

Ce025#1 CVWRLDWALGKL  
Ce033#1 ILWRMATMSASM  
Ce036#1 VAGQHRSLVARL  
Ce039#1 WHWKCFEIVARL  
Ce051#1 GEGAGFNQLAKL  
Ce096#1 DPNEAMQWLARL  
Ce123#1 LGGPAPEFWSSL  
Ce145#5 QPFGSTAMRMSKL  
Ce146#3 AAPVRLVLGHKL  
Ce147#1 GKATGRRLKAKL  
Ce148#2 VVWPLHRLRASL  
Ce149#2 DRKGWTHQPSKL  
Ce150#1 AVSFLSMRRARL  
Ce152#3 GLSDSFYRRSML  
Ce153#1 LWTALLGRTCSL  
Ce154#1 AGSAFSSCRSKL  
Ce155#2 KQWTGKGRSKL  
Ce157#1 RERHSYMQRAKL  
Ce159#1 NLEFWGMMRSKM  
Ce161#2 YGVGRELRQAKL  
Ce167#1 EGKIRTLWCKL  
Ce168#1 DGAHREAPASKL  
Ce169#1 GFRGYMSGLCKL  
Ce171 RKLMSSECRGKL  
Ce172 MRREVLLRAKI  
Ce173#1 TCAREFGRRLRAKV  
Ce175#1 RWRLGWSWRAML  
Ce176 WCWRAPNLGSKL  
Ce179#2 RTEPKLMLRASL  
Ce182#1 RGGGPCQWKAKL  
Ce185#1 GHAPSRPAKARL  
Ce186#1 VSCRCRRLQSRL  
Ce187#1 PDVRRCAGAAKL  
Ce188#1 GSTGWLLQAKL  
Ce190 EHRRSSTSTAKL  
Ce191 VYTVPGTPTCSL  
Ce192 VYGGGVTRGHKL  
Ce193 CHVGSTPWRAAL  
Ce194 CALVGSHIRSRL  
Ce195 QCVHAEVHLARL  
Ce196 LVDPLITRSML  
Ce199#1 RSEEMARSPAKL  
Ce200#1 RLEMARWPRFKM  
Ce201 CQVRGACGMAKL  
Ce202 RSGSMVELRARL  
Ce204#1 NYPQATARVCAKL  
Ce205 CVSDLGHIVHKL  
Ce208 GEKFRDAWRARL  
CeD1#1 IGRRMAPLRSSL  
CeF#1 LELVDPGMRACL  
Hs01 IGLDWLVVISKL  
Hs02 GEYGVGLLWSKL  
Hs03 WGDPLIMPQSKL  
Hs04 SLLALAVQCSKL  
Hs05 RGMWFSSNSKL  
Hs06 ALYQWWALGSKL  
Hs07 GNVGSAVLAACL  
Hs08 KLTVRWAWRAKL  
Hs09 ERVSGRHPQAKL  
Hs10 GVEVLVPGTAKL  
Hs11 GPDPLMSILAKL  
Hs12 PVKCVRKRLSRL  
Hs13 ICDAIFQGGSRSL  
Hs14 PVSWNWVRSRL  
Hs15 IQTGGGRAISRL  
Hs16 RWDIPLRWWSRL  
Hs17 GAMGMLGGVSRL  
Hs18 GEALLGGFLSRL  
Hs19 PPNNIGCTASRL  
Hs20 VDPVIVKVPESRL  
Hs21 LAGHAWKALSRL  
Hs22 ELVDPEGLRSRL  
Hs23 NNGRGKVLSRL  
Hs24 LRHGVNGLWSRL  
Hs25 VRGGYLVRTARL  
Hs26 VAGQHRSLVARL  
Hs27 NGIEQKKWARL

Hs28 CWRDLFQMIARL  
Hs29 AILGGEYIMARL  
Hs30 KEILELVDPARL  
Hs31 SPYTGGSLSCKL  
Hs32 MQIEHSLPFCRL  
Hs33 LVPLYHLTPCRL  
Hs34 FWSYWMVESCRL  
Hs35 VSGWGAGGMSRM  
Hs36 VWREGWCVRARM  
Hs37 PLTPRCILICRM  
Hs38 QDWAGLPLRSHL  
Hs39 KEEWPWGYRAHM  
Hs40 EGLIVMLERGKL  
Hs41 LVDPLYILWGRL  
Hs42 YCYVLRVGGGRL  
Hs43 LVDPFQVMFGRL  
Hs44 MGMTGMLWPRL  
Hs45 KEILELVDPPRL  
Hs46 EQSGDSGLKPKM  
Hs47 PAYRLVAVLANL  
Hs48 WSMELTPIWCNL  
Hs49 SNMSLTFMSPNL  
Hs50 IAMNCVQVKSOL  
Hs51 WARYENIMSSOL  
Hs52 VDPSERRLRSM  
Hs53 LVDPLGLPFSML  
Hs54 TTMRGDTLTSLL  
Hs55 RGMGFPAVRSLL  
Hs56 CRSGLPCLQSL  
Hs57 RAGQGTMWRSLL  
Hs58 VEYMMKWPRALL  
Hs59 SWSRSRLVSALL  
Hs60 GRDRTSVVRCLL  
Hs61 VVGVGVCVKSYL  
Hs62 LGNFVSLCSSL  
Hs63 NGQVRDWCRPSL  
Hs64 RIASNCNLVSAL  
Hs65 MVVRMNPDKCVL  
ADAS\_CAEEL LIDIIGSPHCKL  
ADAS\_DROME PPTSSTPPKAKL  
AK11\_RAT REDGKWAMSCRL  
AMAC\_HUMAN KIIESNKVKASL  
AMAC\_MOUSE RIVESDKLKANL  
AMAC\_RAT RIIESNKLKANL  
AOPP\_HUMAN TCSLAPNIIISOL  
AOPP\_MOUSE TCSLAPNILSOL  
CACP\_COLLII RSLIQSAPKSKL  
CACP\_HUMAN RALLQSHPRAKL  
CACP\_MOUSE RTLLQNHPRAKL  
CAOP\_CAEEL VEKYLKPMTSKL  
CAOP\_HUMAN SYKHLKSLQSKL  
CAOP\_RAT YHKHLKPLQSKL  
CAOQ\_RAT NKSVANRLKSOL  
CATA\_ASCSU KNISNLAKYCKY  
CATA\_BRARE GGASAVAAASKM  
CATA\_CANFA GSHLAAREKANL  
CATA\_CAVPO GSHLSAKEKANL  
CATA\_DROME TEELNLAKSSKF  
CATA\_HUMAN GSHLAAREKANL  
CATA\_MOUSE GSHMAAKGKANL  
CATA\_RANRU SAHVTANDKANL  
CATA\_RAT GSHIAAKGKANL  
DAPT\_HUMAN KTPIGKPATAKL  
DAPT\_MOUSE KKPIGKPATAKL  
DAPT\_RAT KKPIGKPATAKL  
ECH1\_HUMAN NKELKTVIFSKL  
ECH1\_MOUSE KRDTKSIIFSKL  
ECH1\_RAT KKDSKSIIFSKL  
ECHP\_CAVPO WQSLAGLPSSKL  
ECHP\_HUMAN WQSLAGSPSSKL  
ECHP\_RAT WQSLAGPHGSKL  
HAO1\_HUMAN LVRKNPLAVSKI  
HAO1\_MOUSE LVRKNPLAVSKI  
HAO2\_HUMAN EINRNLVQFSRL  
HAO3\_HUMAN EISPDLIQFSRL  
HAO3\_MOUSE EISPDLIQFSRL  
HAO3\_RAT EISPDLIQFSRL

HMGL\_BOVIN TNSKVAQATCKL  
HMGL\_CHICK TNSKVSQAACRL  
HMGL\_HUMAN TSSKVAQATCKL  
HMGL\_MOUSE TSSKVAQATCKL  
HMGL\_RAT TSSKVAQATCKL  
HYES\_HUMAN SDARNPPVVS KM  
HYES\_MOUSE TEVQNPSVTSKI  
HYES\_RAT TEIQNPSVTSKI  
IDHC\_HUMAN ENLKIKLAQAKL  
IDHC\_MICME ENLKAKLAQAKL  
IDHC\_MICOH ENLKAKLAQAKL  
IDHC\_MOUSE ENLKAKLAQAKL  
IDHC\_RAT ENLKAKLAQAKL  
IDI1\_HUMAN NQFVDHEKIYRM  
IDI1\_MESAU SQFVDHEKIH RM  
IDI1\_MOUSE SPFVDHEKI HRL  
IDI1\_RAT SPFVDHEK IH RM  
LUCI\_LUCCR IREILKKPVAKM  
LUCI\_LUCLA IREILKKPVAKM  
LUCI\_PHOPY LIKAKKGGKSKL  
NLTP\_CHICK QNLQLQPGKAKL  
NLTP\_HUMAN QNLQLQPGNAKL  
NLTP\_MOUSE QNLQLQPGKAKL  
NLTP\_RAT QSLQLQPDKAKL  
O55223 MSRFSTLSKAHL  
OXDA\_HUMAN EKKLSRMP PSHL  
OXDA\_MOUSE EKKLSR LPPSHL  
OXDA\_PIG ERNLLTM PPSHL  
OXDA\_RABIT EKKSSRMP PSHL  
OXDA\_RAT EKNLSRMP PSHL  
OXDD\_BOVIN QVLRTPAPKSKL  
OXDD\_HUMAN HALRTPIPKSNL  
P79371 ISRFP SLGKAHL  
PECI\_HUMAN AVVNFLSRKSKL  
PECI\_MOUSE AIMS FVSRKPKL  
PMVK\_HUMAN LENLIEFIR SRL  
PTE1\_HUMAN IRVKPQVSESKL  
PTE1\_MOUSE IRLKPQVSESKL  
PTE2\_HUMAN LGGREGTIPSKV  
PTE2\_MOUSE LDGKKKTIPAKL  
Q27757 LRQMF EKHKSKL  
Q99424 IRPLLQSWRSKL  
SPYA\_CALJA REALQHCPKKKL  
SPYA\_FELCA QEALQRCSR NKL  
SPYA\_HUMAN RAALQHCPKKKL  
SPYA\_MOUSE REALQHCPKNKL  
SPYA\_RABIT REALQHCAQS QL  
SPYA\_RAT REALQHCPKNKL  
URIC\_DROME AQLARKNINSHL  
URIC\_DROPS AQLARKN ISSHL  
URIC\_DROSU AQLARKNLN SHL  
URIC\_DROVI AQLSRKSLKSHL  
URIC\_MOUSE TGTVKRKLPSRL  
URIC\_PAPHA TGTVKRKLSSRL  
URIC\_PIG TGTVKRKLTSRL  
URIC\_RABIT TGTVKRKLSSRL  
URIC\_RAT TGTVRRKLPSRL