Online Supporting Information S1. The benchmark dataset $S = S^+ \cup S^-$, where the positive dataset S^+ contains $N^+ = 731$ SNO sites while the negative dataset S^- contains $N^- = 810$ non-SNO sites. See the text of the paper for further explanation.

Positive dataset \mathbb{S}^+ contains 731 SNO sites and the corresponding peptide sequences

UniProt ID	Site	Sequence (cf. Eq.1 of the paper)
P21980	10	XMAEELVLERCDLELETNGRD
P21980	27	NGRDHHTADL <mark>C</mark> REKLVVRRGQ
P21980	98	WTATVVDQQDCTLSLQLTTPA
P21980	143	GHFILLFNAWCPADAVYLDSE
P21980	230	VGRVVSGMVNCNDDQGVLLGR
P21980	269	DILRRWKNHGCQRVKYGQCWV
P21980	277	HGCQRVKYGQCWVFAAVACTV
P21980	285	GQCWVFAAVACTVLRCLGIPT
P21980	336	DKSEMIWNFHCWVESWMTRPD
P21980	370	TPQEKSEGTYCCGPVPVRAIK
P21980	371	POEKSEGTYCCGPVPVRAIKE
P21980	524	VSYNGILGPECGTKYLLNLNL
P21980	545	EPFSEKSVPLCILYEKYRDCL
P21980	620	QNPLPVALEGCTFTVEGAGLT
P21817	35	ATVLKEQLKLCLAAEGFGNRL
P21817	252	RLVYYEGGAVCTHARSLWRLE
P21817	314	SKAHTKATSFCFRISKEKLDV
P21817	810	KFLPPPGYAPCHEAVLPRERL
P21817	905	VRDDNKRLHPCLVDFHSLPEP
P21817	1039	ATKRSNRDSLCOAVRTLLGYG
P21817	1302	LPVQFHQHFRCTAGATPLAPP
P21817 P21817	1590	QSERKNPAPQCPPRLEMQMLM
P21817	2326	KGYPDIGWNPCGGERYLDFLR
P21817 P21817	2363	VVRLLIRKPECFGPALRGEGG
P21817 P21817	3193	VVKLLIKKFECFGFALKGEGG VEKLRPALGECLARLAAAMPV
P21817 P29474	3635 94	SKQRRRAVVACFRMTPLYNLP
		LSAQAQQDGPCTPRRCLGSLV
P29474	99	QQDGPCTPRRCLGSLVFPRKL
P29474	184	AKQAWRNAPRCVGRIQWGKLQ
P29474	201	GKLQVFDARDCRSAQEMFTYI
P29474	212	RSAQEMFTYICNHIKYATNRG
P29474	661	GLGSRAYPHFCAFARAVDTRL
P29474	802	QYQPGDHIGVCPPNRPGLVEA
P29474	853	GWVRDPRLPPCTLRQALTFFL
P29474	976	DGLGPLHYGVCSTWLSQLKPG
P29474	991	SQLKPGDPVPCFIRGAPSFRL
P29474	1048	QPTPMTLVFGCRCSQLDHLYR
P29474	1050	TPMTLVFGCRCSQLDHLYRDE
P29474	1114	LCLERGHMFVCGDVTMATNVL
P13639	41	GKSTLTDSLVCKAGIIASARA
P13639	67	TDTRKDEQERCITIKSTAISL
P13639	290	PEGKKLPRTFCQLILDPIFKV
P13639	369	PSPVTAQKYRCELLYEGPPDD
P13639	567	CLKDLEEDHACIPIKKSDPVV
P13639	591	ETVSEESNVLCLSKSPNKHNR

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P13639	651	WDVAEARKIWCFGPDGTGPNI
P13639	728	GGQIIPTARR <mark>C</mark> LYASVLTAQP
P17987	76	EVEHPAAKVLCELADLQDKEV
P17987	147	IVNTDELGRDCLINAAKTSMS
P17987	236	PKRIVNAKIACLDFSLQKTKM
P17987	357	QAEEVVQERICDDELILIKNT
P17987	397	EMERSLHDALCVVKRVLESKS
P78527	25	LQETLSAADRCGAALAGHQLI
P78527	478	LAAKGPVLRN <mark>C</mark> ISTVVHQGLI
P78527	795	MQPYYKDILPCLDGYLKTSAL
P78527	1499	KGIAPGDERQCLPSLDLSCKQ
P78527	1507	RQCLPSLDLSCKQLASGLLEL
P78527	2342	ERKNILEESL <mark>C</mark> ELVAKQLKQH
P78527	3187	NIWDDIITNRCFFLSKIEEKL
P78527	3347	IANALSSEPA <mark>C</mark> LAEIEEDKAR
P78527	4045	EKNWYPRQKI <mark>C</mark> YAKRKLAGAN
P10599	62	VIFLEVDVDDCQDVASECEVK
P10599	69	VDDCQDVASECEVKCMPTFQF
P10599	73	QDVASECEVK <mark>C</mark> MPTFQFFKKG
P98170	12	TFNSFEGSKTCVPADINKEEE
P98170	90	VGRHRKVSPNCRFINGFYLEN
P98170	213	CGGKLKNWEPCDRAWSEHRRH
P98170	300	YALGEGDKVKCFHCGGGLTDW
P98170	303	GEGDKVKCFHCGGGLTDWKPS
P98170	327	WEQHAKWYPGCKYLLEQKGQE
P98170	351	NIHLTHSLEECLVRTTEKTPS
Q15149	950	MRGRLPLLAVCDYKQVEVTVH
Q15149	1098	RLMAEREYGSCSHHYQQLLQS
Q15149	1136	LKDIRLQLEACETRTVHRLRL
Q15149	3295	KSHRVPLDVACARGCLDEETS
Q15149	3299	VPLDVACARGCLDEETSRALS
Q15149	3336	PATYGELQQRCRPDQLTGLSL
Q15149	3667 237	NLTYRQLLERCVEDPETGLRL
P10809 P10809	442	YFINTSKGQKCEFQDAYVLLS VEEGIVLGGGCALLRCIPALD
P10809 P10809	447	VLGGGCALLRCIPALDSLTPA
P68366	54	GGDDSFTTFFCETGAGKHVPR
P68366	295	OLSVAEITNACFEPANOMVKC
P68366	347	TKRSIQFVDWCPTGFKVGINY
P68366	376	GDLAKVQRAVCMLSNTTAIAE
P13010	235	YSFSESLRKLCVFKKIERHSI
P13010	249	KIERHSIHWPCRLTIGSNLSI
P13010	296	KEDIQKETVYCLNDDDETEVL
P13010	339	EOMKYKSEGKCFSVLGFCKSS
P13010	346	EGKCFSVLGFCKSSQVQRRFF
P13010	493	PNPRFORLFOCLLHRALHPRE
Q00610	151	MFDRHSSLAG <mark>C</mark> QIINYRTDAK
Q00610	491	RANVPNKVIQCFAETGQVQKI
Q00610	736	DVHFKYIQAACKTGQIKEVER
Q00610	918	YCEKRDPHLACVAYERGQCDL
Q00610	926	LACVAYERGQCDLELINVCNE
Q00610	934	GQCDLELINVCNENSLFKSLS
P00533	190	FQNHLGSCQKCDPSCPNGSCW
P00533	291	PEGKYSFGATCVKKCPRNYVV
P00533	311	VTDHGSCVRACGADSYEMEED
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P00533	329	EEDGVRKCKK <mark>C</mark> EGPCRKVCNG
P00533	470	DVIISGNKNLCYANTINWKKL
P07237	53	YLLVEFYAPWCGHCKALAPEY
P07237	56	VEFYAPWCGHCKALAPEYAKA
P07237	312	LEFFGLKKEECPAVRLITLEE
P07237	397	NVFVEFYAPWCGHCKQLAPIW
P07237	400	VEFYAPWCGH <mark>C</mark> KQLAPIWDKL
P27348	25	QAERYDDMAT <mark>C</mark> MKAVTEQGAE
P27348	94	EKVESELRSICTTVLELLDKY
P27348	134	DYFRYLAEVA <mark>C</mark> GDDRKQTIDN
P00558	50	IKAAVPSIKF <mark>C</mark> LDNGAKSVVL
P00558	99	LGKDVLFLKDCVGPEVEKACA
P00558	108	DCVGPEVEKACANPAAGSVIL
P21333	444	LEARGDSTYR <mark>C</mark> SYQPTMEGVH
P21333	717	LRVQVQDNEG <mark>C</mark> PVEALVKDNG
P21333	810	GQGDVSIGIK <mark>C</mark> APGVVGPAEA
P21333	1260	EPAVDTSGVQ <mark>C</mark> YGPGIEGQGV
P21333	2543	VFVDSLTKATCAPQHGAPGPG
P49327	634	GLSWEECKQR <mark>C</mark> PPGVVPACHN
P49327	642	QRCPPGVVPACHNSKDTVTIS
P49327	1118	EQQVPILEKFCFTPHTEEGCL
P49327	1127	FCFTPHTEEGCLSERAALQEE
P49327	1471	RREPGGNRLRCVLLSNLSSTS
P63244	138	TIKLWNTLGVCKYTVQDESHS
P63244	168	PNSSNPIIVSCGWDKLVKVWN
P63244	182	KLVKVWNLANCKLKTNHIGHT
P63244	249	LCFSPNRYWLCAATGPSIKIW
P06213	1083	NEASVMKGFTCHHVVRLLGVV
P06213	1165	FVHRDLAARNCMVAHDFTVKI
P06213	1261	DGGYLDQPDNCPERVTDLMRM
P06213	1272	PERVTDLMRMCWQFNPKMRPT
P07900	420	KVIRKNLVKKCLELFTELAED
P07900	481	GDEMVSLKDYCTRMKENQKHI
P07900	597	VVSNRLVTSPCCIVTSTYGWT
P07900	598	VSNRLVTSPCCIVTSTYGWTA
P18031	32	DIRHEASDFPCRVAKLPKNKN
P18031	92	ILTQGPLPNTCGHFWEMVWEQ
P18031	215	SPEHGPVVVHCSAGIGRSGTF
P35579	91	SKVEDMAELTCLNEASVLHNL
P35579	569	PKQLKDKADFCIIHYAGKVDY
P35579 P35579	896 917	QEQLQAETELCAEAEELRARL
P46459	21	TAKKQELEEICHDLEARVEEE
P46459 P46459	91	CPTDELSLTNCAVVNEKDFQS SLYTFDKAKOCIGTMTIEIDF
P46459	250	FPPEIVEQMGCKHVKGILLYG
P46459	264	KGILLYGPPGCGKTLLAROIG
P12814	332	LHKPPKVQEKCQLEINFNTLQ
P12814 P12814	480	
P62826	112	PSVNARCQKI <mark>C</mark> DQWDNLGALT PNWHRDLVRVCENIPIVLCGN
P62826	120	RVCENIPIVLCGN RVCENIPIVLCGNKVDIKDRK
P62826 P62736	219	EIVRDIKEKLCYVALDFENEM
P62736	259	VITIGNERFRCPETLFQPSFI
P62736	287	VITIGNERFRCPETLFQPSFI HETTYNSIMKCDIDIRKDLYA
P62736	26 <i>7</i> 376	EAGPSIVHRKCFXXXXXXXXX
P04350	12	REIVHLQAGQCGNQIGAKFWE
104330	12	VETAUDÕYQÕ <mark>C</mark> QNÕTQYVLME

P04350	239	VSATMSGVTTCLRFPGQLNAD
P04350	303	MFDAKNMMAA <mark>C</mark> DPRHGRYLTV
P04350	354	WIPNNVKTAVCDIPPRGLKMA
P55072	105	RLGDVISIQPCPDVKYGKRIH
P55072	535	TLLAKAIANE <mark>C</mark> QANFISIKGP
P55072	572	IFDKARQAAP <mark>C</mark> VLFFDELDSI
P34932	34	ETIANEYSDRCTPACISFGPK
P34932	38	NEYSDRCTPACISFGPKNRSI
P34932	245	KFDEVLVNHFCEEFGKKYKLD
P34932	417	NSPAEEGSSDCEVFSKNHAAP
P31943	34	ADEVQRFFSDCKIQNGAQGIR
P31943	122	FVRLRGLPFGCSKEEIVQFFS
P31943	267	SDRFGRDLNYCFSGMSDHRYG
Q99832	364	GGERYNFFTGCPKAKTCTFIL
Q99832	450	KALEIIPRQLCDNAGFDATNI
P14866	260	SLNGADIYSGCCTLKIEYAKP
P14866	261	LNGADIYSGCCTLKIEYAKPT
P14866	452	NFMFGQKLNVCVSKQPAIMPG
P14866	581	NGPYPYTLKL <mark>C</mark> FSTAQHASXX
P50991	295	NLVKQIKKTG <mark>C</mark> NVLLIQKSIL
P50991	379	${ t GSGKLLKITG}{ t CASPGKTVTIV}$
P50991	410	EAERSIHDALCVIRCLVKKRA
P04075	73	LLTADDRVNPC IGGVILFHET
P04075	240	KPNMVTPGHACTQKFSHEEIA
P04075	339	VKRALANSLA <mark>C</mark> QGKYTPSGQA
P14618	152	ITLDNAYMEK <mark>C</mark> DENILWLDYK
P14618	423	AVGAVEASFKCCSGAIIVLTK
P14618	424	VGAVEASFKCCSGAIIVLTKS
P14618	474	HLYRGIFPVLCKDPVQEAWAE
P36873	127	NFFLLRGNHECASINRIYGFY
P36873	155	NIKLWKTFTDCFNCLPIAAIV
P36873	158	LWKTFTDCFNCLPIAAIVDEK
P36873	245	FLHKHDLDLICRAHQVVEDGY
P15121	299	LLSYNRNWRVCALLSCTSHKD
P15121	304	RNWRVCALLSCTSHKDYPFHE
P31749	224	KYSFQTHDRLCFVMEYANGGE
P31749	296	GHIKITDFGLCKEGIKDGATM
P31749	310	${ t IKDGATMKTF}{ t C}{ t GTPEYLAPEV}$
P35754	8	XXXMAQEFVNCKIQPGKVVVF
P35754	79	TVPRVFIGKDCIGGCSDLVSL
P35754	83	VFIGKDCIGG <mark>C</mark> SDLVSLQQSG
P38646	487	DGQTQVEIKV <mark>C</mark> QGEREMAGDN
P38646	608	EFKDQLPADECNKLKEEISKM
P50570	27	QDAFSSIGQSCHLDLPQIAVV
P50570	86	SKTEHAEFLHCKSKKFTDFDE
P50570	607	YKDLRQIELACDSQEDVDSWK
P50995	294	AIKGVGTDEACLIEILASRSN
P50995	384	TDESKFNAVLCSRSRAHLVAV
Q12879	87	TDPKSLITHVCDLMSGARIHG
Q12879	320	FSYIPEAKAS <mark>C</mark> YGQMERPEVP
Q12879	399	VWPRYKSFSDCEPDDNHLSIV
000429	367	TSELCGGARICYIFHETFGRT
000429	644	RKLSAREQRDCEVIERLIKSY
Q06830	52	FFYPLDFTFVCPTEIIAFSDR
Q06830	173	FQFTDKHGEV <mark>C</mark> PAGWKPGSDT

043175	281	ALVDHENVISCPHLGASTKEA
043175	369	QGTSLKNAGNCLSPAVIVGLL
P36578	96	$\mathtt{RSGQGAFGNM}^{C}\mathtt{RGGRMFAPTK}$
P36578	208	NRRRIQRRGPCIIYNEDNGII
P36578	250	LAPGGHVGRFCIWTESAFRKL
095573	450	LLGGNIRLLLCGGAPLSATTQ
095573	561	FEDENGQRWLCTGDIGEFEPD
095573	573	GDIGEFEPDGCLKIIDRKKDL
P09382	43	LNLGKDSNNLCLHFNPRFNAH
P09382	61	NAHGDANTIVCNSKDGGAWGT
P30050	17	PNEIKVVYLRCTGGEVGATSA
P30050	141	EILGTAQSVGCNVDGRHPHDI
P30050	162	IDDINSGAVECPASXXXXXX
P46782	155	RRVNQAIWLLCTGAREAAFRN
P46782	172	AFRNIKTIAE <mark>C</mark> LADELINAAK
P61247	139	TTDGYLLRLFCVGFTKKRNNQ
P61247	201	DSIGKDIEKACQSIYPLHDVF
P62888	52	KAKLVILANNCPALRKSEIEY
P62888	92	GTACGKYYRVCTLAIIDPGDS
P63167	24	SEEMQQDSVECATQALEKYNI
P63167	56	FDKKYNPTWHCIVGRNFGSYV
P62879	25	RNQIRDARKACGDSTLTQITA
P62879	204	PDGRTFVSGACDASIKLWDVR
P62879	317	VLAGHDNRVSCLGVTDDGMAV
P61081	47	DINELNLPKTCDISFSDPDDL
P61081	65	DDLLNFKLVICPDEGFYKSGK
P11142	574	DEDKQKILDKCNEIINWLDKN
P11142	603	EHQQKELEKVCNPIITKLYQS
P52272	114	LMDAEGKSRGCAVVEFKMEES
P52272	676	WKMLKDKFNECGHVLYADIKM
P52272	694	IKMENGKSKGCGVVKFESPEV
075369	604	IEYNDQNDGSCDVKYWPKEPG
075369	660 1326	AYGPGLEKSGCIVNNLAEFTV
075369	176	SPFKVAVTEGCQPSRVQAQGP
Q92945 Q92945	436	QINKIQQDSGCKVQISPDSGG
P23229	86	EMTFSIPTHKCGLVIGRGGEN RANRTGGLYSCDITARGPCTR
P23229 P23229	131	SOGPGGKVVTCAHRYEKROHV
P23229 P23229	541	PSGICLQVKSCFEYTANPAGY
Q09666	1833	EAEVPDVDLECPDAKLKGPKF
Q09666	1900	GVEVPDVELECPDAKLKGPKF
Q09666	2806	DVSGPKVDVECPDVNIEGPEG
000170	208	AAAKYYDAIA <mark>C</mark> LKNLQMKEQP
000170	238	OOITPLLLNYCOCKLVVEEYY
000170	240	ITPLLLNYCQCKLVVEEYYEV
Q96EK6	113	LIIEHKFIHSCAKRGRVEDVV
Q96EK6	128	RVEDVVVSDECRGKQLGKLLL
Q96EK6	157	KLNCYKITLECLPQNVGFYKK
Q96AG4	48	LPKATILDLSCNKLTTLPSDF
Q96AG4	59	NKLTTLPSDFCGLTHLVKLDL
Q96AG4	131	DPVLAKVAGDCLDEKQCKQCA
Q09161	73	NYKSKILRLLCTVARLLPEKL
Q09161	477	DIVPPTFSALCPANPTCIYKY
Q09161	483	FSALCPANPTCIYKYGDESSN
Q15366	54	GARINISEGNCPERIITLAGP
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Q15366	109	TLRLVVPASQ <mark>C</mark> GSLIGKGGCK
Q15366	158	IAGIPQSIIE <mark>C</mark> VKQICVVMLE
P20618	82	YKLTDKTVIGCSGFHGDCLTL
P20618	89	VIGCSGFHGDCLTLTKIIEAR
P20618	224	DVYTGDALRI <mark>C</mark> IVTKEGIREE
043765	129	ELNPANAVYF C NRAAAYSKLG
043765	148	LGNYAGAVQD <mark>C</mark> ERAICIDPAY
043765	153	GAVQDCERAI <mark>C</mark> IDPAYSKAYG
P07814	92	LEFSATKLSSCDSFTSTINEL
P07814	105	FTSTINELNHCLSLRTYLVGN
P07814	856	PKAKINEAVE <mark>C</mark> LLSLKAQYKE
P50990	148	KAHEILPNLVCCSAKNLRDID
P50990	149	AHEILPNLVCCSAKNLRDIDE
P50990	430	AKQITSYGET <mark>C</mark> PGLEQYAIKK
Q15185	40	NFEKSKLTFS <mark>C</mark> LGGSDNFKHL
Q15185	75	KHKRTDRSIL <mark>C</mark> CLRKGESGQS
Q15185	76	HKRTDRSILC <mark>C</mark> LRKGESGQSW
P55060	344	FEDQNTLTSICEKVIVPNMEF
P55060	387	SDIDTRRRAACDLVRGLCKFF
P55060	939	AQSLHKLSTACPGRVPSMVST
P27695	65	PSGKPATLKICSWNVDGLRAW
P27695	93	WVKEEAPDILCLQETKCSENK
P27695	310	IRSKALGSDHCPITLYLALXX
014920	179	YAKELDQGSLCTSFVGTLQYL
014983	344	RSLPSVETLGCTSVICSDKTG
014983	349	VETLGCTSVICSDKTGTLTTN
015519	254	YKMKSKPLGICLIIDCIGNET
015519	259	KPLGICLIIDCIGNETELLRD
094760	222	LTVPDDIAANCIYLNIPNKGH
094760	274	ELEKVDGLLTCCSVLINKKVD
P00403	196	ATRPGVYYGQCSEICGANHSF
P00403	200	GVYYGQCSEICGANHSFMPIV
P05089	168	DVPGFSWVTPCISAKDIVYIG
P05089	303	VNTAVAITLACFGLAREGNHK
P08670	328	EYRRQVQSLTCEVDALKGTNE
P10415 P10415	158	VAFFEFGGVMCVESVNREMSP
	229	TLLSLALVGACITLGAYLGHK
P19838	61	KQRGFRFRYVCEGPSHGGLPG
P22460	331	ADPFFIVETTCVIWFTFELLV
P22460 P35247	346 35	TFELLVRFFACPSKAGFSRNI TYSHRTMPSACTLVMCSSVES
P35247 P35247	40	TMPSACTLVMCSSVES
P48039	127	NITGIAINRYCYICHSLKYDK
P48039 P48039	130	GIAINRYCYICHSLKYDKLYS
P68104	234	SGTTLLEALDCILPPTRPTDK
P68104	411	IVDMVPGKPMCVESFSDYPPL
Q05586	744	AVLEFEASQKCDLVTTGELFF
Q05586	798	LDKTWVRYOECDSRSNAPATL
Q16665	520	SPEPNSPSEYCFYVDSDMVNE
Q16665	800	SGLPQLTSYDCEVNAPIQGSR
P02730	201	QHSSLETQLFCEQGDGGTEGH
P02730 P02730	317	LLHSLEGFLDCSLVLPPTDAP
P02730 P07355	133	TDEDSLIEIICSRTNQELQEI
Q92769	262	MYQPSAVVLQCGADSLSGDRL
Q92769 Q92769	274	ADSLSGDRLGCFNLTVKGHAK
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Q99836	216	SIASELIEKRCRRMVVVVSDD
Q9UL62	553	ETRAIDEPNNCKGIRCEKQNN
Q9UL62	558	DEPNNCKGIRCEKQNNAFSTL
Q9UQM7	6	XXXXXMATITCTRFTEEYQLF
Q9UQM7	280	WISHRSTVASCMHRQETVDCL
Q9Y272	11	MKLAAMIKKMCPSDSELSIPA
P30046	24	RVPAGLEKRLCAAAASILGKP
P31949	13	KISSPTETERCIESLIAVFQK
P43243	806	IPKTGFYCKLCSLFYTNEEVA
P45880	47	LVKLDVKTKSCSGVEFSTSGS
P45880	76	TGTLETKYKWCEYGLTFTEKW
Q9BUJ2	377	ALYPHVLVKNCAVEFNFGQRA
Q9BUJ2	487	WDVLIQQATQCLNRLIQIAAR
P62070	55	PTIEDSYTKQCVIDDRAARLD
P62070	183	RVIRKFQEQE <mark>C</mark> PPSPEPTRKE
P62701	41	PSTGPHKLRECLPLIIFLRNR
P62701	181	DFIKFDTGNLCMVTGGANLGR
Q9NQC3	1101	YSNSALGHVNCTIKELRRLFL
075390	211	DSMDLIAKLPCVAAKIYRNLY
P04083	324	FYQKMYGISLCQAILDETKGD
P04083	343	GDYEKILVALCGGNXXXXXXX
P46777	76	YARIEGDMIVCAAYAHELPKY
P46777	100	VGLTNYAAAYCTGLLLARRLL
043707	499	HNVNTRCQKICDQWDALGSLT
043707	793	GALGPEEFKACLISLGYDVEN
Q99439	61	QKGLKDGTILCTLMNKLQPGS
Q99439	164	FDDATMKAGQCVIGLQMGTNK
P04843	477	PAAEARMKVACITEQVLTLVN
Q9P2E9	892	KRLDEVSRELCHTQSSHASLR
Q9P2E9	1323	AEFEEAQTSACRLQEELEKLR
P04406	152	NSLKIISNASCTTNCLAPLAK
P04406	247	TANVSVVDLTCRLEKPAKYDD
P20810	408	AAPAPVSEAVCRTSMCSIQSA
P20810 P13489	661 38	IDALSGDLDSCPSTTETSQNT
P13489 P13489	248	QQCQVVRLDDCGLTEARCKDI KLGDVGMAELCPGLLHPSSRL
P13469 P22314	234	VTKDNPGVVTCLDEARHGFES
P22314 P22314	632	ODPPEKSIPICTLKNFPNAIE
Q08211	12	GDVKNFLYAWCGKRKMTPSYE
Q08211 Q08211	1029	NALIHKSSVNCPFSSQDMKYP
P62258	97	OMVETELKLICCDILDVLDKH
P62258	98	MVETELKLICCDILDVLDKHL
P63104	94	EKIETELRDICNDVLSLLEKF
P63104	189	YEILNSPEKACSLAKTAFDEA
P50579	436	SKYLMALKNLCDLGIVDPYPP
P50579	448	LGIVDPYPPLCDIKGSYTAOF
015382	342	VREVFGSGTACQVCPVHRILY
015382	345	VFGSGTACQVCPVHRILYKDR
Q86VP6	571	ATPYIKDLFTCTIKRLKAADI
Q86VP6	1134	FLMLVRLSTLCPSAVLQRLDR
075828	226	RKADRILVNACCPGPVKTDMD
075828	227	KADRILVNACCPGPVKTDMDG
Q16543	308	YESLPEELQKCFDVKDVQMLQ
Q16543	336	PTDAKYHMQRCIDSGLWVPNS
Q92879	61	RSQNPPQSKGCCFVTFYTRKA

Q92879	62	SQNPPQSKGCCFVTFYTRKAA
Q8TCG1	92	QLAVDIETRDCLQNTYNLNSV
Q8TCG1	108	NLNSVLAGVVCRSSHTDSVFL
Q9UBF2	296	APAVSVLQLFCSSPKPALRYA
Q9UBF2	325	AMKHPSAVTACNLDLENLITD
Q9H9Q2	110	TIVSLASRMKCIPYSVLLKDL
Q9H9Q2	240	EMEQQLAERE C PPHAEQRQPT
P17844	200	YCRACRLKST <mark>C</mark> IYGGAPKGPQ
P17844	234	TPGRLIDFLECGKTNLRRTTY
Q9NR33	84	FVETIAKDAY <mark>C</mark> CAQQGKRKTL
Q9NR33	85	VETIAKDAYC <mark>C</mark> AQQGKRKTLQ
P43897	71	FVNCKKALETCGGDLKQAEIW
P43897	240	VLGKYGALVICETSEQKTNLE
P30042	176	FHQAGKPIGLCCIAPVLAAKV
P30042	177	HQAGKPIGLCCIAPVLAAKVL
P10768	176	YKSVSAFAPICNPVLCPWGKK
P10768	181	AFAPICNPVLCPWGKKAFSGY
P21266	39	DTSYEEKRYTCGEAPDYDRSQ
P21266	208	IAAYLQSDQFCKMPINNKMAQ
P49915	449	PGPGLAIRVICAEEPYICKDF
P49915	456	RVICAEEPYICKDFPETNNIL
094927	349	RQVLILGLRRCCLWTELKALH
094927	350	QVLILGLRRCCCLWTELKALHD
Q14103	126	YFSKFGEVVDCTLKLDPITGR
Q14103	226	DNKTNKRRGFCFITFKEEEPV
P61978	184	TQTTIKLFQECCPHSTDRVVL
P61978	185	QTTIKLFQECCPHSTDRVVLI
Q8TEX9	42	LRAPAALPALCDLLASAADPQ
Q8TEX9	708	VFEEVFKLLECPHLNVRKAAH
P40926	212	GKTIIPLISQCTPKVDFPQDQ
P40926 P28482	285 65	CSFVKSQETECTYFSTPLLLG KISPFEHQTYCQRTLREIKIL
P28482	166	LLNTTCDLKICDFGLARVADP
P54886	606	KVTRLVRDSKCEYPAACNALE
P54886	612	RDSKCEYPAACNALETLLIHR
P12004	81	NLTSMSKILKCAGNEDIITLR
P12004	162	SHIGDAVVISCAKDGVKFSAS
029RF7	1079	SKTNEKLYTVCDVALCVINSK
Q29RF7	1073	KLYTVCDVALCVINSKSALCN
Q9Y3B4	74	YEDIFDAKNACDHLSGFNVCN
Q913B4 Q9Y3B4	83	ACDHLSGFNVCNRYLVVLYYN
Q9NPH0	175	NIFRNLESTRCLLAGLFQCQK
Q9NPH0	183	TRCLLAGLFQCQKEGPIIIHT
008752	275	RAKLOPIALS <mark>C</mark> VLNIGACKLK
Q08752	282	ALSCVLNIGACKLKMSNWQGA
P60900	154	DEEQGPQVYK <mark>C</mark> DPAGYYCGFK
P60900	161	VYKCDPAGYY <mark>C</mark> GFKATAAGVK
000231	202	SARTTANAIY <mark>C</mark> PPKLQATLDM
000231	289	YAGRQTEALK <mark>C</mark> VAQASKNRSL
P55036	37	QAQQDAVNIV <mark>C</mark> HSKTRSNPEN
P55036	87	TVQPKGKITFCTGIRVAHLAL
P26599	250	SLDGQNIYNACCTLRIDFSKL
P26599	251	LDGQNIYNACCTLRIDFSKLT
Q06203	339	TPAALAYAGK <mark>C</mark> GLPYVEVLCK
Q06203	348	KCGLPYVEVLCKNRYVGRTFI

Q92600	91	TLTAHQSNRVCNALALLQCVA
Q92600	99	RVCNALALLQCVASHPETRSA
P27635	71	SSEALEAARI <mark>C</mark> ANKYMVKSCG
P27635	195	VAEKRLIPDGCGVKYIPNRGP
P62913	72	IRRNEKIAVH <mark>C</mark> TVRGAKAEEI
P62913	150	SIADKKRRTGCIGAKHRISKE
P50914	42	DQNRALVDGPCTQVRRQAMPF
P50914	54	QVRRQAMPFK <mark>C</mark> MQLTDFILKF
Q8NHW5	27	IIQLLDDYPK <mark>C</mark> FIVGADNVGS
Q8NHW5	119	AAARAGAIAP <mark>C</mark> EVTVPAQNTG
Q8N0X7	499	KVSQFLVDGVCTVANCVGKEL
Q8N0X7	504	LVDGVCTVANCVGKELAPHVK
P49458	39	LKYRHSDGNLCVKVTDDLVCL
P49458	48	LCVKVTDDLVCLVYKTDQAQD
P31948	26	SVGNIDDALQCYSEAIKLDPH
P31948	62	GDYQKAYEDGCKTVDLKPDWG
Q15527	29	RLQTDARKVRCILTGHELPCR
Q15527	38	RCILTGHELPCRLPELQVYTR
P41250	466	IEIVGCADRSCYDLSCHARAT
P41250	471	CADRSCYDLSCHARATKVPLV
P54577	250	DVKKKLKKAFCEPGNVENNGV
P54577	501	LQADFKISEECIAQWKQTNFM
P48643	181	TTLGSKVVNSCHRQMAEIAVN
P48643	407	EAKRSLHDALCVIRNLIRDNR
P49368	173	KAISRWSSLACNIALDAVKMV
P49368	213	KIPGGIIEDSCVLRGVMINKD
095801	367	VKALTPAFLVCVGSSPFCKNF
095801	374	FLVCVGSSPFCKNFLRGRKVY
P09936	152	AHDAVAQEGQCRVDDKVNFHF
P09936	220	GEVRFSAVALCKAAXXXXXXX
P21283 P21283	225 376	VLSEDQDSYLCNVTLFRKAVD
000220	336	YYPYVYYKIDCNLLEFKXXXX
P00750	118	VTVQSPGEAQ <mark>C</mark> LLGPAEAEGS COCPEGFAGK <mark>C</mark> CEIDTRATCY
P00750 P00966	132	GNDQVRFELSCYSLAPQIKVI
P01009	256	KRLGMFNIQHCKKLSSWVLLM
P01009 P01111	118	DVPMVLVGNKCDLPTRTVDTK
P02766	10	XMASHRLLLLCLAGLVFVSEA
P04271	85	MAFVAMVTTACHEFFEHEXXX
P05062	269	RTVPAAVPGICFLSGGMSEED
P05109	42	DDLKKLLETECPOYIRKKGAD
P05976	181	LMAGQEDSNGCINYEAFVKHI
P10242	130	KHLKGRIGKOCRERWHNHLNP
P10620	50	TRKVFANPEDCVAFGKGENAK
P11926	360	YYSSSIWGPTCDGLDRIVERC
P14780	99	ATLKAMRTPRCGVPDLGRFOT
P20337	184	OAFERLVDAICDKMSDSLDTD
P23284	202	KPLKDVIIADCGKIEVEKPFA
P23975	351	RDALLTSSINCITSFVSGFAI
P27816	635	PETVTGTGKKCSLPAEEDSVL
P29350	453	LPHAGPIIVHCSAGIGRTGTI
P30556	289	IVDTAMPITICIAYFNNCLNP
P35557	371	TTDCDIVRRACESVSTRAAHM
P40227	406	RAVKNAIDDGCVVPGAGAVEV
P42574	163	GKPKLFIIQACRGTELDCGIE
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P43235	139	PVKNQGQCGS <mark>C</mark> WAFSSVGALE
P45983	116	IVMELMDANL <mark>C</mark> QVIQMELDHE
P47712	151	CPDLRFSMALCDQEKTFRQQR
P48200	178	CRGQTTCRGSCDSGELGRNSG
P51787	445	EKMLTVPHITCDPPEERRLDH
P68036	86	HPNIDEKGQVCLPVISAENWK
P69905	105	DPVNFKLLSHCLLVTLAAHLP
Q00987	77	YDEKQQHIVYCSNDLLGDLFG
Q04206	38	KQRGMRFRYKCEGRSAGSIPG
Q16623	145	NATQSDYRER <mark>C</mark> KGRIQRQLEI
P18206	85	PPAFIKVENA <mark>C</mark> TKLVQAAQML
P53621	245	NESKAWEVDTCRGHYNNVSCA
Q8WX93	964	TANQEYKVSS <mark>C</mark> EQRLISEIEY
Q99683	869	GKAADIWSLG <mark>C</mark> TIIEMATGKP
Q9GZT4	113	YIVVPQTAPD <mark>C</mark> KKLAIQAYGA
Q9Y696	234	AYSRDEFTNTCPSDKEVEIAY
P23297	86	VVLVAALTVA <mark>C</mark> NNFFWENSXX
P14854	54	MTAKGGDISVCEWYQRVYQSL
P26641	194	FPNTNRWFLTCINQPQFRAVL
P30048	229	FQYVETHGEVCPANWTPDSPT
P31327	600	YALGGLGSGICPNRETLMDLS
P41567	69	LVKAFKKKFACNGTVIEHPEY
P62241	100	LVRTKTLVKNCIVLIDSTPYR
P62280	131	QIGDIVTVGECRPLSKTVRFN
Q8NBI5	226	PLPPNYSYGLCPGNGTTKEEK
P61163	34	GFAGDQIPKYCFPNYVGRPKH
Q8NBU5	137	KGVLLYGPPGCGKTLIAKATA
Q13557	373	NGDFEAYTKICDPGLTAFEPE
Q9NX47	46	RGSTKWVHQACLQRWVDEKQR
P84095	157	KQIHAVRYLECSALQQDGVKE
014979	177	YLSRFGEVVDCTIKTDPVTGR
043390	292	PDDKKKNRGFCFLEYEDHKSA
P30101	244	KFIQENIFGICPHMTEDNKDL
A6NMY6	133	TDEDSLIEIICSRTNQELQEI
015355	13	AYLSQPNTVKCSGDGVGAPRL
P67936	154	EERAEVSELKCGDLEEELKNV
000299	191	LPKLHIVQVVCKKYRGFTIPE
P23396 P30044	97 100	YAEKVATRGLCAIAQAESLRY FGVPGAFTPGCSKTHLPGFVE
P49411	290	ERGILKKGDECELLGHSKNIR
060443	156	QVLEGRNEVLCVLTQKITTMQ
Q9Y277	65	SGNLETKYKVCNYGLTFTQKW
P78347	475	KLRKMVDOLFCKKFAEALGST
007065	100	AAAAASSSASCSRRLGRALNF
Q8TAQ2	145	NCLSRPNIFLCPEIEPKLLGK
Q14498	478	NSAOGNVYVKCPSIAAAIAAV
P16615	560	EWGSGSDTLRCLALATHDNPL
Q14315	1103	GGLGLTVEGPCEAKIECQDNG
P09651	43	HFEQWGTLTDCVVMRDPNTKR
P41091	236	AQLKYNIEVVCEYIVKKIPVP
P42704	930	SARLQWFCDRCVANNQVETLE
P30153	390	VRLNIISNLDCVNEVIGIRQL
P17174	46	LGVGAYRTDDCHPWVLPVVKK
P00505	187	QGYRYYDPKTCGFDFTGAVED
P53396	229	AKVDATADYICKVKWGDIEFP

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Q99798	592	LQILIKVKGKCTTDHISAAGP
P55263	160	SLIANLAAANCYKKEKHLDLE
P05141	160	AEREFRGLGD <mark>C</mark> LVKIYKSDGI
000468	405	CPEPCRFNAVCLSRRGRPRCS
095433	207	ARPVGVKIPT <mark>C</mark> KITLKETFLT
Q12904	161	DVSRLDLRIG <mark>C</mark> IITARKHPDA
Q13155	143	LSLLVLHRLL <mark>C</mark> EHFRVLSTVH
P14550	134	PFPKNADGTICYDSTHYKETW
Q01433	230	QRVTISGEEK <mark>C</mark> GVPFTDLLDA
P39687	123	ENLKSLDLFNCEVTNLNDYRE
Q99873	262	VEDLTFTSPFCLQVKRNDYVH
014744	22	SRVSSGRDLNCVPEIADTLGA
Q9NVM4	262	FSKQVSSSAACHSRRFEPLTS
P08133	114	AISGIGTDEKCLIEILASRTN
P20073	298	TNQEIREIVRCYQSEFGRDLE
P36405	118	AELLEEEKLSCVPVLIFANKQ
Q9H0F7	148	VTSVKVSQLLCLENIKDKPWH
Q6NXE6	484	VAKAALRDLGCHVELRELWTG
P17405	120	ARVGSVAIKLCNLLKIAPPAV
Q99700	301	EGVFKTYSPKCDLVLDAAHEK
095816	162	DQKFQSIVIGCALEDQKKIKR
095861	243	ASAYVFASPGCKKWDTCAPEV
Q01518	93	ERALLVTASQCQQPAENKLSD
Q14444	226	DLLEGKEKPVCGTTYKVLKEI
P42772	74	LLLLHGAEPNCADPATLTRPV
Q4G0N4	393	NRVFSSSRQRCFSSKVCVRSR
Q9P1F3	39	KFGVLFRDDKCANLFEALVGT
Q6P1X6	98	YEAAFPFLSPCGRERNFLRCE
Q9UNE7	199	DDSHVRAQQACIEAKHDKYMA
Q15417	59	QLGLKDGIILCELINKLQPGS
P23528	147	ANCYEEVKDRCTLAEKLGGSA
P53618	248	NPSERARFIRCIYNLLQSSSP
P48444	441	HDSRRNTLEWCLPVIDAKNKS
Q7L5N1 060716	143 394	PSDIHVHKQVCEIIESPLFLK
O96EP5	124	SNAAAYLQHL <mark>C</mark> YRNDKVKTDV KIFVGGIPHN <mark>C</mark> GETELREYFK
P61962	109	RVGETETRLECLLNNNKNSDF
O9BW61	25	NFSRFHADSVCKASNRRPSVY
Q95W01 015523	315	GQQIRDLERGCHLLVATPGRL
P26196	324	AYVTERQKVH <mark>C</mark> LNTLFSRLQI
P60981	147	ANGPEDLNRACIAEKLGGSLI
Q3LXA3	404	ALDRAAGDGDCGTTHSRAARA
043143	190	SLPGPKRGVACTOPRRVAAMS
060610	1227	GPROANRKAGCAVTSLLASEL
Q9UDY4	175	RVSLEEIYSGCTKRMKISRKR
Q99615	175	MDRALEFAPACHRFKILKAEC
Q14181	198	AGNISLKVLGCPEALTGSYKS
Q16555	504	VPRGLYDGPVCEVSVTPKTVT
Q5VYK3	1257	GORTIAALLPCLLDKGMMSTV
Q13144	618	MDSPLDSSRYCALLLPLLKAW
Q13144 Q9BY44	119	LQLYDVKTGTCLKSFIQKKMQ
Q14152	478	ERAIVDAARHCDLQVRIDHTS
Q13347	144	DNNEPYMKIPCNDSKITSAVW
075822	207	KKITNSLTVL <mark>C</mark> SEKQKQEKQS
Q15369	11	MDGEEKTYGGCEGPDAMYVKL

Q9NPA8	40	KELLRAKLIECGWKDQLKAHC
P62495	335	NLDIMRYVLHCQGTEEEKILY
P15170	453	$\mathtt{RPRFVKQDQV}^{ extsf{C}}\mathtt{IARLRTAGTI}$
P13804	53	AATRLGGEVSCLVAGTKCDKV
Q13868	68	GSVERVNKLICVKALKTRYIG
P23610	110	QERDARQRLVCPAAYGEPLQA
Q9NYY8	377	SKVVLDNIHGCPLRIMINILQ
Q7L8L6	670	KAAVPLGGFLCNVADKSGAME
P22087	99	EPHRHEGVFI CRGKEDALVTK
Q9Y3I1	286	RLQLLPESFICKEKLGENVAN
P37268	6	XXXXXMEFVKCLGHPEEFYNL
Q16658	397	VFRGEHGFIGCRKVTGTLDAN
Q9C0B1	104	VSRILIGNPGCTYKYLNTRLF
Q96I24	109	QISRIQAESGCKIQIASESSG
P60520	15	FKEDHSLEHRCVESAKIRAKY
Q92616	1535	AYCAPKQLSSCLPNIVPKLTE
Q9HC38	221	AAAFGRIAFSCPQKELPDLED
Q9H4A6	280	QLLDLDPEVECLKANTNEVLW
P62993	32	GDILKVLNEECDQNWYKAELN
P48507	72	LVREFPDVLECTVSHAVEKIN
P09211	48	TWQEGSLKASCLYGQLPKFQD
Q99714	214	PLLTSLPEKVCNFLASQVPFP
P08397	114	LPPGFTIGAICKRENPHDAVV
Q00839	562	LNTLLQRAPQCLGKFIEIAAR
P00492	106	TVDFIRLKSYCNDQSTGDIKV
P19367	813	ILQQLGLNSTCDDSILVKTVC
Q13907	172	NPDPNEIKSYCYVSKEELKEL
P60842	66	SAIQQRAILPCIKGYDVIAQA
Q9H0C8	190	VISVEKTVKRCLLDTFKHTDE
P12268	140	VFEAKARHGFCGIPITDTGRM
P29218	184	MVLSNMEKLFCIPVHGIRSVG
Q9H2U2	180	VCEIGSKILSCGEVIHVKILG
075153 P54819	753 92	LKDAAAFLLS <mark>C</mark> QIPGLVKDCM LIEKNLETPL <mark>C</mark> KNGFLLDGFP
Q14003	370	TFEFLMRITFCPDKVEFLKSS
P12277	254	GNMKEVFTRFCTGLTQIETLF
P23919	31	TQSRKLVEALCAAGHRAELLR
P05455	232	EMKSLEEKIGCLLKFSGDLDD
Q04760	61	RVLGMTLIQKCDFPIMKFSLY
Q9H9A6	264	KLRFLPEFPSCSLLKELHVGE
095372	56	TIRLPHVKYICPHAPRIPVTL
P23368	441	EEAYTLTEGRCLFASGSPFGP
Q9BQG0	1031	TGPVRPRHQACLLLQKTLSMR
P25205	148	RPKVVRSVHYCPATKKTIERR
P33992	482	AGITTTLNSRCSVLAAANSVF
013257	106	VLERWOFDIECDKTAKDDSAP
P14174	81	AQNRSYSKLL <mark>C</mark> GLLAERLRIS
P45984	163	KPSNIVVKSDCTLKILDFGLA
Q7Z6M4	219	VQQVTKILHSCPSVLREDLGQ
Q9BXJ9	721	CMIRLFNTAVCESKDLSDTVR
Q15274	96	RVAEVRGPAHCLLLGERVALN
060502	896	RILEFYSKLGCFEIAKMEGFP
P55769	30	KKLLDLVQQS <mark>C</mark> NYKQLRKGAN
Q15233	145	RGKQLRVRFACHSASLTVRNL
Q8TAT6	188	GKFVALENIS <mark>C</mark> KIKSGCEGHL
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Q08J23	93	YKSHAKEILHCLKNKYFKELE
075694	974	AFQERLNSYK <mark>C</mark> ITDTLQELVN
P19338	543	FEDAKEALNS <mark>C</mark> NKREIEGRAI
P32322	262	SLLINAVEAS <mark>C</mark> IRTRELQSMA
Q8NC51	11	MPGHLQEGFGCVVTNRFDQLF
Q9Y536	62	CFHRIIPGFMCQGGDFTRHNG
Q4VXU2	339	GGHSKGFGFVCFSSPEEATKA
Q99497	53	PVQCSRDVVI <mark>C</mark> PDASLEDAKK
Q9UHG3	258	WAVEGGNKLVCSGLLQASKSN
Q9Y2S7	143	YYQVLIDARDCPHISQRSQTE
P30086	168	YELRAPVAGT <mark>C</mark> YQAEWDDYVP
P61758	113	TRFLLADNLYCKASVPPTDKM
P18669	153	ADLTEDQLPSCESLKDTIARA
Q9Y237	45	GNAVKVRHIL <mark>C</mark> EKHGKIMEAM
Q9NYY3	26	KMCEQALGKGCGADSKKKRPP
P67775	266	VVTIFSAPNY <mark>C</mark> YRCGNQAAIM
043447	131	GCQFFITCSKCDWLDGKHVVF
Q9Y570	312	FRGLSNLFLSCPIPKLLLLAG
P53041	404	GRSISKRGVSCQFGPDVTKAF
P30041	91	AWSKDINAYNCEEPTEKLPFP
043395	37	EPTVVTAALN <mark>C</mark> VGKGMDKKKA
P62333	193	LARAVASQLDCNFLKVVSSSI
P49721	91	ANFTRRNLADCLRSRTPYHVN
000232	255	KHYRAIYDTP <mark>C</mark> IQAESEKWQQ
Q9UNM6	182	YYKDALRFLGCVDIKDLPVSE
000233	216	TRWAGKGLLGCNIIPLQRXXX
P22102	298	EFNCRFGDPECQVILPLLKSD
P31939	434	AVKYTQSNSV <mark>C</mark> YAKNGQVIGI
P11172	174	QAHGIRLHSVCTLSKMLEILE
P17812	362	VRYHEAWQKL <mark>C</mark> SAHGVLVPGG
P61106	26	IIGDMGVGKS <mark>C</mark> LLHQFTEKKF
P62820	126	NVNKLLVGNK <mark>C</mark> DLTTKKVVDY
P63000	178	VFDEAIRAVL <mark>C</mark> PPPVKKRKRK
P46060	169	GKILAAALTE <mark>C</mark> HRKSSAQGKP
Q86SE5	51	IFSKYGKIVG <mark>C</mark> SVHKGYAFVQ
Q9BQ04	31	LFEQYGKVLE <mark>C</mark> DIIKNYGFVH
P35250	255	FINSENVFKVCDEPHPLLVKE
P40937	238	GKVTEETVYTCTGHPLKSDIA
Q5UIP0	312	DNFALNPDIL <mark>C</mark> SAKRLKLLMQ
Q06587	398	FWKVSRPLELCYAPTKDPKXX
076021	197	SKNLSREINDCIGGTVLNISK
P62829	28	LGLPVGAVIN <mark>C</mark> ADNTGAKNLY
P83731	6	XXXXXMKVELCSFSGYKIYPG
P62910	91	NVKELEVLLMCNKSYCAEIAH
P62987	91	SLRQLAQKYNCDKMICRKCYA
Q96EL3	21	RPVKQVRVQFCPFEKNVESTR
Q5JTH9	317	MLTLLKDLLPCFPEGLVKSCS
Q14684	197	DQNLKFIDPFCKIAAKTKDHT
Q14690	89	SLCEGMRILGCVKEVNELELV
P15880	229	IDDCYTSARGCTATLGNFAKA
P62857	27	VLGRTGSQGQCTQVRVEFMDD
P62753	12	KLNISFPATGCQKLIEVDDER
P82933	330	VDRLGKHDVTCTVSGGGRSAQ
094979	60	LSDPSLDMKSCATFSSSHRYH
Q99961	147	QNFIDPLQNL <mark>C</mark> EKDLKEIQHH

Q9GZT3	48	HFAQFGHVRRCILPFDKETGF
Q7KZF4	440	ETVPAFSERT <mark>C</mark> ATVTIGGINI
095219	172	LLRIASHPILCRDKIFYLFLT
P00441	7	XXXXMATKAV <mark>C</mark> VLKGDGPVQG
Q9UHB9	562	KPLVERFETF <mark>C</mark> LDPSLVTKQA
Q9BXP5	640	YPNEDEMPNRCGIIHVRGPMP
Q05519	79	DSPLPVSSRVCFVKFHDPDSA
015400	28	ISSNIQKITQ <mark>C</mark> SVEIQRTLNQ
Q9Y2Z0	88	NSTAMLRKGI <mark>C</mark> EYHEKNYAAA
P54136	369	DGRKIVFVPGCSIPLTIVKSD
P26639	107	SWKTTPYQIACGISQGLADNT
P52657	98	IKVDKVKIVA <mark>C</mark> DGKNTGSNTT
Q9UI30	33	LRLQATEVRI CPVEFNPNFVA
Q12931	501	RAGTRNIYYLCAPNRHLAEHS
Q15631	225	RGFNKETAAA <mark>C</mark> VEKXXXXXX
Q14166	361	RPGVLLNQFPCENLLTVKDCL
Q9Y6I9	68	GETGRLFTESCSISPKLRSIA
P61088	87	HPNVDKLGRI <mark>C</mark> LDILKDKWSP
Q14139	465	SRLLTFNPTYCALKELNDEER
Q9Y3C8	116	TAKMYRGGKI <mark>C</mark> LTDHFKPLWA
094874	708	THSMLHAPGRCVPQIIAFLNS
Q99536	50	AAAASPPLLR <mark>C</mark> LVLTGFGGYD
Q9NP79	155	ARWKATYIHN <mark>C</mark> LKNGETPQAG
014980	528	VTVIKDLLGL <mark>C</mark> EQKRGKDNKA
Q9BSK1	152	KSQLNDLQKI <mark>C</mark> AGGKPHECSV
P68871	94	GTFATLSELHCDKLHVDPENF

Negative dataset $\,\mathbb{S}^-\,$ contains 810 non-SNO sites and the corresponding peptide sequences

UniProt ID	Site	Sequence (cf. Eq.1 of the paper)
P21333	2378	VHSPSGALEECYVTEIDQDKY
Q6NXE6	201	EADLTCSGIRCVRHACLKHEQ
P34932	13	VVGIDLGFQSCYVAVARAGGI
Q29RF7	971	VKERRAHARQCLLKNISIRRE
Q14166	370	PCENLLTVKDCLASIARRAGG
014980	199	VKSKHLKDSMCNEFSQIFQLC
075369	2431	IEGPSKVKMDCQETPEGYKVM
P48507	46	PSTHSEELHDCIQKTLNEWSS
P17405	431	HIIGHIPPGHCLKSWSWNYYR
P29218	8	XXXMADPWQECMDYAVTLARQ
Q9UL62	808	SFNLGCKKKTCHGPPLIRTMP
P21266	91	LRYIARKHNMCGETEEEKIRV
P21333	1645	RAVPTGDASKCTVTVSIGGHG
P14780	373	SEGRGDGRLWCATTSNFDSDK
Q86VP6	237	SMSTTRTYIQCIAAISRQAGH
P62333	228	MFNYARDHQPCIIFMDEIDAI
Q08211	415	${ t SVVIIRGATGCGKTTQVPQFI}$
P17812	30	ASSVGTILKSCGLHVTSIKID
P12814	774	GTLGPEEFKACLISLGYDIGN
P17844	89	SKEITVRGHNCPKPVLNFYEA
P36873	172	AAIVDEKIFCCHGGLSPDLQS

P48643	253	VEDAKIAILTCPFEPPKPKTK
P16615	70	LLVRILLLAACISFVLAWFEE
P53396	764	SEVQFGHAGACANQASETAVA
000299	89	KIEEFLEAVLCPPRYPKLAAL
Q09161	44	LICKVGEKSACSLESNLEGLA
P14866	151	LVEFEDVLGACNAVNYAADNQ
Q06587	84	VTALRSGNKECPTCRKKLVSK
P00750	519	CQGDSGGPLVCLNDGRMTLVG
P54577	442	RGVESQGMLLCASIEGINRQV
P67775	50	TKESNVQEVRCPVTVCGDVHG
P16615	349	VETLGCTSVICSDKTGTLTTN
000468	844	RGIVTDGRSGCTPCSCDPQGA
P21817	2237	KMVTSCCRFLCYFCRISRQNQ
P49327	1828	IRDGVVRPLKCTVFHGAQVED
Q9H0C8	367	SAADARYEAACNRLANKAVQR
P12004	62	LRSEGFDTYRCDRNLAMGVNL
P22460	564	RKVSGSRGSFCKAGGTLENAD
Q9NYY8	472	TYSSLNHVYKCQNKEQFVEVM
000468	490	AVKNGQAACECLQACSSLYDP
P63000	157	KEIGAVKYLECSALTQRGLKT
Q9BXJ9	214	GLYREALEHLCTYEKQICDKL
P63000	18	VVGDGAVGKTCLLISYTTNAF
P35557	382	ESVSTRAAHMCSAGLAGVINR
Q9NVM4	343	LYLVAHHDDYCVWYSLQRTSP
Q5UIP0	2298	TESVYPPLVNCVAPVDIILPQ
Q8TEX9	566	GEPMRPLAEECCQLGLCDQ
Q8TEX9	797	LKPPGRLAELCGVLKAVLQRK
P50991	252	EKAKIGLIQFCLSAPKTDMDN
Q9NQC3	464	IKDRSGAYITCAPFNPAATES
Q9H4A6	84	REGYTSFWNDCISSGLRGCML
P78527 094979	232 787	GCLKGLSSLLCNFTKSMEEDP
O5JTH9	344	PNIMQLRDRLCRAQGEPVAGH
Q8TCG1	752	MTLSHVLVTACAMQAFHSLFH EKKNKDLOITCDSLNKOIETV
P48039	146	DKLYSSKNSLCYVLLIWLLTL
Q08211	1099	KLQISHEAAACITGLRAAMEA
Q50Z11 O5JTH9	799	KKAYRVLEEVCASPOGPGALF
Q15149	1119	LEQGAQEESRCQRCISELKDI
P30153	447	EFFDEKLNSLCMAWLVDHVYA
P21817	3216	LEPOLNEYNACSVYTTKSPRE
P06213	268	FYLDGRCVETCPPPYYHFQDW
P21333	1353	SPFQVPVTEGCDPSRVRVHGP
P31939	363	ILSKKKNGNYCVLQMDQSYKP
P23610	190	ALOALGEAASCOLLARDYTGA
P53621	693	LQGNHQIVEMCYQRTKNFDKL
P13489	45	LDDCGLTEARCKDISSALRVN
075694	234	GRIFLAGKDGCLYEVAYQAEA
Q99700	852	DSFIENSSSNCTSGSSKPNSP
P06213	315	VIHNNKCIPECPSGYTMNSSN
P00966	331	VYTGFWHSPECEFVRHCIAKS
Q14690	557	GFIIRVKDYGCIVKFYNNVQG
P29218	218	ADAYYEMGIHCWDVAGAGIIV
Q4G0N4	58	GQGQPRELAGCGSRADGGFRP
060716	450	DQDNKIAIKNCDGVPALVRLL
Q06203	503	GLECFEKSGHCTACLTGKYPV

P16615	447	VGEATETALTCLVEKMNVFDT
014983	318	PEGLPAVITTCLALGTRRMAK
P21817	3650	PLYNLPTHRACNMFLESYKAA
P00750	10	XMDAMKRGLCCVLLLCGAVFV
Q9GZT4	269	IFTVTEDEIKCATQLVWERMK
000468	553	GALCEAETGRCVCPSECVALA
Q06587	64	MLKNTMTTKECLHRFCSDCIV
P21817	2436	YAALIDLLGRCAPEMHLIQAG
P16615	417	YDGLVELATICALCNDSALDY
P55072	209	NEVGYDDIGGCRKQLAQIKEM
Q29RF7	583	LELLISPTCSCKQADICVREI
P21817	3170	DVQVSCYRTLCSIYSLGTTKN
Q7L8L6	233	SMLDVYETKCCHQVWEMNMDQ
Q9UBF2	280	AASAIIHLPNCTARELAPAVS
075694	1358	RFTNLCLDAVCGYLVELQSMS
P55263	182	NWMLVEKARVCYIAGFFLTVS
000468	440	AQDGRTYDSDCWRQQAECRQQ
P21817	4877	SEDEDEPDMKCDDMMTCYLFH
Q7L8L6	345	FVMRKIGDLACANIQHLSSRS
P06213	186	KDDNEECGDICPGTAKGKTNC
Q9BQG0	338	DVIRHYGEHVCTAKLPKQFKF
Q7Z6M4	205	NDTVRLLKEKCLFTVQQVTKI
P45880	103	GTEIAIEDQICQGLKLTFDTT
Q09161	231	KPHPQEEYLDCLWAQIQKLKK
P07237	343	ELTAERITEFCHRFLEGKIKP
000468	795	GLAGCPSACQCNPHGSYGGTC
Q9BSK1	303	TGDKPYKCSDCGRTFYFKSDL
P48200	320	VSLTLPEVVGCELTGSSNPFV
Q14181	554 71	LRYFVKDVLGCVCVNPGRLTK
P20618		FSIHTRDSPKCYKLTDKTVIG
P53618 P04350	888 127	LTPEKALSGYCGFMAANLYAR LDVVRKEAESCDCLQGFQLTH
P31948	403	DAKLYSNRAACYTKLLEFOLA
060502	663	MVKSFVQWLGCRSHSSAQFLI
P21817	65	QNVPPDLAICCFVLEQSLSVR
P35754	23	GKVVVFIKPTCPYCRRAQEIL
P78527	1525	LELAFAFGGLCERLVSLLLNP
P09936	201	DTLLKDAAKVCREFTEREOGE
Q9UHB9	344	ERLFESMLSECRDAIQVVREE
P23229	175	DDMDGGDWSFCDGRLRGHEKF
P23610	145	LGOPAAAAALCLELAAALRDL
043143	114	GHAGHTSLPQCINPFTNLPHT
P06213	35	AAGHLYPGEVCPGMDIRNNLT
013144	247	GVEVRYDLLDCHISICSPOVA
043175	295	GASTKEAQSRCGEEIAVQFVD
Q9BUJ2	532	EGFQRKAIVICPTDEDLKDRT
Q9BXJ9	238	ETKGELLLQLCRLEDAADVYR
P31939	325	FGDFVALSDVCDVPTAKIISR
P30044	204	NVEPDGTGLTCSLAPNIISQL
Q12879	745	LNYKAGRDEGCKLVTIGSGYI
P35579	931	EARVEEEEERCOHLOAEKKKM
Q99536	209	GGVGMAAVQLCRTVENVTVFG
P23368	120	IVYTPTVGLACSQYGHIFRRP
000468	296	GSDGADYPGECQLLRRACARQ
Q14139	1002	FLDPIMSTLMCDPVVLPSSRV

99NYY8 506 ENLLDAYSFCLMNYFPLAPF 004760 19 GGLTDERALSCCSDADPSTKD 004760 19 GGLTDERALSCCSDADPSTKD 004760 19 GGLTDERALSCCSDADPSTKD 008339 450 AGRPLFPHVLCHNCAVEFNFG 61962 336 PDWIACYNNCLEILRVXXXX 278527 3912 SHFASSHALICISHWILGIGD 013144 95 ATGVOETFVCCWKAAQIKEH 092600 252 NPRAREALROCLPDOLKDTF 072816 60 CHSAKVKNISCTSTSSTRILT P42704 361 EDVALQILLACPVSKEDGPSV 075528 4 XXXXXXMSSCSRVALVTGAN 0826233 170 QRVGILPROCALYSPETOK 014744 278 GTNHHSSKETCSYLQYLEYLS 932557 213 VNDTWATMISCYYEDHQCEV 090468 1553 SGVGECGDHPCLPNPCHGAP 0914181 375 AVINHDRPOVCILLFOPPLDAR 113489 313 LLCETILLEPGCLESHWYSC 094287 4 XXXXXXMACATRRALAVGS 0141003 6 XXXXXXMACACTRRALAVGS			
Q00839	Q9NYY8	506	ENLLDAVYSFCLMNYFPLAPF
Q00839 450 AGRELPHYLICHNCAVEPNEG P61962 336 PDWIAICYNNCLEILRVXXXX P78527 3912 SHFASSHALICISHWILGIGD Q13144 95 ATGVQETFVPCCKKAAQIKEH Q92600 252 NPRAREALRCLEPOLKDTFF Q718L6 60 CHSAKKVKNICSTFSSRRILT P42704 361 EDVALQILLACPVSKEDGPSV 075828 4 XXXXXXXMSSGSRVALVTGAN P62333 170 QRVGIIFPKGCLLYGPPGTGK 014744 278 GTNHHSBKEFCSYLQYLEVLS P35557 213 VNDTVARMISCYYEDBQCEVG 092616 1446 EGALFAFEMLCTHLGKLFEPY 000468 1553 SGVCECGDHPCLPNPCHGGAP 014181 375 AVINHDRPDVCILFGFFLDAK 014303 6 XXXXXXXMLSSCVUSSFRGRQGA 0143175 225 GLIMDNTFAQCKKGVNVONCA 09Y287 4 XXXXXXMLSSCVUSSFRGRQGA 0143175 225 GLIMDNTFAQCKKGVNVVNCA 09WW4 407 TVLKPDSVCLCVSDGSLLSVL P946459 599 FDDAYKSGLSCVVVDDIEWL </td <td>Q7KZF4</td> <td>837</td> <td>LLNVEHLSAGCPHVTLQFADS</td>	Q7KZF4	837	LLNVEHLSAGCPHVTLQFADS
P61962 336	Q04760	19	GGLTDEAALSCCSDADPSTKD
P78527 3912	Q00839	450	AGRPLFPHVLCHNCAVEFNFG
Q13144 95	P61962	336	PDWIAICYNNCLEILRVXXXX
Q72600 Z52	P78527	3912	SHFASSHALICISHWILGIGD
OT.88L6	Q13144	95	ATGVQETFVFCCWKAAQIKEH
P42704 361	Q92600	252	NPRAREALRQCLPDQLKDTTF
075828 4 XXXXXXMSSCSRVALUTGAN P62333 170 QRVGIIPPKGCLLYGPPGTGK 014744 278 GTMHBEKEFCSYLQYLEYLS 935557 213 VNDTVATMISCYYEDHQCEVG 092616 1446 EGALFAFEMLCTHLGKLFEPY 000468 1553 SGVGECGDHPCLPNPCHGGAP 014181 375 AVINHDRPDVCLIFGPFLDAK 913489 313 LLCETLLEPGGQLESLWVKSC 094287 4 XXXXXXXMAACTARRALAVGS 014003 6 XXXXXXXMSSVCVSSFRRRQGA 043175 225 GLINDNTFAQCKKGVRVVNCA 09NVM4 407 TVLKPDSVCLCVSDGSLLSVL P46459 599 FDDAYKSQLSCVVVDDIERLL 08TCGI 307 SKVLELLLAFCSVQLRHMLT 05VYK3 1503 ADEEKSEKEECNLWTEVWGEN 075369 2057 IQTEDLEDGGTCKUSYPTVG P21817 3304 SALPAGAPPPCTAVTSDLNS 016543 54 QKEKEELDAGCRECKRVABC 09NVM4 366 NERVRQMRPVCDCQAHLLWNR 935250 185 DKIIEPIQSRCAVIRYTKLTD	Q7L8L6	60	
P62333	P42704	361	EDVALQILLACPVSKEDGPSV
014744 278 GTNHHSEKEFCSYLQYLEYLS 935557 213 VNDTVATMISCYYEDHQCEVG 092616 1446 EGALFAPEMLCTHRIGKLFEPY 000468 1553 SGVGECGDHPCLPNPCHGGAP 014181 375 AVINHDRPDVCILFGPFLDAK 113489 313 LLCETLLEPGGQLESLWVKSC 09Y2S7 4 XXXXXXXSSVCSSFRGRQA 043175 225 GLINDNTFAQCKKGVRVVNCA 09NVM4 407 TVLKPDSVCLCVSDGSLLSVL P46459 599 FDDAYKSQLSCVVVDDIERLL 08TCG1 307 SKVLELLLAFCSVTQLRHMLT 05VYK3 1503 ADEKSEKEECNLWTEVWQEN 075369 2057 IQTEDLEDGTCKVSYFPTVPG P21817 3304 SALPAGAPPPCTAVTSDHLINS 016543 54 QKEKEELDRGCRECKRKVAEC Q01518 247 PPPPPVSTISCSYESASRSL Q9NVM4 366 NERVRQMRPVCDCQAHLLWNR P35250 185 DKIIEPIQSRCAVLRYMKLT Q8TEX9 810 LKAVLQRKTACQDTDEEEEE P11172 86 YTALPLATVICSTNQIPMLIR <td>075828</td> <td>4</td> <td>XXXXXXMSSCSRVALVTGAN</td>	075828	4	XXXXXXMSSCSRVALVTGAN
P35557 213	P62333		QRVGIIPPKGCLLYGPPGTGK
Q92616 1446 EGALFAFEMLCTMLGKLFEPY Q00468 1553 SGYGECGDHPCLPNPCHGGAP Q14181 375 AVINHDRPDVCLIFGPFLDAK P13489 313 LLCETLLEPGCQLESLWVKSC Q9Y2S7 4 XXXXXXMASACYSSFRGRGA Q14003 6 XXXXXXMLSSVCVSSFRGRGA Q43175 225 GLLNDNTFAQCKKGVRVVNCA Q9NVM4 407 TULKPDSVCLCVSDGSLLSVL P46459 599 FDDAYKSQLSCVVDDIFERLL Q8TCG1 307 SKVLELLLAFCSVTQLRHMLT Q5VYK3 1503 ADEEKSEKEECNLWTEVWQEN O75369 2057 1QTEDLEDGTCKVSVFPTVPG P21817 3304 SALPAGAPPPCTAVTSDHLNS Q16543 54 QKEKEELDRGCRECKKVAEC Q9NVM4 366 NERVRQMRPVCDCQAHLLWNR P35250 185 DKIIEPIQSRCAVLRYTKLTD Q65379 51 SLHSELMCPICLDMLKNTMTT Q8TEX9 810 LKAVLQRKTACQDTDEEEEEE P11172 86 YTALPILATVICSTNQIPMLIR Q99832 326 EEDLKRTMMACGGSIQTSVNA <td>014744</td> <td>278</td> <td>GTNHHSEKEFCSYLQYLEYLS</td>	014744	278	GTNHHSEKEFCSYLQYLEYLS
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P13489 313	000468		SGVGECGDHPCLPNPCHGGAP
Q9Y2S7 4 XXXXXXMAACTARRALAVGS Q14003 6 XXXXXMLSSVCVSSFRGQGA Q43175 225 GLINDNTFAQCKKGVRVNVCA Q9NVM4 407 TVLKPDSVCLCVSDGSLLSVL P46459 599 FDDAYKSQLSCVVVDDIERLL Q8TCG1 307 SKVLELLLAFCSVTQLRHMLT Q5VYK3 1503 ADBEKSEKEECNLWTEVWQEN Q75369 2057 IQTEDLEDGTCKVSYFPTVPG P21817 3304 SALPAGAPPPCTAVTSDHLNS Q16543 54 QKEKEELDRGCRECKRVAEC Q01518 247 PPPPPVSTISCSYESASRSSL Q9NVM4 366 NERVRQMRPVCDCQAHLLWNR P35250 185 DKILEPIQSRCAVLRYTKLTD Q55TH9 512 GRQAHPVMRKCLQSLCDLRLS Q06587 51 SLHSELMCPICLDMLKNTMTT Q8TEX9 810 LKAVLQRKTACQDTDEEEEEE P11172 86 YTALPLATVICSTNQIPMLIR Q99832 326 EEDLKRTMACGGSIQTSVNA P08397 247 VLHDPETLLRCIAERAFLRHL Q7L5N1 266 LIREAYALCHCLPVLSTDKK	Q14181		
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043175 225 GLLNDNTFAQCKKGVRVVNCA Q9NVM4 407 TVLKPDSVCLCVSDGSLLSVL P46459 599 FDDAYKSQLSCVVVDDIERLL Q8TCG1 307 SKVLELLLAFCSVTQLRHMLT Q5VXK3 1503 ADEEKSEKEECNLWTEVWQEN O75369 2057 IQTEDLEDGTCKVSYFFTVPG P21817 3304 SALPAGAPPPCTAVTSDHLNS Q16543 54 QKEKEELDRGCRECKRKVAEC Q01518 247 PPPPPVSTISCSYESASRSL Q9NVM4 366 NERVRQMRPVCDCQAHLLWNR P35250 185 DKIIEPIQSRCAVLRYTKLTD Q5JTH9 512 GRQAHPVMRKCLQSLCDLRLS Q6587 51 SLHSELMCPICLDMLKNTMTT Q8TEX9 810 LKAVLQRKTACQDTDEEEEEE P11172 86 YTALPLATVICSTNQIPMLIR Q99832 326 EEDLKRTMMACGGSIQTSVNA P08397 247 VLHDPETLLRCLAERAFLRHL Q7L5N1 266 ILREAYALCHCLPVLSTDKFK Q9UBF2 732 SCTMKFTVRDCDPNTGYPDED P00750 430 AQESSVVRTVCLPPADLQLPD </td <td></td> <td>4</td> <td>XXXXXXMAACTARRALAVGS</td>		4	XXXXXXMAACTARRALAVGS
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O75369	Q8TCG1	307	SKVLELLLAFCSVTQLRHMLT
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P21333 478 SPYTVTVGQACNPSACRAVGR P21817 3525 KKMLPIGLNMCAPTDQDLITL P55072 695 ADLTEICQRACKLAIRESIES Q06203 306 DQMVYTVRYRCGQQLAIEAPV Q8TEX9 962 ERHDRVRDNICGALARLLMAS Q92616 1076 LCASSSGDDGCAFAEQEEVDV	Q15149	848	QTQWSWMLQLCCCIEAHLKEN
P21817 3525 KKMLPIGLNMCAPTDQDLITL P55072 695 ADLTEICQRACKLAIRESIES Q06203 306 DQMVYTVRYRCGQQLAIEAPV Q8TEX9 962 ERHDRVRDNICGALARLLMAS Q92616 1076 LCASSSGDDGCAFAEQEEVDV			
P55072 695 ADLTEICQRACKLAIRESIES Q06203 306 DQMVYTVRYRCGQQLAIEAPV Q8TEX9 962 ERHDRVRDNICGALARLLMAS Q92616 1076 LCASSSGDDGCAFAEQEEVDV			_
Q06203 306 DQMVYTVRYRCGQQLAIEAPV Q8TEX9 962 ERHDRVRDNICGALARLLMAS Q92616 1076 LCASSSGDDGCAFAEQEEVDV			KKMLPIGLNMCAPTDQDLITL
Q8TEX9 962 ERHDRVRDNICGALARLLMAS Q92616 1076 LCASSSGDDGCAFAEQEEVDV			_
Q92616 1076 LCASSSGDDGCAFAEQEEVDV	Q06203		DQMVYTVRYRCGQQLAIEAPV
		962	ERHDRVRDNICGALARLLMAS
O75369 783 GEGDVSVGIKCDARVLSEDEE			LCASSSGDDGCAFAEQEEVDV
	075369	783	GEGDVSVGIKCDARVLSEDEE

Q9NVM4	317	EMQWRDHWMQCVYFLPQEEPV
015382	147	PSFDKLELLECIRRLIEVDKD
P19367	834	GVVSRRAAQLCGAGMAAVVDK
Q9BY44	80	LHSFDLLKAVCLEFSPKNTVL
P78527	1742	GTPRFNNYVDCMKKFLDALEL
P07814	123	VGNSLSLADLCVWATLKGNAA
P08397	261	$\mathtt{RAFLRHLEGGCSVPVAVHTAM}$
P06213	219	CWTHSHCQKVCPTICKSHGCT
000468	705	DEDSEDGPCVCDFSCQSVPGS
Q9UHB9	247	VEEISPNIRYCAYNIGDQSAI
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Q00987	374	KTIVNDSRESCVEENDDKITQ
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Q9Y570	347	GKFQMQVLPQCGHAVHEDAPD
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Q8N0X7	123	LYPEFPPKDMCEKLPEPQSFS
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P09211	170	LLIHEVLAPGCLDAFPLLSAY
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Q9NYY3	553	TVHYYAELGQCSVFPATDAPE
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P29350	327	NAKTYIASQGCLEATVNDFWQ
P22314	706	VLQRPQTWADCVTWACHHWHT
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P21817	64	AQNVPPDLAICCFVLEQSLSV
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Q5VYK3	1806	KKLEESKQWECLTSECRVLLI
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P50579 O60443	290 417	VKDATNTGIKCAGIDVRLCDV
P21333	417 1912	CCKLQIIPTLCHLLRALSDDG IEGPSKAEISCTDNODGTCSV
05VYK3	383	TLSLOFVHHICITCPEIKIKP
P62987	99	YNCDKMICRKCYARLHPRAVN
P42574	148	KITNFFRGDRCRSLTGKPKLF
P08133	552	SLETRFMTILCTRSYPHLRRV
000468	210	SPCPSVVAPVCGSDASTYSNE
O9UL62	922	AAOSSECPLACSSSLHCASSI
P27348	237	LWTSDSAGEECDAAEGAENXX
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P31939	575	SAADKVVIEACDELGIILAHT
P48507	114	SSTRSAVDMACSVLGVAQLDS
P37268	147	EKYOTVIADICRRMGIGMAEF
P53621	85	IKVWNYKLRRCLFTLLGHLDY
P10242	347	RPHGDSAPVSCLGEHHSTPSL
P78527	1953	AAYNCAISVICCVFNELKFYO
P14780	468	TPQPTAPPTVCPTGPPTVHPS
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Q9NYY8	265	LRVTQERINECDEICLSVLST
P22314	494	LKNFAMIGLGCGEGGEIIVTD
P18206	1053	KRIRTNLLQVCERIPTISTQL
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Q99683	120	SEALQSLREACETVGATLETL
Q14181	346	DFEQSMVLVACGPYTTSDSIT
Q08752	176	EVKGEKPAKLCVIAECGELKE
Q9H9Q2	157	QRNQLLEVDFCIGRDIRKKDI
P14618	165	NILWLDYKNICKVVEVGSKIY
P45984	213	KENVDIWSVGCIMGELVKGCV
Q9BXJ9	711	IDSSHPWLHECMIRLFNTAVC
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P06213	286	QDWRCVNFSFCQDLHHKCKNS
P23229	94	YSCDITARGPCTRIEFDNDAD
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Q9BQG0	109	LLQSFEDLPLCSILQQIQEKY
P16615	404	EVHKDDKPVNCHQYDGLVELA
Q01433	342	VYTRREPDEHCSEVELPYPDL
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P21333	2102	INTEDLEDGTCRVTYCPTEPG
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Q99497	46	AGLAGKDPVQCSRDVVICPDA
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P78527	1399	VQVMAHLPDVCVNLMKALKMS
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P08397	211	RVGQILHPEECMYAVGQGALG
Q9UL62	178	IPRPHQIRCNCVECVSSSEVD
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P00750	444	ADLQLPDWTECELSGYGKHEA
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P19838	161	ETLEARMTEACIRGYNPGLLV
P30048	101	FFYPLDFTFVCPTEIVAFSDK
P34932	310	TMNRGKFLEMCNDLLARVEPP
P43235	221	MYNPTGKAAKCRGYREIPEGN
092616	907	IKNPFLSLAACVMPSRLKALG
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Q5VYK3	1364	SVLGELVPRLCELIRSGVGLG
	,•	

P31327	516	LGMGGQTALNCGVELFKRGVL
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P06213	495	ALKTNGDQASCENELLKFSYI
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000468	494	GQAACECLQACSSLYDPVCGS
Q99798	410	AKQALAHGLKCKSQFTITPGS
Q8WX93	306	GNPTPRVRWFCEGKELHNTPD
P55072	69	LKGKKRREAVCIVLSDDTCSD
P19367	217	VNDTVGTMMTCGYDDQHCEVG
Q12879	1412	RSSLRSTASYCSRDSRGHNDV
P16615	669	ELNPSAQRDACLNARCFARVE
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Q99615	219	NADALYVRGLCLYYEDCIEKA
P23919	163	NGAFQERALRCFHQLMKDTTL
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Q99683	835	TSKRLAGINPCTETFTGTLQY
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P10242	613	PCSSTWEPASCGKMEEQMTSS
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P78347	121	TLRKTVEDYFCFCYGKALGKS
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Q9NPA8	50	CGWKDQLKAHCKEVIKEKGLE
000468	270	ARSADGLTASCLCPATCRGAP
Q9UBF2	88	QSNDQTLRRMCYLTIKEMATI
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P63244	153	QDESHSEWVSCVRFSPNSSNP
P08133	59	SRSNRQRQEVCQSYKSLYGKD
P68104	111	NMITGTSQADCAVLIVAAGVG
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P14174	57	LMAFGGSSEPCALCSLHSIGK
Q9UQM7	64	HQKLEREARICRLLKHPNIVR
Q5VYK3	1638	NVKYKIVAISCAADILKATKE
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P11172	255	RLMQKKETNLCLSADVSLARE
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P29474 368			
P11926			QKPKTNIAIFCETRAEWMIAA
Q5JTH9			
05JTH9 901 FGSNQEEALQCYLVLIYPGLV 099873 93 LDVGSGTGILCMFAAKAGARK 099798 205 TPNGGGLGCICIGVGGADAVD 277635 8 XXXMGRRPARCYRYCKNKPYP 000220 274 IGSGCGGDPKCMDRVCFWRLG 075369 1087 VEGPCEAKIECSDNGGTCSV P13489 323 CQLESLWVRSCSFTAACCSHF P34932 376 LNADEAVTRGCALQCAILSPA 09C0B1 308 LDDLNATHGUCLAGSQPRFS P23368 428 ALSNPTAQABCTAEEAYTLTE P00966 19 LAYSGGLDTSCILVWLKEQGY Q4VXU2 132 FGRILSCKVACDEBGSGFGFF Q9C0B1 456 QNLRREWHARQSRIARTLPA 075369 450 SPFVVQVGEACNENACRASGR P23396 134 RFIMESCAKGEVVVSGKLRG Q13347 76 HVLTGSADNSCRLWDCETGKO Q9P2E9 933 QSSEAEVRSKCEELSGLHGGL Q9P2E9 933 QSSEAEVRSKCEELSGLHGGL Q9P2E9 933 QSSEAEVRSKCEELSGLHGGL Q9H202 274 ALEVIKSTHCWKFSCOLYHARKON	P11926		PPERIIYANPCKQVSQIKYAA
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000220 274 IGSGCGGDPKCMDRVCFWRLG 075369 1087 VEGPCEAKIESDNGDGTCSV 913489 323 CQLESLWWSSCSFTAACCSHF P31327 761 KNVVSGKTSACFEPSLDYMVT P34932 376 LNADEAVTRGCALQCAILSPA Q900B1 308 LDDLNATHQHEVLAGSQPRFS P23368 428 ALSNPTAQAECTAEEAYTLTE P00966 19 LAYSGGLDTSCILVWLKEQGY Q4VXU2 132 FGNILSCKVACDEBGSRGFFF Q9C0B1 456 QNLRREWHARCQSRIARTLPA 075369 450 SPPVVQUEEACNPNACRASGR P23396 134 RFIMESCAKGCEUVYSCKIRG Q13347 76 HULTGSADNSCRLWDCETGKQ Q06587 332 GASDTGGPDGCGGEGGGAGG Q14315 2369 IAPEDRKDGSCGVSYVVQEPC Q9P2E9 933 QSSEAEVRSKCEELSGLHGQL Q9H2U2 274 ALEVIKSTHQCWKALLMKKCN P05062 158 WRAVLRIADQCPSSLAIQENA P63000 81 SYPQTDVFLICFSLVSPASFE Q94874 641 NEKSIEDFISCLDSAAEACDI <td></td> <td></td> <td></td>			
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P13489 323			
P31327	075369		
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Q9C0B1 308 LDDLNATHQHCVLAGSQPRFS P23368 428 ALSNPTAQAECTAEEAYTLTE P00966 19 LAYSGGLDTSCILWALKEGY Q4VXU2 132 FGNILSCKVACDEHGSRGFGF Q9C0B1 456 QNLRREWHARCQSRIARTLPA 075369 450 SPFVVQVGEACNPNACRASGR P23396 134 RFIMESGAKGCEVVVSGKLRG Q13347 76 HVLTGSADNSCRLWDCETGKQ Q06587 332 GASDTGGPDGCGGEGGAGGG Q14315 2369 IAFEDRKDGSCGVSYVVQEPG Q9P2E9 933 QSSEAEVRSKCEELSGLHGQL P91817 1685 LGNNRVAHALCSHVDQAQLLH Q992U2 274 ALEVIKSTHQCWKALLMKKON P05062 158 WRAVLRIADQCPSSLAIQENA P63000 81 SYPQTDVFLICFSLVSPASFE P04874 641 NEKSIEDFISCLDSAAEACDI P06213 280 PPYYHPQDWRCVNFSFCQDLH P78527 3001 KDFWELASLDCYNHLAEWSL 000468 948 GSDGVTYGNECQLKTIACRQ Q9UQ7 289 SCMHRQETVDCLKKFNARKL			
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Q9UHG3 242 NAFVGAVSLSCSDSGLWAVEG P61962 228 DPQHHPLLRLCWNKQDPNYLA P15170 419 GYNAVLHIHTCIEEVEITALI Q14315 1180 VGEAATFTVDCSEAGEAELTI P14618 358 VANAVLDGADCIMLSGETAKG			
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P15170 419 GYNAVLHIHTCIEEVEITALI Q14315 1180 VGEAATFTVDCSEAGEAELTI P14618 358 VANAVLDGADCIMLSGETAKG	~		
Q14315 1180 VGEAATFTVDCSEAGEAELTI P14618 358 VANAVLDGADCIMLSGETAKG			
P14618 358 VANAVLDGADCIMLSGETAKG			
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U/DD94			
	0/3094	00/	IVISGRHNGICIYFSRIMGNI

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P00533	19	AALLALLAALCPASRALEEKK
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Q9BUJ2	421	GTVGPKSKAECEILMMVGLPA
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P78347	123	RKTVEDYFCFCYGKALGKSTV
Q7KZF4	736	GSYAPRRGEFCIAKFVDGEWY
P53041	240	TLKETEKITVCGDTHGQFYDL
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Q9BUJ2	391	FNFGQRAEPYCSVLPGFTFIQ
P13489	75 127	SNELGDVGVHCVLQGLQTPSC
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P08133	248	FEKLMLAVVKCIRSTPEYFAE
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Q92945	296	IGDPYKVQQACEMVMDILRER
Q16658	19	AVQIQFGLINCGNKYLTAEAF
P23368	274	RFLRKYREKYCTFNDDIQGTA
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P55072	415	GHVGADLAALCSEAALOAIRK
P60981	163	GGSLIVAFEGCPVXXXXXXXX
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000170	78	FKLPVWETIVCTMREGEIAQF
P22314	444	EDKEVLTEDKCLQRQNRYDGQ
092616	414	VHAVSVLALWCNRFTMEVPKK
P78527	1919	NELTKTLIKLCYDAFTENMAG
P49915	523	PIKTVGVOGDCRSYSYVCGIS
P49327	135	ETLVGYSMVGCQRAMMANRLS
P21817	5028	RCWDFFPAGDCFRKQYEDQLS
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043447	128	STNGCQFFITCSKCDWLDGKH
P50579	223	LNAGLAFPTGCSLNNCAAHYT
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043175	48	KEELIAELQDCEGLIVRSATK
095372	135	GALSLYTALTCPHPLAGIVAL
P49327	1558	CSSLRHAQPTCPGAQLCTVYY
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P78527	1455	RSRLAAVVSACKQLHRAGLLH
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P17987	125	TSVISGYRLACKEAVRYINEN
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P17405	385	LRLISLNMNFCSRENFWLLIN
Q8TCG1	445	HIAKILTTVKCTTLIEQQFTY
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P37268	43	QDSLSSSLKTCYKYLNQTSRS
P49327	779	AVLKRGLKPSCTIIPLMKKDH
Q9HC38	171	SDLQKSLNYWCNLLGMKIYEK
P15121	187	KYKPAVNQIECHPYLTQEKLI
Q14315	618	IEGPSQAKIECDDKGDGSCDV
P60842	134	GDYMGASCHACIGGTNVRAEV
P27635	105	HVIRINKMLSCAGADRLQTGM
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P30101	57	LMLVEFFAPWCGHCKRLAPEY
P78527	3403	EEEAQPPSWSCGPAAGVIDAY
P63244	207	TVTVSPDGSLCASGGKDGQAM
P43243	803	DYVIPKTGFYCKLCSLFYTNE
P13489	266	SRLRTLWIWECGITAKGCGDL
P23975	44	LVVKERNGVQCLLAPRDGDAQ
P18206	422	LAEARKIAELCDDPKERDDIL
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Q90L62 P00750	30 / 78	YCWCNSGRAOCHSVPVKSCSE
075694	78 1201	EFADPFKLAECKLAIIHCAGY
P40227		
P40227 095801	499 174	AAEVGVWDNYCVKKQLLHSCT LKHFAEAVNWCDEGLQIDAKE
Q29RF7	956	PLEYMAIFALCAKDPVKERRA
Q29RF / P22460	500	TVGGKIVGSLCAIAGVLTIAL
P48039	100	
P48039 P29350	171	NNGWNLGYLHCQVSGFLMGLS
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P60981	152 46	VIFCLSADKKCIIVEEGKEIL
Q16665	337	
P20073	33 <i>1</i> 327	TKNSQPQCIVCVNYVVSGIIQ GHFERLLVSMCQGNRDENQSI
FZUU/3	341	AUL EVITIA SLICÕGIIANEIIĀST

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P21817	1673	SHTLRLYRAVCALGNNRVAHA
P48444	380	EESFIPLTINCWPSESGNGCD
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P21817	4959	EQVKEDMETKCFICGIGSDYF
Q9C0B1	77	EAFLTLHKHGCLFRDLVRIQG
P53618	189	DFLVNEKDASCKRNAFMMLIH
P36873	39	QLQENEIRGLCLKSREIFLSQ
P11926	217	TFVQAISDARCVFDMGAEVGF
075369	178	GKALGALVDSCAPGLCPDWES
P21333	2582	VGQKSSFTVDCSKAGNNMLLV
P30101	409	IEFYAPWCGHCKNLEPKYKEL
Q13907	86	TFPGCFTNTCCSHPLSNPAEL
P50579	380	GKGVVHDDMECSHYMKNFDVG
Q9NP79	276	PEDFARAQKYCKYAGSALQYE
P26641	166	RVTLADITVVCTLLWLYKQVL
P00533	248	AGCTGPRESDCLVCRKFRDEA
015519	298	GISQILGQFACMPEHRDYDSF
P55060	853	AVGITKLLTECPPMMDTEYTK
P78527	4061	LAGANPAVITCDELLLGHEKA
Q8TCG1	358	LSQPLDGSENCSVLALELFKE
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Q15149	965	VEVTVHKGDECOLVGPAOPSH
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000468	488	TCAVKNGOAACECLOACSSLY
Q7Z6M4	57	SNGGVIEELSCVRSNNYVQEP
Q06203	496	MIOENGNGLECFEKSGHCTAC
P68366	200	TTHTTLEHSDCAFMVDNEAIY
P35579	468	IFDLNSFEQLCINYTNEKLQQ
P29350	382	RAYGPYSVTNCGEHDTTEYKL
P06213	331	MNSSNLLCTPCLGPCPKVCHL
Q5UIP0	613	FFLSLESLVGCVLSGPTSPLA
P21817	3240	LGLPNSVEEMCPDIPVLERLM
Q06203	280	SRSEGNPVAFCIFEYVYFARP
P22314	413	GLAAQEVMKACSGKFMPIMQW
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060502	215	YLGEPETFLFCPTEYCGTFCY
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Q00839	295	DTVVCLDTYNCDLHFKISRDR
095801	160	HLKAIIRGALCHLELKHFAEA
Q08211	612	EDDDANCNLICGDEYGPETRL
Q5UIP0	1077	HQKEVLKTKRCDIPAMYNNLD
Q14684	61	LLKIWKGLFYCMWVQDEPLLQ
075390	101	IRFRGFSIPECQKLLPKAKGG
Q9UBF2	446	LAHLCEFIEDCEHTVLATKIL
Q15417	273	LGRQVYDPKYCAAPTEPVIHN
P21817	2305	LEKVVSYLAGCGLQSCPMLVA
P43235	269	YSKGVYYDESCNSDNLNHAVL
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Q9BY44	132	SFIQKKMQNWCPSWSEDETLC
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P32322	120	AFRPAPRVIRCMTNTPVVVRE
P23368	481	FPGVALAVILCNTRHISDSVF
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014980	920	AAQSFYQTYFCDILQHIFSVV
Q16658	456	DTPVDFFFEFCDYNKVAIKVG
015382	249	LVQQEALKRGCEQVLWLYGPD
075369	416	VEDKGNQVYRCVYKPMQPGPH
P55263	140	YEQNEQPTGTCAACITGDNRS
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P53618	143	LEPLMPAIRACLEHRHSYVRR
Q9NVM4	294	IEMDPEGKIKCTMAPFWAHSD
P78527	1791	SFRRIARRGSCVTQVGLLESV
Q00839	648	GNFTLPEVAECFDEITYVELQ
094927	192	PVVLRDVRTACTLRAQFLQNL
P13489	427	QLVESVRQPGCLLEQLVLYDI
O9NPHO	23	PVGVLTSLAYCLHQRRVALAE
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Q7L8L6	425	AASLPPRVAHCRSKDVAKILW
Q5UIP0	1865	PNENFKTVGPCLGDSKNVSQE
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043143	307	AGKFQIYFDNCPLLTIPGRTH
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Q9NYY8	269	QERINECDEICLSVLSTVLEA
014744	196	WMWWHNFRTLCDYSKRIAVAL
P78527	10	XMAGSGAGVRCSLLRLQETLS
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Q16555	334	GDLQVTGSAHCTFNTAQKAVG
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P26599	23	KRGSDELFSTCVTNGPFIMSS
P43235	210	AYPYVGQEESCMYNPTGKAAK
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P49411	387	VMFSLTWDMACRIILPPEKEL
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Q14684	155	FLDVLMKEVLCPESQSPNGVR
Q14003	646	LPPLPAPGEPCPLAQEEVIEI
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015355	351	LIVANAGDSRCVVSEAGKALD
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060502	618	RSRAAKFEEMCGLVMGMFTRL
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000468	877	SCKPGVAGPKCGQCPDGRALG
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Q7Z6M4	183	EGKLKRVLYCCPEIFTMRQQD
P22102	540	EPLFFLDYFSCGKLDLSVTEA
P55060	630	EAICLSIRITCKANPAAVVNF
015355	495	APDTSGDGTGCDNMTCIIICF
Q5UIP0	731	LVATAEENLCCEELSSKIMSS
000231	257	AITSLKYMLLCKIMLNTPEDV
000251	317	ENVFKKFDGPCDPCQGALPDP
000468	414	VCLSRRGRPRCSCDRVTCDGA
Q6NXE6	127	FCDQCKQDKACRFLAAQKGAY
P21817	603	GRNHKVLDVLCSLCVCNGVAV
P21980	211	PKFLKNAGRDCSRRSSPVYVG
P07355	335	GDYQKALLYLCGGDDXXXXXX
000468	197	AEGPGRASCVCKKSPCPSVVA
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043390	99	SHVQNKSAFLCGVMKTYRQRE
P00750	509	GGPQANLHDACQGDSGGPLVC
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Q14139	490	VHMRGLDKETCLIPAVQEPKF
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075369	1617	RATQTGDASKCLATGPGIAST
P53396	893	PKYSCQFIEMCLMVTADHGPA
Q9NYY8	366	NHRSLILLDECSKVVLDNIHG
000232	244	DQHEGSYLSICKHYRAIYDTP
043707	495	YYDSHNVNTRCQKICDQWDAL
Q16543	183	KYLSDNVHLVCEETANYLVIW
Q9BUJ2	331	DKFAENDVIGCFADFECGNDV
P00533	236	KSPSDCCHNQCAAGCTGPRES
P30153	329	ENVIMSQILPCIKELVSDANQ
000468	172	TPVPPTPPDACRGMLCGFGAV
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P13639	131	VTDGALVVVDCVSGVCVQTET
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075369	991	ILSPSRKVVPCLVTPVTGREN
P78527	1032	DSTLRDFCGRCIREFLKWSIK
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075153	333	TAPOAEHAMDCVRAEDAYTSR
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P05062	202	IPDGDHDLEHCOYVTEKVLAA
P37268	374	STIRTQNLPNCQLISRSHYSP
P13010	157	LDIIIHSLKKCDISLQFFLPF
P21817	2555	EMALALNRYLCLAVLPLITKC
000468	664	TQIEEARAGPCEQAECGSGGS
P53618	235	LVIVELIYKVCHANPSERARF
P06213	911	SRKHFALERGCRLRGLSPGNY
Q6P1X6	130	ADHGPPRLSYCGGGEALAVPF
AOLIVO	130	TEMOLI MEDI COCCUMINIVI I

075153	1196	KTKESSEYLKCLTQQAVALQR
P11926	202	VGVSFHVGSGCTDPETFVQAI
P21980	505	ITNNTAEEYVCRLLLCARTVS
000468	923	VEESGSAHCVCPMLTCPEANA
Q14003	185	GIDETDVEACCWMTYRQHRDA
Q00610	1528	NNRWKQSVELCKKDSLYKDAM
P07237	8	XXXMLRRALLCLAVAALVRAD
Q86VP6	286	IQAFESFVRRCPKEVYPHVST
043143	644	FKQNHESVQWCYDNFINYRSL
P62829	125	SAITGPVAKECADLWPRIASN
Q86VP6	71	NGEVQNLAVKCLGPLVSKVKE
Q9GZT4	2	XXXXXXXXMCAQYCISFADV
Q06587	279	HPLLVEKGEYCQTRYVKTTGN
P47712	220	KALYESGILDCATYVAGLSGS
000468	1843	VPREAAYVCLCPGGFSGPHCE
014983	636	GDNKGTAIAICRRIGIFGENE
P06213	235	SHGCTAEGLCCHSECLGNCSQ
P78527	1954	AYNCAISVICCVFNELKFYQG
Q5VYK3	1390	ASVIVSLTTQCPQDLTPYSGK