

sp|Q63041|A1M_RAT MRRNQQLP---IPVFLLLLLLLLLPRDATAATGKPRYVVLVPSSELYAGVPEKVCVHLNHLNE 56
sp|P06238|A2MG_RAT MGKHLRLSLALLPLLRLLLLLLPTDASAPQKPIYMVMVPSLLHAGTPEKACFLFSHLNE 60
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sp|Q63041|A1M_RAT TVTLNVTLEYGVQYSNLLIDQAVDKDSSYCSFSFTISRPLS-PSALIAVEIKGPTHFFIKK 115
sp|P06238|A2MG_RAT TVAVRVSLBSVRGNQSLFTDLVVDKDLFCHCTSTVPQSSSDEVMFPTVQVKGATHEFRRR 120
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sp|Q63041|A1M_RAT KSMWITKAESPVVFQTDKPIYKPGQTVKFRVVSVDISFRPVNETFPVVYIENPKNRIFQ 175
sp|P06238|A2MG_RAT STVLVKKKESLVAFAQTDKPIYKPGQTVRFRVVSLESHPLNELILPLLQDPKNNRIAQ 180
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sp|Q63041|A1M_RAT WQNVLDLPGHLQLSFPLSVEPALGIYKVVQKDSGKKIEHSFEVKEYVLPKFEVQVKMPK 235
sp|P06238|A2MG_RAT WQNFNLEGLKQLSFLSSEPTQGSYKVVIRTESGRTEVHPFSVEBFVLPKFEVVRTVE 240
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sp|Q63041|A1M_RAT TMAFLEEBLVVTCAGLYTYGKVPVGLVMTMKVCRKYTQSYSNCHGQHSKSICEEFSKQADE 295
sp|P06238|A2MG_RAT TITILEEEMNVSVCGIYTYGKVPVGRVTVNICRKYSNP-SNCFGEBSVAFCEKLSQQLD 299
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sp|Q63041|A1M_RAT KGCFRQVVVKTKVFPQRQKGYDMKIEVEAKIKEDGTGIELTGTGSCEIANTLSKLFKTKAN 355
sp|P06238|A2MG_RAT RGCFSQLVKTKSPQLKRQYEQMLDVHAKIQEBGTGVETGKGLTKITRTITKLSFVNVD 359
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sp|Q63041|A1M_RAT TFRYRPLPFFGQVLLVDEKGGQIPNKNLTVQVNSVRSQFTFTTDEHGLANILIDTNTFTF 415
sp|P06238|A2MG_RAT SHFRQGIPIVFGQVLLVDGRGTPPIYETIFIGADEANLYINTTDDKHLARFSINTDDIMG 419
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sp|Q63041|A1M_RAT SFMGIRVLYKQNNICFDNWWVDEYHTQADHSAARIFSPRSYIQLELVLTGTLACGQTEI 475
sp|P06238|A2MG_RAT TSLTVRAKYKDSNACYGRFWLTEENVEAWHTAYAVFSPRSFLHLSLDPDKLRCDQTEV 479
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sp|Q63041|A1M_RAT RIHFLLEDALDKADKDLTFYYLTKARGSIFNSGSHVLPLEQQKVKGVVSPFIRVPEGMAP 535
sp|P06238|A2MG_RAT QAHYILNNGEAMQELKELVFFYYLMMAKGIVRAGTHVLPKQQQMRGHFSILISMETDLAP 539
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sp|Q63041|A1M_RAT VAKLIVYTIPLNEELIADVQKFDIEKCFANTVNLSPSAQSLPASPDTHLTVKATPLSLCA 595
sp|P06238|A2MG_RAT VARLVLYAILPNGEVVDGTAKEYEINCLANKVDLVFRPNSGLPATRALLSVMASPSQLCG 599
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sp|Q63041|A1M_RAT LTAVDQSVLLKPEAKLSPQSIYNLLPQKAEQGYALGPLPYKGGEN-CIKAEEDITHNGIV 654
sp|P06238|A2MG_RAT LRAVDQSVLLMKPETELASLIYDLPVKDLTGFPQGDQRREDTNGCVKQNDTYINGLIL 659
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sp|Q63041|A1M_RAT YTPKQDLNDNDAYSVFQSIGLKIFTNTRVHKPRYCPMYQAYPPLVYVGEPAQALAMSAIPG 714
sp|P06238|A2MG_RAT YSPVQNTNEEDMYGFLDKMGLKVFNTSNIRKPKVCERLRD-----KGIPA 705
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sp|Q63041|A1M_RAT AGYRSSNIRTSMMMMGASEVAQEVEVRETVRKYFPEWTIWMVPLDLSGDGELPVKVPD 774
sp|P06238|A2MG_RAT AYHLVQSQSHMDAFLESSEPT-----ETRRSYFPEWTIWDLVVDSAGVAEVEVTVPD 758
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sp|Q63041|A1M_RAT TITEWKASAFCLSGTGLGLSSTISHKVFQPFLELTLPYSVVRGEAFILKATVLNLYMPH 834
sp|P06238|A2MG_RAT TITEWKAGAFCLSNDTGLGLSPVQFQAFQPFVVELTMPYSVIRGEAFTLKATVLNLYLPT 818
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sp|Q63041|A1M_RAT CIRIHVSLEMSPDFLAVPVGSHEDSHCICGNERKTVSWAVTPKSLGEVNFATAEALQSP 894
sp|P06238|A2MG_RAT CIRVAVQLEASPDFLAAPEEKRQRSHCICMNRHTASWAVIPKSLGNVNFVSAEALNSK 878
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sp|Q63041|A1M_RAT ELCNGKVAEVPALVQKDTVVKPVIVPEPEGIEKEQTYNTLLCPQDAELQENWTLDPANVV 954
sp|P06238|A2MG_RAT ELCNGEVVVPPEQGKDKTIKSLLVEPEGLNEVTFNSLLCPMAEAVSELIALKLPDVV 938
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sp|Q63041|A1M_RAT EGSARATQSVLGDILGSAMQNLQNLQMPYGCGEQNMVLFVFNIIYVLEYLNETQQLTEAI 1014
sp|P06238|A2MG_RAT EESARASVTVLGDILGSAMQNTQDLKMPYGCGEQNMVLFAPNIYVLDYLNQQLTQTEI 998
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sp|Q63041|A1M_RAT KSKAISYLSGYQRQLNYQHSYDGSYSTFGDRGMRHSQNTWLTAFVLKAFQAQSYIYE 1074
sp|P06238|A2MG_RAT KTKAIAYLNTGYQRQLNYKHRDGSYSTFGDKPGRN-HANTWLTAFVLSFAQARKYIFID 1057
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sp|Q63041|A1M_RAT KTHITNAPNWLMSKQRENGCFQQSGSLLNNAKGGVDEVTLSAYITIALLEMLPVTHS 1134
sp|P06238|A2MG_RAT BVHITQALLWLSQQKDNCGCFRSGSLLNNAKGGVDEVTLSAYITIALLEMSLPVTHP 1117
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sp|Q63041|A1M_RAT VVRNALFCLLETAWASISN-SQESHVYTKALLAYAFALAGNRAKRSEVLESLNKDAVNEE 1193
sp|P06238|A2MG_RAT VVRNALFCLDTAWKARSARGAGGSHVYTKALLAYAFALAGNQDTKKEILKSLDEBAVKEED 1177
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sp|Q63041|A1M_RAT SVHWQRPKNVEENVREMRSFYSYKPRAPSAEVEMTAYVLLAYLTSASSRPTRDLSDDLTT 1253
sp|P06238|A2MG_RAT SVHWTRPQKPSVSV---ALWYQPQAPSAEVEMTAYVLLAYLTPTEPAP---TQEDLTA 1228
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sp|Q63041|A1M_RAT ASKIVKWIISKQNSHGGFSSSTQDVTVALQALSXYGAATFTKSNKEVSVTIESSGTVSGLT 1313
sp|P06238|A2MG_RAT AMLIVKWLTKQNSHGGFSSSTQDVTVALHALSXYGSAATFTTRAKKAAQVTHSSGTFSTKF 1288
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sp|Q63041|A1M_RAT HVNNGNRLLLQEVRLADLPNGYITKVSQSGCVYLQTSLKYNILPEAGEAPFTLKVNTLP 1373
sp|P06238|A2MG_RAT QVNNNNQLLLQVRLTPVPGDYTVKVTGEGCVYLQTSLKYSVLPREE-EFPFTVVVQTLTP 1347
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sp|Q63041|A1M_RAT LNFDKAEHHRKFIHINVSYIGERPNSNMVIVDKVMVSGFIPVKPSVKKLQDQSNIQRT 1433
sp|P06238|A2MG_RAT GTCEDPKAHTSFQIISLNIYSYTGSRSESNMAIADVVMVSGFIPKPTVKMLERSVHVSRT 1407
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sp|Q63041|A1M_RAT VNTNHWLIYIEKLTNQTMGFSFAVEQDIPVKNLKPAPVKVYDYETEFAIEEYSAPFSS 1493
sp|P06238|A2MG_RAT VSNNHVLIYLDKVSNTVNLSTVQQDIPIRDLPKAVVKVYDYEKDEFVAVKYSAPCST 1467
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sp|Q63041|A1M_RAT DSEQGNA 1500
sp|P06238|A2MG_RAT D--YDNA 1472
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