

# PeptideRanker

Your query (ID 171084754113809) has been processed (19.03.2024).

Response:

0.964729	GPPGPPGPPGPPGPPGGGFD
0.930727	GPPGPPGSSSGPQGF
0.909966	GPAGPNGPAGGRGGLGPA
0.899642	GPAGPHGPPGKDGRAGGHG
0.889048	GPRGPPGPPGSSSGPQGF
0.873021	GPSGERGPPGPMGPPGLA
0.846923	GPPGPPGSSSGPQG
0.819015	GPVGPAGKSGDRGESGPAGPA
0.809429	GPVGPAGKSGDRGESGPAGPAG
0.805951	PGPGGPQGEFGAVGPAG
0.791772	MGPRGPPGPPGSSSGPQGF
0.782283	LGPTGPIGTGPMGPR
0.782179	GLPGGPGAVGEPGRLGPA
0.781425	GPAGPRGPSGPA
0.78115	KAADTFNFKTF
0.778331	GPSGPAGARGDKGEAGEAGER
0.776344	GPPGPPGSSSGPQ
0.771791	GPRGPPGPPGSSSGPQG
0.759302	SSYGGANRGMGGGFGAGGGGGSY
0.750221	GASGPAGPRGPSGSAGAA
0.746106	PGPGGPQGEHGQIGPAG
0.740213	GPMGPRGPPGPPGSSSGPQG
0.729904	GEPGKQGPGGPSGERGPPGPM
0.729846	GPSGPAGPSGESGPA
0.727939	SSGGQAQQGPMGPMGPR
0.722902	TGPPGEPGEAGSSSGPMGPR
0.720204	SGESGPAGASGPAGPR
0.698244	SAGGGAGGAGAGAGGGSVARGA
0.693342	GPTGPPGAPGFPGGAGAK
0.691649	GPRGPPGPPGSSSGPQ
0.688174	GDRGDQGVKGADGAGGKDGIRG
0.683052	SGGGQAQMGPMGPMGPRG
0.680767	FTGPPGEPGEAGSSSGPMGPR
0.666412	GPMGPRGPPGPPGSSSGPQ
0.661017	PGPGGPQGEHGQIGPA
0.659388	GESGPAGPAGPNGPAGGRGGLGPA
0.65547	GPAGGQGEKGEGGPAGPN
0.645763	GPMGPMGPRGPPGPPGSSSGPQ
0.644583	GPPGPPGPPGPPGPPGGGFDM
0.639159	MGPRGPPGPPGSSSGPQG
0.639116	PGPGGPQGEHGQIGP
0.638675	GEPGKQGPGGPSGERGPPGPMGPPG
0.630867	SGPAGPAGPNGPAGGRGGLGPA
0.618146	GPAGARGADGSTGPAGPAGPL
0.609093	RGPPGPPGSSSGPQ
0.608775	GPAGAQQEKGEGGPAGPN
0.604673	GFTGPPGEPGEAGSSSGPM
0.595796	GPAGARGADGSTGPAGPAGPLG
0.591814	GFTGPPGEPGEAGSSSGPMGPR
0.581004	GEKGQPGGDGPNGPSGTPGPQ
0.577045	FTGPPGEPGEAGSSSGPM
0.557906	GLTGPAQPQGARGFQGETGIQ
0.557057	MGPRGPPGPPGSSSGPQ

0.547469 FSGLDGA KGETGPAGPKGEAG  
0.535769 GMTGSPGSPGPDGKVGPA  
0.5234 LRGDPGPGGPQGEHQIGPA  
0.521516 WAAFPD VAGNV DYKQ  
0.521043 GPAGSPGDKGETGAPGAVGPS  
0.515051 GHGGLQGMPPGPNGPSGETGSAG  
0.505036 SGESGPAGASGPAGPRGPSGSAG  
0.497159 GPAGARGADGSTGPAGPAGP  
0.483323 NWDDMEKIWHH  
0.482102 GESGPAGASGPAGPRGPSGSAGAA  
0.480255 GSRGEPGNPGPAGAAGPS  
0.475757 FSGLDGA KGESGPAGPKGEAGAS  
0.47384 GTEASGTGNMKFM  
0.473292 SGLDGA KGETGPAGPKGEAGAS  
0.46315 GYTGATGATGLTGPAGPQGARGF  
0.452053 FSDEEEFPDLSLH  
0.451784 EALGDKAKFAGK  
0.435222 GERGEQGAGGPPGFQ  
0.419701 SGESGPAGASGPAGPRGPSGSAGAA  
0.410298 STGEAGATGPAGLRG  
0.409454 FSGLDGA KGETGPAGPKGEAGAS  
0.40943 GARGADGSTGPAGPAGP  
0.404232 AGPAGPAGARGADGSTGPAGPAGPL  
0.403028 GLDASSGELPVVGIR  
0.397305 KEYEPENGKVF  
0.395104 LPYDTPIPGYRNN  
0.394118 ALPYDTPIPGYRNN  
0.379334 GVVGNA GEKGEQGPSGAPGFQ  
0.378089 LEQQVDDLEGSLEQEKKL RMD  
0.377822 ATGATGLTGPAGPQGARGF  
0.368547 LDGA KGESGPAGPKGEAGAS  
0.367079 SMPGPMGPMGPRGPPGPPGSSGPQG  
0.360745 LQMPGPNGPSGETGSA  
0.357719 PANIKWGDAGATY  
0.35649 AGARGADGSTGPAGPAGP  
0.346779 GEEGKRGSTGEVGATGPAGLR  
0.345106 DAGKTFPQHVKSR  
0.33901 GGRSFDNVDTPASAL  
0.334044 HEDYKFGHTK  
0.330889 GSDSLDSAKKEIG  
0.328393 AGGGYDQSGGYDEYR  
0.327242 GQKDSYVGDEAQSKRGIL  
0.32248 DGA KGETGPAGPKGEAGAS  
0.320068 DAGKTFPEHV KSR  
0.31553 YGKDATNVGDEGGFAPNIL  
0.306525 ITGDQLGDLYKSF  
0.302948 GEQIDNLQRVK  
0.301484 IITNWDDMEKIWHH  
0.30112 FDMFDTDGGGDISTK  
0.300186 EMNPPKYDKIEDM  
0.298442 TGPAGPQGARGFQGETGIQ  
0.295713 GFTNDEKIGIY  
0.29555 DEYNKGFLDQL  
0.292514 LDGA KGETGPAGPKGEAGAS  
0.2883 AYE PVWAIGTGKTASPA  
0.288296 AGPAGGGYDQSGGYDEY  
0.283778 AADESTGSVAKRF  
0.278917 GQKDSYVGDEAQSKRGILT  
0.278757 LRGHGGLQGMPPGPNGPSGETGSAG  
0.269936 LDGTENKSKFGAN  
0.269248 AEELKKEQDTSSHLERMKNLE

0.265862 GFTNEEKLGIYK  
0.26388 DFNGDTHSSIFDAGAGIA  
0.263431 IDSELPVDEVFGH  
0.262138 AADGPMKGILGYTEQ  
0.254358 EYDEAGPSIVHRK  
0.253828 SSSSLEKSYELPDGQ  
0.253661 SAETEKELANMK  
0.25276 I IHGSDSLDSANKEIAL  
0.249465 EISDLNRMIQ  
0.249457 ATESFGPGTIEYRP  
0.247899 AEELKKEQDTSSHLERM  
0.247209 GQKDSYVGDEAQSKRGILT  
0.239792 ELDGTENKSKFGAN  
0.233679 GFTNEEKLGIY  
0.233347 LGSIDVNHEDYKFGHTK  
0.231868 EYDESGPSIVHRK  
0.229719 SYEEALDHLETL  
0.225556 GEQIDNLQR  
0.225257 SEATDELAKAM  
0.221956 SIMDLENDKQQADEKIKKK  
0.215944 GEVDRKLAEKDEEMEIQIKRN  
0.215371 AETLWGVQKELTL  
0.212756 VTWYDNEFGYSNR  
0.210672 MLELDGTENKSKFGAN  
0.209331 SGDLEELKNVTNN  
0.208392 KNSYEEALDHL  
0.208336 GEQIDNLQRVKQK  
0.206164 LQSAETEKELANMK  
0.206153 LNDQHLGKTK  
0.202868 GTGEFSEPSVEIEPVEKL  
0.201883 TADQIEDFKEAF  
0.201552 NVLSGGTTMYPGIADRM  
0.199498 VLGNSGLTDVIHM  
0.199461 GVKGERGNSGPAGSAGSQ  
0.196848 NSYEEALDHLETL  
0.194672 ENDKQQADEKIKKKEFETT  
0.191105 GTEFMQDPDVEHL  
0.189142 LGQNPQNKEVAAI  
0.188573 GARQQEGESRLN  
0.188189 EAEASLEHEESKIL  
0.18772 EYDEAGPSIVHR  
0.186978 GQKDSYVGDEAQSKRG  
0.183688 VEELKLFLQ  
0.181834 ELLEGKVLPGVDAL  
0.177117 ELKKEQDTSSHLERM  
0.176656 LLQSAETEKELANMK  
0.176592 GVNHEKYDNSLK  
0.175952 NSYEEALDHL  
0.175926 LSKVEDEQSLGAQL  
0.174546 KDSYVGDEAQSKRG  
0.174138 GADTSADLEETGRVL  
0.173645 KNSYEEALDHL  
0.173594 VGSQVSEANLR  
0.171621 LVTWYDNEFGYSNR  
0.170832 DTSADLEETGRVL  
0.167041 TENGEFGRQL  
0.166978 AEELKKEQDTSSHLERMKK  
0.165729 SYVGDEAQSKRG  
0.165635 IIDQNRDGIIS  
0.162679 SGDLEELKNVTN  
0.158324 DTMDRIIQTM

0.155714	LAEKDEEMEQIKRN
0.155624	LRIDQSILTGE
0.155039	SESLKDAQEKL
0.154054	IAMDGTEGLVRGQ
0.148766	LQSAETEKELANM
0.146696	SKIEDEQSLGAQ
0.146162	VGMGQKDSYVGDEAQSKRG
0.145971	SKVEDEQSLGAQLQ
0.142977	WKQKYEEGQAELEGA
0.142884	ETLWGVQKELTL
0.1421	AKVIHDNYH
0.139815	EAAEGTLEHEEAKIL
0.136843	GVVGNAGEKGEQGPS
0.136801	GQKDSYVGDEAQSKR
0.135484	KVIHDNYH
0.132405	AEELKKEQDTSSHLERMKKN
0.131981	EVKKEVKEEEEVGDWRK
0.130324	GYTGATGATGLTGPAQP
0.130159	GVNHEKYDNSLKVVS
0.129206	LENDKQQADEKIKKK
0.128406	NFSHGTHEYHAE
0.127608	LLEKEAADLITEK
0.127089	DLENDKQQADEKIKKK
0.126999	VDHVNKDIAAK
0.125857	VDRKLAEKDEEMEQIKRN
0.12519	LLSKVEDEQSLGAQ
0.123158	SKVEDEQSLGAQ
0.122966	AEWKQKYEEGQAELE
0.121666	VTGGKSNDDVSGPANQL
0.119758	PTETQSDARSF
0.117163	GVNHEKYDNSLKVVSNA
0.116583	TNEKLQQF
0.113098	TDAEQEQLIADHF
0.112937	LSKVEDEQSLGAQLQ
0.112822	EKDEEMEQIKRN
0.111447	KVEDEQSLGAQLQ
0.107434	EELEEAKKKLAQR
0.107353	EEMEQIKRN
0.104621	VVSSDFNGDTHSS
0.101896	GRTEHTPEGVKWVDTQ
0.101605	EEISERLEEAGGAT
0.101449	LIEDTQDWQPRTGTTQ
0.0990451	LSKVEDEQSLGAQ
0.0964986	GQKDSYVGDEAQSK
0.0945146	DISNADRLGSSEVDQVQ
0.0942321	IAEEADRKYEE
0.0937397	ITGESGAGKTVNTK
0.0932384	TLDDLQSEEDKVNTL
0.0923273	TLDDLQAEEDKVNTL
0.0860746	VGMGQKDSYVGDEAQSK
0.0858503	SDLERTEERAELSEGK
0.0847766	ELEEEELKNVTNN
0.0839209	ISDLTEQIGETGKS
0.0833431	KLKGTEDELDKY
0.0823965	LQNEIENVKGQ
0.0793251	NDENVRQVNDISGQ
0.0744901	LEEEELKNVTNN
0.0718552	EELEEAKKKLAQ
0.0715598	NADRLGSSEVDQVQ
0.0710519	ELKKEQDTSSHL
0.0704079	KRAIEEEVKAK

0.0702853	METGLTPDQVKKN
0.0688986	SELEEEELKTVTN
0.0652111	DENVRQVNDISGQ
0.0651457	KTIDDLEDELYAQ
0.0646284	KTKNDENVRQVNDISGQ
0.0637462	LAKVEKEKHATEN
0.0625328	LIEEEKKEMKQE
0.0623686	ELEEEELKTVTNN
0.0617897	ELEEEELKNVTN
0.060841	EIQLSHSNRQAEEAQ
0.0600667	ENVRQVNDISGQ
0.0559723	TLENGNLVQKQ
0.0553383	KRQLEEEVKAK
0.0538266	TALEEAEGTLEHE
0.0528499	ELEEEELKTVTN
0.0526873	TEQIGETGKSIHE
0.0524878	LEEELKTVTNN
0.0487858	LVEEEELDRAQ
0.0467464	AYKRHSEEAEEAANQH
0.0393565	TKEKKALQEAHQQ
0.0344744	EKAKKTVETEKSEIQ
0.0310724	VEMEKKLEKGEAIDGMIPAQK
0.0287943	LKEVKKEVKEEE