVISGFGDPLICDNQVSTGDTR	91	22	14
TETLLLQAER	68	10	1
FTTLVQDLANAFQQEAQTSGK	74	21	2
LFQQSCPTGLVFSNSCK	75	17	2
AAMALEK	48	7	2
LGGPEAGLGEYLFER	43	15	2
DNLLDTYSADQGDSSEGGTLAR	111	22	5
EVILDLIPYESIVVTR	38	16	5
LDPIQPSDVLSLLDNR	45	16	5
LYTVDLESGLHYLLR	27	15	5
TPDQAILWLWK	55	11	5
ASGSPEPAISWFR	61	13	18
AVDGFTFTEGDKSPDGR	46	17	18
DKLVLPAK	28	8	18
EPSPPSIHGQPSSGK	49	15	18
EVASEIWK	35	8	18
EVVSPQEFK	28	9	18
FQEYILALADVPSSPYGVK	85	19	18
GQHGSSSLHIK	36	11	18
GQTQEATVVLEIYQK	76	15	18

GSNTELTVR		43	9	18
IEIFQTLPVR		56	10	18
IIELSQTTAK		62	10	18
MILEIAPTSDNDFGR	1Met(ox)	45	15	18
NIINSDGGPYVCR		74	13	18
QGEDAEVVCR		58	10	18
SKDKEDQWLEK		54	11	18
SMYLDIEYAPK		77	11	18
SNPPASIHWR		56	10	18
DMDVVSGR		42	8	3
SVPADIFQMQATTR		92	14	3
YPGAYYIFQIK		47	11	3
ITCQGDSLR		27	9	3
NNRPSGIPDR		29	10	3
SELTQDPAVSVALGQTVR		76	18	3
SYELTQPPSVSVSPGQTAR		54	19	1
GVVDSEDLPLNISR		64	14	1
EVQLLESGGGLVQPGGSLR		82	19	1
DTVYLQMDSLR		36	11	2
EVQLVETGGGLIQPGGSLR		91	19	2
QVELVESGGGAVEPGRSLR		36	19	3
QVELVESGGGAVQPGRSLR		27	19	3
QVQLVESGGGAVEPGRSLR		34	19	3
EVQLVESGGDLVQPGR		41	16	1
VTISVDTSR		49	9	1
FFESFGDLSTPDAVMGNPK		117	19	2
SAVTALWGK		47	9	2
AVSVEAAVTPAEPYAR		37	16	6
EPGVTSIEVR		41	10	6
GLHVTAAR		30	8	6
QVAGSVGGNTGVR		37	13	6
VPGPAEGPAEPAAEASDEAERR		41	22	6
VSVLELR		49	7	6
EEEEEMAVVPQGLFR		35	15	3
GEQEHSQQKEEEEEMAVVPQGLFR		89	24	3
SEALAVDGAGKPGAEEAQDPEGK		81	23	3
AEAEENEKETAVSTEDDSHHK		36	21	12
AQSIAYHLK		42	9	12
EKVHENENIGTTEPGEHQEAK		64	21	12
HIQETEWQSQEGK		96	13	12
IYLDEKR		44	7	12
KIYLDEK		40	7	12
KIYLDEKR		68	8	12
KLSENTDFLAPGVSSFTDSNQQESITK		76	27	12
LLAGDHPIDLLLR		54	13	12
SKEESHEQSAEQGK		60	14	12
TGLEAISNHK		54	10	12
TGLEAISNHKETEEK		70	15	12
FYADSVK		27	7	1
EIVMTQSPATLSVSPGER		51	18	1
EVQLVESGGGVVQPGGSLR		33	19	1
IYTYQWR		39	7	2
YALSNSIGPVR		70	11	2
KLVAASQAALGL		86	12	2
LVAASQAALGL		102	11	2
AYLTVLADQATPTNR		54	15	20

DHIVDPR	28	7	20
DLELTDLAER	69	10	20
DNILIECEAK	30	10	20
DQGSYTCVASTELDQDLAK	103	19	20
EFTTPEGVPSAPR	40	13	20
EVAGDTIIFR	40	10	20
FGTALSNR	41	8	20
GMDLLLECIASGVPTPDIAWYK	39	22	20
GPEPESVIGYSGEDYPR	97	17	20
ITNVSEEDSGEYFCLASNK	95	19	20
KEDQGIYTCVATNILGK	32	17	20
LDCPFFGSPIPTLR	60	14	20
LEVKDPTR	48	8	20
LSPYVNYQFR	57	10	20
LTVSWLKDDEPLYIGNR	50	17	20
NLILAPGEDGR	38	11	20
SGTLVIDFR	58	9	20
TSGAPPESNPGDVK	64	14	20
VQAENDFGKGPEPESVIGYSGEDYPR	37	26	20
LEPEDFAVYYCQQYGSSPR	69	19	2
LLMYGASSR	40	9	2
DIQLTQSPSSLSASVGDR	100	18	1
CSVSPTGDLTITNIQR	77	16	4
EGSQNLLFPNQPQQPNSR	30	18	4
GNPQPAVFWQK	31	11	4
QTSGLQATSSWQNLDAK	79	17	4
VTISLDTSK	38	9	1
NYVDWYQQLPGTAPK	54	15	1
AHELGGFTYETQASLSGPYLT 1Met(ox)	44	26	5
EDLYLHSLK	44	9	5
EFWQLLR	35	7	5
LSLEHLNPSIYVGLR	36	15	5
QSISVLAGSTVEDVLK	31	16	5
DIEMTQSPSSLSASVGDR	77	18	4
DIQMTQSPSSLSASVGDR	117	18	4
NIEMTQSPSSLSASVGDR	99	18	4
NIQMTQSPSSLSASVGDR 1Met(ox)	72	18	4
EIVLTESPGTLSLSPGER	59	18	4
EIVLTQSPGTLSLSPGQR	77	18	4
FSGSGSGADFTLTISR	58	16	4
QIVLTESPGTLSLSPGER	44	18	4
ASQSVSNSYLAWYQQKPGQAPR	28	22	2
EIVLTQSPGTLSLSPGER	124	18	2
DIVMTQSPDSLAVSLGER	116	18	3
ESGVPDR	31	7	3
LLIYWASTR	63	9	3
EDDSLTIFGVAER	72	13	5
GPEPESVIGYSGEDLPSAPR	74	20	5
VIAINEVGSSHPSLPSER	80	18	5
YPGSVNSAVLR	79	11	5
YVVGQTPVYVPYEIR	68	15	5
LAGLGLQQLDEGLFSR	93	16	5
LLLLDLSHNSLLALEPGILDTANVEALR	28	28	5
NLHDLDVSDNQLER	52	14	5
SLTLGIEPVSPTSLR	61	15	5
YLQGSSVQLR	34	10	5

ESLLLDTTSLQQR	73	13	3
IEYAPGAGSLALFPGIR	52	17	3
VNDVNEFAPVFVER	42	14	3
ALFQDIK	28	7	2
LGGPEAGLGEYLFER	63	15	2
EASPNPEDGIVR	35	12	10
ELNHLAVDEASGVVYLGAVNALYQLDAK	31	28	10
FGAQLQCVTGPQATR	95	15	10
GSSLHVGSDLLK	30	12	10
IQPETGPLGGGIR	28	13	10
LQLEQQVATGPALDNK	111	16	10
LQLEQQVATGPALDNKK	54	17	10
SFVASNDEGVATVGLVSSTGPGGDR	116	25	10
SPPNVQFTFQQPK	46	13	10
VLYAVFSR	39	8	10
CCVECPPCPAPPVAGPSVFLFPPK	33	24	7
CCVECPPCPAPPVAGPSVFLFPPKPK	36	26	7
FDPWGQGTLVTVSSASTK	67	18	7
GLPAPIEK	38	8	7
KCCVECPPCPAPPVAGPSVFLFPPKPK	29	27	7
STSESTAALGCLVK	75	14	7
VVSVLTVVHQDWLNGK	73	16	7
SASCDALTGACLNCQENSK	89	19	2
VGVIGSICDR	62	10	2
AAAGGPGGAALGEAPPGR	75	18	4
ASGSFVAPVR	66	10	4
EAATSSVLLPLDPGDR	48	16	4
RGNLLGGWK	32	9	4
ELSCESYPIELR	49	12	5
SGEAIIANANYHDTSPYR	92	18	5
VDGTGFVVYDGALFFNK	38	17	5
YLEVQYECVPYK	41	12	5
YSLDFGPLDSR	53	11	5
DQDLNTYSLLAVFAATDGGITR	28	22	6
IFGGVQQLR	58	9	6
LTNTNLLFVVAEKPLCSQCEAGR	29	23	6
QDPTLLWQVFGSATGVTR	51	18	6
RQDPTLLWQVFGSATGVTR	49	19	6
VAGDIESLLDR	72	11	6
FLASVSTVLTSK	84	12	10
KVADALTNAVAHVDDMPNALSALSDLHAHK	79	30	10
LRVDPVNFK	50	9	10
MFLSFPTTK	49	9	10
TYFPHFDLSHGSAQVK	50	16	10
VADALTNAVAHVDDMPNALSALSDLHAHK	30	29	10
VDPVNFK	30	7	10
VGAHAGEYGAEALER	84	15	10
VLSPADK	36	7	10
VLSPADKTNVK	52	11	10
KSDCGEWQWSVCVPTSGDCGLGTR	48	24	4
LTKPKPQAESK	37	11	4
TVTISKPCGK	42	10	4
YQFQAWGECDLNTALK	93	16	4
LVVLDEELEGISPDELK	74	17	3
NTEDLTEEWLR	65	11	3
SESLVVCDVAEDLVEK	69	16	3

DFFTVTDLR		50	9	2
FAFFAGPR		48	8	2
CTELISDIR		30	9	1
ASGVTVNDEVIK	1N-ac	35	12	2
QILVGDIGDTVEDPYTSFVK		78	20	2
IAELLSPGSVDPLTR		67	15	5
LVLVNAVYFR		53	10	5
SCDFLSSFR		47	9	5
SGGGGDIHQGFQSLLTEVNK		36	20	5
TYIGEIFTQILVLPYVGK		57	18	5
NLVVIPK		27	7	3
TTAQVLIR		48	8	3
VAIDVGYR		38	8	3
GVQVETISPGDGR		66	13	1
ATVHIQVNDVNEYAPVFK		66	18	8
EPFTISVWMR	1Met(ox)	40	10	8
FAESFEVTVTK		27	11	8
GVQIQAHPSQLVLTLEGEDLGELDK		61	25	8
ISIKPTCTPGWQGWNNR		48	17	8
LICSELNGR		37	9	8
LTVTAYDCGK		61	10	8
VEAVDADCSPQFSQICSYEIITPDVPFTVDK		85	31	8
CDPVDQCQDSETGTFYQIGDSWEK		61	24	26
DAPIVNK		35	7	26
DSMIWDCTCIGAGR	1Met(ox)	79	14	26
DTLTSRPAQGVVTTLENVSPPR		29	22	26
EESPLLIGQQSTVSDVPR		52	18	26
EYLGAICSCTCFGGQR		91	16	26
FGFCPMAAHEEICTTNEGVMYR		56	22	26
GFNCESKPEAEETCFDK		60	17	26
GFNCESKPEAEETCFDKYTGNTYR		37	24	26
LLCQCLGFGSGHFR		72	14	26
NLQPASEYTVSLVAIK		101	16	26
PGVTEATITGLEPGTEYTIYVIALK		37	25	26
PRPGVTEATITGLEPGTEYTIYVIALK		31	27	26
QAQQMVQPQSPVAVSQSK	1Met(ox)	34	18	26
QDGHLWCSTTSNYEQDQK		36	18	26
RPHETGGYMLECVCLGNGK	1Met(ox)	27	19	26
STATISGLKPGVDYTITVYAVTGR		55	24	26
TEIDKPSQMQVTDVQDNSISVK		65	22	26
TETITGFQVDAVPANGQTPIQR		50	22	26
TNTNVNCPIECFMPLDVQADR		53	21	26
VEYELSEEGDEPQYLDLPSTATSVNIPDLLPG		84	33	26
VREEVVTVGNSVNEGLNQPTDDSCFDPYTV		28	41	26
VTDATETTITISWR		60	14	26
VVTPLSPPTNLHLEANPDTGVLTVSWER		73	28	26
WCGTTQNYDADQK		53	13	26
YSFCTDHTVLVQTR		59	14	26
ELDLNSVLLK		34	10	5
HGTCAAQVDALNSQK		48	15	5
LGIKPSINYYQVADFK		49	16	5
PSINYYQVADFK		57	12	5
VCEDGPVFYPPPK		41	13	5
EVENPQNQLR		44	10	5
NMNVEEMLASEVLGDFLGAVI	2Met(ox)	88	21	5
TPYSDGVLYGSPTAENVGKPTIIEITAYNR		35	30	5

TQFYIDWCK	43	9	5
VPLPINDLK	33	9	5
AAANQMR	56	7	87
ADGSYAAWLSR	85	11	87
AEFQDALEK	60	9	87
AEMADQAAAWLTR	88	13	87
ALEILQEEDLIDEDDIPVR	102	19	87
ASAGLLGAHAAAITAYALTLTK	85	22	87
AVGSGATFSHYYYMILSR	89	18	87
CCQDGVTR	55	8	87
CSVFYGAPSK	50	10	87
DFALLSLQVPLK	77	12	87
DFALLSLQVPLKDAK	60	15	87
DHAVDLIQK	66	9	87
DKGQAGLQR	41	9	87
DVKAAANQMR	28	10	87
EAPKVVEEQESR	62	12	87
ECVGFEAVQEVPVGLVQPASATLYDYYNPEI	83	31	87
ECVGFEAVQEVPVGLVQPASATLYDYYNPEI	50	32	87
EELVYELNPLDHR	74	13	87
EFHLHLR	33	7	87
EGAIHREELVYELNPLDHR	38	19	87
EMSGSPASGIPVK	51	13	87
EPFLSCCQFAESLR	73	14	87
EPFLSCCQFAESLRK	37	15	87
EVYMPSSIFQDDFVIPDISEPGTWK	89	25	87
FACYYPR	52	7	87
FEQLELR	47	7	87
FGLLDEDGK	53	9	87
FGLLDEDGKK	54	10	87
GCGEQTMIYLAPTLAASR	43	18	87
GHLFLQTDQPIYNPGQR	97	17	87
GLCVATPVQLR	52	11	87
GLEEELQFSLGSK	84	13	87
GLQDEDGYR	53	9	87
GPEVQLVAHSPWLK	70	14	87
GQIVFMNR	63	8	87
GSFEFPVGDAVSK	65	13	87
GSSTWLTAFVLK	75	12	87
HLVPGAPFLLQALVR	43	15	87
ITPGKPYILTVPGHLDEMQLDI 1Met(ox)	29	25	87
ITQVLHFTK	50	9	87
KADGSYAAWLSR	97	12	87
KYVLPNFEVK	55	10	87
LELSVDGAK	67	9	87
LGQYASPTAK	69	10	87
LHLETDSLALVALGALDTALYAAGSK	130	26	87
LLLFSPSVVHLGVPLSVGVQLQDVPR	59	26	87
LNMGITDLQGLR	89	12	87
LQETSNWLLSQQQADGSFQDLSPVIHR	104	27	87
LTVAAPPSGGPGFLSIER	64	18	87
LTVAAPPSGGPGFLSIERPDSRPPR	28	25	87
LVNGQSHISLSK	75	12	87
MRPSTDTITVMVENSHGLR	84	19	87
PVAFSVVPTAATAVSLK	67	17	87
PVQGVAYVR	68	9	87

PYILTVPGHLDEMQLDIQAR	73	20	87
QGSFQGGFR	75	9	87
QRVEASISK	29	9	87
RFEQLELR	36	8	87
RGHLFLQTDQPIYNPGQR	54	18	87
SCGLHQLLR	60	9	87
SFFPENWLWR	44	10	87
SHALQLNNR	51	9	87
SHKPLNMGK	38	9	87
SMQGGLVGNDETVALTAFVTI 1Met(ox)	32	39	87
STQDTVIALDALSAYWIASHTTEER	97	25	87
TEQWSTLPPETK	68	12	87
TLEIPGNSDPNMIPDGDFNSY' 1Met(ox)	56	23	87
TTNIQGINLLFSSR	97	14	87
TYNVLDMK	50	8	87
VDFTLSSER	79	9	87
VDVQAGACEGK	67	11	87
VDVQAGACEGKLELSVDGAK	35	20	87
VEASISK	46	7	87
VEYGFQVK	55	8	87
VFALDQK	43	7	87
VGDTLNLNLR	101	10	87
VGLSGMAIADVTLLSGFHALR 1Met(ox)	82	21	87
VHYTVCIWR	42	9	87
VLSLAQEQVGGSPEK	129	15	87
VQQPDCR	36	7	87
VQQPDCREPFLSCCQFAESLR	27	21	87
VTASDPLDTLGSEGALSPGGVASLLR	120	26	87
VVEEQESR	53	8	87
YIYGKPVQGVAYVR	58	14	87
YLDKTEQWSTLPPETK	65	16	87
YVLPNFEVK	52	9	87
YVSHFETEGPHVLLYFDSVPTSR	41	23	87
AEVNGLAAQGK	61	11	15
ALQASALNAWR	85	11	15
DNAGAATEEFIK	73	12	15
DNAGAATEEFIKR	43	13	15
GILAADESVGSMAK 1Met(ox)	84	14	15
GVVPLAGTDGETTTQGLDGLSER	106	23	15
PWALTFSYGR	45	10	15
QVLFSADDR	70	9	15
RAEVNGLAAQGK	63	12	15
TPSALAILENANVLAR	100	16	15
TVPPAVPGVTFLSGGQSEEEASFNLNAINR	27	30	15
VDKGVVPLAGTDGETTTQGLDGLSER	28	26	15
VLAAVYK	51	7	15
YASICQQNGIVPIVEPEILPDGDHDLK	30	27	15
YTPEEIAMATVTALR	93	15	15
FEFGQETSQTLK	67	12	6
GILYVTDTK	45	9	6
LENALYFDR	44	9	6
TYFNLAVDEK	57	10	6
VTETFGTWIR	75	10	6
VTFAFDLLGGK	60	11	6
ASGVPDR	32	7	3
FSGSGSGTDFTLK	94	13	3

VYACEVTHQGLSSPVTK		73	17	3
FSGSIDSSSNSASLTISGLK		77	20	2
SSGSIASNYVQWYQQR		81	16	2
ASQSVSSYLAWYQQKPGQAPR		41	21	2
LLIYDASNR		65	9	2
EGMNIVEAMER	1Met(ox)	28	11	8
FEDENFILK		62	9	8
IIPGFMCQGGDFTR	1Met(ox)	34	14	8
ITIADCGQLE		41	10	8
KITIADCGQLE		82	11	8
SIYGEKFEDENFILK		40	15	8
VNPTVFFDIAVDGEPLGR		57	18	8
VSFELFADK		50	9	8
LDTNYDLLLDQSELR		81	15	2
LEYQACVLGK		40	10	2
DFSLTSSSQTPGATK		68	15	2
GPQLLALVEEVLPR		76	14	2
AIHLDLEEYR		48	10	9
EEILMHLWR		47	9	9
FIIEGMEEAGSVALEELVEKEK		35	22	9
GDGWLTDPYVLTEVDGK		94	17	9
GNSYFMVEVK		53	10	9
GPVLAWINAVSAFR		106	14	9
KPAITYGTR		34	9	9
MFQEIVHK		32	8	9
TVFGTEPDMIR	1Met(ox)	54	11	9
DHFLMDGQVR		56	10	3
ETQQWYTVTHPVPTPR		41	16	3
MLLLQPQAR		43	9	3
VSEPVSAGR		66	9	1
DASGATFTWTPSSGK		65	15	1
ATGIPDR		35	7	11
DSTYSLSSTLTLSK		82	14	11
FSGSGSGTDFTLTISR		129	16	11
HKVYACEVTHQGLSSPVTK		66	19	11
RTVAAPSVFIFPPSDEQLK		63	19	11
SGTASVVCLLNNFYPR		77	16	11
TVAAPSVFIFPPSDEQLK		73	18	11
VDNALQSGNSQESVTEQDSK		123	20	11
VDNALQSGNSQESVTEQDSKDSTYSLSSTL		88	34	11
YLAWYQQK		34	8	11
YLAWYQQKPGQAPR		67	14	11
GLVWVSR		32	7	2
SVTCHVK		27	7	2
DIAPTLTLYVGK		81	12	7
ILGGHLDAK		55	9	7
SCAVAEYGVYVK		77	12	7
SPVGVQPILNEHTFCAGMSK		97	20	7
VMPICLPSK		48	9	7
YQEDTCYGDAGSAFAVHDLEEDTWYATGILS		65	34	7
YVMLPVADQDQCIR	1Met(ox)	77	14	7
ALNPSQTSMSGTLELPNIGAR		66	21	3
DGEIPYYAEVVATNNPDR		56	18	3
LSCSLPGSCEAGPPLTFSWTGNALSPLDPET		30	33	3
IGQESLEFILVQADTPSSPSIDQVEPYSSTAQ		52	48	3
ISVVWNDDSSSTLTIYNANIDDAGIYK		31	27	3

SLDWNAEYEVYVVAENQQGK	111	20	3
FSGSGSGTDFTLTITR	85	16	2
LLIYGASSR	59	9	2
ANDGEWYHVDIQR	48	13	14
AYGLLVATTSR	92	11	14
DGFQGCLASVDLNGR	87	15	14
GNSDRPLNDNQWHNVVITR	29	19	14
GSEYLCYDLSQNPIQSSSDEITLSFK	82	26	14
GYIHYVFDLGNGPNVIK	28	17	14
IYGEVVFK	31	8	14
LELDGGR	34	7	14
LMVNLDCIR	45	9	14
LPDLINDALHR	52	11	14
MGSISFDFR	46	9	14
QLTIFNTQAQIAIGGK	108	16	14
TPFTASGESEILDLEGDMYLGGLPENR	30	27	14
TTSPDGFILFNSGDGNDFIAVELVK	66	25	14
AMDNVTVR	43	8	8
AVGFVSEDEYLEIQGITR	89	18	8
GTGVPVGQK	27	9	8
GTLQCEASAVPSAEFQWYK	68	19	8
QGESATLR	28	8	8
VENRPFLSK	31	9	8
VHLIVQVSPK	51	10	8
VTVNYPPYISEAK	35	13	8
AGLVFPTEVWTALLNYGYVGCIR	42	23	16
CENVATLDPITFETPESFISLPK	79	23	16
DLFIDGQSK	46	9	16
DMTVFSGLFVGGLPPELR 1Met(ox)	77	18	16
FNVGTDDIAIEESNAIINDGK	107	21	16
GKEEYIATFK	51	10	16
IHGVVAFK	45	8	16
ITTQITAGAR	61	10	16
LEFHNIETGIITER	48	14	16
LELDAGR	43	7	16
LQLSFSIFCAEPATLLADTPVNDGAWHSVR	29	30	16
LTVDDQQAMTGQMAGDHTR	41	19	16
NGLMLHTGK	55	9	16
SADYVNLALK	55	10	16
SDLYIGGVAK	63	10	16
VLNMAAENDANIAIVGNVR 1Met(ox)	83	19	16
TPTQEQLLAAAMAAAR	57	16	2
VTAHAEGYTPSAK	53	13	2
DEASSVEVTWPDGK	66	14	13
DGKVDIVYGNWNGPHR	44	16	13
EHGDPLIEELNPGDALEPEGR	31	21	13
FRDIASPK	42	8	13
GDGTFVDAAASAGVDDPHQHGR	51	22	13
GNQGFNNNWLR	45	11	13
GVALADFNR	51	9	13
GVASLFAGR	62	9	13
GVSVGPILSSSASDIFCDNENGPNFLFHNR	30	30	13
LVNIAVDER	43	9	13
NNRWEDILSDEVNVAR	57	16	13
SSPYYALR	44	8	13
WEDILSDEVNVAR	79	13	13

YADLTEDQLPSCESLK		50	16	2
YADLTEDQLPSCESLKDTIAR		29	21	2
VLVTVNYPPTITDVTSAR		40	18	1
AVLGSPR		35	7	16
EACYGDMDGFPGVR		67	14	16
EAEVLVAR		35	8	16
ELEAPSEDNSGR		29	12	16
EYQWIGLNDR		39	10	16
GVVFLYR		35	7	16
IGAHIATPEQLYAAYLGGYEQCDAGWLSDQT		57	33	16
LTLEEAR		29	7	16
NYGVVDPDDLYDVYCYAEDLNGELFLGDPPE		31	32	16
RAVLGSPR		56	8	16
VKGVVFLYR		30	9	16
VKVNEAYR		49	8	16
VKWTFLSR		27	8	16
YAFSFSGAQEACAR		92	14	16
YEVDTVLR		32	8	16
YPIVTPSQR		41	9	16
CTTDFVSLTSHLNSAVDGFDSEFCK		49	25	6
GNLVYHSAVLGISDLMSQR		41	19	6
VEGAWPLIDNNYLSVQVTNVPVVPGSSATAT		44	33	6
VYQAVTDDLPAAFVDGTTSGGDSDAK		118	26	6
YIGTTVFVR		41	9	6
YLTLAIR		42	7	6
DCGATWVVLGHSER		46	14	15
ELASQPDVDGFLVGGASLKPEFVDIINAK		74	29	15
FFVGGNWK		47	8	15
HVFGESDELIGQK		68	13	15
IIYGGSVTGATCK		64	13	15
KFFVGGNWK		33	9	15
KQSLGELIGTLNAAK		30	15	15
SNVSDAVAQSTR		72	12	15
TATPQQAQEVHEK		74	13	15
VAHALAEGLGVIACIGEK		91	18	15
VIADNVKDWSK		50	11	15
VPADTEVVCAPPTAYIDFAR		75	20	15
VTNGAFTGEISPGMIK		75	16	15
VVFEQTK		29	7	15
VVLAYEPVWAIGTGK		76	15	15
ASDSPIDLFYGDFFGDISEAVIQK		31	24	6
DLHIQSHISENRDEVEAVK		31	19	6
ETTEESIKETER		76	12	6
FQNIDFAEEVYTR		75	13	6
IVFLEEASQQEK		72	12	6
LATLGGSQALGLDGEIGNFEVGK		99	23	6
AAVPSGASTGIYEALELR		77	18	13
AVEHINK		32	7	13
DATNVGDEGGFAPNILENK		83	19	13
DYPVVSIEDPFDQDDWGAWQK		75	21	13
FTASAGIQVVGDDLTVTNPK		56	20	13
GNPTVEVDLFTSK		52	13	13
IGAEVYHNLK		62	10	13
KLNVTEQEK		49	9	13
LAMQEFMILPVGAANFR	1Met(ox)	38	17	13
LMIEMDGTENK		27	11	13

SCNCLLLK		28	8	13
VNQIGSVTESLQACK		73	15	13
YISPDQLADLYK		44	12	13
ADLIAYLK		50	8	6
GIIWGEDTLMEYLENPK	1Met(ox)	80	17	6
GIIWGEDTLMEYLENPKK		89	18	6
MIFVGIK		31	7	6
TGPNLHGLFGR		50	11	6
TGQAPGYSYTAANK		66	14	6
ADGDPPPAILWLSPR		47	15	5
AFSGLNSLEQLTLEK		82	15	5
DFPDVLLPNYFTCR		38	14	5
LQEIQLVGGQLAVVEPYAFR		99	20	5
LTVFPDGTLEVR		34	12	5
ADDGRPFPQVIK		28	12	13
ALANSLACQGK		57	11	13
ALQASALK		43	8	13
ELSDIAHR		55	8	13
GILAADESTGSIAK		81	14	13
GVVPLAGTNGETTTQGLDGLSER		94	23	13
IGEHTPSALAIMENANVLAR		42	20	13
LQSIGTENTEENR		72	13	13
PYQYPALTPEQK		60	12	13
QLLLTADDR		50	9	13
SKGGVVGIK		30	9	13
TVPPAVTGITFLSGGQSEEEASINLNAINK		31	30	13
YTPSGQAGAAASESLFVSNHAY		82	22	13
AKLEETITQAR		62	11	32
AQEIFNK		33	7	32
AVEIYIQGK		67	9	32
DSETLKPDNFEESGYTFIAPR		43	21	32
EAGENWQENPETYEDSFYK		81	19	32
EAGENWQENPETYEDSFYKR		45	20	32
EDFASNEVVYYNAK		79	14	32
EEDPSLLWQVFGSATGLAR		113	19	32
FFGEIDPSLMR		42	11	32
FVVTDGGITR		62	10	32
GYYYEIPSIGAIR		54	13	32
IDLYDVR		53	7	32
IIMLFTDGGEER	1Met(ox)	60	12	32
IKPVFIEDANFGR		43	13	32
INTQEYLDVLGR		83	12	32
KTPNNPSCNADLINR		72	15	32
LALEAEK		43	7	32
LEETITQAR		67	9	32
MQEDLVTLAK		55	10	32
NREEDPSLLWQVFGSATGLAR		56	21	32
QLVEIAAR		33	8	32
QSCITEQTQYFFDNDSK		73	17	32
SGPGAYESGIMVSK		88	14	32
SQEPVTLDFLDAELENDIK		105	19	32
SYDYQSVCEPGAAPK		52	15	32
TASGVNQLVDIYEK		91	14	32
TPNNPSCNADLINR		97	14	32
VFTFSVGQHNYDR		64	13	32
VLLDAGFTNELVQNYWSK		107	18	32

VQAAHQWR	29	8	32
YQDLYTVEPNNAR	53	13	32
YYPASPWVDNSR	47	12	32
GFSDCLLK	35	8	2
LGDSMANYPQGLDDK	64	15	2
QVQLVQSGAEVK	66	12	1
AEDTAVYYCVR	38	11	2
WLAWYQQKPGK	50	11	2
KSDVTETLVSGTQLSQLIEGLDR	54	23	7
LTHQIQELTLDTPYYFK	28	17	7
SDEGFYQCIAENDVGNAQAGAQLIILEHAPAT	63	41	7
VETQPEVQLPGPAPNLR	49	17	7
VIGQDVVLPCVASGLPTPTIK	38	21	7
VLPDPEVISDLVFLK	48	15	7
YYTIENLDPSSHYVITLK	51	18	7
DGEDQTQDTELVETRPAGDGTFQK	39	24	2
FIAVGYVDDTQFVR	55	14	2
DTLMISR	45	7	14
EPQVYTLPPSR	77	11	14
EPQVYTLPPSREEMTK	64	16	14
FNWYVDGVEVHNAK	89	14	14
GFYPSDIAVEWESNGQPENNYK	92	22	14
GLEWVANIK	70	9	14
GTTVTVSSASTK	79	12	14
NQVSLTCLVK	60	10	14
NSLYLQMNSLR	71	11	14
STSGGTAALGCLVK	112	14	14
TPEVTCVVVDVSHEDPEVK	104	19	14
TTPPVLDSDGSFFLYSK	82	17	14
VVSVLTVLHQDWLNGKEYK	48	19	14
WQQGNVFSCSVMHEALHNHYTQK	41	23	14
ALPAPIEK	38	8	3
NTLYLQMNSLR	78	11	3
THTCPPCPAPELLGGPSVFLFPPKPK	36	26	3
APWIEQEGPEYWDR	52	14	2
WVAVVVPSGQEQR	43	13	2
FFESFGDLSSPDAVMGNPK	87	19	8
GTFSQLSELHCDK	48	13	8
LLGNVLVCVLAR	52	12	8
TAVNALWGK	66	9	8
VHLTPEEK	44	8	8
VNVDAVGGEALGR	83	13	8
VVAGVANALAHK	71	12	8
VVAGVANALAHKYH	82	14	8
ASGVQVADEVCR 1N-ac	47	12	2
YALYDASFETK	43	11	2
NTLYLQMNSLK	60	11	1
ESATITCLVTGFSPADVFVQWMQR	27	24	2
FTCTVTHTDLPSPLK	40	15	2
VGYVSGWGQSDNFK	49	14	2
VVLHPNYHQVDIGLIK	55	16	2
AGVETTKPSK	64	10	2
ANPTVTLFPPSSEELQANK	63	19	2
DSGLFGQYLLTPAR	52	14	12
ESLFLINGR	55	9	12
FIVSAAADSPWLHVQEITVR	28	20	12

GEIQTLYDLQINSGISDLAFQR		27	22	12
LLVESLFR		58	8	12
NEVGVDEDISSLFIEDSAR		91	19	12
QDLDEDLLGCSPGDLLR		75	17	12
QLLVDSVTDSVLGPNGDVTGTPHTSPDGR		54	29	12
SRPSLQVITEASTGQSQHLIR		27	21	12
TLANILWR		45	8	12
TPFAGVDDFFIPPTNLIINHIR		41	22	12
YIYVAQPALSR		46	11	12
FDHVITNMNNNYEPR		75	15	4
NSLLGMEGANSIFSGFLLFPDI	1Met(ox)	61	24	4
VPGLYYFTYHASSR		63	14	4
VVTFCDYAYNTFQVTTGGMVI	1Met(ox)	52	22	4
AAQVTIQSSGTFSSK		107	15	75
AFQPFFVELTMPYSVIR	1Met(ox)	79	17	75
AGAFCLSEDAGLGISSTASLR		111	21	75
AIGYLNTGYQR		54	11	75
ALLAYAFALAGNQDK		98	15	75
ALLAYAFALAGNQDKR		82	16	75
APVGHFYEPQAPSAEVEMTSYVLLAYLTAQF		37	46	75
ATVLNYLPK		53	9	75
AVDQSVLLMKPDAELSASSV\	1Met(ox)	28	27	75
DLKPAIVK		33	8	75
DLTGFPGPLNDQDDEDCINR		71	20	75
DMYSFLEDMGLK	1Met(ox)	80	12	75
DTVIKPLLVEPEGLEK		76	16	75
EEFPFALGVQTLPQTCDEPK		77	20	75
EQAPHCICANGR		51	12	75
ETTFNSLLCPSGGEVSEELSLK		93	22	75
FEVQVTVPK		70	9	75
FQVDNNNR		48	8	75
FSGQLNSHGCFYQQVK		52	16	75
GGVEDEVTLSAYITIALLEIPLTVTHPVVR		40	30	75
GHFSISIPVK		35	10	75
GHFSISIPVKSDIAPVAR		29	18	75
GPTQEFK		53	7	75
GPTQEFKK		53	8	75
GVPIPNK		27	7	75
HNVYINGITYTPVSSTNEK		104	19	75
HNVYINGITYTPVSSTNEKDM\	1Met(ox)	27	31	75
HYDGSYSTFGER		51	12	75
IAQWQSFQLEGGLK		100	14	75
KDTVIKPLLVEPEGLEK		34	17	75
KLSFYYLIMAK	1Met(ox)	40	11	75
KYSDASDCHGEDSQAFCEK		102	19	75
LHTEAQIQEEGTVVELTGR		143	19	75
LLIYAVLPTGDVIGDSAK		59	18	75
LLLQQVSLPELPGEYSMK		75	18	75
LPPNVVEESAR		71	11	75
LSFYYLIMAK		74	10	75
LVHVEEPHTETVR		71	13	75
LVHVEEPHTETVRK		36	14	75
MCPQLQQYEMHGPEGLR		30	17	75
MVSGFIPLKPTVK	1Met(ox)	44	13	75
NALFCLESAWK		56	11	75
NEDSLVFVQTDK		103	12	75

NQGNTWLTAFVLK		85	13	75
PVPGHVTVSICR		49	12	75
QFSFPLSSEPFQGSYK		78	16	75
QGIPFFGQVR		58	10	75
QQNAQGGFSSTQDTVVALHALSK		66	23	75
QSSEITR		32	7	75
QTVSWAVTPK		63	10	75
RTTVMVK		27	7	75
SASNMAIVDVK		81	11	75
SDIAPVAR		37	8	75
SIYKPGQTVK		31	10	75
SLFTDLEAENDVLHCVAFAVPK		50	22	75
SLNEEAVKK		36	9	75
SSGSLLNNAIK		55	11	75
SSSNEEVMFLTVQVK	1Met(ox)	58	15	75
TAQEGDHGSHVYTK		61	14	75
TEHPFTVEEFVLPK		66	14	75
TEVSSNHVLIYLDK		82	14	75
TGTHGLLVK		34	9	75
VDLSFSPSQSLPASHAHLR		79	19	75
VGFYESDVMGR		96	11	75
VSNQTLSLFFTVLQDVPVR		71	19	75
VSVQLEASPAFLAVPVEK		76	18	75
VTAAPQSVCALR		58	12	75
VTGEGCVYLQTSLK		114	14	75
VVSMDENFHPLNELIPLVYIQDPK		32	24	75
VYDYYETDEFAIAEYNAPCSK		104	21	75
YDVENCLANK		59	10	75
YGAATFTR		43	8	75
YNILPEK		29	7	75
YNILPEKEEFPFALGVQTLPQTCDEPK		33	27	75
YSDASDCHGEDSQAFCEK		84	18	75
GDGELSWEHSDGDIFR		42	16	3
LTCAFPVSVPDSCCR		35	15	3
QPNQCTQCSCSEGNVYCGLK		81	20	3
LLIYDASR		32	8	2
LLIYDASSLESGVPSR		58	16	2
AVQPGETYTYK		48	11	8
DGTDYIEIIPK		70	11	8
DSNMPVDMR	2Met(ox)	33	9	8
HSLVLHK		30	7	8
MDDAVAPGR		48	9	8
SGPESPGSACR		28	11	8
SQHLDNFSNQIGK		63	13	8
VMYTQYEDESFTK		74	13	8
FVSQETGNLYIAK		62	13	4
IEVQFPETVPTAK		44	13	4
ITISEDGNLR		59	10	4
VGGDSAGDLMIR		75	12	4
ANVAVVSGAPLQGLVAR		64	17	3
FGDQVLQINGENCAGWSSDK		88	20	3
SIDNGIFVQLVQANSPASLVGLR		31	23	3
GVALHRPDVYLLPPAR		32	16	4
QIQVSWLR		35	8	4
QVGSGVTTDQVQAEAK		91	16	4
YVTSAPMPEPQAPGR		49	15	4

AEDTAVYYCAK		41	11	5
DGFFGNPR		36	8	5
VFAIPPSFASIFLTK		45	15	5
VSVFVPPR		39	8	5
YAATSQVLLPSK		42	12	5
AKIDQNVEELK		31	11	19
ALVQQMEQLR		81	10	19
DKVNSFFSTFK		58	11	19
EAVEHLQK		46	8	19
ENADSLQASLRPHADELK		44	18	19
IDQNVEELKGR		72	11	19
KLVPFATELHER		84	12	19
LGEVNTYAGDLQK		86	13	19
LKEEIGKELEELR		76	13	19
LNHQLEGLTFQMK		47	13	19
LVPFATELHER		61	11	19
QLTPYAQR		40	8	19
RQLTPYAQR		47	9	19
RVEPYGENFNK		58	11	19
TQVSTQAEQLR		59	11	19
TQVSTQAEQLRR		27	12	19
VEPYGENFNK		29	10	19
VKIDQTVEELR		30	11	19
VKIDQTVEELRR		42	12	19
LLIYAASSLQSGVPSR		43	16	2
NDLGWYQQKPGK		45	12	2
IVTTTGAVFAK		39	11	1
AASLDGFYNSR		71	11	4
EGPWSPESESPMLR	1Met(ox)	30	14	4
SPTHQAALR		38	9	4
VLGQYSGPR		64	9	4
ASQGISSYLAWYQQKPGK		47	18	2
LLIYAASTLQSGVPSR		32	16	2
LLIYGVSSR		34	9	1
DNERPSGIPER		31	11	3
ITCSGDALPK		51	10	3
PGQAPVLVIYK		68	11	3
TLMNLGGLAVAR		46	12	2
YGINTTDIFQTVDLWEGK		40	18	2
LLGELLQDNAK		57	11	3
LVPVLSAK		41	8	3
SYEPLEDPGVK		43	11	3
DTWAFVGR		30	8	2
VEGYGSVCSCKDPTPIEFSPDPLPDNK		27	27	2
DYFPEPVTVSWNSGALTSGVHTFPAVLQSS(		29	49	4
TTPPVLDSDGSFFLYSR		73	17	4
YGPPCPSCPAPEFLGGPSVFLFPPK		27	25	4
YGPPCPSCPAPEFLGGPSVFLFPPKPK		34	27	4
ADLSGITGAR		91	10	15
AKWEMPFDPQDTHQSR		34	16	15
AVLDVFEEGTEASAATAVK		126	19	15
DEELSCTVVELK		73	12	15
DLDSQTMMVLVNYIFFK	1Met(ox)	73	17	15
DYNLNDILLQLGIEEAFTSK		154	20	15
EIGELYLPK		28	9	15
EQLSLLDR		49	8	15

GTHVDLGLASANVDFAFSLYK	56	21	15
ITLLSALVETR	96	11	15
KLINDYVK	54	8	15
LYGSEAFATDFQDSAAAK	111	18	15
NLAVSQVVHK	73	10	15
RLYGSEAFATDFQDSAAAK	62	19	15
WEMPFDPQDTHQSR	46	14	15
DTSISTAYMELSR	61	13	2
SDDTAVYYCAR	62	11	2
AVLTIDEK	52	8	35
DTEEEDFHVDQVTTVK	111	16	35
DTVFALVNYIFFK	72	13	35
ELDRDTVFALVNYIFFK	73	17	35
FLEDVKK	48	7	35
FLENEDR	39	7	35
FNKPFVFLMIEQNTK	69	15	35
GKWERPFEVK	30	10	35
GTEAAGAMFLEAIPMSIPPEVH 2Met(ox)	68	22	35
ITPNLAEFAFSLYR	97	14	35
KLSSWVLLMK	60	10	35
KLYHSEAFTVNFGDTEEAKK	65	20	35
KQINDYVEK	63	9	35
LGMFNIQHCK	59	10	35
LQHLENELTHDIITK	95	15	35
LSITGTYDLK	60	10	35
LSITGTYDLKSVLGQLGITK	30	20	35
LSSWVLLMK 1Met(ox)	65	9	35
LVDKFLEDVK	48	10	35
LVDKFLEDVKK	58	11	35
LYHSEAFTVNFGDTEEAK	93	18	35
LYHSEAFTVNFGDTEEAKK	85	19	35
QINDYVEK	40	8	35
RLGMFNIQHCK	31	11	35
RSASLHLPK	27	9	35
SASLHLPK	60	8	35
SPLFMGK	54	7	35
SVLGQLGITK	64	10	35
TDTSHHDQDHPTFNK	51	15	35
TLNQPDSQLQLTTGNGLFLSEGLK	121	24	35
VFSNGADLSGVTEEAPLK	133	18	35
VFSNGADLSGVTEEAPLKLSK	48	21	35
VVNPTQK	36	7	35
WERPFEVK	43	8	35
WERPFEVKDTEEEDFHVDQVTTVK	27	24	35
AENFFILR	34	8	2
ELLLQPVTISR	43	11	2
DVCDPGNTK	28	9	30
EDFTSLSLVLYSR	47	13	30
ELPEHTVK	27	8	30
ELSSFIDK	37	8	30
ELSSFIDKGQELCADYSENTFTEYK	69	25	30
EVVSLTEACCAEGADPDCYDTR	123	22	30
EYANQFMWEYSTNYGQAPLSLLVSYTK	32	27	30
FPSGTFEQVSQLVK	70	14	30
GQELCADYSENTFTEYK	112	17	30
GQELCADYSENTFTEYKK	105	18	30

HLSLLTTLSNR	76	11	30
HQPQEFPTYVEPTNDEICEAFR	53	22	30
KELSSFIDK	51	9	30
KFPSGTFEQVSQLVK	94	15	30
KLCMAALK	69	8	30
LAQKVPTADLEDVLPLAEDITNILSK	33	26	30
LCDNLSTK	50	8	30
RSDFASNCCSINSPPLYCDSEIDAELK	54	27	30
RTHLPEVFLSK	47	11	30
SCESNSPFPVHPGTAECCTK	77	20	30
SDFASNCCSINSPPLYCDSEIDAELK	92	26	30
SLGECCDVEDSTTCFNAK	122	18	30
SYLSMVGSCCTSASPTVCFLK	78	21	30
TAMDVFVCTYFMPAAQLPELF 1Met(ox)	58	29	30
THLPEVFLSK	44	10	30
VCSQYAAYGEK	47	11	30
VLEPTLK	33	7	30
VMDKYTFELSR	74	11	30
VPTADLEDVLPLAEDITNILSK	95	22	30
YTFELSR	35	7	30
LLIYSNNQR	44	9	4
LLIYSNNQRPSGVPDR	43	16	4
LTVLSQPK	49	8	4
PSGVPDR	33	7	4
AYGILMATTSR	57	11	13
EGFQGCLASVDLNGR	82	15	13
EQGQPFQGQLSGLYYNGLK	84	19	13
GPETLFAGYNLNDNEWHTVR	35	20	13
LAIGFSTVQK	46	10	13
LPDLISDALFCNGQIER	66	17	13
LTVNLDCIR	49	9	13
NGAVSLVINLGSGAFEALVEPVNGK	45	25	13
QLTIFNSQATIIIGGK	39	16	13
SGTISVNTLR	34	10	13
TGSISFDFR	62	9	13
TPYTAPGESEILDLDDELYLGGLPENK	89	27	13
VDSSSGLGDYLELHIHQGK	32	19	13
AVGDKLPECEADDGCPKPPEIAHGYVEHSVI	39	31	22
AVGDKLPECEAVCGK	35	15	22
AVGDKLPECEAVCGKPK	44	17	22
DIAPTLTLYVGKK	62	13	22
DYAEVGR	36	7	22
GSFPWQAK	50	8	22
HYEGSTVPEK	43	10	22
HYEGSTVPEKK	55	11	22
KQLVEIEK	72	8	22
LPECEADDGCPKPPEIAHGYVEHSVR	62	26	22
LPECEAVCGKPK	65	12	22
LRTEGDGVYTLNDK	83	14	22
LRTEGDGVYTLNNEK	96	15	22
NPANPVQR	43	8	22
QKVSVNER	31	8	22
QLVEIEK	45	7	22
TEGDGVYTLNDK	92	12	22
TEGDGVYTLNDKK	62	13	22
VGYVSGWGR	57	9	22

VMPICLPSKDYAEVGR	1Met(ox)	40	16	22
VTSIQDWVQK		72	10	22
VVLHPNYSQVDIGLIK		53	16	22
ITYQPSTGEGNEQTTTIGGR		78	20	4
NLQPDTSYTVTVVPVYTEGDGGR		49	23	4
NVQVYNPTPNSLDVR		62	15	4
VEDIIEAINTFPYR		28	14	4
SWTAADTAAQITQR		99	14	2
WAAVVVPSGEEQR		43	13	2
ATLVCLVSDFYPGAVTVAWK		42	20	2
VGVETTKPSK		47	10	2
ATYFGSIVLLSPAVIDSPLK		73	20	5
FLEQELETITIPDLR		59	15	5
GAFFPLTER		36	9	5
SSVDELVGIDYSLMK	1Met(ox)	56	15	5
VPHDLDMLLR		29	10	5
ASSFLGEK		40	8	3
DSSTWLTAFVLK		60	12	3
LQETSNWLLSQQQADGSFQDPCPVLDR		64	27	3
DAQGQPQAVPVSGDLR		63	16	14
EEPPRPEFLEQPLLGELTVTGVTPDSLR		27	28	14
FLLFGIQDGK		43	10	14
FLLYGLLGK		46	9	14
ILISGLEPSTPYR		31	13	14
LGELWVTDPTPDSLR		46	15	14
LGPLSAEGTTGLAPAGQTSEESRPR		40	25	14
LSQLSVTDVTTSSLR		97	15	14
SGTLYSLTLYGLR		51	13	14
TLSPVLESPR		41	10	14
VSYQLADGGEPQSVQVDGQAR		105	21	14
VVWTAQPDTFAYFQLR		69	16	14
WTVPEGEFDSFVIQYK		69	16	14
YLVTLYGFSDGK		33	12	14
DDHGLFGLDVK		27	11	6
LEENYDNFYTVVTDRPLDR		63	19	6
LTYEIVDGNDDHLFEIDPSSGEIR		42	24	6
NLNYSVPEEQGAGTVIGNIGR		58	21	6
VLENSAPHLLDVDADSGLLYTK		32	22	6
VLGGGGTGGGGGLGGPGGSVPFK		82	23	6
DFMIQGGDFTR		62	11	9
DTNGSQFFITTVK		67	13	9
DVIIADCGK		66	9	9
HYGPGWVSMANAGK		29	14	9
IGDEDVGR		68	8	9
TAWLDGK		28	7	9
TVDNFVALATGEK		69	13	9
VIFGLFGK		37	8	9
VLEGMEVVR		56	9	9
EFTPPVQAAYQK		27	12	7
GTFATLSELHCDK		86	13	7
KVLGAFSDGLAHLDNLK		68	17	7
LLGNVLVCVLAHHFGK		34	16	7
LLVVYPWTQR		67	10	7
VLGAFSDGLAHLDNLK		82	16	7
VNVDEVGGEALGR		87	13	8
ANSFLGEK		53	8	3

LLATLCSAEVCQCAEGK	102	17	3
PVAFSVVPTAAAAVSLK	73	17	3
CLLFSFLPASSINDMEK	36	17	6
EKGEIQNILQK	28	11	6
IYSGILNLSDITK	79	13	6
TSESGTPSSSTPQENTISGYSLLTCK	66	26	6
VLTPDAFVCR	30	10	6
VNIPLVTNEECQK	66	13	6
AVAFQDCPVDLFFVLDTSESVALR	81	24	17
CGPIDLLFVLDSSESIGLQNFEIAK	105	25	17
CPDYTCPITFSSPADITILLDGSASVGSHNFD	55	35	17
DAEEAISQTIDTIVDMIK 1Met(ox)	112	18	17
DTTPLNVLCSPGIQVVSVGIK	80	21	17
DVFDFIPGSDQLNVISCQGLAPSQGR	78	26	17
DVFDFIPGSDQLNVISCQGLAPSQGRPGLSL	34	33	17
ENYAELLEDAFLK	99	13	17
GDEGEAGDPGDDNNDIAPR	42	19	17
LSIIATDHTYR	51	11	17
NLVWNAGALHYSDEVEIIQGLTR	32	23	17
PGLSLVK	44	7	17
SLQWMAGGTFTGEALQYTR 1Met(ox)	79	19	17
TAEYDVAYGESHLFR	77	15	17
VFSVAITPDHLEPR	33	14	17
YLIVVTDGHPLEGYK	37	15	17
YLIVVTDGHPLEGYKEPCGGLEDAVNEAK	33	29	17
SAVQGPPDR	38	9	8
EKYLTWASR	37	9	8
KGDTFSCMVGHEALPLAFTQK	36	21	8
QEPSQGTTTFAVTSILR	90	17	8
SAVQGPPER	52	9	8
TFTCTAAYPESK	49	12	8
VAAEDWK	33	7	8
WLQGSQELPR	76	10	8
KPGESLK	32	7	3
SISTAYLQWSSLK	50	13	3
YSPSFQGQVTISADK	65	15	3
AGGSWDLAVQER	31	12	3
LLNIQTYCAGPAYLK	40	15	3
WLNVQLSPR	65	9	3
ASQGISNSLAWYQQKPGK	58	18	2
LLLYAASR	48	8	2
ITTVAHTEVGPGPESSPVVVR	32	21	2
SGALQIESSEETDQGKYECVATNSAGVR	37	28	2
AEDTAVYYCAR	64	11	4
TPEVTCVVVDVSHEDPEVQFK	78	21	4
TPLGDTTHTCPR	58	12	4
WQQGNIFSCSVMHEALHNR	77	19	4
CPAPELLGGPSVFLFPPK	39	18	3
CPAPELLGGPSVFLFPPKPK	50	20	3
SCDTPPPCPR	53	10	3
AVDGFTFTEGDK	50	12	4
FFTCTAIGEPESIDWYNPQGEK	75	22	4
LTIYNANIEDAGIYR	110	15	4
YDCEAASR	46	8	4
LMIYEVSK	31	8	2
RPSGVPDR	28	8	2

ALDVGSGSGILTACFAR		64	17	3
ELVDDSINNVR		70	11	3
LILPVGPAGGNQMLEQYDK		50	19	3
FYFENLWSR		51	9	2
VTEQLIEAISNGDFESYTK		81	19	2
KGEELEEEWTPTEK		32	14	2
YLSPDATVSTEVR		72	13	2
GLEWVGFIR		58	9	2
SIAYLQMNSLK		34	11	2
AACLLPK		56	7	35
AAFTECCQAADK		61	12	35
ADDKETCFAEEGK		41	13	35
AEFAEVSK		50	8	35
ALVLIAFAQYLQQCPFEDHVK		93	21	35
ATKEQLK		35	7	35
CCAAADPHECYAK		66	13	35
DDNPNLPR		58	8	35
DVFLGMFLYEYAR	1Met(ox)	78	13	35
ECCEKPLLEK		48	10	35
EFNAETFTFHADICTLSEK		117	19	35
FKDLGEENFK		68	10	35
FQNALLVR		73	8	35
HPDYSVVLLLR		63	11	35
HPYFYAPELLFFAK		67	14	35
KQTALVELVK		72	10	35
LCTVATLR		58	8	35
LDELRDEGK		55	9	35
LKCASLQK		40	8	35
LKECCEKPLLEK		57	12	35
LVNEVTEFAK		70	10	35
MPCAEDYLSVVLNQLCVLHEK	1Met(ox)	80	21	35
NECFLQHK		35	8	35
NLGKVGSK		31	8	35
PCFSALEVDETYVPK		81	15	35
PLVEEPQNLIK		44	11	35
QEPERNECFLQHK		33	13	35
QNCELFEQLGEYK		92	13	35
QNCELFEQLGEYKFQNALLVR		46	21	35
RMPCAEDYLSVVLNQLCVLHEK		59	22	35
SLHTLFGDK		57	9	35
SLHTLFGDKLCTVATLR		74	17	35
TCVADESAENCDK		99	13	35
VHTECCHGDLLECADDR		105	17	35
VPQVSTPTLVEVSR		92	14	35
AVWSPEPCTTCLCSDGR		33	17	6
CPQTVIPEGECCPVCSATEQR		65	21	6
FESFSSFPGVESSYNVLPGK		38	20	6
IAPLAWINQENLESIDLSYNK		112	21	6
LPSGCSLSYR		55	10	6
VNENNLQAIDEESLSDLNQLVTLELEGNNLSE		33	44	6
AEGNNQAPGEEEEEEEEATNTHPPASLPSQ		28	31	4
EDSLEAGLPLQVR		67	13	4
RPEDQELESLSAIEAELEK		94	19	4
VAHQLQALR		36	9	4
NGDNVEAPPVYDSYEVEYLPIEGLLSSGK		42	29	3
IGPGDVLTFYDGDDLTAR		48	18	3

EPVCIAACGGVIR	47	13	3
GPEDLDPGAEGAGAQVELLPDR	71	22	2
GPEDLDPGAEGAGAQVELLPDRDPDSDGTK	38	30	2
EPQVYTLPPSRDELTK	38	16	3
GPSVFPLAPSSK	64	12	3
VVSVLTVLHQDWLNGK	89	16	3