

Supplementary data 1. RWP-RK protein sequences used for phylogenetics and motif analyses.

>AtNLP1

MEDDGGSDGGEONGGFSPNSSFGAFADTAMDLMDELLFDGCWLETTDSKSLQTEQSPSASTA
MNDNSPFLCFGENDPSQDNFSNEETERMFPQAQKFLLEAEVGKSWWIAPSASEGPSSVKERLLQAI
GLNEAVQDKDFLVQIWVPIQQEGKSFLTTWAQPHLFNQEYSSLAERYRVSETYNFPADEGMKDFVG
LPGRVFLQKFPEWTPDVRFFRDEYPRIKEAQKCDVRGSLALPVFERGSGTCGVVEIVTTQKMNYR
QELEKMKALEAVDLRSSNLNTPSSEFLQVYSDFYCAALPEIKDFLATICRSYDFPLALSWAPCARQG
KVGRHSDENFSECVSTIDSACSVPDEQSKSFWEACSEHHLLQGEGIVGKAFAETKLFFVPEVATFSKT
NYPLAHHAKISGLHAALAVPLKSGLVEFVLEFFFKAQKEMLKSLCVTLQQDFRSSNLFIK
DLELEVLPVRETMFSENLLCGAETVESLTEIQMQUESSWIAHMKANEKGKDVSLSWEYQKEDPKELS
SGRENSQLDPVPNNVPLEAQQLQQASTPGLRVIGPSTESTGGGNMLSSRRPGEKKRAKTEKTIGL
EVLRQYFAGSLKDAAKSIGVCPTTLKRICRQHGIMRWPSRKIKVGHSLKKLQLVMDSVQGAQGSIQ
DSFYTSFELNSPNMSSNGPSLKSNEQPSHLNAQTDNGIMAENPRSPSSCSKSSGSSNNNENTG
NILVAEDADAVLKRAHSEAQLHNVNQEETKCLARTQSHKTFKEPLVLDNSSPLTGSNTSLRARGAI
VKATFGEARIRFTLLPSWGFAELKQEIAARRFNIDDISWFDLKYLDDDKEWVLLTCEADLVECIDIYRLTQT
HTIKISLNEASQVKLSGSFGNTGLS

>AtNLP2

MEGGRGGGDGNFLPNSNFGVFSDSAMMDMFDELLFDGCWLETTDGKSLQTMGQQVSDSTM
NDNNNNNSYLYGYQYAENLSQDHISNEETGRKFPIPPGFLKIEDLSNQVPDFQSAVMSSAQAEKFLLE
ESEGGRRYWIAPRTSQGPSSVKERLVQAI EGLNEEVQDKDFLIQIWLPIQQEGKNFLTSEQPHFFNP
KYSSLKRYRDVSAYNFLADEDSKESVGLPGRVFLKKLPEWTPDVRFFSEYPRIKEAEQCDVRGSLAL
PVFERGSGTCGVVEIVTTQKMNYRPLEDNICKALESVNLRSSRSLNPPSREFLQVYNEFYAAALPEVS
EFLTVCRVYDLPLALTWAPCARQGKVGSRHSDENFSECVSTVDDACIVPDHQSRHFLEACSEHHLL
QGEGIVGKAFNATKLFFVPEVTTFSKTNYPLAHHAKISGLHAALAVPLKNKFNSSEVFVLEFFFKA
DTEAQQDMLKSLATLQQDFRSNLNFIDKELELEVVFVREEVVFAENPLINAGTGEDMKPLPLEEISQ
EDSSWISHMIKANEKGKGVSLSWEYQKEEPKEEMLTSGWDNNQIGSGHNNFLSEAEQFQKV
TNSGLRIDMDPSFESASFGVGQTLLGSRRPGEKRRRTKTEKTIGLEVLRQYFAGSLKDAAKSIGVCPTTLKR
ICRQHGITRWPSSRKIKVGHSLKKLQLVIDSVQGVQGSIQLDSFYTSFELSSPHMSGTGFKNPNAQT
ENGVSAQGTAAAPKSPPSSCSHSSGSTCCSTGANQSTNTGTSNTVTLMAENASAILKRARSE
RLHTMNQDETKSLSRTLHKTSEHPLFENPPRLPENSSRKLKAGGASKV
KATFGEAKVRFTLLPTWGF
RELQHEIARRFNIDNIAPFDLKYLDDDKEWVLLTCEADLEECIDIYRSSQSRTIKISVHEASQVKLGGSFG
SIGLGPSL

>AtNLP3

MVGPFKKILHAHNQRFPSSSSLDPVVDDSSRKQTRIILSYLLHLSLSLHITYSLCLLHFFFSSNPFP
MEESNNSAVVDFPDNFMDQLLWEECWEEATQHDQALSSPSGLKERVACAMGHQEVGERELLI
QLWVPVETRSGRVLSTEEQPYINTFSQSQLALYRDAASAGYSFAAEVGSEQLVGLPGRVFLRRMPE
WTPDVRFFRKEYPRIGYARRYQVRATLALPLFQGTSGNCVAVMEMVTTHRNEYASQLSTICHALEA
FDLRTSQTSIVPASLKVTSSSSSRTEVASILQGICSSHGLPLAVTWGHQDSSSCLASASYAADHG
SRCFLAACSEHHLLGGEGIAGRAFATKKQCFATDVAIFSKWSYPLSHYAKMFDLHAALAVPILTRGN
RTVQFVLELFFPRDCLDIQTHSLTLASQLKRFQSSPHLMVDDNQIAEVRDAATPPLTQEDPKGKQV
SFSFSSASSLENRKRKTAKEDITLDTLRQHFAGSLKDAAKNIGVCPTTLKRICRQNGISRWPSRKKV
GHSLRKLQVVMDSVEGVQGSLHLASFYSSFPQLQSSSSSFPINTQTVHVPPKSPPSSGSQSSGS
STCCSSEEQQLGGFQKPALSHPQLLSSMHEDQRPVRTSSLPLPSATTPRKAKDGMVKAMFGD

SMLRMSLLPHSRLTDLRREIAKRGFMDDVLSNFSLKYLDDDQEWFVLLCDADLEECIQVYKSSLKET
IRILVHHPLSRPSFGS

>AtNLP4

MEDSFLQSENVVMDADFMGDLGCWLETTDGSEFLNIAPSTSSVSPFDPTSMWSPTQDTSALC
TSGVVSQMYGQDCVERSSLDEFQWNKRWWIGPGGGGSVTERLVQAVEHIKDYTTARGSLIQLWV
PVNRGGKRVLTTKEQPFSHDPLCQRLANYREISVNYHFSAEQDDSKALAGLPGRVFLGKLPEWTPDV
RFFKSEYPRVHHAQDCDVRGTLAIPVFEQGSKICLGIEVVMTTEMVCLRPELESICRALQAVDLRSTE
LPIPPSLKGCDLSYKAALPEIRNLLRCACETHKLPLAQTVSCQQQNKGCRHNDENYIHCVSTIDDA
CYVGDPTVREFHEACSEHHLLKGQGVAGQAFLTNGPCFSSDVSNYKKSEYPLSHHANMYGLHAVA
IRLRCIHTGSADFVLEFFLPKDCDDLEEQRKMLNALSTIMAHVPRSLRTVTDKELEEESEVIEREEIVTPKI
ENASELHGNSPWNASLEEIQRSNNTSNPQNLGVFDGGDKPDGFGLKRGFDYTMDSNVNESSTF
SSGGFSMMAEKKRTKADKTITLDVLRQYFAGSLKDAAKNIGVCPTTLKRICRQHGIQRWPSRKIKKVG
HSLQKIQRVIDSVQGVSGPLIGSFYANFPNLVSQSQEPAKTTPPPPPVLAKSPVSSYSHSSN
SSQCCSSETQLNSGATTDPSTDVGGAALKTSSEIELQSSLDETILTSSLENIPQGTNLLSSQDDDFLR
IKVSYGEEKIRLMRNSRRLRDLLWEIGKRFIEDMSRYDLKYLDENEWVLLCDEDVEECVDVCRTT
PSHTIKLLLQASSHHFPERSSATEYSLWH

>AtNLP5

MENNNSLMPMDPAMDSSFMDGLLLEGCWLETTDASEFLNFSPSTSVAFFDPSSFMWSPTQDTSNSLSQ
MYGQDCPERSSLEDQNQGRDLSTFNRRWWIGPSGHGFVSMERLVQAVTHIKDFTSERGSLIQLW
VPVDRGGKRVLTTKEQPFSHDPMCQRLAHYREISENYQFSTEQEDSDSSRDLVGLPGRVFLGVPE
WTPDVRFFKNEEYPRVQHAQDCDVRGTLAIPVFEQGSQICLGIEVVMTTQMVKLSPDLESICRALQ
AVDLRSTEIPPPSLKGPDFSYQAALPEIRNLLRCACETHKLPLAQTVSCLKQSKTGCRHNDENYIHC
VSTIDDACYVGDPTVREFHEACSEHHLLKGQGVVGEAFLTNGPCFSSDVSSYKKSEYPLSHHATMFG
HGTVAIRLRCIHTGSDFVLEFFLPKNCRDIEEQRKMLNALSTIMAHVPRSLRTVTQKELEEGDSMVS
EVIEKGVTLPKIENTTEVHQSIISTPQNVGLVFDGGTEMGELGSEYGKGVSVNENNTSSASGFNRVTE
KKRTKAENKNITLDVLRQYFAGSLKDAAKSIGVCPTTLKRICRQHGIQRWPSRKIKKVGHSLQKIQRVIDS
VEGVSGHHLPIGSFYASFPNLAAASPEASSLQQQSKITFLSYHSPPAKSPGSSCSHSSCSSETQVIKE
DPTDKTRLVRSRFKETQTTHLSPSSQEDDFLRVKSYEEEKIRFKMRNSHRLKDLLWEIAKRFIEDV
SRYDLKYLDENEWVLLRCDDVEECVDVCRSPGQTIKLLLQLSSSYLPERSSVSGCLS

>AtNLP6

MELDDDDLSGSWPLDQITFASNFKSPVIFSSSEQPFSPLWSFSETSGDVGGELYSAAVAPTRFTDYSV
LLASSESETTKENNQVPSPWGIMPLENPDSYCAIKAKMTQALRYFKESTGQQHVLAQVWAVPKVN
RGRYVLTSGQPFVLGPNSNGLNQYRMVSLTYMFSLDGERDGEGLPGRVFRKKLPEWTPNVQYYSS
KEFSRLGHALHYNVQGTLALPVFEPSRQLCVGVVELIMTSPKINYAPEVEVKCKALEAVNLKTSEILNHE
TTQICNEGRQNALAEILEILTVCETYKLPLAQTVVPCRHRSLAFGGFKKSCSSFDGSCMGKVCMS
TSDLAVYVVDAHVGFRDACAEEHHLQKGQGVAGRAFQSGNLFCRDFVTRFCKTDYPLVHYARMF
KLTSFAVCLKSTYTGDDEYVLEFFLPPAITDKSEQDCLLGSSLQTMKQHYSSLKVSETELCENNMSLE
VVEASEDMVYSKLEPIRIHHPAQISKDYLELNAPEQKVSLNSDFMENNEVDDGVERFQTLPIPEAK
TVKKSERKRGKTEKTISLEVQQYFAGSLKDAAKSLGVCPTTMKRICRQHGISRWPSSRKINKVNRSLTRI
KHVIDSVQGADGSNLTSLSRPWPHQIPPIDIQLAKNCPPSTSPLSNLQDVKIERDAEDSAGSSTS
RASCKVNPLICTRFLPTHNQEPESRQVALDDSDSSKNMTNFWAHLTCQDTASPTILQHKLVSIKATY
REDIIRFKISPESVSITELKQQVAKRLKLETAAFELKYLDDREWVSVSCDADLSECLDTAAKANTL
RLSVHDVTFNFGSSCESSEETMMCL

>AtNLP7

MCEPDDNSARNGVTQPSRSRELLMDVDDLDGSWPLDQIPYLSSNRMISPIFVSSSEQPCSP
WAFAFDGGNGFHATSGGDDEKISSVGVPSFRLAEYPLFLPYSSPSAAENTTEKHNSQFPSPLMSL
VPPENTDNYCVIKERMTQALRYFKESTEQHVLAQVWAPVRKNGRDLTTLGQPFVLNPNGNGLNQ
YRMISLTYMFSVDSESDVELGLPGRVFRQLPEWTPNVQYYSSKEFSRLDHALHYNVRGTLALPVNP
SGQSCIGVVELIMTSEKIHYAPEVDVKVCALEAVNLKSSEILDHQTTQICNESRQNALAEILEVLTVC
THNLPLAQTWVPCQHGSQLANGGLKKNTSFDGSCMGQICMSTTDMACYVVDAHWGFRDAC
LEHHLQKGQGVAGRAFLNGGSCFCRDITKFCKTQYPLVHYALMFKLTTCFASLQSSYTGDDSYILEFF
LPSSITDDQEQQDQLLGSILVTMKEHFQSLRVASGVDFGEDDDKLSFEIIQALPDKKVHSKIESIRVPFGF
KSNATEMLIPQPVVQSSDPVNEKINVATVNGVVKEKKTEKKRGKTEKTISLDVLQQYFTGSLKDAAK
SLGVCPTTMKRICRHGISRWPSRKIKKVNRSITKLKRVIESVQGTDGGLTSMAVSSIPWTHGQTSA
QPLNSPNGSKPPELPNTNNSPNHSSDHSPNEPNGSPELPPSNGHRSRTVDESAGTPSHGSCDG
NQLDEPKVPNQDPLFTVGSPGLFPYSDHDVSAASFAMPNRLGSIDHFRGMLIEDAGSSKDLR
NLCPTAAFDDKFQDTNWMMNNNNNNLYAPPKEEAIANVACEPSGSEMRTVTIKASYKDDIIRFRI
SSGSGIMELKDEVAKRLKVDAGTFDIKYLDDNEWVLIACDADLQECLEIPRSSRTKIVRLLVHDVTTN
LGSSCESTGEL

>AtNLP8

MENPFASREKGFGNYSDFPTEQMDGLSSNFGSGVRNLISDDMFNPSELNFDSLAAWCNSPSAT
DILFAQYGLSNSQPMMPFGAFTSFHVADPKATSLTRSFYDLESSYYGEERSSAQEMNSQFHRSDDDEL
SGKRRKVVNQKIGFPNVLNCTIPRSLSHSLDEKMLKALSLFMESSGSSEGILAQVWTPIKTDQYLLST
CDQAYLLDPRFSQYREVSRRFTFAAEANQCSFPGLPGRVFISGVPEWTSNVMYKTDEYLRMKHAID
NEVRGSIAPILEASGTSCCAVMELVTSKEKPNFDMEMDSVCRALQAVNLRTAAIPRPQYLSQRDA
LAEIQDVLRTVCHAHKLPLALA WIPCRKDQSIRVSGQKSGENCILCIEETACYVNDMEMEGFVHACLE
HCLREKEIVGKAFISNQPFSSDVKAYDISEYPIVQHARKYGLNAAVAIKLRSTYTGEDDYILELFLPVS
MKGSLEQQLLDSLSGTMQRICRTLRTVSEVGSTKEGTKPGFRSSDMNSNPQTTSSENFTQISLDSE
FNSTRSMFSGMSSDKENSITVSQGTLEQDVSKARTPEKKSTTEKNVSLQHQFSGSLKDAAKSLG
GETSAYFQAWVYFFCP TTLKRICRHGIMRWPSRKINKVNRSLRKIQTVDLQVQGVEGLKFDSATGE
FIAVRPFIQEIDTQKGLSSLNDAHARRSQEDMPDDTSFKLQEAKSVDNAIKLEEDTTMNQARPGSF
MEVNASGQPWAWMAKESGLNGSEGIKSVCNLSSVEISDGMDPTIRCSGSIVEPNQSMSCSISDSSN
GSGAVLRGSSSTS MEDWNQMRTHNSNSSES GTTLIVKASYREDTVRFKFEPSVGCPQLYKEVGKRF
KLQDGSQLKYLDDDEEWVMLVTDSDLQECLEILHGMGKHSVKFLVRDLSAPLGSSGSNGYLTG
>AtNLP9

MENPSASRDNKGF CFPDIPVEEMDGWVKNLISEEDMFSSSTSELNFESFASWCNSPSAADILFTQ
YGLSTSQSIIIPFGGLEGSYACEKRPLDCTS VPRSLSHSLDEKMLKALSLFMEFSGEGILAQFWTPIKTGD
QYMLSTCDQAYLLDLSRLSGYREASRRFTSAEANQCSYPGLPGRVFISGVPEWTSNVMYKTAEYLR
MKHALDNEVRGSIAIPVLEASGSSCAVLELVTCREKPNFDVEMNSVCRALQAVNLQTSTIPRRQYLS
SNQKEALAEIRDVLRACVYAHRLPLALA WIPCSYSKGANDELVKVYKNSKECSLLCIEETSCVNDM
EMEGFVNACLEHHLREGQGIVGKALISNKPSFSSDVKTFDICEYPLVQHARKFGLNAAVATKLRSTFT
GDNDYILEFLPVSMKGSSEQQLLDSLSGTMQRLCRTLKVSDAESIDGTEFGSRSEMTNLPQATV
SVGSFHTTFLDTDVNSTRSTSNISSNKRNE MAGSQGTLQQEISGARRLEKKKSSTEKNVSLNVLQQY
FSGSLKDAAKSLGVCPTTLKRICRHGIMRWPSRKINKVNRSLRKIQTVDLQVQGVEGLKFDSVTGE
FVAVGPFIQEFGTQKSLSSHDEDALAR SQGDMDEDVSVEPLEVKSHDGGV KLEEDVETNHQAGPG
SLKKPWTWISKQSGLIYSDDTDIGRSEEVNKDKE DLCVRRCLSSVALAGDGMNTRIERGNGTVEPN
HSISSSMSDSSNNSGAVLLGSSSASLEQNWNQIRTHNNSGESGSSSTLVKATYREDTVRFKLDPYVV
GCSQLYREVAKRFKLQEGAFQLKYLDDDEEWVMLVTDSDLHEC FIELNGMRKHTVKFLVRDIPNTAM

GSSAGSNGYLGTT

>AtRKD1

MKSFCCKLEYDQVFGKENNSFSFLNHSSLYSHQSELANPFFELEDEMPSATSSNCFTSASSFLALPDLE
PISIVSHEADILSVYGSASWAEETMFVSDFAKKSETTTKRRRCREECFSSCSVSKTLSKETISLYFYMPIT
QAARELNIGLTLKKRCRELGIKRWPHRKLMQLQKLISNVKELEKMEGEENEDKLRNALEKL
EKEKKTIEKLPLDKFEDTKRLRQACFKANHKRKRRSGMSTPITSSSSASASSSSYSSVSGFER

>AtRKD2

MADHTTKEQKSFSLAHSPSFHDHSSLSYPLFDWEEDLLAQENSGSQAFPTTSLPLDLEPLSEDVL
NSYSSASWNETEQNRGDGASSEKKRENGTVKETTKRKINERHREHSVRIISDTTYTSSAPTLSKET
VSRYFYMPITQAAIALNVGLLKKRCRELGIIRRPHRKLMNLISNVKELQKMEGEENAELQDAL
EMLEKEKRTIEDLPDLEFKDKTKRLRQACFKANHKKRSLKSDQSQVPSCSSGSVPSDESV
DEAGMESDEEMKYLLCGFSSEFTSGL

>AtRKD3

MADQRPLMTWLEANNYESFLQEDIFSFLDQSLFVDPHSSIDPFKDFQTQNWFSLQDSIVNHISTTFA
ADHTFLASLDLEAISSTFSDLISGGWWNENNGNNYNNQVEPNLDEISRTNTMGDPNMEQILHEDVNT
MKEKTSQKRIIMKRRYREDGVINNMSREMMKQYFYMPITKAAKELNIGVTLLKKRCRELGIPRWPWRK
LTSLNALIANLKDLLGNTKGRTPSKLRNALELLEMEKKMIEEVPDLEFGDKTKRLRQACFKAKYKRRRL
FSSSS

>AtRKD4

MSSSKHSSVFNYSALFLFLQQMDQNSLHHLDSPKIENEYEPDSLVDMLDKLPLDSLLDMEDLKPN
AGLHFQFHYNFEDFFENIEVDNTIPSIDIHLLTQEPIYFSSDSSSSPLAIQNDGLISNVKVEKVTVKK
NLKKKRQDKLEMSEIKQFFDRPIMKAAKELNVGLTVLKKRCRELGIYRWPHRKLSNLNSLIKNLKNVGM
EEEVKNLEEHRFIIEQEPDAELSDGTKKLRQACFKANYKRRKSLGDDYY

>AtRKD5

MVDQGFFTLKKEKKNILIKLFYVKILGFCFCDSWSSDMAHSLTSLAVFQSVIRKEMVRSLHVYESV
EIREFWFKSKSCYVEKKAKPLFRSEDFFRPEISEGSVFGTWRCIFVFRFNHSLPRFPTLLCLSRNP
LEDIPNLANELKFISELKPSKIYEEEQCSSTEGLYNSDLPKPRKLVLQDLNCLPDSETESEESVNE
KTEHSEFENDKTEQESDAKTEILKKKRTPSRHVAELSLEELSKYFDLTIVEASRNLKVGTVLKK
CREFGIPRWPWRKIKSLCLIHDLQREAQKQEKNEAAAMAVAKKQEKLETEKRNIVKRPFMEIGET
KKFRQENFKKRHRASRAKKNQESLVTSST

>AtRWP-RK-1

MAGDDPKSSAPKDDGYEDPYINDPDFGFGPTDEDLINVMADINPNLMDMSFTEHTPPSVLPTLPSD
VDETHLFHELDLSISNYDSKDEFETLSFVEDGFTIPLDMFEQPNDHNDNNDVPGEIGNDNGKNVVG
DSTRLELSTRVFGGANGLDFEIGGSATMPVPVTTISDGVHACNCCLLRELVHLKEGEERSKLDIYGGI
GFICHAILLIQLLAPDSMQRQPLIFHLQHLMEEVKFIEDCYQRVANGLQLQDTNAAFYQAMSAN
SIFNQPPPMLPSSTDVPLSLLSPNEDHDVPLSLLSPEEALDIQPVPPHVELREEPTTRLPKQREKRQTP
LAAQVISLVSQFNI

>OsNLP1* in 2005 (OsNLP2)

MEQKPSPPPPRSDEEDGLMGCGMGGTDIAGGDLDLMEFLLATPGFDLSEFWHPGAASPFSPLF
DIGSSVTLTTPAPAAGEDDRDEAEMPSRGGGLEVSPAHRGWTQFQEVAVEPTVKERLRRALE
RIASQSQAQRGDGELLQVWVPTRIGDRQVLTCGQPFWLDRRNQRLANYRTVSMKYQFSADE
SARADLGLPGRVFVGRVPEWTPDVRYFSTEYPRVQHAQYFDIRGSVALPVEPRSRACLGVVELVMT
TQKVNVSAEINICNALKEVDLRSSDVSSDPRSKVVDASYRAIIPEIMDVLRAVCDTHNLPLAQTWIPCI

CQAKRGSRHSDESYKHCVSTVDEACYVRDCSVLGFHQACSEHHLRGEVVGRAFGTNEPCFSPDIT
TYSKTQYPLSHHAKLFGLRAAVAIQLRSVKTGSLDFVLEFFLPMKCINTEEQRAMLNSLSNTIQQVCYT
LRVVPKELVNDGPFEISQPTRPEFYAKSVHEDLDELCSGINVPGRTSLEASEEVSSWIASLVDAQNKK
GGKGEIDVDLPGFSKQDDEGFSVTAGWHTSPVMAPDGSMFSGFKRHEDYDVKENTC **SSDPSNSN**
SDKAVEKRRTKTEKTVSLQDLRKHFAGSLKEAAKNLGVCPTTLKRICRQHGINRWPSRKIKVGHSLLK
LQMVIDSVHGPEGTVQLSSLYENFTKTTWSERELQGDVHFPAEQNFQLEPSVPDRPCEGRFTSHTS
GSNSISPSCSQSSNSSLGCSSVPKTQQQHGSAPQLAVKEEISMDENQCSTLIKSASHAEALQMFVEE
RPTMLFRSQSQVLLSEHKPIENMSNVQKARSDSLKIKAIYGEERCIFRLQPSWGFQRLKEEIVKRGFISQ
DTHVDLKYLDESEWVLLTCADLLECIDVYKSSSNQTVRILVNPsiQPLLNASFGQTGLS

>OsNLP2* in 2005 (**OsNLP6**)

MDMPTPSNRAGCNGNTGGTMGPSDDPYGAAAMNLDCYSEIYSPSVADQLFSLLNDPAAHRMFA
MWPSMGSSPCAAGTSEDMPLDAYSGLGEAVEEPSQIMSVNPTEAEEKTGSSGELGSDDGAHQGSS
MVPRSVVGSSLADRMLMAISLFRESLGSGALAQVWMPVEQEGHVVLSTCEQPFLDQVLAGYREVS
RHVFVSAKEEPGLQPGPLPGRVFISGVPEWTSSVLYNRPEYLRMEHALHHEIRGSLAMPIYDPSKDSCC
AVFELVTRKEKPDFSAEMDNVCNALQAVNLKATKGSSNQKFYTENQKFAFTEILDVLRAICHAMLPL
ALTWVPTSNGIDGGYVVGKDGASFQSQGKTIIRIHESACYVNDGKMQGFLQACARRHLEKGQGIAGR
ALKSNLPFFSPDIREYSIEDYPLAHARKFSLHAAVAIRLRSTYTGNDDYILEFFLPVSCKGSGEQQMILL
NNLSSTMQRICKSLRTVYEAEVNVNAGTAAVFRKNNECLPTGHTESSSHGDQSITGASFEDTSLA
NKPGVMEPELAEQVQPSSIGHAEKKRSTAENISLDVLRKYFSGSLKDAAKSLGVCPTTLKRICRHHGIS
RWPSRKINKVNRSLLKIQTVINSHGVDRSLQYDPATGSLVPVSLPEKLTFCGDLPTPSVGKTVEE
NSDLKSEEGCSLPDGSQRQSCQLQISDVKKSNEDEFHIGSGNSDFYGANATAKSNSEVTQGPLCPTG
AFSALHLKGTDCTNPSSLRPSSESTRNQIVGRNSPSIQQEDLMDLDNHEAEDKDHMHPSTGMDT
SSSGSASSHPTFKQNTRSALKDAASPALTVKATYNGDTVRFKFLPSMGWYHLEEIAKRFKLPTGAYQ
LKYKDDDEWVILANDSDLQECVDVLDSIGSRIVKLQVRDLPICVSSSGSSTCLQLAAHSS

> OsNLP3* in 2005 (**OsNLP1**)

MEVDPSSSLPGAGEGGGGGGGGGGDWPFDSLTSLLFSSVSASPQPLPASSSSWLTTPSPLWLFD
RQLLPLDMGAPAAPATAPPAEAAAEEVHRTRSGNSDTSKRVHQINSKWQFHLSIDDNTDSSCLF
KERLTQALRYFKESTDQHLLQVWAPVKSGDRYVLTSGQPFVLDQSQSIGLLQYRAVSMMMYMFSD
GENAGEGLPGRVYKQKVPEWTPNVQYYSTEYPRLNHAISYNVHGTVALPVFDPSVQNCIAVVELI
MTSKKINYAGEVDVKCKALEAVNLKSTEILDHPNVQICNEGRQSALEIETLVVCEEHKLPLAQTV
PCKYRSVLAHGGGVKKSCSFDSMCIGEVCMSTSDVAFHVIDAHMWGFRDACVEHHLQKGQGV
GKAFIYRRPCFSKDISQFCLEYPLVHYARMFGLAGCFAICLQSMYTGDDDYILEFFLPPNCRNEDDQ
NALLESILARMKKCLRTLKVVGNGDTNEVCLQISNVLIETEDLKTNVHFENSEGCFRESPESNGSQRV
HEVDNDGNKVSIMSERHLLADDNSQNNNGASVGRPNGSGASDSLHKSNKPPERRGKAEKTIISDVL
QQYFSGSLKNAAKSLGVCPTTMKRICRQHGISRWRPSRKINKVNRSLSKLQVIESVQGSDAAFNLTSIT
GPLPIVGPSSDSQNLEKASPNKVAELSNLAVEGDRDSSLQKPIENDNLAILMSQQGFIDANNLQLE
ADKASHSRSSSGEGSINSRTSEASCHGSPANQTFCVCKPIASTFAEPQLIPEAFTKEPFQEPA
LSRMLIE DSGSSKDLKNLFTSAVDQFLARSSNLALMQNSGTVTIKASFKE DIVRFRPCSGSVTALKDEVAKRLR
MDVGMFDIKYLDDDHEWVKLACNADLEECMEISGHSHVIRLLVSDVA AHLGSSCGSSG

>OsNLP4 (**OsNLP4**)

MEEGDPQPSISLARTPSEGAAAVALDLLEQLLSADNAWLEVAANTSRSRPNFFATPSNCLTDASVAT
TTPANSWWIQPAGASTSVRERFDQALAYIRETQSDADVLVQLWVPVKNDGQLVLTSGQPFTLDQ
RSNSLIQFREVSTKYQFSADVAGSSPGLPGRVFIGRLPEWSPDVRYFTSYEYPRINHAQYLDVHGT
GLPVFERGNYSCLGVIELIMTKQKLNFTSELNTICSLQAVNLTSTEVSSIPRAKLN
SASYKDALPEILEV

LRAACITHKLPLAQTWVTCAQQGKGRSRHSDENYKYCISTIDAACYVNEPRMOSFHEACSEHHLLRG
QGVAGKAFTTNQPCFLPDIGSSTKLEYPLSHAKIFNLKGAVAIRLRCRTGIADFVLEFFLPTDCEVL
EEQKAVLDSLSGTMRSVCQTLRVVTDKEMEDEAMREMNELNFSPRGKNKVEELSGDNTRGDREE
ASWTLVGTQSQKGSDLAELLTHGMLSHGGHGSSQAGDQTSKEGSKVKRRTKTEKTVSLQVLRQYFA
GSLDAAKSLGVCPPTLKRICRHGIRWPSRKIKKVDHSLRKLQQIIDS VHGAETAFQLNTLYKDLTN
TSVSSDNNLSGSVTVPLANQNNLDFEMHQHHRLSSNIPSTS LSHSSCSQSSDSSPSCSGGATKHSPQ
VGADQVRSGCLPQHSPVQTLQTEA SINEHFSGQEAPIDLQDV AEKANGEQHMSQSPSSPKQTA
NVGMRVKVTFGSEKVRFRLKPECDFQELKQEISKR LSIADMINS LIVKYLDDSEWVLMTCDADLHECF
HVYKLADIQT KISVHLAASPTTRITIGHTGFS

>OsNLP5 (**OsNLP6**)

MDMLMLSSALDGLDSYTDLVAGSSVADSIFSALT CFPSSQERLLHVSTPVGSNSRQDDSDVSITKEGT
TTRRGDCAAGLASGE PVAAGIGSVPKPF DGVTLTERMLRALAMLKEAS GGE AILVQVWMPVRNGEQ
HVLTTSDQPFLDQKLTGYREVS RQFTSAEGPGLFPGLPGRVFM SGMP EWT S NVMYYHGSE YLRV
DYARRHEVRGSLAMPVF NSSGGSCAVLEV M TREKDNCLEMVN VSNAL QSVQLSTVNAWRHS
QSYSRDQKLALMEIFDVLQAVCQA HLLPLA LA WIPVCSK RDVL V SVEYGANFGKRNKEVLCIEESAC Y
VNDTRMRDFVQVCAEHPLEKGQGVAGNAYLSNNPFFSDV K DYDMHAYPLVN HARKFGLHAAVA
IRLQSTYTKNDDYVLEFFLPVLCKGGEQQLLDSISATMRRVCKSLRTVSDAELKEDVTRKPSNENRS
GTRCPSPVNLIYSGREIDVSNETKTNTPLEYQIEGIDEQLSDTKSTNKLIKCSNASDGEKRRSTEKS VSL
SVLQQYFSGSLKDAAKSIGVCPTTLKRICRHGIRWPSRKIKKVNRS LKKIQNVISSVHGVEGVLYDP
STGCLVSSVSPSIEPVLMNVEHKGS DPLPIESELPHLN FEPDCDAYRREHAGQDV LHKLQN KQNGEIN
FDMDDGELFRNSHSTRTLSGAF CEDMPNGLYVAREMTCVAKTGTRTERLEHKPSSRDSFSAPQEYR
MESETDKSNKNSKQSLPSSSM TDCSTSSGETFKSIKS QSANESNKVVVKASYKNDTIRFKLLPSMKY
EQLLEEAKRLKLSIGLQLKYK DDEGDWVILASDADLQECL EILDTRRLRILK LQVQD VVCPIGSSSGSC
SMLRP

>OsNLP6 (**OsNLP7**)

MERVVGDFNLLLQRGGAAGERGSEGGGGSPGTEAAA AVVKQRIARALRLYK EAAGDGGGGWM
VQVWAPARDGARRVLATRGQPFVLA SQCHRLFQYRTVSLTRVFPVGGAAA ADEQGLPARAFDATA
PEWTPNVQCYGSGEYARISYALIYDIQGSLALPILD PDDASSPLAVLELVTTAPLLRVSGEVANLCNALR
AVSLRGAGICNRRAECYSLAVNQIVHRD ATTRAAMA EVSELLITVCEAHKLPLAQ TWVRCWSCGGGG
EDTEKA ALTTAGFHLAAGADARGFRD ACV EHHLQRGQGLVGT AARAPGGGRLCADVARCSKDD
YPLAHYAGMYGLAGCLVLAELSAAAMADA AAAA TAGDEEDCVVLEFLPPDCTGVAEQKA A VDAVS
ATIKQCSGNLKAIVISNLDDFLDTMADGDHQLRHEMDDLGDDQRC SDEEDLQLLEN TNIGELNIHN
ADQIRNEDPTS QVGKNKTKRGKA EKSVTLEELQKHFSGSLKDAARSLGAGY SVDYLHQNP DVY AQIN
TQPLKV GSPGSMASQGHFARSAR CGNGDGVTIKASHRGDIIRFRVPCSAGVAAVKA VAKRLSL
DAGAFDV KYLDDHEWVLLSCAD FQE CLDV VPALPSMSVTARSGSGAA APVV RL MVQEADN I
GSSCASSD

>OsRKD1 in 2005 (**OsRWP-RK1**)

MCAAPMASTAQPQQQPQQQQEQQPVAAA AVPTPAPPASEAQPKPTRVSLSYEEISKLFSLPIAEA
ASILGVCTS VLKRICRHGIVRWPYRKL VSGKGDDTKNAEREKAGLLEISKVAKQKALSASGLSTVSP
GAFQGVAKSQGSSKAGQVSPPGKQNVLGGSAI SYGTQTKGIP TYMDDFKYGFPSGLSLQTMKW
WGTD SHTETTPAKDDNGEAPESANEASKGMTDELDWGADEAE AEA DADS AITTEPSAQLCSLRRK
AVDDGRKLLTGKCGGLECLRLNKRQKM ALAQVFGASLPEQLRSKLG

>OsRKD3 2 in 2005 (**OsRKD2**)

MEMHECCYYGGSIGGDWLNLPLAIPPPCSSSSSWSSQLLLSDHDDVLLHSAGDHGGA VAGIGGA

CMTADLVVRDEEMEMAAGYLPVAASAAAAADVHDHYMYQQFQLEPDQFVSTLPAVAVAATAG
GGSHDDELLRMPFTIDLDCAFADARDVVGVGEPKPSPQHTLAAIALPAVGGGAHHFGTQDDD
VKFDVTKQRNDAALAGDDLSMVIVESYEMGMRRHAAEQQEQQPKIITSAAATLPLPLPPPPPPP
RVTRSRRDGSSAATAGGKTRLDHIGFEDLRRYFYMPITKAAREMNVGLTVLKKRCRELGVARWPHRK
MKSLSLILNEMGSKGMSSAAMRREALENCCALMERNPAVELTERTKKLQRACFKENYKRRRAAA
VDVLDDHCFSAAGHCHRHHHQQLALPPPPAAAADHRRRDFFGY

>OsRKD4 in 2005 (**OsRKD5**)

MDAAAAAVTLSALAVFASTLDHGAVRSVHGYKVYGRGGRRWERWVEREFLTPASCREVPAPVA
PPRILPAEWGRGPAYREGQVVAAGAWRCILAFDSAAPPRTPPPVLSPFLNPRLMCVPSLYNDLEKVF
RFQNVEKIPKLMQCDSEEKSSWDARDKSSDEVHASESDSDDLQSSEEEKPTVQKQRRANKKIAS
ITLVDIAQYFHLPIREASRTLKGVSILKRKCRQYNIPRWPHRKIKSLDSLIQDLEYVIDGDDHDDTGDD
VQQEKKQTAEEKQEAIMLTRRKQMLETEKETIQQIPAMDLKETKQFREDVFKRRYRAKKDLAND

>OsRKD5 (**OsRKD2**)

MRHNRSLAALLRAGRYGAARRLFDALPARSVVTWNSLLAGLARRPDARAAREFFDAMPVRDAVSW
NTLLAAYSASPHPDHLAARRLFDEMPQRDVVTWNTLLGAYARRGLMDEARRLFDEMPQRNAAS
WNTMVTGFFAAGQVVKALDVFDAMPAKDSASLSTMVGFTKNGMLHEAEELLTKRLSVTDMDKAV
DAYNTLIVAYGQAGRFSDAKRLFDMIPKGQYQHNMLKRKGFERNVWSNSMMICYIKAGDVCSAR
ALFNEMPDKDLVSWNTMISGYTQASDMKESEKLFWEMPPDTVSWNLIIQGFMQKGEAEHARGFF
DRMPERTISWNTMISGYEKNGNYISSVKLFSKMLEVGEIPDRHTFSSVLAACASIPMLGLGAQIHQLV
EKSFPDTAISNALITMYSRCGALNDAEAIFKQMIHTKKDLVSWNALIGCYEHGRATKALQLFKEMRR
AKVMPHTITFVSLAACVNAGLVSEGRMVFDTMVHEYGIVARIEHYAALVNLIGRHGQLDALEVINS
MPMAPDRSVWGAFLGACTAKKNEPLAQMAAKELSTINPDSSAPYVLIHNLHAHEGKWGSAAVRE
EMERQGIYKQPGYSWIDLEGKMHVFISGDTWHPNAQEIFSVLEDWQWHNPMSLEMNQLTQVKLIA
KRNNQIMDKQATATKMHKATNIRHIYSKFKVDEQQNTSWWPYCTSLWPDSYLLEEAFLSSLSFPSF
HPQPVYSTVMQSNVLQDELGVIFEDDVLYWDEMEQSENKVEKSEKGLPLYYGDENGAASKIMRD
DVRSEEKALTfelvsqyfympitqaarelnvgltlkkcrelgiprwhrkmkslqtlinnvqlqea
skanneeqlrmlveumlqeerrlleqkpyvqleektrklrqacfkanckrrllaleagep

>OsRKD6 (**OsRKD2**)

MEKWRFYGGHQFQSYSFHEEENLFQDWSDLYLLLGEDEPFTHHFSTSVDHSNFVQDELYTLFDGDILS
IWGDMKEDAYHRSDKDGGEKEEKLHHEKAMELQLQRLPSGRQSGEKTTFELVSQYFCLPKIQAAQE
LNVGLTLLKRRCRVLCIGPRWPHRKVKSLETIKNVQELGMETGQDEDNTRNAVEMLQQTKKLIEQSPD
AKLDDWTKMLRQACFKENYKRRLLAIEG

>OsRKD7 (无)

MAMS RATCSTRPRISRCSPSRRWFVPAAAIESHPTAIHVVEREWPQVQISRFDPSCPPPARMLP
GGQQQQPHPAPGGLDLRSDEIHALKQRRRPPNPQSSSPPEGLLGAANPDRPDDDELAILSSFRDE
AAHGDGGGVARVQVLPVEQDANLPSLPLRHGQLDCSRCHLVRHVMHAVLPYVKFSLKLVWEE
AFHRIRRMYVISVRSNVYSLSKWTQEWAIFIARNIDTMRNNTNGQLLDSGYSNLVESVRTNVNPH
TAVEVNLLQTIMSAPSADHHQNAADQVAAPAAQPSAAPPVALPPKAAPRKARKDRDYASMLVA
EEFYVAATSRPVPSNDVEILESSHVSQQQDGGRAIYPSLQARRGKTKQEVPRRANKDVLEYLSLARKE
TEKEINTLSSFDGIYRNDGTLSYLMTEVRRLNKIWLQKNAPSTLSSRLLASVKEIDDICKVEKGRLYAQF
ISALKKLCRKMKDDGGSAPSANN

>OsRKD8 (**OsRWP-RK2**)

MAGDGGNGAADGGGEHELEDFPSPLLDNFVLELLDSLGVDDDGAPPAPAPVQMSVDGAG
GGRINKPGSSVDDLVDWSTAFCGNNC SKIDGEYGGASTSSAAPALAPPQEDYCSGCQVLREVHNS

GLEITKLCIHHGVASGEFYHAILDVYRVSASAPAPALAHHSIINFRGRGYDWVKQYLTEYALRRAGGGF
AVVQDSLSAFHDALCTTMAPCSSHVGDDDAHRRASSSSAAAAEERTNGNGDHGQLVVHNAAVLP
MLESSRCLVAADQAATTNNNGDERRLVLDTTAIQPPASGCLHMAFPSKSYQSSRPTLPSRYQE
CTCNPVAYEMDGVRSSLAPICWPELLGYNVTNREKTQQLQLSDIAPYFELPIAKAAKKDICATALKGIC
RKHGVLWPYRKVRSIDRQIATLRRSGNGDATRNEIETLIASRRRIVAGLDQ

>OsRKD9 (**OsRWP-RK2**)

MATGGSSSSNVPSGGAAGGGAGSGKLPMPVPGSIECLRKRCRQLRVLIHVNDHRKAVVVLHAGED
GKPDHVLVQNVSPEGEASVQSTYRIDVSGETPESQAEMLNDWYTSFRMDTTGVLYDSDQNVYGV
RGHPGGDVPRSLAILPPAPKKNQHGKAPATESNSSLVEEPLLVQTDQPAIGKRKKFTFPDQRKRVK
TMTKKDLESYFHTQKSAAHIGLSIGTTALKNLCRANDLPRWPYRQIASLDNKFNNNLKKQITGWNLG
KAVQGVTKAFLRKEKEFYQKIMSSMPEQLQSIDEIVNSLPEADDIDIEDDEDNDVIEDNDDNS
DEN

>OsRKD10 (无)

MAGEHRYDSNGDSRYMLDDMEFLPLFGEAEAATTLPLPVLPDVEPLPTAPAPAATAHVEPAPAAD
FGLGNPVLAFLGFDVLDFFPELNQSPPPPPAMNAGGYTHQVQASPPVMHHQQQQQPLAPL
AAHGFAGQPAPATTMAPSGGDDGLFLAAPSRDAPCSPVMFNFMDFNVDMDGVDMDDVLMWA
DQDTGAAAGGDTAPPVVVDEYADFVPFQAGDLDCSNCHLVREMMHANASRTIYFLVHATGVGS
FQHAIVDRRTATGAEGLHFPGRQLLYFDLTNHTIESASDFIASNVEKLKNDDTGHFLDTGYNFGA
VRTDMANSHTAMEMNMLHTIVSAPFENVTTDAASPPAAQFIGAPPAAELPAPVPAPAPPAHEQN
AVVATLLFKVEEFYAAANSRPAAKRADVKILESSQVTQQAGGSSAATATMPSMVDRKRKRAQATPS
RMAPHEVIQYLRTAVETDKELETNNFFKVCDCGEDKALITFSVEQIRSIKKIGRIINKPVTAMSSRRM
ARFIDEIDTIKEEKARVFEEIIKILKNPDKRENDGSSGSNRKNVGGSSGGKKTVGGSSGSKKKNVGRP
SAKKAQK

>**MeNLP1**

MPEQEDDKSKENQTVGGDKGETLMDLDDLDTSWPLDQISYLSNPLSPLLLSSSDQPCSPLWAFSDA
DDDRLLPVSSASHATTPLRFSDPYPIFLACAPNSVAENHTENDDKRKFPSPLLGLMPFDNPDGYCVIKE
RMTLALRKFKESTEQHVLAQVWAPTKNGGRYVLTSGQPFVIDPHSNGLHQYRMVSLMYMFSVDG
DCEGELGLPGRVFRQKLPEWTPNVQYYSSKEYSRRDHALHYNVQGTALPVFEPFGQSCVGVL
TSQKINYGPEVNKVCKALEAVNRSSEILDHPSTQICNEGRRNALAEILEILTVCETHKLALAQTWVPC
MHRSVLAFDGGLKKSCTSFDGSCNGQCMSTTDVAFYVVDPMWGFREACLEHHLQKGQGVAGR
AFLSHNACFCPDITQFCKTEYPLVHYARMFGLSGSFAICLRSSYTGNDNYVLEFFWPLSISDNYEQN
LGSLVATMKQHFQLKVASGMDLEEEGFVIIQTSMSGRDLRLECIRIPQSPKSPPNLNTSPKEGKT
AQLDPLKHHLMVNLDADVNGGNISQAEGSHFPTSLPENKGNNKPSERKRGKTEKSISLEV
LQYFAG
SLKDAAKSLGVCPTTMKRICRQHGISRWSRKINKVNRSLSKLKRVIESVQGAEGAFDLTPL
SPLPVTV
GSISWPSNLNGCNQQNSLNSKLEPIDEKNGSPICNAAGNDQGGGEAQLLGGRILSREEAVLQNG
FSPKIGLVSDFRKAGNASGEESTGPTSHGSCQGPANDSAPANDSPVSPVHEKCKNAGNTPELAFQ
TKGKLNLSAAYSVPDALVATEAQAPLGEMLIKDAGSSKDLRNLCPSVADTILEERVPESSWTNYPGP
FPTTEAVAALEHAMTQGTRQEMMTITIKAAYREDIIRFRISLSSGIVELKEEVAKRLKLEV
GTFDIKYLD
DHEWVLIACDADLQECIDISRSGSSIIRLSVHDVNTNLGSSCESNGE

>**MeNLP2**

MAEQEDDKSKDNQSVAGNKGETLMDLDDLDTSWPLDQIYFLSNPLSPLLLSSSDQPCSPLWAFSD
ADDERLAASSSHATHPLRSLDYPPIFLTCNPSSLAESQTENDDKRKFPSPLLGLMPFDNPDGYCIKER
MTQALRNFKESTEQHVLAQIWAPVKNGARYVLTSGQPFVIDPHSNGLHQYRMVSLTYIFSLDGEFE
CEPGLPGRVFRQKLPEWTPNVQYYSSKEYSRRDHALHYNVQGTALPVFEPFGQSCVGVIELIMTSQK

INYGPEVDVKCKALKAVNLKSSEILDHPSTQICNEGRKNALAEILEILTVCETHKLALAQTWIPCMHRS
VLAFFGGLKKSCSFDGSCNGQCMSTTDVAFYVVDPMWGFREASLEHHLKGQGVGRAFLSH
NACFCPDITQFCKTEYPLVHYARMGLTGCFAICLRSSHTGDDEYILEFFLPPGISDFQQKALLGSLLA
TMKQHFQSLKVASGMDEEEGFVEIIQTSVSGKLDLRLECIQTPESPESLPNVNTSPKEGQMAQVDP
VVNLDDVNNGGSTGDAEGGHFPTSPENPKGNKKPSERKRGKTEKTISLEVLQQYFSGSLKDAAKSLG
VCPTTMKRICRQHGISRPSRKINKVNRSLSKLKVIESVQGAEGTFDLTPIATSPLPVAFGSISWPSNL
NGCNQQNSPNSRSSEPNERNGFICKVAGSEEQSGSENQLPGGRVISHEELFLQNCFSPEIGQGSS
RHKAGNVSREESAGTPTSHGSCQGSLANDIAPAKVASASPQEQQNEVGTIPELSFQPKGKPNLALSAL
YIPDAFVATEAQAPFRERLIEDAGSSKDLRNLCPSVADTMILEERVPEYSWTNHPGPEFPTKQSMAAL
EHATPRVTARQEMRFVTIKATYREDIIRFRISVGSGITELKEEVAKRVKLEVGSFDIYLDLDDHEWVLIAC
DADLHECIDISRLSGGNIIRLSIHDVNTNLGSSCESNGELLGCTYGKV

>MeNLP3

MLGTRADSAMDFFDCMDKLLLEGCWLETIDGSEFFNPSPFSSAAFVDASFQWPTSEINNCCLASSPSK
RSNREEQISLLPRNSALNEAHGRSLINAEDALDHGVGSADRLGYNTTEGSEVSRRWWIGPRTNPGPKT
SVRDRLIRALGYIREFTKDKDVLIQIWVVPDKGGRVLTTHDQRFALVPNCQRLANYRDISTNYQFSVE
EDSKDMVGLPGRVFLGKVPWEWTPDVQFFRSDEYPRVDHAQQYDVRGTLALPVFEQGSSTCLGVIEV
VTTKQKIKYHPELESVCMALKAVDLQSSEVPSLQHVKMCDMSYQAVLPEIHEVLRSACTHKLPLAQT
WVPCIQQGKGGCRHSNNYYRCVSTIDLACYVHDGVQAFHEACSEHHLLKGQGVAGEAFLTNQP
CFSSDITSYRKTEYPLSHHARMFGLHAAVAIRLRLHSGAADFVLEFFLPAYCTDPEKQKEMLTSLSITV
QQVCQSLRVVTDKEEETGCPISELLASSVDTPPKEEMLVTPQPSVSYGGESLSHADRLTGQQSGSV
VSSCQKKKQKVLEDQISVDCWQNQEDYSLKVSVESGGDSAVAEGSFSSVFKGKKGEKRRTKAEKSITL
QLVLQRYFAGSLKDAAKSIGVCPTTLKRICRQHGIKRWPSRKIKVGHSLQKLQLVIDSVQGASGSFQIG
SFYTNFPELASPNLSRSNLNSTLKQSDHPEPSSIQSERYFSSQAATPKSPSSCSQSSNSSHCISSGTQL
HASAVTTPTSEDPVLGENDNCVLRARSDAELRASIQEQQNLLPRSQSQKSLGEQPNLGNLLPLPNN
SSSIIRDFTQRVKVTYGNENKIRFRMPNNWELKDLLQEIARRFNIDDIKYELKYLDDESEWVLLCDD
DLEECIDVCQSSQTHTIKFLLQVSPHLFDRSMNSRGLS

>MeNLP4

MDHGASALNSTWTFSSTTMDLDVMDKPLSQGYCLENNDRANSLEPEPFTSHVLYDHSEYSSFED
LYHQIYRKEMKPLFENIPSAYHGIEGAETAKSFVQSESFTLEGSNSGSRWLIDPSYSVKQRLMLAIKYLKE
YVEDSDVLIQIWVPTQIGGQNVLTIDQPYSLNPSCSKLASYRNVSKTFQLTAEENLKELAGLPGRVFL
GKLPEWTPDVRFFRSDECPRKSYAKMYNISGCLALPVFEQDNGVCLAVVEIVTTQKISYSLELEIVGRA
LEAVDLKSSQDFPLGIKACNESYQISILEISDILQFVCKTHGPLAMTWALRDRQGEVEHWQFSKKYD
YCLSTVDSACYVADSDLLGFHEACSEHYLFPNQGIVGKAFTKKQWFATDITSFSKANYPLSHHARMF
NLRAALAVPLCNIYTGLIEFVLELFPWDCKDIEEQKHFWDMLSIVMQQACKSFHVIVDKEIDEVSEQ
MVVASDGRYNREGIQLVELVAPSNESSQEESSTPHEDSQNKDKFSISWECPKEEDEFKVITHWE
NDEMGLYQEPVFSDQKATQSSMPKPGVDIAENFSNGRHCSSSEKASDKRQTKIEKTITLQILRQYF
AGSLKDAAKSIGVCPTTLKRICRQHGGINRWPSRKIKVHDHSRKLQLIVNSIHGAEGLIQIDSFYKSFEL
SSSKFSGNDPFSSSRMSENCSILKPQPRSLFSPKGKVPKSRSSCSQNSGSTDEEKRLLTINGLNTGCS
LAIDDPVLMKAKSKPELHNLNQEEKEPKVLKSSLHKT LHGNTTGHSLQDGMOVFRVKATFGEENIR
FSFLPNWDFRDLQQEIAKRFKIDDFSRIDLKYLIDQEPVLLCDADLEECIDLLRLSQCNTIKISLHQAS
KSNLGSSFRIKGLF

>MeNLP5

MEDGGFKTDSVSRNLPGAAMDLDLFMDELLYDGCWLTDGLNISQPGLOSSSTVNDSSQYFPFVDTN
LHQQINQDGMENNFPENPPLSNPKVEQLADDQDHKSVKITTSSVSAGFPNEGSELRSRFWIEPRA

DPGPTSSVKVRLMHAIIRLKECTKDREVLVQIWVPTKKEGHVLTTDQPCFVSLKSESLARYRYVSEN
YHFPVEGDSKESLGLPGRVFLGKLPESTPDVRRNDEYPRKTYAKQYNISGSLAVPVFERGTGTCGV
VEVITTRYINYRSELETICKALEAFDLRSSQDFCOPPSVKACKEFCQAAVPEISEILGVCETHKLPLALT
ARCFQQGKGGCRHFDEKFANCISTVDSACFVDDRECLVFHAACSEQFLSGQGIVGRAFVTNKQCF
ASDVTAFSKTDYPLSHAKVGLHAAVAPILRSTHDGSADFVLEFLPKDCRDPEEHKQMQUELLPIAQ
QACRSLQVVMMDKELEASSLNEPYPLGSSWFAQMLEAQKKGKRVCVSDNPEEPKEEFKVATHWDE
GLEKLYEGQVFAESGSQLQQNSGRKDSTEGGNNSFAGQQSLGSRKTEKRRTKTEKTISLEVLRQYFA
GSLKDAAKSIGVCPTTLKRICRQHGISRWPSRKLKKKTFINDE

>MeNLP6

MENLFSLKEKGKGWTSRAQADGMAPSDGGAKNSISEDIFNSFSEMFNFDTYAGWCTSPSATDQ
MFASYGSSSFQSTPYSTFDALSFAEQSCLTSVGGNALNAAGTSYSSGDKMAVQQVNVYASDLMDA
DDLCAKERTGAQRQIEEMANCMISRPVGFSLDEKMLRALSLRESADGGILAQVWVPMRRGDQYILT
TTEQPYLDDQSLAGYREVSRTYTFSAEVTPLPLGPLGRVFISRVPEWTSSVYYIAEYLRGQHARNHK
VQGSIALPIFEPDVNCACAVLELTVKEKPNFDSEMENTVRLALQAVNLRSTAPPQLLPQSLSRNQRAA
LAEITDVLRAVCHAHSPLALTWIPCHYMEEAFDEVMKVRVREGNSRSGKFVLCIERTACYVNDREM
QGFVHACSEHYIEEGQGIAGKALQSNHPFFSDVKAYDITKYPLVHHARKYGLNAAVAIRLSTYTGD
DDYILEFFLPVNMGSSEQQLLNNLSGTMQRICKSLRTVSDAELKGGEGLSAVDFQKGAISSFPLSV
SITSSQTSSEAVLNLTDKVPLDASSSKYDAMKSDGPHEQSFNASRRQPEKKRSTAENVSLGVHQY
FSGSLKDAAKSIGVCPTTLKRICRQHGISRWPSRKINKVNRSLLKIQTVDLQVQVEGLKFDPPTGGF
VAAGSIVQEFDQSQRIFLSPDKNLPARNSQSATEEAVSVPSVPCIDGSNSAVKVEEDEFCMDTCGEVLM
KSSIPVIDCSEDSKSIATDAEICQKGRLCGPWAMDNASAFAKGTKGSLNSGSAKVDNSDTHFVSR
SSCSLGAAEVLDTKEEGDDVMVEHNQPTCSSMTESSNGGSMIHGSASSSPSFEEKHLVKCDDGGS
KISVKATYREDTVRFKFEPAGCFQLYEEVAKRFKLQNGTFQLKYLDDEEWVMLVTDSDLQECIEILDY
VGKRSVKFLVRDSVFTMGSSSSNCFLGGSS

>MeNLP7

MEDHILSPGTMLGAHADSTMFDYMDKLLLEGFRLESIDGSEFFNPSSSSAACVDSSFLWPASEIS
NGNFASSRSRNQEDEHILLPRIAPPNEAHGRSLVTAQAHGEDIGSAYRLGNTVDGSEVSRRWWIGPR
TSPGSKTSVRDRRLIRALGYIKDFTKDNNILIQIWVVDKGGRRVLTTHDQCFALVNCQRLANYRHVSI
NYQFLAEEDSKHMVGLPGRVFLGKPEWTPDVRFFRSDEYPRVHDAQLCDVRGTLALPFEQGSRT
CLGVIEVVMTKQKIKYHPELERICKALEAVNLQSSEVPSLQHVACDTSYQAVLPEIHEVLKSACETHKL
PLAQTVWPCIQQGKGGSRHSDENYYRCVSTDHACYVRDSGVQAFQEACSEHHLLKGQGVAGEAF
LTNQPCFTSDLTSYGKTEPLSHARMFGLHAAVIRLSVHTGTADFVLEFFPADCKDPEEQKKML
TSLSIIIQEVCQSLRVVTDKELEEENDLPISEVLALSYGTPDEEMFRVTQPYSESYHGDNLWSDSFPGI
QSGSAVSSFQIEKQKLLIHEKSVDRCQTQEDYSPKVSVEYGGDSAIAEGSFSSVCVSKTGEKRRTKGE
KTITLQVLRQYFAGSLKDAAKSIGVCPTTLKRICRQHGIKRWPSRKIKVGHSLRKLQVVIDSVQGASG
AFQIDSFTKFPELASNLSRSTPLSTSQQLGHPESSSIQPEGGIFSSQVAAPKSPSSSSQSSSSHCCS
SGTEQHASTLTIPTCKDMLGENSGNCVLGVKSDAELHVSIQEEQNLPRSQSQSLWEQPNLGNL
PPLAKSSNRICQEIDGQRVKVTYGNFKIRFRMPNNWELKDLLQEIARRFNIDDIHYDLKYLDDSEW
VLLTCDDDLEECIDLCQSSQSHTIKLLLQISPILLDRSTNSRGLS

>MeRKD1

TKDNERKRSAPLELEEIRKHFDVPITKEAKEMKVGLTILKRSRELKIMRWPHRKIRSLKSLINNIKEMGLT
NEIMMLEEHQRLEKKPDMEFSETTKKLQRQAFKANYKKRCLVAHHI

>MeRKD2

MDFSQDYSLTALRIFQNTINREFIRSLHVYRLKDGTDEVEREFLSDDGPYVEMVANPLLRLDRFRVLE

LFEGQVIGVWHCIFAFNAHHSPHLSRIPSLLSISRNPKLKSPTLANDLQLIFKLISRTADEEPLQFLSEEK
CRMIGCFQSQRNLPVLDQDLNCLPYSVATSQVPKSQQIEPSEPASVMAKKKKRAATEDIARIALEDL
VKYFDLPIAEASRNLKVGTVLKRKCREFGIPRWRHKIKSLDSLRLNLQEEAERQKQENEDAAMAVAK
RQKMLEREKESIERKPFMEIQSETKFRQDVFKRRHRARALKTQGLRVSQASA

>MeRKD3

MDSKITVELMNFPKNDTDCFLMDEYFERLSELSPLESFLEFDSLSPYDTNFGFEDYLLENGHPLSWDV
DIPIDAKPLRVFTSIDNTVKYDDDFTGTSSNEEKRVIMGRKRSAPLEMEEIRKHFDKPITKAAKEMKV
GLTVLKKRCRELSIMRWPHRKIKSLKSLRNVKEGLTNEAVMLEEHQLLEKMPDMELNDSTKKLRQ
AIFKDNYKKRRCLEAH

>MeRKD4

MADPRYVVYPYHDPHDSPINYEILNFLRDANPTLENLPSTDEPTSLHVSSDDIVNPMLSLEDPMIWDIV
NQPNSGGQSQGEGPSLNERRATENLEQRYLDDGKPISWPPPAMPFQCTCCQVLREIIHTDGNC
TKLEIHGRLGIICHAVLEIKDQVMYEAQQADQTEGGNEAERGSRSTLAQQRERTGRLTLKDFEAYFHL
PIEEAAKIMNLCPVVKKICRRYGMTRWPHRKIKSIQRQISNLRANSNWNDPGERARAQAEIQRLEVEI
SNICSGVTN

>MeRKD5

MGLTLLKKRCRELGIRRWPHRKLMQLQTLIKNVQEMKKVEGKESEEKLREAIEILESERKMLEEIPDLQLE
HKTKRLRQACFKSNYKKRKLMMMSRPSSSGAIDYYGQINEDEDEEIKSLLADPVCHPTMMF

>MeRKD6

MANWQNELATQESFLAAYPLIESFVPDPFCASLEIENSTIHDQDVNGLGDWNEFSTLFDPQKQLL
SCDSARGSSRTEPFEEEKAKKCREEKVNNISKALSRQTISMFYMPITQAAKELNVGLLLKKRCREL
GIRRWPHRKLMQLQTLIKNVQEMKKVEGEESEKKLREAIEILERERKMVEETPDMQLENKTKRLRQACF
KANYKKRRLMGMDKPSSSSSTITNKMMNEDEENIEEYEEIISLLADPTSHTNMLF