

CLUSTAL 2.1 MULTIPLE SEQUENCE ALIGNMENT

File: /media/morpheus/disk1/fst/pep_msa/GSADate: Tue Feb 1 12:16:05 2022

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                                :           :           *
Pavo_muticus      -----MLLH-----LAAEFDLRRDVVP-----17
Pavo_cristatus    -----MLLH-----LAAEFDLRRDVVP-----17
Gallus_gallus     -----MLLQ-----LAAEFDLQRDVVP-----17
Anolis_carolinensis MAALPQGALLPEAMPPLLR-----LSARFEPAKEVAP-----31
Mus_musculus      -----MALR-----LVTHFDVLEADVLP-----S18
Rattus_norvegicus -----MALR-----LVTHFDVLEADVLP-----S18
Homo_sapiens      -----MALR-----LVADFDLGKDVLP-----WLRAQ22
Pan_troglodytes   -----MALR-----LVADFDLGKDVLP-----WLRAQ22
Macaca_mulatta    -----MALR-----LVADFDLGKDVLP-----WLRAQ22
Callithrix_jacchus -----MMMEEDETENVPVWGSGTGRWCSSLLQMLLFLLCLPSGGGANEQAR46
Canis_lupus_familiaris -----MALR-----LVCFDFDLRKDVLP-----GLRAR22
Equus_caballus    -----MALR-----LAADFDLRKDVLP-----GLRA-21
Bos_taurus        -----MALR-----LIADFDLEKDVLP-----WLRVQ22
Heterocephalus_glaber -----MALR-----LVTFDFDLAADALP-----17
Rhinatrema_bivittatum -----MVLRL-----FHAAFDLRRDVLP-----17
Xenopus_tropicalis -----MVLD-----FCASFSLQDVMP-----17
1.....10.....20.....30.....40.....50.....60

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                                * :*:*. . :*:*. . . :*:*. . :*:
Pavo_muticus      --WLAAR--GWAEALGTSSENSALYIVNVERNGKIIYTWKGNQSTHIGLYDLQTKENEH73
Pavo_cristatus    --WLAAR--GWAEALGTSSENSALYIVNVERNGKIIYTWKGNQSTHIGLYDLQTKENEH73
Gallus_gallus     --WLAAR--GWAVALDTSEKSSALYIVNVERNGKIIYTWKGNQSTHIGLYDLQTKENEH73
Anolis_carolinensis --WLAAGPGDSGVAEAVDNYLETLLHIVNVERNGNIIYTWKGNQGYTHIGLYDHHTEKNOH89
Mus_musculus      LLTQAATTDEGDRAVLETTYGSLRVLNRIERNGNIIYTYKDNKGNAVFGLYDCQTRONEH78
Rattus_norvegicus LLVQAATADEGDEG--AETTLGSLRVLNRIERNGDIITYKDNKGNAVFGIFDCQTRONEH76
Homo_sapiens      RAVSEASGAGSGGADVLENDYESLHVLNVERNGNIIYTYKDDKGNVVFGLYDCQTRONEL82
Pan_troglodytes   RAVSEASGAGSGGADVLENDYESLHVLNVERNGNIIYTYKDDKGNVIFGLYDCQTRONEL82
Macaca_mulatta    RAVSEASEAGSGGADVLENDYESLRLVNVERNGNIIYTYKDDKGNVIFGLYDCQTRONEL82
Callithrix_jacchus RQARAISENSEKQDVVENDYESLRLVNRIERNGNIIYTYKDDKGNVVFGLYDCQTRONEL106
Canis_lupus_familiaris RAASLASGARGGGPGILEDNYESLRLVNVERNGNIIYTYKDDKGNVAFGLYDCRTRONEH82
Equus_caballus    -----AAGARGGGGAGILENNYESLRLVNVERNGNIIYTYKDDKGNVVFGLFDCQTRONEH76
Bos_taurus        LAASAAAGARGGGPGVLENNYECLRLVNVERNRNIIYTYKDNKGNAVFGLYDYQTKONEH82
Heterocephalus_glaber -----ALRAARAGGADVLENTYESLHLLNVERNGNIIYTYKDGKGNVVFGLYDCQTRONEH73
Rhinatrema_bivittatum -----WLRPPGENDVFEKISETLHIINVERSRVTLYTWKGIIQGYTHIGLYDLEAKONEH71
Xenopus_tropicalis -----WILAREASEISKRNSRTLRIVNVERNGRVLFTWKGGDGLTINIGLYDPDLRONEM71
.....70.....80.....90.....100.....110.....120

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**::*:*. . :*:*. . :*:*. . :*:*. . :*:*. . :*:
Pavo_muticus      LYTFEKDLRIISCSINSERTLLAVSFROYTEERVTLQLQSVSKYLTLLEIHPINNVRV133
Pavo_cristatus    LYTFEKDLRIISCSINSERTLLAVSFROYTEERVTLQLQSVSKYLTLLEIHPINNVRV133
Gallus_gallus     LYTFEKDLRIISCSINSERTLLAVSFROYTEERVTLQLQSVSKYLALLIEIHPINNVRV133
Anolis_carolinensis LYTFEKDLQVISCSVNSEKTLAVSFLOSAKEERVNLVFPVSKCLTLLEIHPVNNVKV149
Mus_musculus      LYTFEKDMQAVSCSVNSERTVLAASFQYTTTEGVKN-DLQPGSKCLTLLEIHPVNNVKV137
Rattus_norvegicus LYTFEKDMQAVSCSVNSERTVLAASFQYTTTEGVKN-DLQPGSKCLTLLEIHPVNNVTV134
Homo_sapiens      LYTFEKDLQVFSVNSERTLLAASLVQSTKEGKRN-ELQPGSKCLTLLEIHPVNNVKV141
Pan_troglodytes   LYTFEKDLQVFSVNSERTLLAASLVQSTKEGKRN-ELQPGSKCLTLLEIHPVNNVKV141
Macaca_mulatta    LYTFEKDLQVFSVNSERTLLAASLVQSTKEGKRN-ELQPGSKCLTLLEIHPVNNVKV141
Callithrix_jacchus LYAFEKDLQVFSVNSERTLLAASLVQSTKEGKRN-ELQPGSKCLTLLEIHPVNNVKV165
Canis_lupus_familiaris LYTFEKDLHVISCSINSERTLLAASLVHSAKEGRKN-ELQPGSKCLTLLEIHPVNNVKV141
Equus_caballus    LYTFEKDLQVSCSVNRERTLLAASLVQSTKEGRRN-ELQPGSKCLTLLEIHPVNNVKV135
Bos_taurus        LYTFEKDLQVSCSVNKEKTLATSLVQAAKEGRSN-ELQPGSKCLTLLEIHPINNVKV141
Heterocephalus_glaber LYTFEKDLQVISCSVNNERTLLAASLVQSTKEGRRN-ELQPGSKCLTLLEIHPVNNVKV132
Rhinatrema_bivittatum LYTFEKDLQIISCSINHRTLLAVSYFQSAKEGVNEPLRPVSRCLTLLEIHPFNNVKV131
Xenopus_tropicalis LYSFDKDLLIISCSVNCESLLALSVCNYSVKEHGYE-PLRTVSKYLALLIEIKPVNNVRV130
.....130.....140.....150.....160.....170.....180

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***** : ****:* :.: ***:***: * *: : . *:: :.:*
LKAVDSCVRVQFLYPVEGRNTSTESH-LLLVS190EDKYIEQFDIHVVAEEEHRVVI200QNSGQL
LKAVDSCVRVQFLYPVEGRNTSTESH-LLLVS210EDKYIEQFDIHVVAEEEHRVVI220QNSGQL
LKAVDSCVRVQFLYPVEDRNSSTESH-LLLVS230EDKYIEQFDIH-VAAEEHRVVI240QNSGQL
LKAVDSCIRVQFLYPVAETHSFLESR-LLLISEDKYAEKVDIR-VVRDGGQVVIANS190SQL
LKAVDSCVWVQFLYPQAESHLLPQNH-LLLISEEKYIERFHIQITREDGGRVVI200RNSSHL
LKAVDSCVWVQFLYPQAESHLLAQNH-LLLISEEKYIERFHIQITREDGGRVVI210RNSSHL
LKAVD220SYIWVQFLYPHIESHPLPENH-LLLISEEKYIEQFRIHVAQEDGGRVVI230IKNSGHL
LKAVD240SYIWVQFLYPHIESHPLPENH-LLLISEEKYIEQFRIHVAQEDGGRVVI190IKNSGHL
LKAVD210SYIWVQFLYPHVESHPLPENH-LLLISEEKYIEQLRIHVAQEDGGRVVI220IKNSGHL
LKAVD230SYIWVQFLYPHVESHPLPENHLLLISEEKYIEKFRVRVTQEDGGRVVI240IKNSGHL
LKAVD190SSIWVQFLYPQVESHPPPENH-LLLISEEKYIEKFHIHVIQEDGNKVVL200RD210SGHL
LKAVD220SYIWVQFLPHPHVESNPLPENH-LLLISEEKYIEQFHIQVIQEDGNGVVI230IKNSGHL
LKAVD240SYIWVQFLYPHVESCPQPKNH-LLLISEEKYIEQFHIQVVQEDGGRVVI190IKNSGHL
LKAVDSCVWVQFLYPHGASHPLPQSH-LLLVS210EEKYIEQFHIQVTKEDGGRVVI220IKNSGHL
LKAVDSCVRVQFLYP-AEMVQFPQSR-LLLISEDRYIEQFHISDVTEEGCGVLL230QHCGQL
LKAVDYDIRVQFLYPTIEDVCPFPESH-LLLVS240EEKYIEQFHVILTVEGDSVVIKNSGKL
.....190.....200.....210.....220.....230.....240

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: * : * : : : * . * * * * * : * : : * * . * : * * : . : . : * * : : .
 PRARVADDLIWAQWDMTEQRLFYIVPKESRSTLKCVOFYFPDENFNSTLESQLDISVNDKR
 PRARVADDLIWAQWDMTEQRLFYIVPKESRSTLKCVOFYFPDENFNSTLESQLDISVNDKR
 PRARVADDLIWAQWDMTEQRLFYIVPKESRSTILRCVOFYFPDENFNSTLESQLDISVNDKQ
 SREIADDLIWAQWDMMEQRLFYIVPKESLDTLNCIOFFPDKNFKLTLEAPLDI SLADIA
 PRDLRAEDFVWAQWDLSEORLYYIELKESRSILKCIQFRADESFNLMFEMPLDITLTGLR
 PRERIAEDFVWAQWDVSEQRHYIELQESRSILKCVQFWADESFTIMFEMPLDISLSGLR
 PRDRIAEDFVWAQWDMSEORLYYIDLKKSRSILKCIQFYADESYNLMFEVPLDISLSNSG
 PRDRIAEDFVWAQWDMSEORLYYIDLKKSRSILKCIQFYADERYNLMFEVPLDISLSNSG
 PRDRIAEDFVWAQWDMSEORLYYIDLKKSRSILKCIQFYADESYNLMFEVPLDVLSLSNSG
 QRDRIAENFVWAQWDMSEORLYYIDLKKSRSILKCIQFYADESYNLMFEVPLDISLSNSG
 PRERVAEDFVWAQWDMSEORLYYIVLKKSRSILKCIQFSANEKFNLMFEAPLDITLSASG
 PRERVAEDFVWAQWDTSEORLYYVELKKSRSILKCIQFYADENFNLMFEAPLDIALSDSG
 PRERIAEDFVWAQWDMSEORLYYIDLKKSRSVLKCIQFYAAEHFNLMFEAPLDISLSDSG
 RRDRIAEDFVWAQWDMREORLYYIDLKKSRIILKCIQFNADESFTLMFETPLDIPLSGSG
 PKDRVAEDFIWAQWDILGQRLFYIISKDSKPLLKCIQFYFPDQNFKELLEVPDLALADTG
 PRERIAEDFVFWQWDMLHORLFYIIPKHSVCVLHCIOFYHEDHFKIIFEVSLEIIFVYQA
250.....260.....270.....280.....290.....300

. : * : . : : : : * . * : : * : : * : *
 VKLVNFGYDGDCKDQDVAS-KSLNLQVFTSKAGGLCVCCSLASDIPGEITYCIYFLHKGFS
 VKLVNFGYDGDCKDQDVAS-KSLNLQVFTSKAGGLCVCCSLTSDIPGEITYCIYFLHKGFS
 VKLVNFGYNCDERDVPS-KSLNLQVFTSKAGGLCVCCSLPSDIPGEISYSIYFLHKGFS
 LKPMNLDYNDCHQDREIIP-KPLNIRVITNETGGCLCIYSLIPITSEEVTVSVFLHKGYS
 FKLNVNFGYDYRDREKLC-NQPSLCIFTNHTGSLCMCYSPKSDSREEITYSVFYLHKGYS
 FKLNVNFGYDYRDQAKLC-HQPSLCIFTNHTGSLCVCYSPKSDSWEKITYSVFYFLHKGYS
 FKLNVNFGCDYHQYRDKFS-KHLTLCVFTNHTGSLCVCYSPKCAWQITYSVFYIHKGHS
 FKLNVNFGCDYHQDRDKFS-KHLTLCVFTNHTGSLCVCYSPKCAWQITYSVFYIHKGHS
 FKLNVNFGCDYHQDREKLS-KHLTLCVFTNRTGSLCVCYSPKCAWQITYSVFYIHKGHS
 FKLNVNFGCDLQDQGNLS-KHLTLCVFTNHTGSLCVCYSPKFDSEWIKITYSVFYFHKGHS
 FNLNVNFGCTDLQDQEKFS-EHPTLCVFTNHTGSLCVCYSPKLDSEWIKITYSVFYFHKGHS
 FKLNVNFGYSDLQDKEELS-EHLTLCVFTNHTGSLCVCYCPNFDSEWIKITYSVFYFHKGHS
 FKLNVNFGCGYHEEQEKLS-KHLTLYVFTNHTGSLCACYSQKFDSWEKITYSVFYFHEGYR
 LRLNVNFGVDTYQDRSP-----NVHVFAKSEGGGLCLCYSGPSD-AAEVKYSVAFLHRGCS
 VCLINLGFDPYEVKEQESS-TSLNIQVFTDNTGGCLCLFYLPFKDAKEVKYMVIFLHRGCS
310.....320.....330.....340.....350.....360

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Pavo_muticus      KTFTVSLERPETPQLK-----EVAFLNLDYYVAAAYLPGQFLHLLNIQHPDLLCYSLF 363
Pavo_cristatus    KTFTVSLERPETPQLK-----EVAFLNLDYYVAAAYLPGQFLHLLNIQHPDLLCYSLF 363
Gallus_gallus     KTFTASLERPETPQLK-----EVAFLNLDYYVAAAYLPGQFLHLLNIHHPDLLCYSLF 362
Anolis_carolinensis KTYTVALDEMNSLEGN-----DLTFLNLDYYVAIYLPGHFLHLLNTRHPDLMCYSFF 378
Mus_musculus      KIFTAAPGSADSQVTNGADSQVTDGIAFLNLGYFVAVYSPGHFLHLLNIQHPDLVCHSLF 375
Rattus_norvegicus KTFTVAPGSTDSQVAN-----GVTFNLNLGYFVAVYSPCRFLHLLNIRHPDLICHSLF 364
Homo_sapiens      KTFTTSLENVGSHMTK-----GITFLNLDYYVAVYLPGHFFHLLNVQHPDLICHNLF 371
Pan_troglodytes   KTFTTSLENVGSHVTK-----GITFLNLDYYVAVYLPGHFFHLLNVQHPDLICHNLF 371
Macaca_mulatta    KTFTTSLENVGSHVTK-----GITFLNLDYYVAVYLPGHFFHLLNVQHPDLICHSLF 371
Callithrix_jacchus KTFTTSLKNVGSMTK-----GITFLNLDYYVGVYLPGHFFHLLNVQHPDLICHNLF 395
Canis_lupus_familiaris KTFTAALGSVDSLVTK-----GLTFLNLDYYVAVYLPGHFFHLLNIQHPDLICHSLF 371
Equus_caballus    KTFTAALGSADSRVTK-----GITFLNLDYFVAVYLPGHFFHLLNIQHPDLVCHSLF 365
Bos_taurus        KTFTTTLGSVDSHVTK-----GITFLNLDYYVAVYLPGHFFHLLNIQHPDLICHSLF 371
Heterocephalus_glaber KTFTAAPGSGGSHVTK-----DITFLNLDYYVAVYLPGHFFHLLNIQHPDLICHSLF 362
Rhinatrema_bivittatum KTFFAVAITGPLSGHNR-----DVLFLANLDYYVVVYLPGHFLHLLNTQHPDLMCYHLE 354
Xenopus_tropicalis KTFKVALSAEEMKQLK-----KMSFINLGGYVAVYLPDHFLHLINTRHPDLMCYHLE 361
.....370.....380.....390.....400.....410.....420

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*: : .: .: * : : .: .: * : : * : : * : : *
Pavo_muticus      LTGEDARIDILPNCIS-IQSPPLVSTVLDCIGRMVAVSISDSALLKFLQNSKRDSERLAAL 422
Pavo_cristatus    LTGEDARIDILPNCIS-IQSPPLVSTVLDCIGRMVAVSISDSALLKFLQNSKRDSERLAAL 422
Gallus_gallus     LTGEDARIDMLPNCIS-IQSPPLVSTVLDCIGRLYAMSISDSALLKFLQNSKRDSERLAAL 421
Anolis_carolinensis LTGGEAKINGLHAST-IIISPLKSTVFDIRSTGELFTTIEINKEALFQFLWNKCDTYKLAAL 437
Mus_musculus      LTGNNKIAAVLPPSP-LQSLPGSLVLDCCSGKVYRVTLDSYLLRFLWNAHLDCERMAAL 434
Rattus_norvegicus LTGNNKTAAVLPPSP-LQSLPGSLVLDCCSGKVYRATLDSYLMGFLWNAQLDCEKMAAL 423
Homo_sapiens      LTGNNEIDMLPHCP-LQSLSGSLVLDCCSGKLYRALLSQSLLQLLQNTCLDCEKMAAL 430
Pan_troglodytes   LTGNNEIDMLPHCP-LQSLSGSLVLDCCSGKLYRALLSQSLLQLLQNTCLDCEKMAAL 430
Macaca_mulatta    LTGNNEIDMLPHCP-LQSLSGSLVLDCCSGKLYRALLSQSLLQLLQNTCLDCEKMAAL 430
Callithrix_jacchus LTGNNEIDMLPHSP-LQSLSGSLVLDYCSGKLYRAMLSQSLLQLLQNTCLDCEKMATL 454
Canis_lupus_familiaris LTGNNEVDMLPHSP-LQSLSGSLVLDWCSGKLYRALLNQSYLLQFLWNTQLDCEKMAVL 430
Equus_caballus    LTGNNEVDMLPHGP-LQSLSGSLVLDWCSGRLYRADLNRSYLLEFLWDARLDWEKMAVL 424
Bos_taurus        LTENSEVIDMLPHSP-LQSLSGSLVLDSSGKLYRVLLNQSYLVEFLRSARLDCEKMAAL 430
Heterocephalus_glaber LTGNNSMADVLQSA-LRPLSGSLVLDWGLGKLYRATLNPRLLOFLGAAQLDCDRMATL 421
Rhinatrema_bivittatum LTGEAARLDVLQHSPTQALLEGALLDCCSGIMFTVNINQSVLKLWDSLEDCQKLAAL 414
Xenopus_tropicalis LADTDARIGEICCDGPVQSVLMSVIEYCQGILFSVSLNHLNLLKYLWTCKRDCERLAVL 421
.....430.....440.....450.....460.....470.....480

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** * : : * : : * * * : * * : * *
Pavo_muticus      HCALLCVSSTTDLEMK-----IIWWISENLSTCHSFDPPQEFIIASL 464
Pavo_cristatus    HCALLCVSSTTDLEMK-----IIWWISENLSTCHSFDPPQEFIIASL 464
Gallus_gallus     HCALLCVRSTTDLEMK-----IIWWISENLSTCHSFDPIQEFIIASL 463
Anolis_carolinensis HCLLLHIGSTREIESQ-----VIQWILDKTSICFTFDPMQEFIIASS 479
Mus_musculus      HCILSCSQDPGFPEEQ-----IIQWISEHVSACHSFDLIQEFIIASS 476
Rattus_norvegicus HCALSCSDSPGFP-EQ-----IVQWVSERVSACHSFDLIQEFIIASS 464
Homo_sapiens      HCALYCGQGAQFLEAQ-----IIQWISENVSACHSFDLIQEFIIASS 472
Pan_troglodytes   HCALYCGQGAQFLEGOWRQRADLNTNRVLSDFLKIIQWISENVSACHSFDLIQEFIIASS 490
Macaca_mulatta    HCALYCGQGTRFLEGO-----IIQWISENVSACHSFDLIQEFIIASS 472
Callithrix_jacchus HCALSCGHGAQLLEGO-----IIQWISENVSACHSFDLIQEFIIASS 496
Canis_lupus_familiaris HCVLSCGRDPRFLEAK-----IIQWISENISACHSFDLIQEFIIASS 472
Equus_caballus    HCVLSCRQDTRFLEAK-----IIQWISENISACHSFDLIQEFIIASS 466
Bos_taurus        HCALSHGRDPRRLEAK-----IIQWISENISACHSFDLIQEFIIASS 472
Heterocephalus_glaber HCLLACGRELQPLEAQ-----IIQWISFNSTCHSFDLIQEFIIASS 463
Rhinatrema_bivittatum HCALLHFENPMHWETO-----IIQWICDCASTCATFNPIQEFIVASL 456
Xenopus_tropicalis HCFLLHLDQKPLWETE-----IIEWICENLSVCQMFDPQEFIIASL 463
.....490.....500.....510.....520.....530.....540

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Pavo_muticus      YCRMCPETNNLDKLLPYTSLLDWTGVIPGVACATDIISLPVLEIQNSKGFWEKLDNSNLES 524
Pavo_cristatus    YCRMCPETNNLDKLLPYTSLLDWTGVIPGVACATDIISLPVLEIQNSKGFWEKLDNSNLES 524
Gallus_gallus     YCRMCPETNNLDKLLPYTSLLDWTGVIPGVACATDIISLPVLEIQNSKGFWEKLDNSNLES 523
Anolis_carolinensis YWTMCLDVTNLDKLLPYTPLLQWNEDFPGIICRTQIISMPALKVQHCKDFWEKLSMSLEC 539
Mus_musculus      YWSVYAELDDMGMLLQYSSVLTWNTEIPGIKFTTEELPLPLMKVYGLKGYWAKLNSNLEY 536
Rattus_norvegicus YWSVYPGLDDVDLLPYSSVLTWDEIPGMKLVTEELPLPLMKVYSLKGYWAKLNSNLEY 524
Homo_sapiens      YWSVYSETSNMDKLLPHSSVLTWNTEIPGITLVTEEDIALPLMKVLSFKGYWEKLSNLEY 532
Pan_troglodytes   YWSVYSETSNMDKLLPHSSVLTWNTEIPGITLVTEEDIALPLMKVLSFKGYWEKLSNLEY 550
Macaca_mulatta    YWSVYSETSNMDKLLPHSSVLTWNTEIPGITLVTEEDIALPFMKVLNFKGYWEKLSNLEY 532
Callithrix_jacchus YWSVYSETSNVDKLLPHSSVLTWNTEIPGITLVTEEDIALPLMKVHSFKGYWETLNSNLEN 556
Canis_lupus_familiaris YWSIYPETSNIDKLLPYSSVLTWNTEIPGITLVTEETILPFMKVHSFKGYWEKLSNLEY 532
Equus_caballus    YWSIYPETGNMDKLLPYSSVLTWNTEIPGITLVTEEDIALPLMRVHSFKGYWEKLHSNLEY 526
Bos_taurus        YWSIYPETSNMDKLLPYSSVLTWDEIPGITLVTEETILPLPLMKVHSFKGYWEKLSNLEY 532
Heterocephalus_glaber YWSVYPETSNMDMLLQYSSVLTWDEIPGITLVTEETILPLPLMKVHSFQGYWEKLSNLEH 523
Rhinatrema_bivittatum YRRTIPEASNLSKLLPYTSVPCWNMAIPGIMCTTDIALPIVKKRGLGGFWENLHSGLEY 516
Xenopus_tropicalis YRRMSLETIYLDKLLPYTSLPFWNQIGIEVLWSSDITDMPILKIGMFKGFWEKLFHSELEY 523
.....550.....560.....570.....580.....590.....600

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*:  :  .  :*:  .  :
Pavo_muticus      VKYAEPHLHYHNNVLRREWRNLSEETE-----ERRATAYLRNIFENA 566
Pavo_cristatus    VKYAEPHLHYHNNVLRREWRNLSEETE-----ERRATAYLRNIFENA 566
Gallus_gallus     VKYAEPHLHYHNNVLRREWRNLSEETE-----ERRATAYLRNIFENA 565
Anolis_carolinensis VKYSEPYLHFNYKMLQKEDNLLSEER-----TEERTTYMKNIFGNA 581
Mus_musculus      IKYTKPHLHYHNSVVRREWHNLISEER-----TGKRRSTMYVRNILENA 580
Rattus_norvegicus IKYTKPHLHYHNSVVRREWHNLISEER-----TGKRRSTMYVRNILDNA 568
Homo_sapiens      VKYAKPHFHYNNNSVVRREWHNLISEEK-----TGKRRSAAYVRNILDNA 576
Pan_troglodytes   VKYAKPHFHYNNNSVVRREWHNLISEEK-----TGKRRSAAYVRNILDNA 594
Macaca_mulatta    VKYAKPHFHYNNIVVRREWHNLISEEK-----TGKRRSTAYVRTILDNA 576
Callithrix_jacchus VKYAKPHFHYNNNSVVRREWHNLISEEK-----TGKRRSTAYVRNILDNA 600
Canis_lupus_familiaris VKCSKPCLLYNNSMVKREWHSLISEEK-----TGRRRSMVYVRNIFDNA 576
Equus_caballus    VKYSEPHSHYSNSVVRRWHSLSISEEK-----TGKRRSMAYVRNILDNA 570
Bos_taurus        VKYSKPHLHYNNNSVVRREWHNLISEEK-----TGKRRSTVYVRNILDNA 576
Heterocephalus_glaber VKYATPPSRYTNSVVRREWHNLVAELGCGWGORLRSPTVWSQETGKRRPTVYVRNILHSA 583
Rhinatrema_bivittatum VKYAEPCHFHK--TLRRDWWKLSLSELN-----TEDKRTTVYHKNILDNA 558
Xenopus_tropicalis MKHT---HHNNPVHRRDWCKQIADVD-----TQEWNRNIYQRNILENA 563
.....610.....620.....630.....640.....650.....660

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Pavo_muticus      KKVLSHLDAWDSEERLVPLFQEDDYQQLLMGLMVAQLKDHLMRHLQYVGKKKIDQIVLD 626
Pavo_cristatus    KKVLSHLDAWDSEERLVPLFQEDDYQQLLMGLMVAQLKDHLMRHLQYVGKKKIDQIVLD 626
Gallus_gallus     KKVLSHLDAWDSEERLVPLFQEDDYQQLLMGLMVAQLKDHLMRHLQYVGKKKIDQIVLD 625
Anolis_carolinensis KRVFSSLSNMSNVEGRVLPFFQEDDNQQLLTGLMVAQLKDHLARHLQYIGTKNMEQIAVD 641
Mus_musculus      MKVIASMETRTLEPRLIPFLQEDDRHORLLMGLMVSELRDHLLRHLQGEVKKKIEQMVDL 640
Rattus_norvegicus VKVISNMEMKTFEPRLIPLLOEDDRHORLLMGLMVSELRDHLLRHLQGEVKKKIEQMVDL 628
Homo_sapiens      VKVISNLEARNLGPRLTPLLQEDDSHORLLMGLMVSELKDHFRLRHLQGEVKKKIEQMVDL 636
Pan_troglodytes   VKVISNLEARNLGPRLTPLLQEDDSHORLLMGLMVSELKDHFRLRHLQGEVKKKIEQMVDL 654
Macaca_mulatta    IKVISNLEARNLGPRLTPLLQEDDSHORLLMGLMVSELKDHFRLRHLQGEVKKKIEQMVDL 636
Callithrix_jacchus IKVISNLEARNLEPRLTPLLQEDDSHORLLMGLMVSELRDHFLRHLQGEVKKKIEQMVDL 660
Canis_lupus_familiaris MKVISNLEARNLEPRLTPLFQEDDYHORLLIGLMVSELREHLLRHLQGEVKKKIEQMVDL 636
Equus_caballus    IKVISKLEARHLEPRLTPLFQEDDNHORLLVGLMVSELKDHLRLRHLRGVEKKRIEQMVDL 630
Bos_taurus        IKVISNVEAKNLEPRLTPLFQEDDTHQOLLIGLMVSELREHLLRHLQGEVKKRIEQMVDL 636
Heterocephalus_glaber VKVISNVEARHLEPRLAPLLQEDDTRHLLVGLMVSELKDHLRLYLGVEVKKKIEQMAVD 643
Rhinatrema_bivittatum KTVLSHLCTCNSEHRIAPLFQEDDYQKELIGLIMVRLKDHLTRHLPYLRKNIIDKIALD 618
Xenopus_tropicalis KKVILNMETWSSDQRIVPLHQEDDYLOKDLMLWMLTIKLDHLSQPLLHVGNKNNIDKIVLD 623
.....670.....680.....690.....700.....710.....720

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CLUSTAL 2.1 MULTIPLE SEQUENCE ALIGNMENT

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*  :: * *:  ::: * : :  : :  :  *::* :: ::: :
Pavo_muticus      YVANLLNLVHRIMKEVWKIYQLHSCIFCFD-----ERGSEAEFRVFHIMSRILEAANGMC 681
Pavo_cristatus    YVANLLNLVHRIMKEVWKIYQLHSCIFCFD-----ERGSEAEFRVFHIMSRILEAANGMC 681
Gallus_gallus     YVANLLNLVHRIMKEVWKIYQLHSCIFCFD-----ERGSEAEFRVFHIMSRILEAANGMC 680
Anolis_carolinensis YVSKLLRLIWQIMESVWKKYNLQSHTFSIK-----GQNSQEVAAAFHIMCRILQATNGMC 696
Mus_musculus      YISKLLDLIWCLLETSSWRKHSMPVLVHLN-----SHCSAADFEVFHLMTRILDAASSLC 695
Rattus_norvegicus YISKLLDLVWCLETSSWRKHSVHPVWLHLN-----EHGSPADFEVFHLMTRILDAASSLC 683
Homo_sapiens      YISKLLDLICHIVETNWRKHNLSWVLHFN-----SRGSAAEFAVFHIMTRILEATNSLF 691
Pan_troglodytes   YISKLLDLICHIVETSSWRKHNLSWVLHFN-----SRGSAAEFAVFHIMTRILEATNSLF 709
Macaca_mulatta    YISKLLDLICHILETSWRKHNLSWVLHFN-----SRGSAAEFAVFHIMTRILEATNSLF 691
Callithrix_jacchus YISKLLDLICHILETSWRKHNLSWVLHFN-----SRGSAAEFAVFHIMTRILEATNSLF 715
Canis_lupus_familiaris YISKLLDLICQILETSWRTHHLHPVWLHFN-----RRASAAEFVVFHIMTRILEATMSLF 690
Equus_caballus    YISKQLDLICQILEASWRKHNLSWVLHFN-----RRASAAEFVVFHIMTRILEATHSLF 685
Bos_taurus        YVSKLLDLICQILEASWRKHNLSWVLHFN-----RQASAAEFVVFHIMTRILEATNTSLF 691
Heterocephalus_glaber YTSKLLHLICCIETSSWRKHSLSWALRLD-----GSGSAAEFAVFHLMTRILEATSSLF 698
Rhinatrema_bivittatum YVAKLLDLIGLIMETVWRKHDLSRVFRLD-----GKGSEDEFMFHIMTRVLEAANGLC 673
Xenopus_tropicalis YISKQLDLICLILEVWKKYNLDLYAFSFLFICSSSRGCSDDYSAFYMMCHISESAIKLG 683
.....730.....740.....750.....760.....770.....780

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*****::*  ***:***  .::: ** *: * ***  . :*:***:.. ..*:** *:
Pavo_muticus      MPLPPGFHTLHLGLGVRCPLPLHTLLHYIDNGVLHLTETCVRKLLKDLDDNEKNEKLKFSI 741
Pavo_cristatus    MPLPPGFHTLHLGLGVRCPLPLHTLLHYIDNGVLHLTETCVRKLLKDLDDNEKNEKLKFSI 741
Gallus_gallus     MPLPPGFHSLHLGLGVRCPLPLHTLLHYIDNGVLHLTETCVRKLLKDLDDNEKNEKLKFSI 740
Anolis_carolinensis MPLPPGFHTLLMGLGVRCPLPLHTLLHYIDNGVLHLTEMNAIKLLKELDDTVKNEKLKLSI 756
Mus_musculus      LPLPPGFHSLHTILGVHCLPLYSLLHYIDNGVLLLTETAVTRLMKDLDNNEKNEKLKFSI 755
Rattus_norvegicus FPLPPGFHSLHTILGVHCLPLYNLLHYIDNGVLLLTETAVTRLMKDLDNNEKNEKLKFSI 743
Homo_sapiens      LPLPPGFHTLHTILGVQCLPLHNLHLCIDSGVLLLTETAVIRLMKDLDNTEKNEKLKFSI 751
Pan_troglodytes   LPLPPGFHTLHTILGVQCLPLHNLHLCIDSGVLLLTETAVIRLMKDLDNTEKNEKLKFSI 769
Macaca_mulatta    LPLPPGFHTLHTILGVHCLPLHNLHLCIDSGVLLLTETAVIRLMKDLDNTEKNEKLKFSI 751
Callithrix_jacchus LPLPPGFHTLHTILGVHCLPLHNLHLCIDSGVLLLTETAVTRLMKDLDNTEKNEKLKFSI 775
Canis_lupus_familiaris LPLPPGFHTLHTILGVHCLPLHNLHLCIDSGVLLLTETAVIRLMKDLDNNEKNEKLKFSI 750
Equus_caballus    LPLPPGFHTLHTVLGVRCPLPLHNLHLCIDSGVLLLTETAVTRLMKDLDNTEKSEKLKFSI 745
Bos_taurus        LPLPPGFHTLHMLGVRCPLPLHNLHLCIDSGVLLLTETAVTRLMKDLDNTEKNEKLKFSI 751
Heterocephalus_glaber LPLPPGFHTLHTILGVHCLPLHSLNLYIDNGVLLLTETAVTRLLKDLDDNEKNEKLKFSV 758
Rhinatrema_bivittatum APLPPGFQTLHTVLGVRCPLPLHTLLHYIDNGVLQLTEACVTRLLKDLDDTKNNEKLKFSI 733
Xenopus_tropicalis MPLPPGFQTLHLVLGVRCPLPLGNNLHYIDAGILQLTEAFVAVKLLKELDDNESNEKLKYSI 743
.....790.....800.....810.....820.....830.....840

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Pavo_muticus      VTRLPEVTLDALGLKARHFDHPINANFLARKYVKLLLEKLGNRQCSRP-VPE--RHPVP 798
Pavo_cristatus    VTRLPEVTLDALGLKARHFDHPINANFLARKYVKLLLEKLGNRQCSRP-VPE--RHPVP 798
Gallus_gallus     VTRLPEVTLDALGLKARHFDHPVNAFLARKYVKLLLEKLGNRQCSRP-VPE--RHPVC 797
Anolis_carolinensis LTRLP----EEIGHKICHIWSPARNAVARNYVKFLLKCRNKQQRML-VVD--GLSAR 809
Mus_musculus      IVRLP----PLIGQKVCRLWDHPMSSNIISRNHVARLLKNYR-KEPRNS-MID--KSSFP 807
Rattus_norvegicus IVRLP----PLIGQKICRLWDHPMSSNIISRNHVTRLLQNYK-KQPRNS-MIN--KSSFS 795
Homo_sapiens      IVRLP----PLIGQKICRLWDHPMSSNIISRNHVTRLLQNYK-KQPRNS-MIN--KSSFS 803
Pan_troglodytes   IVRLP----PLIGQKICRLWDHPMSSNIISRNHVTRLLQNYK-KQPRNS-MIN--KSSFS 821
Macaca_mulatta    ILRLP----PLIGQKICRLWDHPMSSNIISRNHVTRLLQNYK-KQPRNS-MID--KSSFS 803
Callithrix_jacchus IVRLP----PLIGQKICRLWDHPMSSNIISRNHVTRLLQNYK-KQPRNS-MID--KSSFS 827
Canis_lupus_familiaris IVRLP----PHIGQKICRLWDHPMSSNIISRNHVTRLLQNYK-KQPRNS-MID--KSPGS 802
Equus_caballus    IVRLP----PLIGQKICRLWDHPMSSNIISRNHVTRLLQNYK-KQPRNS-MID--KSSYS 797
Bos_taurus        IMRLP----PLTGQKICRLWDHPVSSNIISRNHVTRLLQNYK-KQPRNS-MID--KSSFS 803
Heterocephalus_glaber IVRLP----PLIGQKIYRLWDHPVSSNVIARNHVTRLLQNYK-KQPRNS-MIN--KSSFS 810
Rhinatrema_bivittatum VTRLP----EEFCQKIYQVWDHPVSSFIARSYVRLLLVKLAKRPLEQR-SVSDRRLSVC 788
Xenopus_tropicalis IVRLP----ERICQNIHRLWDHAISSNCCIARKYVEQLLCKLKKRECHLAGQAACSQSALS 799
.....850.....860.....870.....880.....890.....900

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CLUSTAL 2.1 MULTIPLE SEQUENCE ALIGNMENT

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Pavo_muticus      VEFLLPNYLTINVLAETIEN--QGVHPYEKQDHINVRFVEEAALKHTMMLLGLRYS 850
Pavo_cristatus    VEFLLPNYLTINVLAETIEN--QGVHPYEKQDHINVRFVEEAALKHTMMLLGLRYS 850
Gallus_gallus     VEFLLPNYLTINVLAETIES--QGVHLYEKQDHINVRFVEEAALKHTMMLLGLRYS 849
Anolis_carolinensis IEFLLPNYLTINILAEVEC--QGWAASVQPENMNVKLVEELALKHTTMLLGL--- 858
Mus_musculus      VEFLLPNYFIEILMGLESSNQALYGFEGHDNVD AEFVEEAALKHTTMLLGL--- 858
Rattus_norvegicus VEFLLPNYFIEILMHLESSNQALHGFEGHDNVD AEFVEEAALKHTTSLGL--- 846
Homo_sapiens      VEFLLPNYFIEILTDIESSNQALYPFEGHDNVD AEFVEEAALKHTTAMLLGL--- 854
Pan_troglodytes   VEFLLPNYFIEILTDIESSNQALYPFEGHDNVD AEFVEEAALKHTTAMLLGL--- 872
Macaca_mulatta    VEFLLPNYFIEILTDIESSNQALYPFEGHDNVD AEFVEEAALKHTTMLLGL--- 854
Callithrix_jacchus VEFLLPNYFIEILTDIESTNQALYPFEGHDNVD AKFVEEAALKHTTMLLGL--- 878
Canis_lupus_familiaris VEFLLPNYFIEILTDIESSNQALYAFEGHDNVD AKFVEEAALKHTTMLLGL--- 853
Equus_caballus    VEFLLPNYFTGILTDIESSRNLNAFEGHDNVD AKFVEEAALKHTTVLLGL--- 848
Bos_taurus        VEFLLPNYFIHILTDIESSNPALYAFEGHDNVD AKFVEEAALKHTTAMLLGL--- 854
Heterocephalus_glaber VDFLLPNYLIIEILTSLESSTQALNDFEGHDNVD AEFVEEAALKHTTMLLGL--- 861
Rhinatrema_bivittatum LEFLPLNYLVTVLSEVEER--ALNPFE-EDNVD MKFVEETALKQTTILLGL--- 836
Xenopus_tropicalis MNFLPLNYLIKMLSTIEE--QALNPFE-EDNIDA AFLEEIALKQTTVLLRLHKS 850
.....910.....920.....930.....940.....950....

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