

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

>AtNLP1

MEDDGGSDGGEONGGFSPNSSFGAFADTAMDLMDELLFDGCWLETTDSKSLQTEQSPSASTA
MNDNSPFLCFGENDPSQDNFSNEETERMFPQAQKFLLEAEVGKSWWIAPSASEGPSSVKERLLQAI
GLNEAVQDKDFLVQIWVPIQQEGKSFLTTWAQPHLFNQEYSSLAERYRVSETYNFPADEGMKDFVG
LPGRVFLQKFPEWTPDVRFFRDEYPRIKEAQKCDVRGSLALPVFERGSGTCGVVEIVTTQKMNYR
QELEKMKALEAVDLRSSNLNTPSSEFLQVYSDFYCAALPEIKDFLATICRSYDFPLALSWAPCARQG
KVGRHSDENFSECVSTIDSACSVPDEQSKSFWEACSEHHLLQGEGIVGKAFAETKLFFVPEVATFSKT
NYPLAHHAKISGLHAALAVPLKSGLVEFVLEFFFKAQCLDEAQMLKSLCVTLQQDFRSSNLFIK
DLELEVLPVRETMFSENLLCGAETVESLTEIQMQUESSWIAHMIKANEKGKDVSLSWEYQKEDPKELS
SGRENSQLDPVPNNVPLEAQQLQQASTPGLRVDIGPSTESTGGGNMLSSRRPGEKKRAKTEKTIGL
EVLRQYFAGSLKDAAKSIGVCPTTLKRICRQHGIMRWPSRKIKVGHSLKKLQLVMDSVQGAQGSIQ
DSFYTSFELNSPNMSSNGPSLKSNEQPSHLNAQTDNGIMAENPRSPSSCSKSSGSSNNNENTG
NILVAEDADAVLKRAHSEAQLHNVNQEETKCLARTQSHKTFKEPLVLDNSSPLTGSNTSLRARGAI
VKATFGEARIRFTLLPSWGFAELKQEIRRFNIDDISWFDLKYLDDDKEWVLLTCEADLVECIDIYRLTQT
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>AtNLP2

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NDNNNNNSYLYGYQYAENLSQDHISNEETGRKFPIPPGFLKIEDLSNQVPDFQSAVMSSAQAEKFLLE
ESEGGRRYWIAPRTSQGPSSVKERLVQAI EGLNEEVQDKDFLIQIWLPIQQEGKNFLTSEQPHFFNP
KYSSLKRYRDVSAYNFLADEDSKESVGLPGRVFLKKLPEWTPDVRFFSEYPRIKEAEQCDVRGSLAL
PVFERGSGTCGVVEIVTTQKMNYRPLEDNICKALESVNLRSSRSLNPPSREFLQVYNEFYAALPEVS
EFLTVCRVYDLPLALTWAPCARQGVKGSRHSDENFSECVSTVDDACIVPDHQSRHFLEACSEHHLL
QGEGIVGKAFNATKLFFVPEVTTFSKTNYPLAHHAKISGLHAALAVPLKNKFNSSEVFVLEFFFKA
CL DTEAQQDMLKSLATLQQDFRSNLNFIDKELELEVVFVREEVVFAENPLINAGTGEDMKPLPLEEISQ
EDSSWISHMIKANEKGKGVLSWEYQKEEPKEEMLTSGWDNNQIGSGHNNFLSEAEQFQKV
TNNSG LRIDMDPSFESASFGVGQTLLGSRRPGEKRRRTKTEKTIGLEVLRQYFAGSLKDAAKSIGVCPTTLKR
ICR QHGITRWP
SRKIKVGHSLKKLQLVIDSVQGVQGSIQLDSFYTSFELSSPHMSGTGFKNPNAQT
ENGVSAQGTAAAPKSPPSSCSHSSG
SCTGANQSTNTGTTNTVTLMAENASAILKRARSEV
RLHTMNQDET
KSLRTSHKTFSEHPLFENPPRLPENSSRKLKAGGASKV
KATFGEAKVRFTLLPTWGF
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SIGLGPSL

>AtNLP3

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LQEVMGERELLI
QLWVPVETRSGRVLSTEEQPYINTFSQSQLALYR
DASAGYSFAAEVGSEQLVGLPGRVFLRRMPE
WTPDVRFFRKEYPRIGYARRYQVR
ATLALPLFQGTSGNCVAVMEMVT
THRNLEYASQLSTICHALEA
FDLRTSQTSIVPASLKV
TSSSSSRTEVASILQGICSSHGLPLAVTWGH
HQDSSSCLSALISASYAADHG
SRCFLAACSEHHLLGG
EGIAGRAFATKKQCFATDVAIFSKWSYPLSHYAKMFDL
HAALAVPILTRGN
RTVQFVLELFPRDCL
DIQTHSLT
LASQLKLFQSSPHLMVDDNQIA
EEVRDAATPPLTQEDPKGKQV
SFSFSSASSLENRK
RKT
KA
EK
DIT
LDT
LRQHFAGSLKDA
AKNIGVC
PTTLK
RICRQ
NGIS
RWPSR
KIKV
GHSLRK
LQV
VMD
SVE
GV
QG
SLH
LAS
FYSS
FP
QL
QSS
SS
SF
P
NPT
QTV
HV
PP
K
SPP
SS
GS
QSS
SG
STCC
SSEE
QQL
GGF
QK
PAL
SHP
QLL
TSS
MH
EDQ
RP
VR
VT
SSL
P
L
PS
ATT
PR
KA
K
DGM
K
VK
AM
FGD

SMLRMSLLPHSRLTDLRREIAKRGFMDDVLSNFSLKYLDDDQEWFVLLCDADLEECIQVYKSSLKET
IRILVHHPLSRPSFGS

>AtNLP4

MEDSFLQSENVVMDADFMGDLGCWLETTDGSEFLNIAPSTSSVSPFDPTSMWSPTQDTSALC
TSGVVSQMYGQDCVERSSLDEFQWNKRWWIGPGGGGSVTERLVQAVEHIKDYTTARGSLIQLWV
PVNRGGKRVLTTKEQPFSHDPLCQRLANYREISVNYHFSAEQDDSKALAGLPGRVFLGKLPEWTPDV
RFFKSEYPVHQAQDCDVRGTLAIPVFEQGSKICLGIEVVMTTEMVCLRPELESICRALQAVDLRSTE
LPIPPSLKGCDLSYKAALPEIRNLLRCACETHKLPLAQTVSCQQQNKGCRHNDENYIHCVSTIDDA
CYVGDPTVREFHEACSEHHLLKGQGVAGQAFNTNGPCFSSDVSNYKKSEYPLSHHANMYGLHAVA
IRLRCIHTGSADFVLEFFLPKDCDDLEEQRKMLNALSTIMAHVPRSLRTVTDKELEEESEVIEREEIVTPKI
ENASELHGNSPWNASLEEIQRSNNTSNPQNLGVFDGGDKPDGFGLKRGFDYTMDSNVNESSTF
SSGGFSMMAEKKRTKADKTITLDVLRQYFAGSLKDAAKNIGVCPTTLKRICRQHGIQRWPSRKIKKVG
HSLQKIQRVIDSQGVSGPLIGSFYANFPNLVSQSQEPSSQAKTTPPPPPVQLAKSPVSSYSHSSN
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>AtNLP5

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VPVDRGGKRVLTTKEQPFSHDPMCQRLAHYREISENYQFSTEQEDSDSSRDLVGLPGRVFLGVPE
WTPDVRFFKNEEYPRVQHAQDCDVRGTLAIPVFEQGSQICLGIEVVMTTQMVKLSPDLESICRALQ
AVDLRSTEIPPPSLKGPDFSYQAALPEIRNLLRCACETHKLPLAQTVSCLKQSKTGCRHNDENYIHC
VSTIDDACYVGDPTVREFHEACSEHHLLKGQGVGEAFLTNGPCFSSDVSSYKKSEYPLSHHATMFG
HGTVAIRLRCIHTGSDFVLEFFLPKNCRDIEEQRKMLNALSTIMAHVPRSLRTVTQKELEEGDSMVS
EVIEKGVTLPKIENTTEVHQSIQPNVGLVFDGGTEMGELGSEYGKGVSVNENNTSSASGFNRVTE
KKRTKAENKNITLDVLRQYFAGSLKDAAKSIGVCPTTLKRICRQHGIQRWPSRKIKKVGHSLQKIQRVIDS
VEGVSGHHLPIGSFYASFPNLAAASPEASSLQQQSKITFLSYHSPPAKSPGSSCSHSSCSSETQVIKE
DPTDKTRLVRSRFKETQTTHLSPSSQEDDFLRVKSYEEEKIRFKMRNSHRLKDLLWEIAKRFIEDV
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>AtNLP6

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RGRYVLTTSGQPFVLGPNSNGLNQYRMVSLTYMFSLDGERDGEGLPGRVFRKKLPEWTPNVQYYSS
KEFSRLGHALHYNVQGTLALPVFEPSRQLCVGVVELIMTSPKINYAPEVEVKCKALEAVNLKTSEILNHE
TTQICNEGRQNALAEILEILTVCETYKLPLAQTVVPCRHRSLAFGGFKKSCSSFDGSCMGKVCMS
TSDLAVYVVDAHVGFRDACAEEHLQKGQGVAGRAFQSGNLFCRDFVTRFCKTDYPLVHYARMF
KLTSFAVCLKSTYTGDDEYVLEFFLPPAITDKSEQDCLLGSSLQTMQHYSSLKVSETELCENNMSLE
VVEASEDMVYSKLEPIRIHHPAQISKDYLELNAPEQKVSLNSDFMENNEVDDGVERFQTLPIPEAK
TVKKSERKRGKTEKTISLEVQQYFAGSLKDAAKSLGVCPPTMKRICRQHGISRWPSSRKINKVNRSLTRI
KHVIDSVQGADGSNLTSLSRPWPHQIPPIDIQLAKNCPPSTSPLSNLQDVKIERDAEDSAGSSTS
RASCKVNPLICTRFLPTHNQEPESRQVALDDSDSSKNMTNFWAHLTCQDTASPTILQHKLVSIKATY
REDIIRFKISPESVSITELKQQVAKRLKLETAAFELKYLDDREWVSVSCDADLSECLDTAAKANTL
RLSVHDVTFNFGSSCESSEETMMCL

>AtNLP7

MCEPDDNSARNGVTQPSRSRELLMDVDDLDGSWPLDQIPYLSSNRMISPIFVSSSEQPCSP
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VPPENTDNYCVIKERMTQALRYFKESTEQHVLAQVWAPVRKNGRDLTTLGQPFVLNPNGNGLNQ
YRMISLTYMFSVDSESDVELGLPGRVFRQLPEWTPNVQYYSSKEFSRLDHALHYNVRGTLALPVNP
SGQSCIGVVELIMTSEKIHYAPEVDVKVCALEAVNLKSSEILDHQTTQICNESRQNALAEILEVLTVC
THNLPLAQTWVPCQHGSQLANGGLKKNTSFDGSCMGQICMSTTDMACYVVDAHWGFRDAC
LEHHLQKGQGVAGRAFLNGGSCFCRDITKFCKTQYPLVHYALMFKLTTCFASLQSSYTGDDSYILEFF
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KSNATEMLIPQPVVQSSDPVNEKINVATVNGVVKEKKTEKKRGKTEKTISLDVLQQYFTGSLKDAAK
SLGVCPTTMKRICRHGISRWPSRKIKKVNRSITKLKRVIESVQGTDGGLTSMAVSSIPWTHGQTSA
QPLNSPNSKPKPELPNTNNSPNHWSSDHSPNEPNGSPELPPSNGHRSRTVDESAGTPSHGSCDG
NQLDEPKVPNQDPLFTVGSPGLFPYSDHDVSAASFAMPNRLGSIDHFRGMLIEDAGSSKDLR
NLCPTAAFDDKFQDTNWMMNNNNNNLYAPPKEEAIANVACEPSGSEMRTVTIKASYKDDIIRFRI
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>AtNLP8

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

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QYMLSTCDQAYLLDLSRLSGYREASRRFTSAEANQCSYPGLPGRVFISGVPEWTSNVMYKTAEYLR
MKHALDNEVRGSIAIPVLEASGSSCAVLELVTCREKPNFDVEMNSVCRALQAVNLQTSTIPRRQYLS
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GDNDYILEFLPVSMKGSSEQQLLDSLSGTMQRLCRTLKVSDAESIDGTEFGSRSEMTNLPQATV
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FSGSLKDAAKSLGVCPTTLKRICRHGIMRWPSRKINKVNRSLRKIQTVDLQVQGVEGLKFDSVTGE
FVAVGPFIQEFGTQKSLSSHDEDALAR SQGDMDEDV SVEPLEVKSHDGGV KLEEDVETNHQAGPG
SLKKPWTWISKQSGLIYSDDTDIGRSEEVNKDKE DLCVRRCLSSVALAGDGMNTRIERGNGTVEPN
HSISSSMSDSSNNSGAVLLGSSSASLEQNWNQIRTHNNSGESGSSSTLVKATYREDTVRFKLDPYVV
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GSSAGSNGYLGTT

>AtRKD1

MKSFCCKLEYDQVFGKENNSFSFLNHSSLYSHQSELANPFFELEDEMPSATSSNCFTSASSFLALPDLE
PISIVSHEADILSVYGSASWAEETMFVSDFAKKSETTTKRRRCREECFSSCSVSKTLSKETISLYFYMPIT
QAARELNIGLTLKKRCRELGIKRWPHRKLMQLQKLISNVKELEKMEGEENEDKLRNALEKL
EKEKKTIEKLPLDKFEDTKRLRQACFKANHKRKRRSGMSTPITSSSSASASSSSYSSVSGFER

>AtRKD2

MADHTTKEQKSFSLAHSPSFHDHSSLSYPLFDWEEDLLAQENSGSQAFPTTSLPLDLEPLSEDVL
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VSRYFYMPITQAAIALNVGLLKKRCRELGIIRRPHRKLMNLISNVKELQKMEGEENAELQDAL
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>AtRKD3

MADQRPLMTWLEANNYESFLQEDIFSFLDQSLFVDPHSSIDPFKDFQTQNWFSLQDSIVNHISTTFA
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MKEKTSQKRIIMKRRYREDGVINNMSREMMKQYFYMPITKAAKELNIGVTLLKKRCRELGIPRWPWRK
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FSSSS

>AtRKD4

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NLKKKRQDKLEMSEIKQFFDRPIMKAAKELNVGLTVLKKRCRELGIYRWPHRKLSNLNSLIKNLKNVGM
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>AtRKD5

MVDQGFFTLKKEKKNILIKLFYVKILGFCFCDSWSSDMAHSLTSLAVFQSVIRKEMVRSLHVYESV
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LEDIPNLANELKFISELKPSKIYEEEQCSSTEGLYNSDLPKPRKLVLQDLNCLPDSETESEESVNE
KTEHSEFENDKTEQESDAKTEILKKKRTPSRHVAELSLEELSKYFDLTIVEASRNLKVGTVLKK
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KKFRQENFKKRHRASRAKKNQESLVTSST

>AtRWP-RK-1

MAGDDPKSSAPKDDGYEDPYINDPDFGFGPTDEDLINVMADINPNLMDMSFTEHTPPSVLPTLPSD
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DSTRLELSTRVFGGANGLDFEIGGSATMPVPVTTISDGVHACNCCLLRELVHLKEGEERSKLDIYGGI
GFICHAILLIQLLAPDSMQRQPLIFHLQHLMEEVKFIEDCYQRVANGLQLQDTNAAFYQAMSAN
SIFNQPPPMLPSSTDVPLSLLSPNEDHDVPLSLLSPEEALDIQPVPPHVELREEPTTRLPKQREKRQTP
LAAQVISLVSQFNI

>OsNLP1* in 2005 (OsNLP2)

MEQKPSPPPPRSDEEDGLMGCGMGGTDIAGGDLDLMEFLLATPGFDLSEFWHPGAASPFSPLF
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RIASQSQAQRGDGELLQVWWPTRIGDRQVLTCGQPFWLDRRNQRLANYRTVSMKYQFSADE
SARADLGLPGRVFVGRVPEWTPDVRYFSTEYPRVQHAQYFDIRGSVALPVEPRSRACLGVVELVMT
TQKVNVSAEINICNALKEVDLRSSDVSSDPRSKVVDASYRAIIPEIMDVLRAVCDTHNLPLAQTWIPCI

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

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

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

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

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

>OsNLP4 (**OsNLP4**)

MEEGDPQPSISLARTPSEGAAAVALDLLEQLLSADNAWLEVAANTSRSRPNFFATPSNCLTDASVAT
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RSNSLIQFREVSTKYQFSADVAGSSPGLPGRVFIGRLPEWSPDVRYFTSYEYPRINHAQYLDVHGT
GLPVFERGNYSCLGVIELIMTKQKLNFTSELNTICSLQAVNLTSTEVSSIPRAKLN
SASYKDALPEILEV

LRAACITHKLPLAQTWVTCAQQGKGRSRHSDENYKYCISTIDAACYVNEPRMOSFHEACSEHHLLRG
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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 LAWIPVCSK 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 QVGKNKTRGKA EKV TLEELQKHFSGSLKDAARSLGAGY SV DYLHQNP DVY AQIN
TQPLKVGSPGSMASQGHFARSARCGNGDGVTIKASHRGDIIRFRVPCSAGVAAVKA VAKRLSL
DAGAFDV KYLDDHEWVLLSCAD FQE CLDV VPALPSMSVTARSGSGAAAPVV VRLMVQE VADNI
GSSCASSD

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

MCAAPMASTAQPQQQPQQQQEQQPVAAA AVPTPAPPASEAQPKPTRVSLSYEEISKLFSLPIAEA
ASILGVCTS VLKRICRHGIVRWPYRKL VSGKGDDTKNAEREKAGLLEISKVAKQKALSASGLSTVSP
GAFQGVAKSQGSSKAGQVSPPGKQNVLGGSA ILSYGTQTKGIP TYMDDFKYGFPSGLSLQTMKW
WGTD SHTETTPAKDDNGEAPESANEASKGMTDELDWGADEAE AEA DADS AITTEPSAQLCSLRRK
AVDDGRKLLTGKSCGGLECLRLNKRQKM ALAQVFGASLPEQLRSKLG

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

MEMHECCYYGGSIGGDWLNLPLAIPPPCSSSSSWSSQLLLSDHDDVLLHSAGDHGGA VAGIGGA

CMTADLVRDEEMEMAAGYLPVAASAAAAADVHDHYMYQQFQLEPDQFVSTLPAVAVAATAG
GGSHDDELLRMPFTIDLDCAFADARDVVGVGEPKPSPQHTLAAIALPAVGGGAHHFGTQDDD
VKFDVTKQRNDAALAGDDLSMVIVESYEMGMRRHAAEQQEQQPKIITSAAATLPLPLPPPPPPP
RVTRSRRDGSSAATAGGKTRLDHIGFEDLRRYFYMPITKAAREMNVGLTVLKKRCRELGVARWPHRK
MKSLSLILNEMGSKGMSSAAMRREALENCCALMERNPAVELTERTKKLQRACFKENYKRRRAAA
VDVLDDHCFSAAGHCHRHHHQQLALPPPPAAAADHRRRDFFGY

>OsRKD4 in 2005 (OsRKD5)

MDAAAAAVTLSALAVFASTLDHGAVRSVHGYKVYGRGGRRWERWVEREFLTPASCREVPAPVA
PPRILPAEWGRGPAYREGQVVAAGAWRCILAFDSAAPPRTPPPVLSPFLNPRLMCVPSLYNDLEKVF
RFQNVEKIPKLMQCDSEEKSSWDARDKSSDEVHASESDSDDLQSSEEEKPTVQKQRRANKKHIAS
ITLVDIAQYFHLPIREASRTLKGVSILKRKCRQYNIPRWPHRKIKSLDSLIQDLEYVIDGDDHDDTGDD
VQQEKKQTAEEKQEAIMALTRRKQMLETEKETIQQIPAMDLKETKQFREDVFKRRYRAKKDLAND

>OsRKD5 (OsRKD2)

MRHNRSLAALLRAGRYGAARRLFDALPARSVVTWNSLLAGLARRPDARAAREFFDAMPVRDAVSW
NTLLAAYSASPHPDHLAARRLFDEMPQRDVVTWNTLLGAYARRGLMDEARRLFDEMPQRNAAS
WNTMVTGFFAAGQVVKALDVFDAMPAKDSASLSTMVGFTKNGMLHEAEELLTKRLSVTDMDKAV
DAYNTLIVAYGQAGRFSDAKRLFDMIPKGQYQHNMLKRKGFERNVWSNSMMICYIKAGDVCSAR
ALFNEMPDKDLVSWNTMISGYTQASDMKESEKLFWEMPPDTVSWNLIIQGFMQKGEAEHARGFF
DRMPERTISWNTMISGYEKNGNYISSVKLFSKMLEVGEIPDRHTFSSVLAACASIPMLGLGAQIHQLV
EKSFPDTAISNALITMYSRCGALNDAEAIFKQMIHTKKDLVSWNALIGCYEHGRATKALQLFKEMRR
AKVMPHTITFVSLAACVNAGLVSEGMRVFDTMVHEYGIVARIEHYAALVNLIGRHGQLDALEVINS
MPMAPDRSVWGAFLGACTAKKNEPLAQMAAKELSTINPDSSAPYVLIHNLHAHEGKWGSAAVRE
EMERQGIYKQPGYSWIDLEGKMHVFISGDTWHPNAQEIFSVLEDWQWHNPMSLEMNQLTQVKLIA
KRNNQIMDKQATATKMHKATNIRHIYSKFKVDEQQNTSWWPYCTSLWPDSYLLEEAFLSSLSFPSF
HPQPVYSTVMQSNVLQDELGVIFEDDVLYWDEMEQSENKVEKSEKGLPLYYGDENGAASKIMRD
DVRSEEKALTfelvsqyfympitqaarelnvgltllkkcrelgiprwhrkmkslqtlinnvqlqea
skanneeqlrmlveumlqeerrlleqkpyvqleektrklrqacfkanckrrllaleagep

>OsRKD6 (OsRKD2)

MEKWRFYGGHQFQSYSFHEEENLFQDWSDLYLLLGEDEPFTHHFSTSVDHSNFVQDELYTLFDGDILS
IWGDMKEDAYHRSDKDGGEKEEKLHHEKAMELQLQRLPSGRQSGEKTTFELVSQYFCLPIKQAAQE
LNVGLTLLKRRCRVLCIGPRWPHRKVKSLETIKNVQELGMETGQDEDNTRNAVEMLQQTKKLIEQSPD
AKLDDWTKMLRQACFKENYKRRLLAIEG

>OsRKD7 (无)

MAMS RATCSTRPRISRCSPSRRWFVPAAAIESHPTAIHVVEREWPQVQISRFDPSCPPPARMLP
GGQQQQPHPAPGGLDLRSDEIHALKQRRRPPNPQSSSPPEGLLGAANPDRPDDDELAILSSFRDE
AAHGDGGGVARVQVLPVEQDANLPSLPLRHGQLDCSRCHLVRHVMHAVLPYVKFSLKLVWEE
AFHRIRRMYVISVRSNVYSLSKWTQEWAIFIARNIDTMRNNTNGQLLDSGYSNLVESVRTNVNPH
TADEVNLLQTIMSAPSADHHQNAADQVAAPAAQPSAAPPVALPPKAAPRKARKDRDYASMLVA
EEFYVAATSRPVPSNSDVEILESSHVSQQQDGGRAIYPSLQARRGKTKQEVPRRANKDVLEYLSLARKE
TEKEINTLSSFDGIYRNDGTLSYLMTEVRRLNKIWLQKNAPSTLSSRLLASVKEIDDICKVEKGRLYAQF
ISALKKLCRKMKDDGGSAPSANN

>OsRKD8 (OsRWP-RK2)

MAGDGGNGAADGGGEHELEDFPSPLLDNFVLELLDSLGVDDDGAPPAPAPVQMSVDGAG
GGRINKPGSSVDDLVDWSTAFCNNCSKIDGEYGGASTSSAAPALAPPQEDYCSGCQVLREVHNS

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
MHRSVLAFLDGLKKSCTSFDGSCNGQCMSTTDVAFYVVDPMWGFREACLEHHLQKGQGVAGR
AFLSHNACFCPDITQFCKTEYPLVHYARMFGLSGSFAICLRSSYTGNDNYVLEFFWPLSISDNYEQN
LGSLVATMKQHFQLKVASGMDLEEEGFVIIQTSMSGRDLRLECIRIPQSPKSPPNLNTSPKEGKT
AQLDPLKHHLMVNLDADVNGGNISQAEGSHFPTSLPENKGNNKPSERKRGKTEKSISLEV
LQYFAG
SLKDAAKSLGVCPTTMKRICRQHGISRWRPSRKINKVNRSLSKLKRVIESVQGAEGAFDLTPL
SPLPVTV
GSISWPSNLNGCNQQNSLNSKLEPIDEKNGSPICNAAGNDQGGGEAQLLGGRILSREEAVLQNG
FSPKIGLVSDFRKAGNASGEESTGPTSHGSCQGPANDSAPANDSPVSPVHEKCKNAGNTPELAFQ
TKGKLNLSAAYSVPDALVATEAQAPLGEMLIKDAGSSKDLRNLCPSVADTILEERVPESSWTNYPGP
FPTTEAVAALEHAMTQGTRQEMMTITIKAAYREDIIRFRISLSSGIVELKEEVAKRLKLEV
GTFDIKYLD
DHEWVLIACDADLQECIDISRSGSSIIRLSVHDVNTNLGSSCESNGE

>**MeNLP2**

MAEQEDDKSKDNQSVAGNKGETLMDLDDLDTSWPLDQIYFLSNPLSPLLLSSSDQPCSPLWAFSD
ADDERLAASSSHATHPLRSLDYPPIFLTCNPSSLAESQTENDDKRKFPSPLLGLMPFDNPDGYCIKER
MTQALRNFKESTEQHVLAQIWAPVKNGARYVLTSGQPFVIDPHSNGLHQYRMVSLTYIFSLDGEFE
CEPGLPGRVFRQKLPEWTPNVQYYSSKEYSRRDHALHYNVQGTALPVFEPFGQSCVGVIELIMTSQK

INYGPEVDVKVCKALKAVNLKSSEILDHPSTQICNEGRKNALAEILEILTVCETHKLALAQTWIPCMHRS
VLAFFGGLKKSCSFDGSCNGQCMSTTDVAFYVVDPMWGFREASLEHHLKGQGVGRAFLSH
NACFCPDITQFCKTEYPLVHYARMGLTGCFAICLRSSHTGDDEYILEFFLPPGISDFQQKALLGSLLA
TMKQHFQSLKVASGMDEEEGFVEIIQTSVSGKLDLRLECIQTPESPESLPNVNTSPKEGQMAQVDP
VVNLDDVNNGGSTGDAEGGHFPTSPENPKGNKKPSERKRGKTEKTISLEVLQQYFSGSLKDAAKSLG
VCPTTMKRICRQHGISRPSRKINKVNRSLSKLKVIESVQGAEGTFDLTPIATSPLPVAFGSISWPSNL
NGCNQQNSPNSRSSEPNERNGFICKVAGSEEQSGSENQLPGGRVISHEELFLQNCFSPEIGQGSS
RHKAGNVSREESAGTPTSHGSCQGSLANDIAPAKVASASPQEQQNEVGTIPELSFQPKGKPNLALSAL
YIPDAFVATEAQAPFRERLIEDAGSSKDLRNLCPSVADTMILEERVPEYSWTNHPGPEFPTKQSMAAL
EHATPRVTARQEMRFVTIKATYREDIIRFRISVGSGITELKEEVAKRVKLEVGSFDIYLDLDDHEWVLIAC
DADLHECIDISRLSGGNIIRLSIHVDVNTNLGSSCESNGELLGCTYGKV

>MeNLP3

MLGTRADSAMDFFDCMDKLLLEGCWLETIDGSEFFNPSPFSSAAFVDASFQWPTSEINNCCLASSPSK
RSNREEQISLLPRNSALNEAHGRSLINAEDALDHGVGSADRLGYNTTEGSEVSRRWWIGPRTNPGPKT
SVRDRLIRALGYIREFTKDKDVLIQIWVVPDKGGRVLTTHDQRFALVPNCQRLANYRDISTNYQFSVE
EDSKDMVGLPGRVFLGKVPWEWTPDVQFFRSDEYPRVDHAQQYDVRGTLALPVFEQGSSTCLGVIEV
VTTKQKIKYHPELESVCMALKAVDLQSSEVPSLQHVKMCDMSYQAVLPEIHEVLRSACTHKLPLAQT
WVPCIQQGKGCCRHSNNYYRCVSTIDLACYVHDGVQAFHEACSEHHLLKGQGVAGEAFLTNQP
CFSSDITSYRKTEYPLSHARMFGLHAAVAIRLRLHSGAADFVLEFFLPAYCTDPEKQKEMLTSLSITV
QQVCQSLRVVTDKEEETGCPISELLASSVDTPPKEEMLVTPQPSVSYGGESLSHADRLTGQQSGSV
VSSCQKKKQKVLEDQISVDCWQNQEDYSLKVSVESGGDSAVAEGSFSSVFKGKKGEKRRTKAEKSITL
QLVLRQYFAGSLKDAAKSIGVCPTTLKRICRQHGIKRWPSRKIKVGHSLQKLQLVIDSVQGASGSFQIG
SFYTNFPELASPNLSRSNLFSTLKQSDHPEPSSIQSERYFSSQAATPKSPSSCSQSSNSSHCISSGTQL
HASAVTTPTSEDPVLGENDNCVLRARSDAELRASIQEQQNLLPRSQSQKSLGEQPNLGNLLPLPNN
SSSIIRDFTQRVKVTYGNENKIRFRMPNNWELKDLLQEIAARRFNIDDIKYELKYLDDESEWVLLCDD
DLEECIDVCQSSQTHTIKFLLQVSPHLFDRSMNSRGLS

>MeNLP4

MDHGASALNSTWTFSSTTMDLDVMDKPLSQGYCLENNDRANSLEPEPFTSHVLYDHSEYSSFED
LYHQIYRKEMKPLFENIPSAYHGIEGAETAKSFVQSESFTLEGSNSGSRWLIDPSYSVKQRLMLAIKYLKE
YVEDSDVLIQIWVPTQIGGQNVLTIDQPYSLNPSCSKLASYRNVSKTFQLTAEENLKELAGLPGRVFL
GKLPEWTPDVRFFRSDECPRKSYAKMYNISGCLALPVFEQDNGVCLAVVEIVTTQKISYSLELEIVGRA
LEAVDLKSSQDFPLGIKACNESYQISILEISDILQFVCKTHGPLAMTWALRDRQGEVEHWQFSKKYD
YCLSTVDSACYVADSDLLGFHEACSEHYLFPNQGIVGKAFTKKQWFATDITSFSKANYPLSHARMF
NLRAALAVPLCNIYTGLIEFVLELFPWDCKDIEEQKHFWDMLSIVMQQACKSFHVIVDKEIDEVSEQ
MVVASDGRYNREGIQLVELVAPSNESSQEESSTPHEDSQNKDKFSISWECPKEEDEFKVITHWE
NDEMGLYQEPVFSDFKATQSSMPKPGVDIAENFSNGRHCSSSEKASDKRQTKEKTTLQILRQYF
AGSLKDAAKSIGVCPTTLKRICRQHGGINRWPSRKIKVHDHSRKLQLIVNSIHGAEGLIQIDSFYKSFEL
SSSKFSGNDPFSSSRMSENCSILKPQPRSLFSPKGKVPKSRSSCSQNSGSTDEEKRLLTINGLNTGCS
LAIDDPVLMKAKSKPELHNLNQEEKEPKVLKSSLHKT LHGNTTGHSLQDGMOVFRVKATFGEENIR
FSFLPNWDFRDLQQEIAKRFKIDDFSRIDLKYLIDQEPVLLCDADLEECIDLLRLSQCNTIKISLHQAS
KSNLGSSFRIKGLF

>MeNLP5

MEDGGFKTDSVSRNLPGAAMDLDLFMDELLYDGCWLTDGLNISQPGLOSSSTVNDSSQYFPFVDTN
LHQQINQDGMENNFPENPPLSNPKVEQLADDQDHKSVKITTSSVSAGFPNEGSELRSRFWIEPRA

DPGPTSSVKVRLMHAIIRLKECTKDREVLVQIWVPTKKEGHVLTTDQPCFVSLKSESLARYRYVSEN
YHFPVEGDSKESLGLPGRVFLGKLPESTPDVRRNDEYPRKTYAKQYNISGSLAVPVFERGTGTCGV
VEVITTRYINYRSELETICKALEAFDLRSSQDFCOPPSVKACKEFCQAAVPEISEILGVCETHKLPLALT
ARCFQQGKGGCRHFDEKFANCISTVDSACFVDDRECLVFHAACSEQFLSGQGIVGRAFVTNKQCF
ASDVTAFSKTDYPLSHAKVGLHAAVAPILRSTHDGSADFVLEFLPKDCRDPEEHKQMQUELLPIAQ
QACRSLQVVMMDKELEASSLNEPYPLGSSWFAQMLEAQKKGKRVCVSDNPEEPKEEFKVATHWDE
GLEKLYEGQVFAESGSQLQQNSGRKDSTEGGNNSFAGQQSLGSRKTGEKRRTKTEKTISLEVLRQYFA
GSLKDAAKSIGVCPTTLKRICRQHGISRWPSRKLKKKTFINDE

>MeNLP6

MENLFSLKEKGKGWTSRAQADGMAPSDGGAKNSISEDIFNSFSEMFNFDTYAGWCTSPSATDQ
MFASYGSSSFQSTPYSTFDALSFAEQSCLTSVGGNALNAAGTSYSSGDKMAVQQVNVYASDLMDA
DDLCAKERTGAQRQIEEMANCMISRPVGFSLDEKMLRALSLRESADGGILAQVWVPMRRGDQYILT
TTEQPYLDDQSLAGYREVSRTYTFSAEVTPLPLGPLGRVFISRVPEWTSSVYYIAEYLRGQHARNHK
VQGSIALPIFEPDVNCACAVLELTVKEKPNFDSEMENTVRLALQAVNLRSTAPPQLLPQSLRNQRAA
LAEITDVLRAVCHAHSPLALTWIPCHYMEEAFDEVMKVRVREGNSRSGKFVLCIERTACYVNDREM
QGFVHACSEHYIEEGQGIAGKALQSNHPFFSDVKAYDITKYPLVHHARKYGLNAAVAIRLRSTYTGD
DDYILEFFLPVNMGSSEQQLLNNLSGTMQRICKSLRTVSDAELKGGEGLSAVDFQKGAISSFPLSV
SITSSQTSSEAVLNLTVDKPLDASSSKYDAMKSDGPHEQSFNASRRQPEKKRSTAENVSLGVHQY
FSGSLKDAAKSIGVCPTTLKRICRQHGISRWPSRKINKVNRSLLKIQTVDLQVQVEGLKFDPPTGGF
VAAGSIVQEFDQSQRIFLSPDKNLPARNSQSATEEAVSVPSVPCIDGSNSAVKVEEDEFCMDTCGEVLM
KSSIPVIDCSEDSKSIATDAEICQKGRLCGPWAMDNASAFAKGTKGSLNSGSAKVDNSDTHFVSR
SSCSLGAAEVLDTKEEGDDVMVEHNQPTCSSMTESSNGGSMIHGSASSSPSFEEKHLVKCDDGGS
KISVKATYREDTVRFKFEPAGCFQLYEEVAKRFKLQNGTFQLKYLDDEEWVMLVTDSDLQECIEILDY
VGKRSVKFLVRDSVFTMGSSSSNCFLGGSS

>MeNLP7

MEDHILSPGTMLGAHADSTMFDYMDKLLLEGFRLESIDGSEFFNPSSSSAACVDSSFLWPASEIS
NGNFASSRSNQEDEHILLPRIAPPNEAHGRSLVTAQAHGEDIGSAYRLGNTVDGSEVSRRWWIGPR
TSPGSKTSVRDRLIRALGYIKDFTKDNNILIQIWVVDKGGRRVLTTHDQCFALVNCQRLANYRHVSI
NYQFLAEEDSKHMVGLPGRVFLGKPEWTPDVRFFRSDEYPRVHDAQLCDVRGTLALPFEQGSRT
CLGVIEVVMTKQKIKYHPELERICKALEAVNLQSSEVPSLQHVACDTSYQAVLPEIHEVLKSACETHKL
PLAQTVWPCIQQGKGGSRHSDENYYRCVSTDHACYVRDSGVQAFQEACSEHHLLKGQGVAGEAF
LTNQPCFTSDLTSYGKTEPLSHARMFGLHAAVIRLSVHTGTADFVLEFFPADCKDPEEQKKML
TSLSIIIQEVCQSLRVVTDKELEEENDLPISEVLALSYGTPDEEMFRVTQPYSESYHGDNLWSDSFPGI
QSGSAVSSFQIEKQLLIHEKSVDRCQTQEDYSPKVSVEYGGDSAIAEGSFSSVCVSKTGEKRRTKGE
KTITLQVLRQYFAGSLKDAAKSIGVCPTTLKRICRQHGIKRWPSRKIKVGHSLRKLVVIDSVQGASG
AFQIDSFTKFPELASNLSRSTPLSTSQQLGHPESSSIQPEGGIFSSQVAAPKSPSSSSQSSSSHCCS
SGTEQHASTLTIPTCKDMLGENSGNCVKGKSDAELHVSIQEEQNLPRSQSQSLWEQPNLGNL
PPLAKSSNRICQEIDGQRVKVTYGNFKIRFRMPNNWELKDLLQEIARRFNIDDIHYDLKYLDDSEW
VLLTCDDDLEECIDLCQSSQSHTIKLLLQISPILLDRSTNSRGLS

>MeRKD1

TKDNERKRSAPLELEEIRKHFDVPITKEAKEMKVGLTILKRSRELKIMRWPHRKIRSLKSLINNIKEMGLT
NEIMMLEEHQRLEKKPDMEFSETTKKLRQAIFKANYKKRCLVAHHI

>MeRKD2

MDFSQDYSLTALRIFQNTINREFIRSLHVYRLKDGTDEVEREFLSDDGPYVEMVANPLLRLDRFRVLE

LFEGQVIGVWHCIFAFNAHHSPHLSRIPSLLSISRNPKLKSPTLANDLQLIFKLISRTADEEPLQFLSEEK
CRMIGCFQSQRNLPVLDQDLNCLPYSVATSQVPKSQQIEPSEPASVMAKKKKRAATEDIARIALEDL
VKYFDLPIAEASRNLKVGTVLKRKCREFGIPRWRHKIKSLDSLRLNLQEEAERQKQENEDAAMAVAK
RQKMLEREKESIERKPFMEIQSETKFRQDVFKRRHRARALKTQGLRVSQASA

>MeRKD3

MDSKITVELMNFPKNDTDCFLMDEYFERLSELSPLESFLEFDSLSPYDTNFGFEDYLLENGHPLSWDV
DIPIDAKPLRVFTSIDNTVKYDDDFTGTSSNEEKRVIMGRKRSAPLEMEEIRKHFDKPITKAAKEMKV
GLTVLKKRCRELSIMRWPHRKIKSLKSLRNVKEGLTNEAVMLEEHQLLEKMPDMELNDSTKLRQ
AIFKDNYKKRRCLEAH

>MeRKD4

MADPRYVVYPYHDPHDSPINYEILNFLRDANPTLENLPSTDEPTSLHVSSDDIVNPMLSLEDPMIWDIV
NQPNSGGQSQGEGPSLNERRATENLEQRYLDDGKPISWPPPAMPFQCTCCQVLREIIHTDGNC
TKLEIHGRLGIICHAVLEIKDQVMYEAQQADQTEGGNEAERGSRSTLAQQRERTGRLTLKDFEAYFHL
PIEEAAKIMNLCPVVKKICRRYGMTRWPHRKIKSIQRQISNLRANSNWNDPGERARAQAEIQRLEVEI
SNICSGVTN

>MeRKD5

MGLTLLKKRCRELGIRRWPHRKLMQLQTLIKNVQEMKKVEGKESEEKLREAIEILESERKMLEEIPDLQLE
HKTKRLRQACFKSNYKKRKLMMMSRPSSSGAIDYYGQINEDEDEEIKSLLADPVCHPTMMF

>MeRKD6

MANWQNELATQESFLAAYPLIESFVPDPFCASLEIENSTIHDQDVNGLGDWNEFSTLFDPQKQLL
SCDSARGSSRTEPFEEEKAKKCREEKVNNISKALSRQTISMFYMPITQAAKELNVGLLLKKRCREL
GIRRWPHRKLMQLQTLIKNVQEMKKVEGEESEKKLREAIEILERERKMVEETPDMQLENKTKRLRQACF
KANYKKRRLMGMDKPSSSSSTITNKMMNEDEENIEEYEEIISLLADPTSHTNMLF