

Supporting Information Table 1. Experimentally Measured and Predicted NRTs of All 834 peptides

Peptide seq	NRT		Peptide seq	NRT	
	meas	pred		meas	pred
ASAQLETIK	0.431	0.437	FGEIEEVELGRIQK	0.566	0.601
YFPDATILALTTNEK	0.671	0.666	GDPVYLK	0.448	0.454
MVEVNACLK	0.504	0.496	IVVFNNSVLGFVAMEMK	0.774	0.774
QHEFSHATGELTALLSAIK	0.674	0.623	NGVEERK	0.127	0.090
LDEAGVVRMIGPNCPGVITPGECK	0.586	0.611	TCPK	0.121	0.133
VNEDLGLLSEEK	0.547	0.561	FFPAEANGGVK	0.497	0.509
NHLNMHFVSNVDGTHIAEV K	0.587	0.621	GFSGEDATPALEGADVVVLISAGVARK	0.679	0.683
GFTSEITVTSGNGK	0.500	0.480	ERVTEAK	0.410	0.391
DAASFAPLNPAHLIGIEEALK	0.651	0.661	IIQIDINPASIGAHSK	0.596	0.584
LGELLEALK	0.608	0.597	NYDPRATVMRETCHEVLK	0.527	0.534
RAMDVYCHRLAK	0.440	0.443	IYAYLSRGLCGR	0.545	0.541
ITDAYAENPQIANLLAPYF K	0.745	0.754	LLPWIDGLLDAGEK	0.753	0.737
QEAAAPAAAPAPAAGVK	0.438	0.460	LLAWLLETLK	0.679	0.682
QVIDASHAEKG	0.371	0.372	IIASVAEK	0.420	0.455
PLSPETWQHKLK	0.529	0.530	HLINK	0.171	0.194
SIREAGVQEADFLANVDK	0.581	0.596	VLGVK	0.386	0.400
VGIVSRSGTILTYEAVK	0.526	0.562	DSVSYGVVK	0.452	0.436
IAAGDTNSLGDSTLADPGVVEK	0.550	0.590	NGLACITPISALNQPGK	0.622	0.600
AMQEVLQQFAHVVK	0.599	0.575	ELGTK	0.141	0.140
LASAK	0.123	0.172	ARHLVDLYQQQQVKE	0.477	0.503
GVNLPGVSIALPALAEK	0.698	0.656	FLCIAK	0.510	0.493
VTGQALTVNFK	0.431	0.449	AGQTFTFTTDK	0.501	0.508
VIVVTDGGERILGLGDQGIGGMGIPIGK	0.700	0.682	LSPTLAMYRAK	0.505	0.523
RALIVTDRFLFNNGYADQITSVLK	0.687	0.755	LPYITPEGSEEHTYLHAQRQK	0.572	0.581
APVIVQFSNGGASFIAGK	0.635	0.648	LIQLMNETVDGDYQTFK	0.630	0.667
TDITELEAFRK	0.538	0.525	NINPTQTAAWQALQK	0.575	0.561
TRNAIIFSPHPRAK	0.436	0.465	VMWGSRWDELLRK	0.608	0.626
DWGYQLAREEFGGELIDGGPWLK	0.729	0.742	AVLNRGVSVVVLPGDVALK	0.629	0.672
LMPEFWQFPTVSMGLGPIGAIYQAK	0.812	0.827	EYLPASYHEGSK	0.443	0.454
ESAPAAAAPAAQPALAARSEK	0.468	0.468	GVHEGHVAAEVIAGK	0.444	0.427
ELLEDPTTRLDDVGLCGR	0.733	0.731	RGLCGR	0.242	0.209
SVLCIGGSWLVPADEALEAGDYDRITK	0.723	0.736	MRTLGTAAACPYYHIAFVIGG TSAETNLK	0.642	0.687
LVSWYDNETGYSNK	0.540	0.556	MVGGVTPGK	0.410	0.426
VHPNDDVNK	0.264	0.294	NLDDK	0.136	0.135
RPFK	0.144	0.158	EYAPAEDEPGVVSVEIYQYYK	0.664	0.638
MVPCDFIAPAIATHNPLSDHH QK	0.629	0.622	HQFAQSLNYEIAK	0.518	0.511
APNPSVAGRATVIFPDNLNTGNTTYK	0.660	0.653	HVTWRGEAIPGSLFDALYFFHNYQALLAK	0.880	0.815
RMQAAGAQAYLVNTGWNGTAK	0.555	0.564	EDGIYVTMEGK	0.513	0.509
EGTRPAVVIPTNEELVIAQDASRLTAGLCGR	0.720	0.700	IINELEGIFEGAGWNVIK	0.756	0.782
IAFGCDHVGFILK	0.632	0.651	LRIMPMIIISVEEVRALRK	0.698	0.728
ESGLLGLTEVTSDCRYVEDNYATK	0.648	0.623	ANSIGHGAPGMADMIAEVWLWDFLK	0.826	0.740
QLIPQLK	0.539	0.502	TAGVIGTK	0.374	0.387
GVVPQLVK	0.524	0.475	YIPEELRK	0.440	0.458
HFSTTPAEK	0.355	0.357	LMAELEGLCGR	0.583	0.609
LNERHYGALQGLNK	0.454	0.453	ALLSMAIRAAK	0.549	0.503
IAEAAVVGIPHNK	0.537	0.532	FCQEEDK	0.344	0.358
YYDELPTEGNEHGQAFRDVE LEK	0.552	0.572	IAELAGFSVPENTK	0.581	0.583
TTLSTDTPK	0.371	0.361	LEPLSLQDFGALLEEQSK	0.762	0.729
MLNK	0.163	0.204	ERAADVIRDIGK	0.382	0.352
AIEGTDRSSLRILGSVGEPINPEAWEYW WK	0.727	0.758	DWRGRRGASQNIIIPSSSTGAAK	0.490	0.468
QTYCGPIGAEMYHITSTEK	0.581	0.558	GFDSPREFYVGRLTIEGIAL GAAFYPK	0.747	0.708
PARVVMEK	0.369	0.412	AVAAVNGPIAQALIGK	0.622	0.593
DSILEAIDAGIK	0.662	0.588	LAVNFGAEILK	0.634	0.606
FSATFDDQMVLVDYSK	0.625	0.651	LNAK	0.124	0.149
VIPSIAYTEPEVAWVGLTEK	0.708	0.723	AEAPAAAAPAAK	0.365	0.367
TRTQQIEELQK	0.422	0.456	FNLMLETK	0.577	0.574
IFMGLATIFLERDLALIEINPLVITK	0.897	0.868	EAAPAAAAPAAAAAK	0.404	0.432
VYYDLLEQRKK	0.497	0.494	DFEDAVEK	0.451	0.443
QRSLYIPYAGPVILLEPFLNK	0.767	0.742	AGVYDK	0.199	0.203
HFDEMKG	0.330	0.319	DVTIADLFAK	0.658	0.590
AGRGLCGR	0.310	0.320	HYFDPK	0.387	0.355
YVALQFLRNDSIAAK	0.606	0.619	PGVITGDDVQK	0.438	0.474
YIALRCAGFNNVNDLDAAK	0.595	0.628	FPLPVVEIPMARSavarqlvk	0.703	0.693
RYYLGNADEIAAK	0.498	0.492	LAPSLTLGCGSWGGNSISENVGPK	0.631	0.643
NRITEETLAK	0.418	0.410	LQTLGLTQGTVVTISAEGED EQK	0.607	0.647
LLAYNCSPSFNWQK	0.609	0.617	IYQLK	0.426	0.436
HFAALSTNAK	0.415	0.429	VLPAVAMLEERAK	0.617	0.573
QMNDIEHQNLVGVSNHRTLE FAK	0.545	0.581	GDVLYDEVMERMDHFMDWLAK	0.783	0.734
LDNLVVFVINCNLQRLDGPVTGNGK	0.730	0.706	EEAHGAPLGEEEVALARQK	0.497	0.498
EGVFHTEWLGLCGR	0.647	0.653	GIANSILIK	0.552	0.515
TLGEFIVEK	0.556	0.565	IGVAMILRILK	0.610	0.610
LSEDAFDDQCTGANPRYPLISELK	0.629	0.651	LRGSVPECTLAQLGAAK	0.583	0.569
RISTVPEAVERMRSRVAK	0.500	0.512	DITLAMDCAASEFYK	0.661	0.657
NDSFRLMGFGHRVYK	0.532	0.542	AAYSSGK	0.133	0.149
RLLTTCNIPVPSDVRVATEFSETATPLK	0.656	0.678	GMNTAVGDEGGYAPNLGSNAEALAVIAEAVK	0.727	0.719

Peptide seq	NRT		Peptide seq	NRT	
	meas	pred		meas	pred
QWRRGFAVTPPELTK	0.543	0.561	DIADYEHNLMRRLRQLIAQS WHTDEIRK	0.683	0.618
VNNIGEI AAGADMVFAGSAIFDQPDYK	0.794	0.778	PLFSSHMIDLSEESLQENIEICSK	0.688	0.715
VNPGSI LSAEMMLRHMGWTEAADLIVK	0.868	0.822	EIGPLATPDVLHWTDSL PK	0.693	0.664
MRGSHHHHHHHTDPALRAIK	0.343	0.339	ADLWLA EYYDQRILVDK	0.650	0.697
ASMEVPSPQAGIVK	0.530	0.532	VQQPEFAAAK	0.451	0.449
GNDFFQQIQLTQC TRCVK	0.618	0.606	GAIVGMTGYGESAPADK	0.513	0.530
NDDYLILD AVNNQVYVNPTN EVIDK	0.683	0.699	YTSSV VIDESVIQGK	0.618	0.615
SAGLCGR	0.354	0.356	LRSPVDEAK	0.395	0.439
FPNGVK	0.404	0.400	VEPEALTLLAQQAFHDASFMLRPAHQK	0.778	0.752
ALAEHGIVFGEPK	0.527	0.548	RLQELAK	0.360	0.385
GAITIATREK	0.419	0.408	VAAEQLSLITVEGDK	0.514	0.550
ALAAGADV VHFDVMDNHYVPNL TIGPMVLK	0.719	0.788	DEEHINQNVLNK	0.368	0.349
ESWQDLPQNK	0.480	0.464	VINDNFGHIEGLMTTVHATTATQK	0.733	0.703
DHPIYYAGPAK	0.442	0.473	HYGRPM DIEWAK	0.505	0.501
IPSINVSK	0.481	0.446	AGTHIDSHGAPLGDAEIALTREQLGWK	0.601	0.591
QILLDTYYGRDYVEGETAAK	0.584	0.626	LEQAK	0.129	0.171
VNVGDK	0.158	0.188	VVMTGPSK	0.389	0.426
IRGIPADLED RR VEITGPVERK	0.534	0.551	TYSINEGNYIK	0.509	0.477
YNQLIRIEE ALGEK	0.623	0.585	RAVEEARAVTDK	0.379	0.349
FTGANTLEV EGENGK	0.492	0.505	RVFIRADLN VPVK	0.534	0.566
VVSDYLA K	0.460	0.476	IRMVYAPTQE HGK	0.447	0.451
DIALGEEFVN K	0.568	0.566	QYLIAPSILSADFARLGEDTAK	0.719	0.719
DQRVAASK	0.133	0.135	IEVEDFPAFILV DDK	0.733	0.740
TICGLCGR	0.470	0.474	PIAQPTGGDADPQTTIVRANSSTTAAEPLK	0.518	0.514
SFPQLK	0.485	0.439	GQRVWTGGGDEETLSK	0.464	0.493
HRLDVQFDNVEEAIAMSAMYAANHLK	0.751	0.703	VIHDISEMNRRIE PGDVLVTD MTDPDWEPIMK	0.698	0.722
AQREYASFTQEQV DK	0.463	0.470	LAVNIGLEILK	0.676	0.663
YPLADYALTPDMAIVDANLVMDMPK	0.794	0.799	GYINSLGALTGGQALQ QAK	0.610	0.613
TPSDVLAVHLLL K	0.644	0.636	MVQAPIFHVNADDPEAVAFVTRLA LDFRNTFK	0.813	0.797
DGH利VNGK	0.398	0.401	DDLLEVAMELENIALN DPYFIEK	0.886	0.825
RVVMCSG	0.334	0.376	AGIALNDNFVK	0.559	0.533
AGEGEAVRVDLNRP MK	0.476	0.473	VIAAGANVVR MNFSHGS PEDHK	0.515	0.546
VFEGNRPTNSILLREITPFS LGALIALYEHK	0.775	0.763	AETLAK	0.198	0.184
IIDLSAIQDEV ILVAADLTP SETAQLNLK	0.853	0.843	YINELQANPAK	0.469	0.476
LDVILLMSVNPFGGGQSFIPQTLDK	0.785	0.787	LVLV LNCGSSLK	0.605	0.606
ELHRDVCRFANTLLELGIK	0.675	0.663	FIRDHIAK	0.387	0.436
LAREAVEGAK	0.388	0.394	LYPNVDFYSGIILK	0.695	0.693
RPQDRVALP DVPK	0.493	0.473	SGFINK	0.390	0.396
PSLLMCK	0.509	0.503	GPSIMP GGQK	0.442	0.428
GLCGR	0.251	0.256	SATPAQAQAVHK	0.364	0.358
LNGSRH MVH ELV S NLRK	0.486	0.480	SEFAENDAYV HATPLIRRLAREFGVNLK	0.697	0.671
LEQEK	0.124	0.167	EQQLNSEN FVAFNL TERMQLIGGTWYG GEMK	0.755	0.745
HLAK	0.118	0.141	DIFTK	0.430	0.425
QAQIEK	0.381	0.384	VTS DARIRASLPTIELALK	0.615	0.641
FAAYK	0.382	0.344	VALDPLTGPMYQGREL AFK	0.646	0.678
TEEVCAR QIDA VLK	0.555	0.539	VSRAIA FAVGK	0.494	0.469
STNRPTSYR IGS LADF HRNLLK	0.624	0.624	ISPRFTIAASFGNVHGVYK	0.600	0.614
TANGCEAV C IFVN DDGS RPV LEELK	0.667	0.671	GNR SQVT DACK	0.338	0.321
LGSQFHIPH GLAN ALLICNV IRYN AND NPTK	0.692	0.690	SQSSNDVPTAMHV ALL ALRK	0.698	0.654
GNRSQQV TDACHK	0.338	0.282	FAALEAAGVK	0.522	0.490
HGGFYLG SIGG PAVLAQGS IK	0.666	0.654	DLAN AIRAL SMDA VQK	0.676	0.597
DPATLDYV VP FK	0.626	0.615	GVLG YT EDDVV STDFN GEV CTS VFD AK	0.681	0.684
EHGIRRYGAHGT SHF YV TQEAK	0.434	0.458	LIDDAWAWAK	0.563	0.581
RVIVRLSDFK	0.508	0.508	YVGLK	0.398	0.393
VIDEMRSELAK	0.475	0.506	SLISLK	0.505	0.462
YINIGVAVDTPN GLV VP FK	0.724	0.720	VAEETPH LIHK	0.426	0.446
IARPRQLY TGYEK	0.455	0.478	SLR RIDRSN VREK	0.361	0.362
VDMALVG DJK	0.573	0.579	PFIYQAPF PMGK	0.639	0.636
PFLNAYS RLLIK	0.626	0.617	ALLLK	0.462	0.483
LVAMGGIGHTS CLY TDQDNQPARV SYFGQK	0.594	0.614	LLAARIQ QQD IDN GTP DFIS ETAS IRD ADWK	0.680	0.720
AAQEIVNSGK	0.377	0.384	VREATR LAQEK	0.364	0.366
TFTTQETM TNAHS RDWFLK	0.575	0.633	IIRNTNF EDAK	0.425	0.470
TQT GALIM IFDSADGA ADAAPA QAEK	0.685	0.722	HGER FMERYA PNAK	0.439	0.420
LHGYLPSR QPNF TEK	0.489	0.529	AVT GLK	0.486	0.465
LTQLISAAQK	0.498	0.509	VLP IT HGA AVT GQGV VQETLN MSK	0.652	0.679
AGGVK	0.118	0.111	MW RLLH GESK	0.480	0.486
PMDPYV VEE DPGV K	0.539	0.547	GLCGR	0.253	0.256
IGNEK	0.123	0.123	ASMEV PAPF AGT VK	0.586	0.573
GLVTRQLSGK	0.432	0.417	GPEIRT MK	0.390	0.387
YLANK	0.205	0.210	SSNTET FVAIR VD IDN WRWAGV PFY LRTG K	0.727	0.729
PVVG YIAGV TAPK	0.564	0.561	IVIRPLP GLPV IRDLV VD MGQ FYAQ YEK	0.840	0.809
VFMVDRFGLL TDK	0.637	0.688	GRIL RED VQAY VK	0.488	0.488
MPTSLIH ALA LTK	0.624	0.621	TQSWQ EA AW SEV FS YAE DRF NL PR GTIK	0.775	0.776
GLT DAA QQV VAA VEGK	0.670	0.543	EFVESLET PRR ILL MVK	0.650	0.700
RAEA APAAT GGGI PGML PWPK	0.628	0.623	TYLYQETK	0.447	0.454
QAIAMP SGLCGR	0.546	0.536	RFSLEG GDALIPMLK	0.665	0.671

Peptide seq	NRT		Peptide seq	NRT	
	meas	pred		meas	pred
LQDLAK	0.366	0.424	HIPEFVRRAK	0.426	0.422
ISGANLSRNWVMIPHVTDFK	0.598	0.645	GSDYIGDQDAIEYMCK	0.583	0.562
ENNFALPAVNCVGTDSINAVLETAAK	0.726	0.688	SQSTVGAGDSMVGAMTLK	0.563	0.551
IFDFVK	0.570	0.547	NLRSRSLMATQAWLEARLK	0.651	0.631
E LAESEGAIERK	0.421	0.426	AQQSWNEK	0.349	0.357
AVQDVILK	0.497	0.507	ILDPRNTYASPEQWQE K	0.540	0.555
VFLNIGDK	0.539	0.529	FIEEK	0.354	0.421
IANDVRWLASGPRCGIGEISIPENEPGSSIMPGK	0.656	0.667	ISWMEIYTGEK	0.630	0.608
GLGEAK	0.166	0.152	IVSYAQGFSQLRAASEEYNW DLNYGEIAK	0.690	0.759
RLVTVQT DANGQPVNQILVEAATDIAK	0.688	0.681	IVTTLGPATDRDNNE LK	0.494	0.542
TFSEAIISGEWK	0.605	0.597	ACIGIITNPVNTTVAIAAEVLK	0.826	0.726
NHFASEYYNAYK	0.528	0.497	LPLGSRPAK	0.401	0.441
SLYEA DLVDEAK	0.542	0.561	QYGEAF EK	0.432	0.414
LT LNGDTA VEL DVLK	0.643	0.681	NHTTIFSEWYALDLVK	0.711	0.693
LN VPENPII PYIEGD GIGV DVT PAMLK	0.762	0.765	NGALNAAIIVGQPAYK	0.529	0.518
DIWPSAQEIARAVEQV STEMFRK	0.839	0.681	GLDDLAK	0.444	0.430
WTQS ALDLGEPLS LITESV FARYISSLK	0.871	0.792	GNDF QQIVNK	0.596	0.539
VV NGLK	0.365	0.397	TVRGQYTAGFAQGK	0.441	0.439
LGRHVAILGDLQGPK	0.531	0.549	DGAYVREHFFGK	0.501	0.482
STQVYQGDVWLPAETLDLIREYRVAIK	0.758	0.767	MTD L DLAGK	0.487	0.519
IEDVPQEQ RQDIFSLTNEEVQ ELAK	0.677	0.671	E PARPMVAIVGGSK	0.495	0.505
LLSNF FAQTE ALA FGK	0.745	0.746	TVRS LADIGEALK	0.571	0.533
L REGITATD LVLT VTQMLRK	0.769	0.696	R IYK	0.137	0.150
VFMAD FEDS LAPDWNK	0.676	0.714	ENRFTG WYD VDLSEK	0.589	0.585
RIVLPEGD EPR TVK	0.478	0.509	VLSGPQA QPAGDK	0.410	0.453
FEALATC DALVQAHGALK	0.606	0.657	SMFSGEK	0.396	0.397
YGDEQVK	0.341	0.340	APIVHALRGK	0.408	0.427
IEVEDFP AFILV DDK	0.732	0.740	RLGGDTAK	0.142	0.145
EVLESRLPGILELSRTFAHVDPVK	0.697	0.684	VLPELNGK	0.478	0.471
QYP IVSIED GLD ESD WDGFAY QTK	0.709	0.681	DIGAQYIIIGH SERRTYHK	0.501	0.508
PTQE QYDEFK	0.456	0.471	VIDGQINLR DAVNGT ISYTNEAGK	0.605	0.597
EELAIP P QIK	0.551	0.546	RISIK	0.302	0.246
DED GN YLV D VIL DEA N K	0.730	0.636	VTA EAK	0.298	0.305
LGG GLS AEA L TEK	0.525	0.532	GTGV GGR LTRED VEK	0.420	0.423
DDET LSK	0.231	0.195	TEEVIA EN PGK	0.429	0.433
WV D SITE A WAMD NDAPK	0.645	0.672	TQE QQVVAAC DK	0.421	0.429
EITST DDFY RL GK	0.535	0.538	ERMD NGEG LPQYIK	0.510	0.502
MAVA ESGM GIVE DK	0.534	0.571	I ELLPYHE LGK	0.578	0.594
TIIGFGSP NK	0.521	0.495	QVD ILGK	0.472	0.451
VNPET TLFLV ASK	0.637	0.618	TENTR VS YPI HID NIVK	0.573	0.589
LVP YYTVK	0.517	0.535	VG IDYK	0.428	0.412
VOLLGSGSILRH VREA AEILAK	0.686	0.669	RWI QRIES GRAT FN SEEK	0.479	0.498
EGL TLK	0.438	0.397	FAD VACAG PLLA E DAL GK	0.748	0.730
VE MK	0.147	0.170	TNW LR GLC GR	0.566	0.501
TYLYQETK	0.439	0.454	VIC QGFT GSQ GT FH SEQ AIA YG TK	0.561	0.611
IRV STFK	0.431	0.431	EDEK	0.115	0.092
MRAV QEQV A SEK	0.400	0.399	IQLVG DDL FVT NT K	0.633	0.653
GQAIYAYV TL NHGE E PSELYA EVRN W VRK	0.665	0.670	ATD LFLK	0.520	0.531
FICI A DASK	0.514	0.507	ELPY MN FPK	0.597	0.566
R IQNAGT EV VEA K	0.424	0.411	DDTE YY LLT SEH VS E FEG Q EIL K	0.681	0.703
LT VLD SLSK	0.549	0.545	APYNGRK	0.142	0.123
LLPS LYQ LEK	0.614	0.628	QENV YY YITL NEN YHMP AM PEG AE EG IRK	0.671	0.653
GQAGLCGR	0.365	0.391	ISYISTGG AF L FVEG K	0.708	0.697
MVINALNANVK	0.539	0.515	LDLFANEK	0.538	0.519
VNPRLA GLC GR	0.480	0.481	TV TYD FERL M DGAK	0.617	0.576
DAL GEH I LER VET IRK	0.588	0.544	RPDLMIDGPLQY DA VMAD VAK	0.696	0.692
GCPHLC PW CAN PESIS GK	0.586	0.586	GMPLY EHIA EL NGTPG K	0.587	0.578
MI AVTT SGTE V TPAV VT DDA TGQK	0.648	0.627	WV AMGEY K	0.512	0.517
IRV TAER DPAN LK	0.443	0.454	EAIFE GHIM LLEDE ELE QEI IALIK	0.842	0.850
HPELT DMV IFREN SEDI YAG IEWK	0.685	0.733	PAI GV GAG NT PVV IDE TADIK	0.604	0.631
MRTLGTAACPPYHIAF VIGG TS AET NLK	0.649	0.687	SRAFA DIVK	0.480	0.450
IFRAA ALA AADARI PLAK	0.593	0.563	E LA VK	0.331	0.261
EDEIV IDR K	0.426	0.446	TLT QTL NEK	0.440	0.452
VV ITAT QM LDS MIK	0.698	0.686	TDID DV TQL AK	0.532	0.524
QAGLCGR	0.361	0.395	GLC GR	0.259	0.256
LYNDAGISN DRIL IK	0.551	0.613	VLD LIA HIS K	0.573	0.554
LFPF GGFT AEN IVA K	0.762	0.743	MRA DK	0.120	0.104
LEHNP GQYI P QEL RQAGE GEAVK	0.557	0.540	ADEQILDIG DASA QELAE I LK	0.754	0.737
FLK	0.300	0.336	SRFLG K	0.360	0.357
GVLPTC QDGTG TAI IV GK	0.586	0.597	EPIPV IPTCH YM MG GI PK	0.641	0.628
YAA LCD VFM DAF GT A HRAQ AS THIG IK	0.671	0.681	MAQQ QGV AVK	0.388	0.434
AGL VDILG ASGA EN VQGE VQK	0.623	0.627	FNQIGSLT ET LAA I K	0.688	0.672
TYH YYSL PLA AK	0.554	0.541	GAN FDK	0.205	0.198
QEA ALGAGA AH HSE AL NFIV N TILA QK	0.758	0.754	LDGV K	0.317	0.255
QTVY AFLG DGE MDE PESK	0.631	0.589	LEG GND VSL K	0.439	0.456
QLFARYGLPAPVGYACTP REAEEA ASK	0.611	0.652	VMF GLC GR	0.563	0.547

Peptide seq	NRT		Peptide seq	NRT	
	meas	pred		meas	pred
GEQQAMQVATRAK	0.424	0.395	YAPFEIPSEIYAQWDAK	0.716	0.687
FNIDADK	0.443	0.436	ARDIVAGIASEEDEIVVFEGCEHAK	0.655	0.677
DSMGAIDVPADK	0.496	0.500	YTDTTPAGAALVAAGPK	0.533	0.551
FLLDANL GK	0.584	0.585	SLECVEYPELGMEAIW K	0.708	0.722
TPELNLFK	0.581	0.532	GETCPNELVGLCGR	0.555	0.563
NFRQLHSK	0.290	0.350	LIFDK	0.480	0.496
VRRALYLGK	0.413	0.437	ELLGLCGR	0.547	0.526
YDSTHGRFDGTVEVK	0.452	0.464	YLQQVNESFGFELEFFDFLLTEK	0.877	0.850
LIITITEGIPTLDMLTVK	0.774	0.786	LSNPVRLVK	0.480	0.470
SLGDITRLPK	0.519	0.487	LEDLLAAK	0.520	0.526
SLRNYGITAPIDVHLMVK	0.621	0.644	HNLPHNSLN FVFHGGSGSTAQEIK	0.544	0.581
ALLHVAK	0.419	0.433	GVLHTTGGYLVYAA LTFK	0.684	0.696
YYDPRVWL RAGQTSMIARLEK	0.652	0.646	EISTTI A FVRALNVMLK	0.750	0.723
TSAEISLT TGPVVPVIVVK	0.685	0.676	SIGTLSAFEQN ALEGMLDTLK	0.794	0.749
NRITINYFAMPSTFGAICK	0.672	0.670	DVL MEEIVANYHANTK	0.625	0.592
NYPDRVLPLDRQAVTM DK	0.496	0.511	QEQTPLAVIAGK	0.568	0.511
HMTADAAAHEVIEGQASALE ELDDEYLK	0.674	0.714	TGNTPDGRRAGAPFGPGANPMHG RDQK	0.445	0.446
VLNNGDLGENK	0.418	0.440	AEFI EK	0.415	0.439
MSYVEGLLSSNQK	0.579	0.536	RIDK	0.118	0.108
GDVA VFFGLSGT GK	0.640	0.602	NPNTVSVEHVVVLK	0.546	0.530
RSDIEIVAIN DLLDADYMAYMLK	0.806	0.800	ESDELI AK	0.417	0.437
RAIHTLWNVLN DEDQAWLPVEK	0.777	0.790	PFHYQAPFPLK	0.580	0.586
AGGYLT DGT ELHD TNFARIAEAC GTGIRVEK	0.644	0.676	AAGLSQYEHLIDDAI AWGK	0.720	0.690
QILT AEAEVFLTEL VTHFTPQRNK	0.869	0.751	IAAVAEDGEPCV TYIAGDA GHYVK	0.579	0.646
ERIELTRNSFGGAK	0.472	0.449	DIQK	0.130	0.142
LVLVRHGESQWNK	0.463	0.504	YPETAA LVADWTDEQI WALN RGHD PK	0.699	0.690
EIVIK	0.416	0.438	QELRDEGK	0.156	0.224
IMWV MYEH PETHFEE ALRFMDIRK	0.735	0.805	VDLNRPMK	0.418	0.426
VVDA AVEK	0.379	0.427	GQIEGA VSSSDASTEK	0.420	0.423
TGSLIMIFEVEGA AAPAAAPAK	0.700	0.709	QVAAILHDPEASENDK	0.486	0.501
GFSLYMLRAI SGRDEVIELAK	0.755	0.757	VEAEQSLITVEGDK	0.514	0.549
QLRTN VP HIF AIGDIV QPMLAHK	0.657	0.693	LAVYSTK	0.417	0.435
QGIRCIVDSSGEA LSA ALAIGNIELVK	0.783	0.750	GVFTDPGFTS ACESK	0.629	0.619
ILIGEV TVVDESEPF AHEK	0.625	0.678	TARILINTPASQGGIGDLYNFK	0.631	0.670
ITLQ NGK	0.377	0.394	NSTAM LTTFNEVN MK	0.593	0.598
LTSLRQ YTTVVADTGDIAAMK	0.620	0.653	DVMPEV NAVLEK	0.589	0.574
VLL ENLLR WQD GNSV TEEDI HALAGWL K	0.782	0.831	ELGLK	0.403	0.344
VTTIIQGGTSSVTALT GSTEESQ FGLCGR	0.650	0.676	GMFSMM NYLLPLK	0.777	0.735
EAIRM GS E VFHH LAK	0.520	0.509	GEAASNPEVIA RTLRK	0.493	0.458
LAQ LLRY SSNIALMC GSG CAGAH K	0.617	0.644	SAMEPVWNGK	0.490	0.472
LSYTGEVK	0.438	0.433	EAILEAQSVK	0.474	0.477
AITDV VNI GIGG SDSL GPY MV TEAL RPY K	0.738	0.724	AGAGT DAAI DLSK	0.482	0.471
VVNSK	0.118	0.158	MPNLLPQTK	0.630	0.593
TLNAEIVFVMSQG TDTP EQLK	0.691	0.683	ADAFAVIVK	0.561	0.515
IFRAG CIIR A QFL QK	0.606	0.659	AYPQEAAEFT RRMK	0.501	0.469
AAGVETEV FFEV EADPTLSIVRK	0.689	0.723	GVNNTYIEDN LRY SQA ALDMYK	0.643	0.652
PGNV VLPTL RDSQ EYVSK	0.613	0.597	LCDEVLF DLK	0.647	0.660
GGNL NTNEELA QT FTEW NN ELSS YLIDITK	0.803	0.799	LRLLYECNPMAFLAEQAGGK	0.713	0.725
RHG IRLGF MFSYV K	0.612	0.588	IMIDL DGTE NK	0.542	0.561
QLAN RILPELK	0.554	0.551	LFGV TLDIIR SNTFVA ELK	0.736	0.776
VREIAAK	0.248	0.289	YLERMSK	0.379	0.395
GLP ADVV PGD ILL DDGRVQLK	0.707	0.714	AATEG TIPLIP GISTV SELMLG MDY GLK	0.856	0.801
HFAATGK	0.167	0.180	NIAHQVK	0.480	0.304
AATYEQIK	0.413	0.407	LGLCGR	0.443	0.451
ELAN AIRAL SMDA VQK	0.615	0.592	AELAK	0.145	0.174
RAVASV LMSK	0.459	0.451	ANEAYLQGQLGNPK	0.496	0.484
IYADK	0.169	0.253	GLAASLMK	0.507	0.474
TVLK	0.175	0.211	INREGI WI EK	0.506	0.517
LWGAQ TQRS LEHFRISTEK	0.553	0.578	EQQQ DVITR TFTQS MGLCGR	0.621	0.602
DYLDGV DVAEGELV VLEN VR FNK	0.740	0.731	NYTPYEGDES FLAGATE ATT TLW DK	0.713	0.688
IRTYGLCGR	0.459	0.469	LADSGLN II AAK	0.558	0.557
YVALQFLRN SEIAAK	0.612	0.608	CSEVV VYFK	0.561	0.521
TEIDK	0.154	0.148	SSSV ETMPD LPLK	0.575	0.566
DNT PMFVK	0.443	0.483	GTLGQD VIDIR TLGSK	0.596	0.558
VDFSK	0.342	0.345	VMVTSHLGRP TEGEY NEEF SL PVV NYLK	0.701	0.723
IGR THLQ DATPLT LQ EIS GWV AMLEH NLK	0.832	0.748	HLEC VAYPELG MEAI W K	0.677	0.698
TILWNGPVG VFEFP NFRK	0.721	0.720	ILDWIK	0.586	0.570
QQQDLV NDALN VEG LCGR	0.627	0.624	SLLR HPLA VSS LEE LANG TFLPAIGE IDEL DP K	0.774	0.793
FAII DAVNG EY LSLG LAEC FHLPEARIK	0.777	0.752	EHLG TG DVK	0.357	0.343
SMLL QRNAGL CGR	0.510	0.543	LFIDNFDK	0.592	0.594
LLTK	0.254	0.294	GYGM GDAA EGK	0.410	0.396
EDIRAF AENW LGK	0.646	0.583	AA DIVLQ AAIA AGA PK	0.626	0.588
ELAGVAG CAVAI APPEM YIDMAK	0.707	0.741	TAAILL DTK	0.513	0.535
LHG GEPANFLD VGG GATK	0.547	0.543	SLFK	0.391	0.369
AGQLNPDT RIIGV GRADW DK	0.551	0.555	EEGYSDFAY TS VLK	0.665	0.647
GLNP TRAIGH IK	0.470	0.432	AAEIAK	0.175	0.197
VLEVQGMK	0.470	0.485	HAI VAITEHM CDV DELA HFIEK	0.672	0.697

Peptide seq	NRT		Peptide seq	NRT	
	meas	pred		meas	pred
MVHNGIEYGDMQLIAEAYSL LK	0.772	0.762	CTNPQWK	0.405	0.398
LDAPLIVVATQGGK	0.619	0.616	AELGIPK	0.468	0.439
LLEEK	0.351	0.349	IADQLIVGGGIANTFIAAQGHDVGK	0.661	0.712
VGINGFGRIGRIVFRAAQK	0.606	0.598	VVPAQGK	0.393	0.410
EYAEVFEGTAEWK	0.589	0.594	ELALQSGLAHK	0.483	0.471
RMGHAGAIIAAGGK	0.398	0.394	NTNTITGVIVNK	0.502	0.462
ALVAGGVRVLEVTLRTECAVDAIRAIK	0.729	0.716	SGHPGAPMGMADIAEVLWRDFLK	0.802	0.716
AGIAK	0.138	0.160	QDAAYRESVLPK	0.462	0.456
EVNTGTNLPQAQIDLYAVDGD EYK	0.636	0.630	EVNTGTNLPQAQIDLYAVDGEYK	0.637	0.630
HGGFYLGSIQGPAAVLAQQSIK	0.669	0.658	AAAEGEMK	0.205	0.204
TGLTGEEK	0.448	0.452	ECDLAGAIK	0.485	0.459
ALFRQMLGEEMK	0.568	0.619	EVNVPDINGDEVEVTEVMVK	0.648	0.630
GVLPTCQDTGTAIIVGK	0.589	0.597	AFAASNELEVNAUTH	0.493	0.469
YIVRDDTRDFTFWADK	0.592	0.634	VNVGDK	0.158	0.188
QTVAAYIAK	0.477	0.448	LNAPVDEQGRTRPDLSEIFDDSSK	0.585	0.539
VVREALETFMK	0.555	0.569	FAVLK	0.470	0.464
SVRNIYDYYK	0.500	0.474	GSGPYFYLPK	0.581	0.550
IYGDRQRAMAAGEK	0.410	0.438	LEHAVPMAK	0.415	0.450
RVSDMEPK	0.357	0.341	SLGIHVFDLNEVDSLGIYVDGADEINGHMQMIK	0.741	0.778
GMEGAINAK	0.407	0.392	TIVK	0.165	0.224
GHIELSRELMITISK	0.675	0.642	DHPIYYAGPAK	0.451	0.473
QRWPDVLLQFEDFAQK	0.712	0.698	HQNILYTAPTAIRALMAEGDK	0.688	0.667
ASEVDEALQRFAISDGVLVDVVAK	0.759	0.748	PVEELNIHTCHLGNNGGSVSAIRNGK	0.609	0.618
FTEGAFK	0.451	0.460	SIEK	0.127	0.136
VLAEQALAQPPTDELMTLVN K	0.685	0.737	NTSFAPGNVSIK	0.515	0.477
EVTVEDLMK	0.567	0.526	EISSHDSTSNTGLINRYK	0.465	0.469
ESAWNEK	0.373	0.364	VFQVAK	0.421	0.436
ELVEFAGK	0.499	0.488	ISADQVDQEVEFLSGRAK	0.671	0.547
NLVQQVAK	0.457	0.412	TIASFQQQLSDMGYK	0.613	0.599
SQNGAAMSFGRSTFLDVYIERDLK	0.684	0.663	ELLIEAQNLGLGQAQFGGK	0.663	0.717
LHTYRDAIPTQSVLTTSNVYVGK	0.644	0.638	TQLPSGESELSLYDIAPVTGVAVDLSHIPTAVK	0.728	0.725
IIRDINK	0.332	0.398	FLREEMGVK	0.467	0.503
DGYTFVSHHQEVGTGYFDK	0.563	0.547	YVLAGEGNK	0.419	0.422
AGETFGEEK	0.374	0.379	AAGYELGK	0.409	0.382
DASRYSSTISDPDTNVK	0.456	0.481	RAENMLWHK	0.433	0.424
HEIVAHLVERGVEVIDK	0.567	0.574	TLESAGVK	0.377	0.387
GGGAALTREK	0.241	0.243	ELSAVNRELTQPDVVRK	0.540	0.557
SIYFRRGSLPLIDEVITDGHK	0.634	0.647	PAPEGATMHWYHAPQPVVTPEEEELRK	0.560	0.606
MPTMAAMCYK	0.545	0.547	FAIDQEK	0.439	0.443
GDWLDCAK	0.514	0.478	LTGMAFRVPTPNVSVDLTVRLEK	0.699	0.703
ETMERVK	0.276	0.209	QGDLCILDGK	0.533	0.521
AQETK	0.117	0.090	VLPVTQEAIQK	0.501	0.545
VFTEVTGGPLSNNK	0.560	0.566	AFQEELNAIDVGLCGCR	0.707	0.717
ALLTPGK	0.425	0.448	GILEQYGHK	0.454	0.409
WDEVGVDVVAEATGLFLTDETARK	0.777	0.744	VAVLGAAGGIGQALALLK	0.830	0.793
AFTSEEFTHFLEELTK	0.753	0.689	RFLSELTAEGLERYLGAK	0.665	0.654
RRLIGDDEHGWDGGVNFNFECCYAK	0.616	0.591	NVDDALK	0.395	0.346
DPQTEAIVMIGEIGGSAEEAAAYIK	0.804	0.726	QQSNNDRAQQIVDATDK	0.445	0.465
RVVVSGLGPQGALGVSENCIQVVPPVK	0.649	0.658	HK	0.349	0.090
TESEEMLAK	0.418	0.436	FGANAILAVSLANAK	0.172	0.558
AVTARVAVEAGIADYWYK	0.626	0.626	INRQGIWIEK	0.489	0.509
YLEHRGLK	0.369	0.382	LGLCGR	0.443	0.451
MRGSHHHHHHTDPALRASK	0.114	0.140	SVNDVK	0.150	0.180
RGAFAMGGMAAFIPSX	0.623	0.616	EAYELVAPILK	0.627	0.620
NQPELSEDTIK	0.465	0.463	AAALALGYLSEAEDFSWVRPEQMVGSMK	0.755	0.798
PNPAVLICRVRGLHLPEK	0.617	0.606	TQGAAAFEGAVIAYEPVVAIGTGT	0.715	0.753
ATVILAHTIK	0.484	0.504	MLEISSVK	0.499	0.511
NGVMIVNTSRGALIDSQAAIEALK	0.655	0.715	AVQRSADLISIGPMLQGMRK	0.621	0.616
NNQHDVAIVRIEQLYPFPHK	0.614	0.600	GIASMHCSANVGEK	0.448	0.431
QAFLIRELNSK	0.544	0.532	NYLVEK	0.410	0.405
STLRALLPLVEEK	0.625	0.616	MNIDTDTQWATWEGVLNYYK	0.729	0.719
LASTWEGIRAAELEK	0.593	0.607	LGVLGFEDHERNLAAFRGK	0.601	0.645
HITAGAK	0.133	0.141	YAGQDIVSNASCNTCLAPLAK	0.604	0.632
IILSDKK	0.429	0.489	VEAEQSLITVEGDK	0.514	0.549
PGMDRSIDLNVNAGIVK	0.593	0.593	DAEVLVLEGLVPTRK	0.617	0.604
HNLQITK	0.372	0.352	GGIYLYPSTASHPDGK	0.509	0.526
ASMEVPAPFAGVVK	0.626	0.589	IVCTIGPK	0.466	0.486
ITPVVFIMK	0.637	0.642	LREVRRRIDESGFDIRLEVDGGVK	0.547	0.563
AMIEAGAAA VHFEDQLASVK	0.641	0.668	LETIEGSK	0.395	0.421
LASTWQGIRAAEQLK	0.574	0.603	TAHQLVLSK	0.416	0.431
LVQQFTK	0.433	0.478	GEMPSDFDAK	0.467	0.471
IMDATQARDVVK	0.467	0.490	LGWHHPPFEIPK	0.567	0.578
VINQLTGGLAGMAK	0.572	0.602	PVDRIVPDFAAAGASIIHFPEASEHVDRTLQLIK	0.694	0.728
LGLGAQFGGK	0.517	0.502	AVAAALEQMPRFNSSLSEDGQRLTLK	0.620	0.687
IGSLGMDVYENERDLFFEDK	0.667	0.696	DAGVMAASWAQYQAQDALIK	0.651	0.713
PYLDK	0.368	0.338	QAFSRFSVVPAGVICHQVNLEYLGK	0.645	0.673
AHPQLAEFFRRMSGGGLPK	0.552	0.533	LYGLCGR	0.496	0.486

Peptide seq	NRT	
	meas	pred
EHVEYDNPYDVGMTGLIGFSSGFHTMMNADTLVLLGTQFPYRAFYPTDAK	0.841	0.842
HTIPANIADRCLINPQQYEAMYQQSINVPDFWGEQGK	0.698	0.719
MRGSHHHHHHTDPALRAMVRIYTLTLAPSLDSATITPQIYPEGK	0.656	0.622
QRQVMVYTHDSIGLGEDGPTHQPVEQASLRVTPNMSTWRPCDQVESAVAWK	0.650	0.670
LQESSPLPVLGAVPWSDLIATRAIDMARHLNATIINEGDIINTRVK	0.882	0.827
ALIRRARQLNRNAVITAQMMESMITNPMPTRAEVMDVANAVLDGTDAVMLSAETAAGQYPSETVAAMARVCLGAEK	0.900	0.870
HIEEPLPHVAELALGGTAVGTLNTHPEYARRVADELAUTCAPVTAPNK	0.746	0.719
NMYPSGDGARRDEDGYYWITGRVDDVLNVSGHRLGTAEIESALVAHPK	0.700	0.687
RRLLSFSNHGSMAAMPQALGAQATEPERQVAVMCDDGFSMLMGDFLSSVQMK	0.797	0.790
TTVTRHTMIHEQITRLFHAFRRDHSHPMAVMCGITGALAAYFYHDSLVDVNPRHREIAAFRLLSK	0.753	0.717
TLAAQFNRPGLDIVDHYTAFMMDGCMMEGISHEVCSSLAGTLK	0.759	0.786
LRCTAPVFEPGGGINVARAIAHGGSATAIFPAGGATGEHLVSSLADENVPVATVEAK	0.776	0.805
ASAIRQADEVLAQHDDEFPLAQTGSGTQSNNMNEVLANRASELLGGVRGMERK	0.740	0.750
RWVSLPGEHHGVDFAPNSE EAIILELRAEIGHVSAERV LSGPGLVNLYRAIVK	0.825	0.818
QIADDYQQALRDVVAYAVQN GIPVPTSAAVAYYDSYRAA VLPANLIQAQRDYFGAHTYK	0.823	0.853
VQEYVANYINADWIESLTATRSERSRLSPPAFRYQLTELARK	0.706	0.709
DDIFFRSTGGGVTLSGGEVLMQAEFATRFLQLRLWGVSCAJETAGDAPASK	0.843	0.858
VVRVPAYDPEAWEAHGMMMTLNRRRIHRYQTRDANSLEGLTGFTMYGK	0.772	0.770
NAMPILLRNECISFNDDIQGTAAVTGTIAASRAAGGQLSEK	0.814	0.778
GVTAAITMTESGRTALMTRSRISSGLPAMSRHERTLNLTYRGVTPVHFDSDANDGVAASAEAVNLLRK	0.763	0.837
GDIIIDGGNTFFQDTIRRN RLSAEGFNFIGTGVSGEEG ALK	0.712	0.724
LAENASLEEMVRFGVAAGSAATLQNQGTRLCSHDDTQK	0.683	0.686
EGIHTCLDTNGFVRRYDPVI DELLEVTDLVMLDLK	0.899	0.825
YSYEASLMAHLDRDVIRTMACCGIAGLSVAADSLSAIK	0.783	0.745
VVLEVPAASADGILDADVLEDEGTTVTSRQLGRLREGNSAGK	0.749	0.739
ERILDIIPETLHQRRSFFVGNDHMVEDVERFIREFPDAGLCGR	0.765	0.779
EVPEAIVGAGTVLNPQQLAEVTEAGAQFAISPGLTEPLLK	0.817	0.844
TNLAGLMDGYFHHASIEIEGGQHQLNVNMNREMlldamenpeK	0.713	0.765
SDVPQGAAILGAISGAHHVHQMAEHYGVPLHDCAK	0.644	0.652
TFDNGVICASEQS VVVVVDSYDA VRERFATHGGYLLQGK	0.797	0.742
DWTRQNLHVHVVEASGEQYRFVMPGAALNEDEFRQLEEQVLEIESGAILVISGSLPPGVK	0.878	0.853
AGGGSATLSMGQAAARFGLSLVRALQGEQGVVECAYVEGDGQYARFFSQPLLLGK	0.866	0.843
GGVNVAACTGINSYNTIPVVEEPYEPGNLELERIRS AIRWNAIMTVLASK	0.729	0.782
GSAFSEMEERRNFNLGLPEVETIEEQERAWIQYQGFK	0.895	0.847
VAPEALTLLARQAFHDASFMRPAHQQQVADILRDPEASE NDK	0.744	0.782
HNPQNPSPWADRDRFVLSNGHGSMLIYSLHLLHTGYDLPMEELK	0.759	0.765
DLIGWIDQPSVELSNALMHHDPI DINLILATGGPGMVK	0.809	0.825
WGFAVVENSLWQGPVNPYLRELNEQLEENLYK	0.869	0.837
MRGSHHHHHHTDPALRASV IGHFESCGTVDPGIRFI TFFQGCLMRCLYCHNRDTWD THGGK	0.736	0.645
GQRVWTGGDEAALARGVYN TYIEDNLRYSNQAPLDMYK	0.675	0.674
VERAEAVCSQDAMDDII LASDVMVARGDGLVEIGDPELVGIQK	0.828	0.809
LDLSYSETFNQTHLADAYERLLLETMRGQALFVRRDEVEEAWK	0.801	0.856
ALQAIAGPSQVRCPTGGIS PAN YRDYALK	0.705	0.717
QTAFSQYDRPQARRYAEIADHGLSAPGDRTAAK	0.545	0.550
TPGHPEVGYTAGVETTGPLGQGIANAVGMAIAEK	0.708	0.633
YSMPVPMNIINGCEHADNNVDI QEFMIPQVPGAK	0.741	0.737
VIFLTADAFGVLPVSR LTADQTQYHFSLSGFTAK	0.771	0.837
YIGAYTALMDGRDAVVF TGGIGENAAMVREL SLGK	0.762	0.830
GYLMSGDLVITVQGDVMSTVG STNTTRILTVEGLCGR	0.748	0.759
NAHADREIAYRPARVLMQDFTGVPAVVDLAAMREAVK	0.697	0.717
LAQFTSLQADLENGVNLEQTIRLREEIAEQHRLGQMK	0.685	0.725
HCAVGIEPNRERINQLNESMLVTALNTHIGYDK	0.785	0.739
RLFVVDACFGANPDTRL SVR FITEVAWQAHFVK	0.762	0.768
DTRAIIDAILNGSLDNAETFTLPMFNLA IPT ELPGVDTK	0.894	0.851
EAEPEIYN AIRRDALLENVTVREDGTIDFDDGSK	0.666	0.661
TPEGYASGSLGPTTAGRMDS YVDQLQAQGGSMIMLAK	0.687	0.651
LPVREFDAVVIGAGGAGM RAALQISQSGQT CALLSK	0.743	0.799
EYVHDIPVYLVHDPNGLLG SGAHLRQTLGHILGLCGR	0.748	0.757
PVN DLSR GALVDDIVYTI ALTAQSAQQQQGLCGR	0.877	0.793
IVFLGAGSAGCGIAEMI SQTQREGLSEEARQK	0.740	0.757
EVV TYRHF MNASGGGV T ASG GEAILQAEFVR DWFRACK	0.886	0.750
VWIRYVVPGWSDDDS AHR LGFTRDMGNVEK	0.658	0.676
MRGSHHHHHHTDPALRAA VTNAELNALVERVK	0.587	0.542
ATLLIETLP AVFQMD EILHAL RDHIVGLNCGRWDYI FSYIK	0.892	0.870
SLEAN NGHD GTWIAH PGLAD TAMA VFN DIL GSRK	0.762	0.716
YGVERQDGPTALILSRQNL AQQERTEEQLANIA RG GYVLK	0.663	0.670
QQIGVVGMA VMGRN LALNIE SRGYTVSIFN RSREK	0.717	0.707
TAALIGV DYLAVSF PRCGED LN YARR LARDAG CDAK	0.713	0.679
LAGTERGITEPTP ITSA CF GAFLS LHPTQYAEVLVK	0.753	0.783
ASQNAIEAFGPLLPEFLGG SADLAPS NLTL WSGSK	0.814	0.817
TVINF DN AIIAAGSRP IQLPFIP HEDPRI WDSTD ALELK	0.759	0.825
REAEGSHIMLGAQNV DLNLSGAFTGETSAAMLK	0.664	0.749