

P1 (g11108)
LDSDGNGKVVYNITGGNEGAFKIDPDTGVIVKVRSLTTSASTFALDVEAKDKGNPPRSNIAKVNLNVFLPDGPPKFVV
KPVVEEVTEGIKANNRVMVVAATSEALTYEVVSGNEDGLFRINPSTGEIMVTRELDYEEARERRLVVRVMDTRDRSDQV
TVILOIKNINDNAPTFPGEVNGYVERKVEDDFQIGDPVARLSAFDGDAGDKITYTLSEANASFTINEEGFLIAKKPRKE
FSSPVTFDLTAKDNGSPRESKVNVLVLSYRPEQQHPVREYREDKEVGSVVAIVPRYFPGGTLSIVYPRKANFTVDN
SGRVRMTPFDFAEQLYLTVREQEPPLSRTNDVDEINVIDINDNPKVMKVEFFGRVNTNSRPGTSAYQLKAEDKD
GGLSGRIGYQMVSRGVPFAINPLTEVVETGGKLEDRGGYNVTVFDFGVPRQFGDSVYLDIKTVNFKPQFSESSYNFEV
FENDLPGK**VIGVNVATSASGARLGFSIVQGDTDNKFSIESTGELVNSKLDREKTSMYNLKVRTTEQIPRGYSNDVDVRI**
SVKNANEYYPYFPLTVYEKSINEDVGSGASVLSITAYDCDCTGCVCKNAELKYSLEGTSVFKIDEVTGLISVGDTPLDY
TQRHILFK**VVVVEDFGEKKFSSRSYVRRSVQNKNDEEPKFQRGEYSIDIAEDAINTNKALAAILARDGDGDSSLYTITSGAS**
GIFDINRNTGVLSLVQSVLGKK**QYTLTVRATDTAPNPKFDEVRVVINIEDINDNRPVFTCPKEVTVAENKPTGERVTO**
VTAKDSDNGRNGEVEYMLVTGGERLFEINNSTGLITTITSLDRETADRHTLIVQAEDGGHGRNEAERLLSYCIEVKAD
VNDNPVFFFTRQYYGSVFHKAPVGTTVLTVSATDDDTGNQKVYSLVGSNDKFEVDGATGSIRTIADLSNFQQTULLKI
RASNQATPTATKTSSSETTVEINVVNQOPPKFQPSSVYTAEIQEDVEVGTAVVTVKAVSQVDSKNGIAYSLVK**SYPSEAO**
NFKVDPGTGVISTASTLNYEQOSSYTLQIRARENNDLYATCSVNIRLKDVNDDTPFKLEEYSARVPEETVGGFNIVT
KADDRTLLSGEVSYSLGASQKFRIVASNGTIFSQEKFRENQTKHTVFTATDKGTPPLQAKVAVSVVVVDQNDPEPLF
PNRKYTANQEDAPIGTSIDDLTATDADIGDNARLEYFISGGEQSQLFRMETVYSPENKGILYNGKLDYETKPTYTIKV
TATDRKNSDTDVVIAVTKSWAH*

P2 (g11187)
MIVRDNDDDDDDDDDDDDDDDDDGVDDDDDDDDFSDNEEMLSFEVDELEEKDVGNDVDTM**QRHSVDSFDDVDFT**
FTKVTQAKYDGLPVTNVNLSATLPSSSSLEIMVYLFRAGKVTFGNETFRVEKGTIKFNIRISNWDFCDGSRFDCDEGK
IGEFLDLKLKIKSKDSPAIESDDRKAACRPICEDEDSDDGVDDPTSDNDDDDCPSIYDIGGSEMLLNKGVTDDD
KYTALPEGYPKFEVEDGEKKFVFRIPFKKSVLDPNVGRVNGVNSLAHANTGLAFLLFIAAFIVTNYV

P3 (g12510)
MSDATFILMKNAVTOIMDQYGTDRIHYSVIIFGSIVTTQVTFDRRFPEKNELIROVARYGKV**DGNPNLSAALEEKRIYE**
LQSVRPNARKVLVILDAPSASNTTAVKKAVTELDRKDVLVIGVGISSANENDLIGITKDDRHLINATLNKPYQEVAR
IIEAIFYGINTAGFSQWGDWSACSKSCRSGVAGTQERRICEIPRGCVGPTVETRECNIQNCEDCGEIGPLEGSYE
SSSAQPPRF**AKLNTTNPGVNKQGWCP**TAEADPEQYLOQINL

P4 (g9861)
MVVSOSITTVQESAPMEEFFNTOITEYEATLKSSADE**EGLOQEEILNLQTKLAKLQGDYDRAILDLAATKKCDLKVKK**
FEVQLQNSQKQRDDLKRQLEKLRDDLKKSREALSSAQTEAVNNKRAADRLRDEAEGKRNQIIRNLEQKNGSLEDQIEQLKO
ELSHSEMSAEVEAK**RAKDLENELRSSERRVAE**QEAIIESLRSTNEQRELAISSNRV**STLESQNGKMSTRIRELENELV**
RCHARIHDLENEFESVNTKCVQLOSSYDSSVSEVDGLRSRVGGHQSKVNELESRSVTSKERDDLKLKRISPLEKTLAELR
LTLERSENTCDGFEREIDLKAKNSNLETQLASANKLTQLRADIADAEDKDSNQLREITALQTOLGEAQRNLDFKNDN
VRKQKVIDDLRNTIRGLEAKIHAQENDVKOLEMQDMAEEEKEAMHDELLDNNRKLAEQEEESLRQAOQDKLDNVEKELIDS
QKNNEKLENTLENTEKEKRDFFENR**VQLQONLTR**VEAEELRDALKAVLEEDAKQHSAVVSKLTDEIDRLKGLLAEAQRO
IDAYRRESDSKDATITALKSQISDLKENDDLRAELSKVKTQLAAVEEEKRKLQIEHNTKNEEINRLTGEIDITVKRNEE
LQSECIVVADKVQSVESSYQDFMSLAGEQDGLLEGEKEKNAQLKKENNALKLKITSLEKKDRSQEVDQLRADLDAANEEI
PKLQDRIQQLVGENDDLSRIDLVVQSNLSQKENDLEAAEKDKEKAEDDVHIAQGKIDSLESQLAALRKQKVQLKDNLKDA
NNKIIKAKESNDLQAKVSQOEKIIIEGLKNGNTKKENIRELTDKIKKLQNELADAKREVSAAKLTAEGLRSDDKDLERK
LADQIKRIEALASEIKDLKAQRNSMETDLSNKTHKLALIESOLNSIKKERENMKEEVKDLNDRKLNLKEENLSFOTEIRR
LEKELEYKNKDNQDKDAVIERLKATKATVDEELLKKELMTQNLDAENKTKALENSQRQTNLSMNKLEVTIONLNR
EREERDRVGGSSHDSQYIEIDTLYKASQDRENRTKRELQAVKARLEVETDYYKIQQESVDRRAKVNLSLEKKVRLQAND
RDKKEKQAIRSDLEAASNQVADLEVAYDNEAKEKDSFKQLODALEKLAKARKQNLELEQTIVENKIQLEEAEKEMTEKE
NAINNLASS**KLEGQLDAAKSGLDKAQRE**IDELKSEDASKQNEILELHNKINELEARIVKLTGERDRALDDTAINADKI
ANLEQQIGELVSERTLLERIEDLKEISSLREERMEVNQKOLTDLOGNISFLTREVENRDKVITDLRNKEELQEKEIDNL
RKNLRRLESENESALQKISELRKAYESSSQKTIMIESKSDSHRDKAKRLDEENVSLKRRITELEITHNVNEEKQELQDK
LLRAQETIGNLESAPPMESSIVVETETNVEAAAGDIEIEFLKGDLNLKDKNAQLEKHLQTLDSNLKAQMAKQDQT
LKDNEDTILDERRIQELTIRLKTDQTDAPGIVSSLRNQKDSLERNVRMKELEAVKASQVDTQKRRADVENQLOQIK
RKLNEADFKLNMANQENLHQELLRAEKNRAIEAERESALQEQAQLRGGLNETNERLNRAKVDLANLKTQKRDLTGDL
EAVKDRRDDKESTINWHLGNERQLDEIASLKRKLDNMRDANEKLSHTVDNLNAQKQEREKIADEHHM

TLLRDEGEKLRRDLATANASVADFKKKYFNALKDIERLQTELNTKIQRIASLENRIKSLENAKSLLRNEISVHKNTISDQ
KVKLEASRRSDRIPTEITVTKVVGPSIQLSAPDMFQFTDDSVPSKKYKDLETSYKVLREKEDEGKDAATLKNRLAKLQE
LDETNRVKNAVEGDSMWKAKRISOLESSELNDLQSKDMVDSDEALGWKNKIPHLEQDLHAANNKIVRLEAEIETYEKMMKD
YSDDLKAKERVAQLESKVEDDQEKEIGQRKELMEAYKHEADLEAATKSCHNSKSELNGVIMTLRDKIGALENELOKEAK
QAHAVRGTLQDMKIREDDLKRKIAKLOKQLREVQQYDSQVQDWSERKYTIDGLRKENADYIATVEKLRRELGMYKNDLD
NMATEKRDQLQNELNELTPKMEQLETRLHKALEDRTFQIELDDTKRALDEVGRRESETVIROVSGGLEAGVDMFPELSN
DLESTKNERDRLASDVSSWQNKHDTLQNNYDNLYADKKRLEEKLD**SMQK**TMNQLDHNNQTLTSDKNRQTVELESAKRKIA
DLLSELDDAKRDKDALKVEYEIIQKKITKFEAEYEIHICHRHDYDDSGAYAKLEEAYKRSQERENEVQKELLVCYKTIGE
LKHQLENAHETMERLT**TVTYNTETIRSATLREEVVSYQRLSEEEKFEINRGANEDMENELTVLQRMEELSRENSSIQE**
AKALVLVEVTNQRNKIARLEQQLEDCESEKKSLRLQLTAMHSKGDAEAHEISKQISQLSEQSLTFOREKQNLERQLSQKH
AETISLQRENDALKSQEYSSSTEKQIESFFEMGGDADLGPSAESTRFEFDNDFEGGGTTVKRTTVKVRSSKKILLSK

P5 (g11674)

MQGKPNSNRVSQYKLSYSDDNRVYTPITDQVKKKNPINGGWEWTAWSACDPGCGNLRSRTRTCTNPVPVFGLDC**
VGANTITQFCPPNCAVDGQWSNWSAWSPCSKTCGPGSTWRTRDCVNPPPSNGGAFCKGQAFQLGSCNDGACPEATTAA
TTTEVTTAATTTEATTTEATTTEATTTEAPTTAATTTEATTTEATTTEAPTTAATTTEATTTEATTTEAPTTA
ATTTEATTTEATTTEATTTEAPTTAAPTNTAPTGIPLSDIDITFALSATSANSNATYTFMQNTLKFIDKYRANQIQ
YSILVFGNVERVVTNFNSFPPTANOLKAAIDAQPALPGPVLEDALKETFKI**F**N**E**S**Q**R**P**A**K**K****A**L**V**V**I**T**D****N****N****S****G****D****K****K**
TLSNAVKTVEEQGVLVISVAVGGV****

P6 (g11666) see g4601

MCKAGFAGDDAPRAVFPSIVGRPRHQGMVGMQKDSYVGDEAQSKRGILT**LYPIEHGIVTNWDDMEKIWHHTFYNELR
VAPEEHPVLLTEAPLNPKANREKMTQSLRLCLSTPLVEQLVSSLTAEMVVSHTVPIYEGYALPHAILRLDLAGRD****LTDY******

LMKILTERGYSFTTAEREIVRDIKEKLCYVALDFEQEMOTAASSSSLEKSYELPDGQVITIGNERFRAPEAMFQPSFLG
MESAGIHETTYNSIMKCDVDIRKDLYANTVLSGGSTM**FPGIADRMQKEITALAPPTMKIKIIAPP**PERKYSVWIGGSILAS
LSTFQQMWISEQYDESGPSIVHRKCF

P7 (g4601) see g1166 and g13727

MCKAGFAGDDAPRAVFPSIVGRPRHQGMVGMQKDSYVGDEAQSKRGILT**LYPIEHGIVTNWDDMEKIMFETFNSPAM
YVAIQAVLSLYASGRTTGIVFDSDGDGVSHVPIYEGYALPHAILRLDLAGRDLTDYLMKILTERGYSFTTAEREIVRDI
KEKLCYVALDFEQEMOTAASSSSLEKSYELPDGQVITIGNERFRAPEAMFQPSFLGMEAGIHETTYNSIMKCDVDIRKD
LYANTVLSGGSTM**FPGIADRMQKEITALAPPTMKIKIIAPP**PERKYSVWIGGSILASLSTFQQMWISEQYDESGPSIVHR
KCF**

P8 (g9654)

MDKCSNLITDTYFLTVGKPRIKLPKKSDQEWKFHDWKVKAAGQKIKNVGQHGDMPVIRWCVTSPCEMKCER
MAMEFNYKISPRMQSCVMASCKESCMEIQRGNADIMTAEGDEITYAGKMHYLKPIMAETDDHDHVPNYDTKFKFQKP
DTLKHFSVALIKKSTAGIKSFKDLSSKKSCHPGVDSKAGFK**SPVCSLIK**KNVISSVGNFESAGEEFFKESCLPGVQHER
FNPNTNPESLCNLCKQGDKDDEFQCILSDKEPYYGCDGALKCLREGVGDVAFFHTDILMKYDTFSTEFDIVCEKDF
PLTWANLMKPECHLTEEYPQVLMVNGEKSKNWIKSCTDALMDASTKFKSPTSGGFDMFDSTQYCAKDDSDKGSEMKNMG
FKVDTNLMKNKDLIFKDCSSSLMLIPEDKRTYEKFLGVEMLGWVKAQELEPK**PRAFIC****CVVGDDQMTKCQDMITKFQKIT**
EVKKVSWGCIEAPSKQECVEMVANGLADLVDLNATEMFVAGMDFNLAPEVNAHKDKDAMTESITIGVMLKDGSDFLGDM
QGFMCNAGVDKLSGFHMGWLLANGTVPRRTTPLESIGKYFGESCPGIDKLDMTSHPLLSHTLNWRGMLKDICKRWE
DWYDAPLWRQWQDKEDGKTVDTAMH**KIEDFQSPEF**KTMIAMGFSNPRMRKYHTPDIRQVLEKMPWEWMIGWVWHKPED
WTEEQWALFQOMWAERSWDQEDLVSYVEKWTLDWETFKRWMMNAEFQDYSTKDKSLHQDTF**IRLSLMCSRY**YLLNEWKN
LQYDDIDMKQCREKLCNTCAGKGDKKCSMDSNNEDYYGYKGALKCLKQKDGNNVAFIDAYSFENIIQEDSYKNNDFVLLCP
DGKTVDTGADSYEKCNFGRIPSRTLVTNCMHDGIWRWVTKALLVAQEKLPOQVQFQDCIFSKNMESFTSIPFVNQTYQV
RLGPMFLRAMESLMO**PPVKE**KAVKYHTVDCCLGEECDGDVEDRLACDECLEGDDVCLENCWDGEPIVMEEMPISHIIPKIPL
KMPLKMOVGGVKKITLWKPCKRHSHQMMGKMHMGAMEFGMKGHMHDDMRMSGLIMKGSSPVMHQRWSMMRGKVFPHPK
MSAKPTKFGMKHMMKFGPMMKSGSDKLHMMSSKVMEMHPFKSMPKPFMDEPIHGGKMANHFMETGPLMHSYKSLR**GEFY**
NVLFARDGTCNSIVHDLSCFGERREIRMKAGRVGQTTITVPMCPKPTIFKNTTALFTCDNGASFAKPVSLPLKCAYAPCK
PGFWLPDWSLDQYWQGDNSHIGGDHWGSKAWWKNRQGNDNWFLAARDENRVQKKSSVGKQ

P9 (g10811)

MRDISVIVYSILCYFCLLRTAIGQTFDVRVNLEEGKADAGQPVHSFPLPASNEVYTFPAQDADSKAALTLFOISEQGVVT
TTKPIDFEIGKKNYYDLVAVRDRGDKEGGVPLSIRISITDTNNFGPTFPSNLYHGRVKENSPVDTIVLGLENCFAEDRD
TEGSITYSISGGNEKGYFKAEGRTVNNINRVFLVLKTTNIKIVRDVNTPKIIITVRAFDGRRFTKVSITVLDVNNNDP
KFDKDTYTATINEDTPLMTSVLVRATDADVGNTGGIYYYISGGSPWFSVDAITGVMKVVSRLPNQAAVDLRVTARDGR
DGTRSHEAVVKVNINLIADYPPADTASPGANTPPVPEGSYSANVREDFPVGAABLVIHAVDRDPGRNSRIQYRLSGDN
AFRINPDNGVVSRLTNLDYSLQNSYDLNVIAQDQAQNPLSAEASLKITVQOVDKNRFAPEFQAPNKYQRQASIPENRAIN
TKVGSSIATDSDGSQRPEGQVVSIASGSGLPYFKIDANTGDLRSVAVLDKEKOPQYNNLIEARDKSLYPLKSNLVYMI
AVTGVEDNFPDFSQPVYSAKVPGNAPRGTFVTAIRAIQDQDGPVTYSIDNAGSAFTIQADSGVITARPLDPANEERNFV
LSVRATAANKDSRAQVIVNVVSKADSPPTFVNLPYKATVPENLGQIDSLLCIAAKDSQGRFSATGSFDYEAFGGVYQVQ
ASSGPQQIATESVTIAVTDTKDPKFSKGSYKFVIDEDTSAGTTITVTNQDGSSGGLVIEDADTTITNFECTIENIKSLE
IENHFTIRLVTSNSNKNAQNSNNVRRRVEVSQVYAPGTTFIQFDPASYSKDISENAPTDTNVFVTAKGPNISRYSIV
GGNINNAFKITQSGQVQVANSLDRETQATYKLVIRAFKDGTKLASEVTTTINVLDVNDVTPKITFMETQPKELADEDYSP
KGSFVVVKVRSYPLAM

P10 (g11107)

MVLQAVDLDGINAVIHYELQDNAGGKFKIDQDTGLITTLEREGPENEFTVQVKATNR**QAGAPLLSGTVTATIKVSD**
GNDQSPVFDPLVYKANPENALPGYLVTKVAIDKDVGPNALEYTITAGNDPYEFYIDPRNGR**ILVSGLLDFDKGKKT**
NLTVTVDGRNPPKSAPRPALVYINVLDANDNAPIFVPAEYNKKVSESVPGDTVLLVTAVDKDTGTNAQFQFSITEGDD
ANMFAVRSNANNGSIGEYTLLQOLDRETVPRYNLTVTATDGGLOGIAVVHLTTIDINDNGPWFKPRYYEGTVKAGINPS
QGQPVVTLEAYDPDEAINGPPFTSFLVNTDLSNRFKITKATDKTANMAAFGEFNRDVKQSWEVQVKAEDSGNPTMDNTF
VYIDVEDDRNRNEPFDGKLTIIVNAYNRF**FAGGIIGK**AYRDNDYNGDQDKYMSMDVQSLTLPETGDITAENIPVGRY
TFSVKVEETKQRPGNFPKIVTSDTVIVQSNTAAIQQSVAVLILEMRKTSFFVADFYTKVQGVLANILTGGDTQILIF
SIQKAPINRVPLSALFGVEIYLSVKSAGGIMDRLRVLNQLIONKQRLLEGLKIGSIGIDACAYEQQVGICKNVVKASS
AFTVVGSDYQGPATESSLTVVAMNVNLQAEYTSIIILPEINCTTGTCPFHGGTCHNAVPKGIICECGREYRGPECOSTTR
TFKGNSYLWLEKLAAYERSTVSLQFMTGTADGLLLYQGPVSEGSNNGLPDVMAVMMVGGYIKLVLISLGPHPNAPLELFMN
RGDRLDDROWHTVEVIRELKKVTVRIDLCSKAQVVEDDGQVLEKRAACEIEGEVWGSNVYLNGFGPLQIGGVENKLNDAK
ISYAGFQGCIRNIKDNDVMYDLLNPVSEKNTDLGXAFVNSSGNAILRTADGNIISSGIGSGLAGYSSDAELMGTMITVG
RSGQVHLSSSYMKMYSFGMRAVVGSSGRVFVGRATTSTRTIIRSISYRSTPIGVKSGGVNVRVPIPTNYDPSGGSIV
ISSTKGGSVVGSSGGKGSAGGGSGASDSSGASAGGGVVVDGGAVAGGGGVVGGSGVVGGTVGHQYVTGATVLG
GASSADAYQOINAYGTSLLGGWTFLGATPNGAKEIEVIGDFGGCTASNYSNGVLDNDPSIEARRONVEFGCPCSSGFCLH
GGTCVDAVPPYCMCPCPGWTGPRCETVVTSPNP

P11 (g13727) see g11666 and g4601

SLVTOTVKASELDEKGVMVMGQKDSYVGDEAQSKRGILTLYKPIEHGIVTNWDDMEKIWHHTFYNELRVAPEEHPVLLT
EAPLNPKANREKMTQIMFETFNCPAMYVAIQAVLSLYASGRTTGIVLDSDGVSHTVPIYEGYALPHAILRLDLAGRDLT
DYLMKILTERGSFTTAEREIVRDIKEKLCYVALDFDQELOQTAASSSSLEKSYELPDQVITIGNERFRCPEAMFQPSF
LGMEAISGIHETTYNSIMKCDVDIRKDLYANTVLSGGSTMYSGIADRMQ**KETALAPSTMKIKIIAPP**KYSVWIGGSIL
ASLSTFQQMWISKQEYDESGPAIVHRKCF

P12 (g2385)

MFIFRHYQKVLLISQLPLTLQDETENKCSQTTITEAKDHLRLNGGGPLGTYNENYMAFTLDEVRTNFKNLLTLQPGNF
SALAAWAPLSFPATLESGACVKDGNCEGCPVVKDLGANYYPRFLNEVNCKSSQICGQGLGECRTSVKSQRF
RDPSTD LSTCNKDLEEEYYDMEMCCQCMLL*

P13 (g6918)

MGYNAGDPSNLRFYNIPSSGTVEAHLVDER**LGNTNREGVWFFRLDVNPILDILPAKKCYTWIQQQDNIQVPAIRPC**CTL
D QAVADKNFYVDYNQVVAKRSNLTLCAYSVRDLTS**RWAQQCCYTEIPGGGRVLSVGPTEGGTPFLIGIPGLPIITD**LEAHS
YCCNASILCARYY**QLRPSDTCSR**YSSRREALIWDPHFVTLNDNKAYTFNGLGEYTMISIDNVGFELQARTKKTSGR**GLGT**
FSAAVAKEANTAAVEGRINSQGELEVVFVAGIQQNILTVAPERGLISGSNITLIRETAGSISALFPGSISISFSRVSDALA
TTFNGPIEFVNHTKGLLGTWNDDPSDDFTTPDGTVLPADATPOQIHYDFGLKWOIHASQLFTYQAHENPSTFIDL
SYVP MFIDNITWTNESFRROAESVCGNNTHCLFDAVTVDCSFKGKSTKELEENNQVNKL

P14 (g9951)

IIPDIDVGF A VRDEEVESVGLFTLMRDTISTINHRYGTERVRYTVILYGNINTTVQFSNRNIDHQELISTVTELGRVP
AIDLEAKLTAAEALFRSLPSRAGSKVFIVLTDVTAGNESTITSAGASLVREGILVFSVNRS E DGTNVVTVTQIDYLNI
PTFTTDRSVVIAETIIRKASS

P15 (g1532)

MVVRNEKDDDDDRDGKDGDKDDSDSKGDIDDKNDKDSL YDKDKFDDKDDKDDKDNKDDDDDDDED FSDDNE
NLLSFELDEIEEKDADDNEVDDKHSVDSFDDVKFTFGVNVKSTLDGIHVTTVNLSTYLDQKASLEIIIVYLHEAGSV
FISNWNFCDGDESSGECSES KIGEFLDL KIKSKGSPKEVDDDRKKVCNSKDDDDDDKNEDEDDND DDECPIIYD
VGGDSEMLLNKGVSRAHL

P16 (g11702)

ISYLCMAKF DSDQKT ALMIPAKKVGEVDNWDDKVLAKMCNLL EALPVREMLKL ASDVAIKDMLSNLNLSEGQLRVLA
V ALDLLGSPDKWNKDYIKELGNMVAGLMPSELRQIGAQVVKDSLQFLKGIDFDLDQLSKLDKDDVVNL SKAIDGFLSSDV
G KMTQAVVIAAFPEMKVAKEVAMPVLREFIKKFEEDPSSAGKSIAOLGDFAVALSRGEVDAESVDKVITNLDKLGVLPWD
TKTTLIKKIRT KWGDFNGTDADDSDSPNWGF LNMKRLGRLALGIAKEELRDLPIRGIEDVIDVLGKEKDWD RGQ

P17 (g12472)

CTNLPPIFIYKVYMF KFDSTHGRFKGTVEAKDGKL VINGNPISVHAKRD PSEIOWGADGAQYVVESTGVFTTVEKASAHL
KGGA KRVIIISAPSADAPMFVMGVNEDKYDPSKMTIVR NM

P18 (g810)

LMEAGVIGSHGQA ALRRAEAVHRRASALATIHHHNMAASSVPEAIQIVNLAALKVAQLMEDGLIGHNGLPVTKRVVVATP
IGEGNVQTQYHRTVVKAQVTKM NIGVAILRDAQS VAVGVRGVSGQAAVRLAMDNEQGTAHALILHHPTMEPLAPGQ EYN
LKRAMWGSVQMAAGQPGVNGRPVQNRVVEEHKGQDLAPLH PMAERSVWAKRSPSSAKNKHQVWMAGGLSGIHESA
ALHVEVENDQEHEPVLIHHHLVTDQDVLAALWRLNNAKHNRVQWMANFLRGLNGNNARRVEEELKPGPGGVIVLLLPLV
GRPVLEILS RLEIVAPMSAQLM VYQGTGKVGRSVAGHAVAENNQEVYV SVAH GRPMVGNVVLATIQRQGSAKPRPVQLMA
DGLNGPQGH AMLYVAMENATEQEPNAYTTLAATPTPTATAVPTPTIDPNIPKIDLVFAISATSVSSRSYELMKNTIKRF
IDRYGVNSIHYSIIIVYGDQVVRVINFNRTFPSSANELKTAIDNQLALSGGPVLINALQEAYRVFKE SVGRPGAKRVLVVI
ADENSGSSPSFLSRAVRPLEDLGV LVISVGVGDRISRSELNIISP NLLDVISARLNINPSLLAVRIMERILRLNFPDV DV
GFAISAASANSDEIFSLMKQI INTI IDRYGVSKVRF SIIYGSRV TTRFTFDNAPITQEE LIKAVNGTKV TGD PDEKA
LEEA EKLFTKSSRPNATKVFVVLTD FVGAGDDNSL IANAVRLRKSGVLILSVGFGQ QVNAIGNQMTKVVITQSDYI RVPD
FTT QRPV VIAETIMFKALQKGKVHVYECAAYNHA

P19 (g20041)

KTENTATIINEDGVIK LGPWEPPSHGIISLLFKTPYPKGTILFNGDTKD YFHLEIINETSIRLEYNIGNGVSTIDLSLA
NEKELNDRKWHAVTVYRNMLQFGLKLDQEKHENPLFKEKE LNVGDKLN VGTNPRDVTEGFVGCIRRLVSNHCSSRL

P20 (g6066)

CDFEKKSFCAWLNVPNGNKS LGDDFDWTLGSGSTPSWQTGPSSDHTT GSAIGTYAFIETSSPRKRGEIARLRSKMF AST
TGKCMSSWFHMYGATVANLNIYM KQVGKLESIVWNLN GSQGNVWKKA EVTITSRTSYQ MIFEGVTGRDFQSDIAIDDILF
NEG CIGLCS VNSQQRVDCGYVGITKATCVGLRCCWDDSVNVPYCFYHPSKIVATLESRQLS VSLGDAAGIILSRTF
LGASTDRAGQHPSPPPQPRLQRGHHPNGTAISSKATVTGTTQKRTISTGGDSL VVHLL EPVQLRI IQQEVKKFAFEGIR
GOSYQGDIALDDIRVLDGSCPLL RNYMYIET SAPRKPGDYARLESPTH S VTDNGKCLVFWYHMGSGIGRLNVYIKRGN
SLGSRVWTASGNQGNRWLRGMITVQSPNTQWKVRNSVCRT

P21 (g18277)

MMAILEHQPNHGNLMDKITLKDVSCTLNDLGEFNATHLWMKAPLDGCLTNHSTSED TITYMNSIIAETRASAGSVLISR
EFQAEFPFKCSYPRSAIMSVASFSPREKVIYTRTAEGNFTFTMDMYETDQYLT PYD SYPVEKKLNEKMHL EVKVKS NDS
KLVLIPERCWATPSQDPKDSKFFAFIDKGCGEDDTLVF NYEENAVQRF TLDAFRFLQESTGSTVYLHCEVEACRKGDNES
RCAKGCIQDNSRKRRLSETDSTIMETVSVGPLTAEDIQPOQKQ

P22 (g19762)

MVGAGGANVG VGGGNMGGGLGAGVGGITGGGGVGAGGIAGGGTMGGMVGGVGMTSGGT VGGGLTGGMVAGGGIVG
GGVGGNMGGGGSIGGGIGGVTGAGGNIGGGVG VGGGNVGGVGGVGSIGGNIGGGVGAEGGIGGGVGGMGGMGGI GT
GVGIGGGGVGGGIGGGANMGGAGAETVGGAKGMEGISVHF CRPCFDGHQELCDYCPSEFQDD DEVEEGYKRHHHHHKH

DKIHLQLGNDPRHPERKRPFAKVSGDYEVKQFKQAGPSKGYERHDEHGYRADTFSNRNDDVHPTTNPDTDYYGEYKFDD
SKMTVEDSGKYSEENMKTQ

P23 (g1057)

MFGHLRCSVSASTNYNNFNVTEGLPNGTFVGNIGNQSYDIVAITPRDRDDLNLDPISMNITTKIVLDRKLSAYIIDLV
RLNPFKVVTFFINVTDINDNIPTFLVGGYDFEFYEGNPVGARVTAVDDDLGSYAVQNSIISGNTDNMFALTEYVDAVGR
LGAKIELQPGKQLDREQRDLYSLTVSASDGTPQGTGFLVNITVLDVNDNPFVNPKYYNPNTENTKQFTPIVEVYAT
DKDIGTYGKVMYSVIHDSTSMDPFGHFALRTNEKGIVNLAKLDREKQDSYSLRVLAHNGSGYYYKRLAYVTINILDVN
DNSPEVSYTFYLDDAQVYEDASVGTKVARWVTDKDAGKNAEIESVTIQGGNGHLSIAYDAANKVHIIISLAKELDREKO
PTFRVYIYAKDRGDPQQTQDGFLNVGDINDNPPKFDQDFIAVVSERATIGTSVVVAHATDKDIGSNAKLVYNISYTS
QATLYKTWFQIDQATGEIRTSARLDREKFARLEFLVTVDGVPALSANCTVIVNITDANDNDPIFTKAVYFSTVNESA
NNTSVLQLANDIDSGINGIVHYSLATQHRRVPFSIDPNTGLLSTSGQIDFEAQSLYVINVVASDGGRLDKSILNISVV
DENDNYPVINPIIYDVSVYENLTIDVVATVMAKDNDGILSQINFASNGNDDGLFFINSSTGVITLIKPLDRETKDFH
QLEVOATDGGGLKSNSAVIQTVLDVNDPEPVFEPSSYKFAIDENSDVNTLLGSVYASSKDLGTNAQIYYSISDGNIHN
VFTINSTGTIFTQNNIDHERSPQLLLTVQAKDGGSPPLYGFANVSVTVIDLNDNAPSFESEGVIDVNILEDIKIGEVFYNV
TAQDPDGLFGRVRYTFLSNPNNTFQMSSTGALTVTHLIDFEGPRQYLIIVLAEDGGSPPLNSTATFRVTVDVNDHTP
KFSNKSYSTHVDEHTPAGTNILNVSATDADTGNNALISYSFQSGVDTSMFGLRSNGWLFITRQLDREQQEYVYRFTVVATD
KGHIPKSSTADVIVYIDDINDHNPKFTQQAYIFSVNENQPNRTFVGQVPASDKDAGANAKISYTPEPPSLEFAIDHNTGI
IRTNQVLNREQRSSYILTVKANDHGTSPTDHTSVTVQDINDNAPTFKKPFYEKAVPEDIPVGSSVLTVEASDPDAGS
NGMITYFFASNPVGFTIDARTGVITTARLDREKHDRYTLSIGAKDHGNPQSHVVFTIHVLDVNDNSPRFVNNSIFVN
IPEKQAI GTVVTVTAKDEDTGNNNGKVRYTIDQGNGGKVFIEINETSGVIALRKPLDFEKSKYQLRIVARDQGQNSQSSF
LYVTVVVLDSDNRPTFEMNPVTVQQLQEGVPLNTVTTIKAHDYDSGQNSWIRYSIDSQSPGPAFKVDPSTGVVQTIGS
IDREAVDEYTLNIRATDQAFTESERLSSTLTIFIIVSDVDNKPIFVSPNETYVMEDEGFDPVITITAIDNDLGSNAKV
KYRIEHDENRDSFRKFALNAATGLLKLGVNLDHETTPKYVLNISVTDGGTPQFSFQLLTVFVVDVNDNGPEFNQSLFLG
NVSENQPVGTPVMAVSAYLDDSGENGALTYSPRGNVNQKFALNGSSGVIYTNSTLDREEKDEYTLTVYAMDNVYPRRGV
TCTVKIKVLDVNDHRPVFIPSRLNLTVLEHRPPDFHTLRAQDPDIGHQNGRIRYIIQHGNEDAKFSVGEFTGVLSTTAEL
DRERKSRYYLVIEARNVVVPFYNASIYVTIFVGDKNDNSPKFVNSSYSVNIRELTKGTPILRAQANDIDSGKNGEITFS
LSSDALGIFRIDSKTGTIYTLQEFDDYITAAYNFLCRATDKGVAPRETSVGVTVSIIIDENNHPVDAIPYEATIVSGSS
VPQNELVRVHAEDKDSKSIHMTYRLDNSSQFFQLESNSGVMRVKQGVSSI PDKVYVLNIIADDGGNLTRGRGIVEIIIVG
SVSPNRPIFVNITPSFVSIPEPKSLEVARVLANSNDVVSIVDGNSGNAFLVNSQTGIVTVQNPQLDFERLDFRL
RIIVSRKSQPSRNDYLTLHVNLTDVNDNPKVFYPANISVQLLEDDGSQSSFTVRSVANLTTDKDSGNQEVTYKIISG
NVDDKFAINAQLGLLTTKGFLDREKHAVYDLVVNATDKGTPSRSSVSHVVNIVDVNDNIPAFSGPYLVNIKEDLRVGSV
VKRVVATDKDEHPKLMYSGNDKTIYDVFRIDQLTGDIALLESLDYERNTSYELNVTCDGRYKSHTTVNVKDNDNA
PEFLESSYQATLSEETDQGTSILKVS AEDRDDGSNRELTYAFVTSSDLFRLNNTSGVITYTAKKIEASDRDSFYFVLVSAV
DRGPVQHQAFVMVQIRINRKPMFKEPAYRARIAEDTLPGTTVLTVAADDGMNVAKISYKFKFGNSEGLFRIGRRSGII
KVNKGGLDYESEKNYLMGVEARDESANKSVVVEVNITISDVNDNPVFPDSEYVKEVYENISIGTLLRVNATDKDSAN
GRLVFSILSGNDKQSFRINSRTGEIVTIKRLDHEIKEHTLSIQAIDE

P24 (g15888)

HEVYCFSCOEQLPITEHEEEAKGSFYFKVFGKELFYNYFTYDDVKELEDGVIRVSNDLKDLLKKGNWNITRVLRPIEV
KIRQPSCVGLPLDFSVDGVFLRLNTTVKAEVQPRFFIPLPDQVTLSGKLNI

P25 (g11220)

MPQGRKTITLEVEPSDSIENVTKI QDKEGIPPDQQLIFAGKQLEDGRTLS DYNIQKESTLHLVLRLRGGMQIFVKTLT
GKTITLEVEPSDSIENVTKI QDKEGIPPDQQLIFAGKQLEDGRTLS DYNIQKESTLHLVLRLRGGMQIFVKTLT GKTITLE
TLEVEPSDSIENVTKI QDKEGIPPDQQLIFAGKQLEDGRTLS DYNIQKESTLHLVLRLRGGMQIFVKTLT GKTITLE
EPSDSIENVTKI QDKEGIPPDQQLIFAGKQLEDGRTLS DYNIQKESTLHLVLRLRGGMQIFVKTLT GKTITLE
SIENVTKI QDKEGIPPDQQLIFAGKQLEDGRTLS DYNIQKESTLHLVLRLRGGMQIFVKTLT GKTITLE
VKTQI QDKEGIPPDQQLIFAGKQLEDGRTLS DYNIQKESTLHLVLRLRGGMQIFVKTLT GKTITLE

P26 (g1441)

SNTWLMYRKPVQGSNDWS SLYRSSELGFEIGEVTEVRPNKGCLTVNERYPKLFKITFMNDRRHVR LATVVYIHDA
SKNPVEWTRETVMNTDQVTLFQRMIEDIEPQKGYEEKWRYISDLPLS LSSDNVPENDLPHHHMKIVFVTPVAGKLCLR
PKRTKESVFSVGRHHTNLSEL CYGKQILGFDVCLSGRYISPHTLHPPAFCPCPTRLNISVQPSRNPAQKIQFKFGRGDR

FDEYHKTWRAGIYLOGTNPEKKIEADVDYDVQNQRTDVTVDIHNMYPWLRKVCIIQVISDNVVEFDLRFGENCLSDYEV
Q

P27 (g18472)

MLTFQVDSMEEKDSNGNVVGLTSQKKHSWLSSENPNQKFTFSPLNNFTTFQRLPVKRINLSVSLAGPQANLEIQVLLFLK
SGKIEFGNETFNRSGTFKFNIKVSDWQFCGTNAEVCKNSTTGANEIGQFLDIGMSIGSVAEEPEEVDDDQIPGEDICAP
EEPDEEHECPRIFNMGGNSEMMLNKG

P28 (g11651)

MKLAVAVLLVCVAAPQMIILTIVLIIILSSTVFGSLPSFEYGEAFEKAKEIGQEVVREVGRHAEEKNSLLESGEQAMEKL
RQLKEANRKAGEIQETLEESGDQAIKEKLRELKEQAGQRAEKLKGNAEEILEEQKKSLOETGEQAVEKLTQLKEQASNRA
EEMKNTFHEKRKEAMEKLSNLKEQASQRAEEVKGNVEEVVEETKKTFHEKREEAMEKLRNLKGNAEEMLEEOKKSLQETR
EQAVEKLTQLGEOASNRTEEMKNTFHEKREEAMEKLRNLKEQASQRAEEVKGNIEEMVEETKKKFHEKREEAMEKLRNLK
EEASQRAEQVKGNVEEMYEENRKALQDKREQAMEKLLELKQALKSKEILKEKKQOLKEKMNQMWDMDMKQKAGTNKEQLO
EAFRQKKQEMKEYWKELENEQAGESKEKFRKIWRENKENWRNKAEEVSVAHSFRYLSFYASLDKYLPACKDPVYM

P29 (g13377)

MREIVHLQAGQCGNQIGAKFWEVISDEHGDPTGTYHGSDLQLERINVYYNEATGGKYVPRAVLVDLEPGTMDSVRSGP
FGQIFRPDNFVFGOSGAGNNWAKGHYTEGAELVDSVLDVVRKEAESCDCLOGFOLTHSLGGGTGSGMGTLLISKIREEYP
DRIMNTFSVVPSPKVSDTVEPYNATLSVHQLVENTDETYCIDNEALYDICFRTLKLTTPTYGDLNHLVSATMSGVTCL
RFPGQYRALTVPELTQOMFDAKNMMAACDPRHGRYLTVAAMFRGRMSMKEVDEQMLNVQNKNSSYFVEWIPNNVKTAVCD
IPPRGLKMSATFVGNSTAIQELFKRISEQFTAMFRRKAFLHWYTGEGMDEMFT

P30 (g11056)

MMGIKGDTVFSQSQGQSSSEVMVNLTGVSATLKWSIRQLPMIYNGNANLSCKAGAVGDFDPTMAKASADYETNCQSSSS
SKFQSCAVGDLSTMLGDLDRNQQPTTLSDPKLMIPISGPHSGMKGALVLYDGDMPKACALIGPTQPMKTFVAVFKGPVAG
FVYFRQVDENSDFVFVDFLFFVNGANSKKNFQWQINQGVVSHDASDPSSYCKNVQTFNPKNMGDVSNCNTIHGKCAIGD
MTSKHGKIEVSLATEMQSSTKAIFIIDTNLPLSGDDSVQGETLLLYPSEPTQAIACAKIVRLEPKTVKVSFNGSIQDGVTG
HLMFTQSSPFDATTAKIELTGLNKAEGYHIHAYSPPPMOLLPNESCTGLIAGDHWNPFNVDIKTSPPAGTGTDEEYEV
GDLGGKYGTFLNLSSFTKTVMDNFNLPFMGRNSIQGRSLVIHKQKIKDNNMRWVCSDLILEVDEGITYFMKATANIRGPAV
KGVVVL

P31 (g20420)

YVVRLGGNDKIKELLLGQDLNDGKPHRVDVVKDLVINVTLDGDSSQEKGQITTKYKLEIDVAIYVGAPSFOGLLLGV
KSNFYFVGCLIDVKFRIVRGTPVREIQLLKKDVVESVKVDMAKDRCSPLAFQPYTFSKPDSEYTFVIDKKTSIAGSFKFR
TYQKDGIILKQGNGQNGFTIKFGGSGVILSLVIKGTPTEVTLPTDNEPRVDAGNWIAKFTLSATDIKLEGYNNELPVT
KTPTTAVGNFFHPDVT**AGGFFIGCMRD**LQVNGKDFKPINGVSKVKGVEFDRCNITDLCVFVPCNGATCTTDGKGLKCDC
TGTGYKGDVCQLGKYLTVGCIRN

P32 (g5540)

LPKEINVFDAVGLGKKVFPGVSTTGQNNRSPAYHFTKQTKDMIASQNAFEEVDKLIHNSSDFIVSAYAKLMANEYGNP
IVSISSKDGNQLYFLLQISYTPQPYPKDAQLRLGQLFEVYQEDIGEITSFRFRGDLQDVFKVGSPLSDCQRIAKASYCRC
SISCOCTDVGKK**DCVVDGLRYADGKRWT**KDKCNICHCKWGQVMCTYTCQVCKDNGQTYMEGETWKSSNDSCHTCCKCEDG
KVKCSPPVCAKDCSGK**SGDLVTLKGQCC**PICKEDQCKGTGKVKFCGCDVTCNFHDNDACQSCTEGCYCPEGRVLNDQG
KCMASARCGCLYQGKRYKVRK

P33 (g8985)

MTRLGDTFIRQLHEKGEDFTNLWAFSEFLESVDKGVPDTKTIPVGLKKMKGIRNAAIEMLVYAGINYKKFKVEAIN
ELLTERLRWVHANLAKDSKVVNFRLPDGMIKRPKRKFVIRHSSLPLDIRPGIVTEESGKYFPEQLAAWISAVRRE
MDDCGFKDGHLVHVHEKGLVDSLQSLANGANGIWAQSMIIQGASMGAASSTVLMNLVRLGNKKVLKKYNCTALREA
AQEICRATTGQEPYPLQPIYGERALDMVFGMPK**LGINEFDLAKF**GKPLMRMTTLASAEMIITRLKNIFGEDPQFTIE
RGTRMKEVMLEDLHKNIKEEYMSAAGLAQLFDRSGGQLTGKMAVDVLAQDEPRKAHAQVILAEVRAMDEWDLRDGKRDDQ
LEFDAFYNGFLAPYFGCYRCDETQALKAMDDEDGTVDWNEFAVYLVKWAMRQYPETKTAED**LLSVAFR**KGLIPAMQDE

P34 (g1714)

RYMYIETSSPRVGETARLVSDFKIADGHNWCLKFWYHMYGNSVGALKVKLKMPFNPSKPYYNTLRTTQYNRGDSWYF
DQVRINSADDFEIVFEASVGTSYDADIAIDDIADVTKGYCPSPTPTPNACAVKCKSGKCVSSTMCDVFNDGVGDNTD
EQNCGDCTFETGTALGHYMYVEASQGDSWDLARFESTWLQOSASTCVVSFWYHMFGSGIGTLYGYVKVGQTYTRLFEWG
NKGNKWIQGKLYVGRHSGPFKVIFEAERSYNVFGDIAIDDVSFENCTLPIVSSCKRGQHRCARGSCIDPRLCDYTDDC
GDNSDEVNCYSYNYRCSFERSLCQWNQLSDDDFDWTRNQGSTASYGTGPMFDHTLGTSAGSYLYAEASWPRKKGDKARLA
SGFLSALPQGDTSCKLRFYFHMFGPDIGSLNVYTRPCNGCAETLVYTRSGNVGNFWDRAEVSLPSTVPFQVVIESVRGSS
YQGDISIDDLMSMTQYCRSYNGPIPTAPPPTAAPTVACPGFRFKCSNGPCIDWGYVCDYRDDCGDGSDEVNCGPCNFES
GLCNWKDNSVGRYQWSRNKGSTVATGTGPSIDHTCGNAS

P35 (g7349)

LHLVFFNKKYETFONAVDPDGLAVLGVLITATCPGNRILGNFAKKLTKVVEEGANTNATATDSIKLNYLMPYNNKQGDE
DDEEDDIASDDPEAVEEEEVDAKKKIKYYTYKGSLLTPPCYESVTWIVFKDKIKISNTOLNKFRKLKAQFGGAP

P36 (g13890)

HYMYIETSSPRKQGDKARLVSDFSPITIRGRCVKFYWYHMYGASIGTLRILEKTGPGNKSESTIWELSGNFGDQWYSGQA
PITSSSPYQVVFEAMRGRSVSGDIAIDDVTFSTSCKCSVVPSLAVPPTPPPTPPPPIINNCTFEGGFCSWTNLNGDDFDWT
RSRGSTASWSTGPTVDHTLGTSGYVYIETSSPRSPNDKAQLOSGLIQPTTVTSGRCLFWFHMGWRHVDTLNVYRKDS
SSRV