

CLUSTAL 2.1 MULTIPLE SEQUENCE ALIGNMENT

File: /media/morpheus/disk1/fst/pep_msa/WD036pep Tue Feb 1 14:53:52 2022

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Species	Sequence	Length
Pavo_muticus	-----	
Pavo_cristatus	-----	
Gallus_gallus	-----	
Anas_platyrhynchos	-----	
Chelonia_mydas	-----	
Mus_musculus	-----	
Rattus_norvegicus	-----	
Homo_sapiens	-----	
Pan_troglodytes	-----MCCTEGSLRKRDSSQRAPEAVLCLQLWQRTIVPLDTLKGLGTCFP-----	43
Macaca_mulatta	-----MTVPLDTLKGLDTCFP-----	16
Callithrix_jacchus	-----MCCTAGSLRKRDSSQRAQEAVLNQLRPRIVKLHTLKGLVTCFL-----	43
Heterocephalus_glaber	-----	
Canis_lupus_familiaris	-----	
Equus_caballus	LAEPKSRPDCGLRTELFFSCTENGLRRRDWPRAPEALVNLOPGARIAPRDARRGQLPAFR	60
Bos_taurus	-----	
Latimeria_chalumnae	-----	
Rhinatrema_bivittatum	-----	
Xenopus_tropicalis	-----	

Pavo_muticus	-----NLLNFS-----	6
Pavo_cristatus	-----NLLNFS-----	6
Gallus_gallus	-----MAAER-----AALFAGFRALGRYS SHVPHVRLRYHGRHREFYVATAV	41
Anas_platyrhynchos	-----MAAGGGGGGGGGGLFAGFRALGRYS GHVPHVRLRYHGRHREFYVATAV	48
Chelonia_mydas	-----MSRAGSAVRAAPGGGSALFSGFRALGLYSNHLPHVRLRYHQHREFYELITAA	51
Mus_musculus	-----MAEMESAVEGRTASVLFAGFRALGLFSNEVPHVVRYSAALKRRFYVTTCV	49
Rattus_norvegicus	-----MESAVEGRTASVLFAGFRALGLFSNDIPHVVRYNALKRRFYVTTCV	46
Homo_sapiens	-----MERASEERTASALFAGFRALGLFSNDIPHVVRFSAALKRRFYVTTCV	46
Pan_troglodytes	-SGPELRGAGIAAAMERASERRTASALFAGFRALGLFSNDIPHVVRFSAALKRRFYVTTCV	102
Macaca_mulatta	-SGPELSGAGIAAAMERASERRTASALFAGFRALGLFSNDIPHVVRFSAALKRRFYVTTCV	75
Callithrix_jacchus	PSGPERRGAGIAAAMEQASERRTASALFAGFRALGLFSNDIPHVVRFSAALKRRFCVTTTCV	103
Heterocephalus_glaber	-----MEGGAEGRTASALFAGFRALGLFSNDISHVVRFSAALKRRFYVTTCV	46
Canis_lupus_familiaris	-----MERGAG---SALFAGFRALGLFSTDVAHVLRFSALKRRFFVTTTCV	42
Equus_caballus	-SGLAPRGARTAAAMEGGAGSRTASALFAGFRALGLFSNDVPHVVRFNALKRRFYVTTCV	119
Bos_taurus	-----MERGPGETASALFAGFRALGLFSSDIPHAVRFSAALKRRFYVTTCV	46
Latimeria_chalumnae	-----MSSNKKGSGIFLGFALGLYSNHLAHVVRYHGRHREFYLLTAA	43
Rhinatrema_bivittatum	-----MSGD--GSVLFAGFRALGLYSGHLPHVRLRYHKKHREFYVATAV	41
Xenopus_tropicalis	-----MAEENVARGSVLFTGYRSLGVYSNHLPHVVRYHKKHKEFYLVTSV	45
70.....80.....90.....100.....110.....120	

		* ..*: : * *:*** *:*.:. . *.**.*:*** :.***	
Pavo_muticus	-----VFAGNALLQDITCLAADRMLIFASYG	VDVFHAIARNKEVVHTYEGHK	52
Pavo_cristatus	-----VFAGNALLQDITCLAADRMLIFASYG	VDVFHAIARNKEVVHTYEGHK	52
Gallus_gallus	GRSVHTYNVKKLGIVAVSNALLQDITCLAADRMLIFASYGN	VDVFHAIARNKEVVHTYEGHK	101
Anas_platyrhynchos	GRSVHTYNVKKLGIVAVSNALLQDITCLAADRMLIFASYG	VDVFHAVARNKEVVHTYKGHK	108
Chelonia_mydas	GRSFHTYNVKKLGIVAVSNALLQDITCLAADRMLIFASYGN	LLHFAFARNKEVVHTYDGHK	111
Mus_musculus	GKSFHTYDVQKLSLVAVSNSVPODICCMAADGRLVFAAYGN	VFSAFARNKEIVHTFKGHK	109
Rattus_norvegicus	GKSFHTYDVQKLSLVAVSNSVPODICCMAADGRLVFAAYGN	VFSAFARNKEIVHTFKGHK	106
Homo_sapiens	GKSFHTYDVQKLSLVAVSNSVPODICCMAADGRLVFAAYGN	VFSAFARNKEIVHTFKGHK	106
Pan_troglodytes	GKSFHTYDVQKLSLVAVSNSVPODICCMAADGRLVFAAYGN	VFSAFARNKEIVHTFKGHK	162
Macaca_mulatta	GKSFHTYDVQKLSLVAVSNSVPODICCMAADGRLVFAAYGN	VFSAFARNKEIVHTFKGHK	135
Callithrix_jacchus	GKSFHTYDVQKLSLVAVSNSVPODICCMAADGRLVFAAYGN	VFSAFARNKEIVHTFKGHK	163
Heterocephalus_glaber	GKSFHTYDVQKLSLVAVSNSVPODICCMAADGRLVFAAYGN	VFSAFARNKEVVHTFKGHK	106
Canis_lupus_familiaris	GKSFHTYDVQKLSLVAVSNSLPDICCMAADGRLVFAAYGN	VFSAFARNKEVVHTYKGHK	102
Equus_caballus	GKSFHTYDVQKLSLVAVSNSLPDICCMAADGRLVFAAYGN	VFSAFARNKEVVHTFKGHK	179
Bos_taurus	GKSFHTYDVQKLSLVAVSNSLPDICCMAADGRLVFAAYGN	VFSAFARNKEVVHTFKGHK	106
Latimeria_chalumnae	GKSFHTYNVKKLGIVAVSNALPEDITCLAADRMLVFASHGN	LVKAFARSKEVVHTYKGHS	103
Rhinatrema_bivittatum	GRSFHTYNVKKLGIVAVSNALPEDITCLAADRMLVFVSHG	KHLKAFARNKEVVHTYEGHN	101
Xenopus_tropicalis	GKSFHTYNVKKLGIVAVSNALPEISCLAADRMLVFAAHSN	LLKAFARNKEVVHTYQGE	105
130.....140.....150.....160.....170.....180		

CLUSTAL 2.1 MULTIPLE SEQUENCE ALIGNMENT

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Pavo_muticus      ARIHLLQPFGDHIIISVDVANVLIVWDIQSEEEYLQVVFDKATFAVSAILHPSTYLNKILL 112
Pavo_cristatus    ARIHLLQPFGDHIIISVDVANVLIVWDIQSEEEYLQVVFDKATFAVSAILHPSTYLNKILL 112
Gallus_gallus     ARIHLLQPFGDHIIISVDVANVLIVWDIQSEEEYLQVVFDKATFAVSAILHPSTYLNKILL 161
Anas_platyrhynchos ARIHLLQPFGDHIIISVDVANVLIVWDIQSEEEYLQVVFDKATFAVSAILHPSTYLNKILL 168
Chelonia_mydas    GEIRLLQPFGDHVISVDIDNVLIVWDVQSEEEYLQLTFDKAVFAVSAIMHPSTYLNKILL 171
Mus_musculus      AEIHLLOPFGDHVISVDTDVSLIIWHIYSEEEYLQLTFDKSVFKISTILHPSTYLNKVLL 169
Rattus_norvegicus AEIHLLOPFGDHVISVDTDVSLIIWHIYSEEEYLQLTFDKSVFKISTILHPSTYLNKVLL 166
Homo_sapiens      AEIHFLQPFGDHIIISVDTDGILIIWHIYSEEEYLQLTFDKSVFKISAILHPSTYLNKILL 166
Pan_troglodytes   AEIHLLOPFGDHIIISVDTDGILIIWHIYSEEEYLQLTFDKSVFKISAILHPSTYLNKILL 222
Macaca_mulatta    AEIHLLOPFGDHIIISVDTDVSLIIWHIYSEEEYLQLTFDKSVFKISAILHPSTYLNKILL 195
Callithrix_jacchus AEIHLLOPFGDHIIISVDTDVSLIIWHIYSEEEYLQLTFDKTVFKISAILHPSTYLNKILL 223
Heterocephalus_glaber AEIQLLOPFGDHIIISVDTDVSLIIWHIYSEEEYLQLAFDKLVFKISAILHPSTYLNKILL 166
Canis_lupus_familiaris AEIRLLQPFGDHIIISVDTDVSLIIWHIYSEEEYLQLTFDKSVFKISAILHPSTYLNKILL 162
Equus_caballus    AEIHLLOPFGDHVISVDSVSLIIWHIYSEEEYLQLTFDKSVFKISTILHPSTYLNKILL 239
Bos_taurus        AEIHLLOPFGDHVISVDTDVSLIIWHIYSEEEYLQLTFDKSVFKISAILHPSTYLNKILL 166
Latimeria_chalumnae AEVCLLLPFGDHVISVDRDNVLIWDVQSEEEYLQLNFDKTVFGISAIMHPSTYLNKVLL 163
Rhinatrema_bivittatum TEIHLLOPFGDHVISVDLDNLLIIWEVQTEEEYLQLNFDKVKVFTVSAIMHPSTYLNKILL 161
Xenopus_tropicalis AEIHLVQPFGDHVISVDTSNVLMIWDIKSEEEYLQLTFDKTNFAISAIMHPSTYLNKILL 165
.....190.....200.....210.....220.....230.....240

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Pavo_muticus      GSEQGGLQLWNIRSNKLLYSFPGWSLG-VTTLAQAPAVDVIAVGLVSGHIIHNKIFDET 171
Pavo_cristatus    GSEQGGLQLWNIRSNKLLYSFPGWSLG-VTTLAQAPAVDVIAVGLVSGHIIHNKIFDET 171
Gallus_gallus     GSEQGGLQLWNIRSNKLLYSFPGWGLG-VTTLAQAPAVDVIAVGLVSGHIIHNKIFDET 220
Anas_platyrhynchos GSEQGGLQLWNIRSNKLLYSFPGWGLG-VTTLAQAPAVDVVAVGLVSGHIIHNKIFDET 227
Chelonia_mydas    GSEQGSLQLWNVKS NKLLYSFPGWGLGGVTTLEQAPAVDVVAIGLVSGHIIHNKIFDET 231
Mus_musculus      GSEQGSLQLWNIRSNKLLYTFPGWKVG-VTALQAPAVDVVAIGLVSGQVVIHNKIFDET 228
Rattus_norvegicus GSEQGSLQLWNIRSNKLLYTFPGWKVG-VTALQAPAVDVVAIGLVSGQVVIHNKIFDET 225
Homo_sapiens      GSEQGSLQLWNVKS NKLLYTFPGWKVG-VTALQAPAVDVVAIGLVSGQVVIHNKIFDET 225
Pan_troglodytes   GSEQGSLQLWNVKS NKLLYTFPGWKVG-VTALQAPAVDVVAIGLVSGQVVIHNKIFDET 281
Macaca_mulatta    GSEQGSLQLWNVKS NKLLYTFPGWKVG-VTALQAPAVDVVAIGLVSGQVVIHNKIFDET 254
Callithrix_jacchus GSEQGSLQLWNVKS NKLLYTFPGWKVG-VTALQAPAVDVVAIGLVSGQVVIHNKIFDET 282
Heterocephalus_glaber GSEQGSLQLWNVKS NKLLYTFPGWKVG-VTALQAPAVDVVAIGLVSGQVVIHNKIFDET 225
Canis_lupus_familiaris GSEQGTLQLWNIRSNKLLYTFPGWKVG-VTALQAPAVDVVAIGLVSGQVVIHNKIFDET 221
Equus_caballus    GSEQGSLQLWNVKS NKLLYTFPGWKVG-VTALQAPAVDVVAIGLVSGQVVIHNKIFDET 298
Bos_taurus        GSEQGSLQLWNVKS NKLLYTFAGWKVG-VTALQAPAVDVVAIGLVSGQVVIHNKIFDET 225
Latimeria_chalumnae GSQQGRLQLWNVKS NKLLYTFPGWGS-VTVLQAPAVDVVAIGLVSGQVVIHNKIFDET 222
Rhinatrema_bivittatum GSQQGSLQLWNIRSNKLLYTLQGWGS-ITTLQAPALDVVAIGLVSGQVVIHNKIFDET 220
Xenopus_tropicalis GSQQGSLQLWNIRSNKLLYTFQGWGS-VTVLQAPAVDVVAIGLVTCGKIIHNKIFDET 224
.....250.....260.....270.....280.....290.....300

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~::~: *****:~::~:~** * :~::~: *****:~::~:~** .:~::~:*****:~**
Pavo_muticus      LMKFQQDWGPITAIISFRTDGHPVMAAGSPVGHIALWDLEKKLMSQMRNAHSTAVAGMSF 231
Pavo_cristatus    LMKFQQDWGPITAIISFRTDGHPVMAAGSPVGHIALWDLEKKLMSQMRNAHSTAVAGMSF 231
Gallus_gallus     LMKFQQDWGPITAIISFRTDGHPVMAAGSPVGHIALWDLEKKLMSQMRNAHSTAVAGMSF 280
Anas_platyrhynchos LMKFQQDWGPITAIISFRTDGHPVMAAGSPVGHIALWDLEKKLISQMRNAHSTAVAGMSF 287
Chelonia_mydas    LMKFQQDWGPITAIISFRTDGHPVMAAGSPIGHIALWDLEKKLISQMRNAHSTAVAGMSF 291
Mus_musculus      LMKFHQDWGPITISISFRTDGHPVMAAGSPCGHIGLWDLEDKKLINQMRNAHSTAVAGMSF 288
Rattus_norvegicus LMKFHQDWGPITISISFRTDGHPVMAAGSPCGHIGLWDLEDKKLINQMRNAHSTAVAGMSF 285
Homo_sapiens      LMKFRQDWGPITISISFRTDGHPVMAAGSPCGHIGLWDLEDKKLINQMRNAHSTAVAGMSF 285
Pan_troglodytes   LMKFRQDWGPITISISFRTDGHPVMAAGSPCGHIGLWDLEDKKLINQMRNAHSTAVAGMSF 341
Macaca_mulatta    LMKFRQDWGPITISISFRTDGHPVMAAGSPCGHIGLWDLEDKKLINQMRNAHSTAVAGMSF 314
Callithrix_jacchus LMKFRQDWGPITISISFRTDGHPVMAAGSPCGHIGLWDLEDKRLINQMRNAHSTAVAGMSF 342
Heterocephalus_glaber LMKFRQDWGPITISISFRTDGHPVMAAGSPCGHIGLWDLEDKKLINQMRNVHSTAVAGMSF 285
Canis_lupus_familiaris LMKFRQDWGPITISISFRTDGHPVMAAGSPCGHIGLWDLEDKKLINQMRNAHSTAVAGMSF 281
Equus_caballus    LMKFRQDWGPITISISFRTDGHPVMAAGSPCGHIGLWDLEDKKLINQMRNAHSTAVAGMSF 358
Bos_taurus        LMKFRQDWGPITISISFRTDGHPVMAAGSPCGHIGLWDLEDKKLINQMRNAHSTAVAGMSF 285
Latimeria_chalumnae LMKFQQDWGPVTAISFRTDGHPVMAAGSPVGHIGLWDLEKKLAGQMRDVHSTAVAGMSF 282
Rhinatrema_bivittatum LMKFQQDWGPITISISFRTDGHPVMAAGSPVGHIGLWDLEKKLAGQMRDAHSTAVAGMSF 280
Xenopus_tropicalis LMTFQQDWGPITISLISFRTDGHPVMAAGSPVGHIGLWDLEKKLAGQMRDAHSTAVAGMSF 284
.....310.....320.....330.....340.....350.....360

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Pavo_muticus      VPGEPLLIITNGADNAIRVWIFDGPGGTGRVLRSRMGHSAPPTKIRYHGONGEQILSAGQD 291
Pavo_cristatus    VPGEPLLIITNGADNAIRVWIFDGPGGTGRVLRSRMGHSAPPTKIRYHGONGEQILSAGQD 291
Gallus_gallus     VPGEPLLIITNGADNAIRVWIFDGPGGTGRVLRSRMGHSAPPTKIRYHGONGEQILSAGQD 340
Anas_platyrhynchos VPGEPLLIITNGADNAIRVWIFDGPGGTGRVLRSRMGHSAPPTKIRYHGONGEQILSAGQD 347
Chelonia_mydas    VQGEPLLVITNGADNAIKIWIFDGPGGXGRFVRCRMGHSAPPTKIRYHGOMRQOILSAGQD 351
Mus_musculus      LHREPLLVITNGADNALRIWIFDGPAGEGRLLRFRMGHSASLTKIRYYGONGQQOILSASQD 348
Rattus_norvegicus LHREPLLVITNGADNALRIWIFDGPAGEGRLLRFRMGHSASLTKIRYYGONGQQOILSASQD 345
Homo_sapiens      LHREPLLVITNGADNALRIWIFDGPAGEGRLLRFRMGHSAPLTNIRYYGONGQQOILSASQD 345
Pan_troglodytes   LHREPLLVITNGADNALRIWIFDGPAGEGRLLRFRMGHSAPLTNIRYYGONGQQOILSASQD 401
Macaca_mulatta    LHREPLLVITNGADNALRIWIFDGPAGEGRLLRFRMGHSAPLTNIRYYGONGQQOILSASQD 374
Callithrix_jacchus LHREPLLVITNGADNALRIWIFDGPAGEGRLLRFRMGHSAPLTNIRYYGONGQQOILSASQD 402
Heterocephalus_glaber LHREPLLVITNGADNAVRIWIFDGPAGEGRLLRFRMGHSAPLTNIRYYGONGQQOILSASQD 345
Canis_lupus_familiaris LHREPLLVITNGADNALRIWIFDGPAGEGRLLRFRMGHSAPLTNIRYYGONGQQOILSASQD 341
Equus_caballus    LHREPLLVITNGADNALRIWIFDGPAGEGRLLRFRMGHSAPLTNIRYYGONGQQOILSASQD 418
Bos_taurus        LHREPLLVITNGADNALRIWIFDGPAGEGRLLRFRMGHSAPLTNIRYYGONGQQOILSASQD 345
Latimeria_chalumnae VHGEPLLIITNGADNAIRVWIFDTPGGGGGRLLRCRMGHSAPPNKIKYHGONGQQOILSAGQD 342
Rhinatrema_bivittatum VQGEPLLIITNGADNALRVWIFDGPGGSGRLLRFRMGHSAPPTKIRYHGONGEQILSAGQD 340
Xenopus_tropicalis VHGEPLLIITNAADNAIRVWIFDGPGGGGRLLRFRMGHSAPPTFIRHHGADGKQILSAGQD 344
.....370.....380.....390.....400.....410.....420

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*****:*:::*.***: * : * * : * : * : * : *
Pavo_muticus      GTLQSFSTVHERFNKSLGH-----GSINKKSKKKGLQHDT-MALPPITAFASEVA 341
Pavo_cristatus    GTLQSFSTVHERFNKSLGH-----GSINKKSKKKGLQHDT-MALPPITAFASEVA 341
Gallus_gallus     GTLQSFSTVHERFNKSLGR-----GSINKKSKKKGLQHDT-MALPPITAFASEVA 390
Anas_platyrhynchos GTLQSFSTVHERFNKSLGR-----GSINKKSKKKGLQHDT-MALPPITAFASEVA 397
Chelonia_mydas    GTLQSFSTVHERFNKSLGR-----GSINKKSKKKGLQYDT-MALPPITAFATEIA 401
Mus_musculus      GTLQSFSTVHEKFNKSLGH-----GLVNKKIVKRKGLQNTMSVRLPPIITQFAAEEA 399
Rattus_norvegicus GTLQSFSTVHEKFNKSLGH-----GLVNKKIVKRKGLQNTMSVRLPPIITQFAAEEA 396
Homo_sapiens      GTLQSFSTVHEKFNKSLGH-----GLINKKRVKRKGLQNTMSVRLPPIITKFAAEEA 396
Pan_troglodytes   GTLQSFSTVHEKFNKSLGH-----GLINKKRVKRKGLQNTMSVRLPPIITKFAAEEA 452
Macaca_mulatta    GTLQSFSTVHEKFNKSLGH-----GLINKKRVKRKGLQNTMSVRLPPIITKFAAEEA 425
Callithrix_jacchus GTLQSFSTVHEKFNKSLGH-----GLINKKRVKRKGLQNTMSVRLPPIITKFAAEEA 453
Heterocephalus_glaber GTLQSFSTVHEKFNKSLGH-----GLINKKRVKRKGLQNTMSVRLPPIITKFAAEEA 396
Canis_lupus_familiaris GTLQSFSTIHEKFNKSLGH-----GLINKKRVKRKGLQNTMSVRLPPIITKFAAEEA 392
Equus_caballus    GTLQSFSTIHEKFNKSLGH-----GLINKKRVKRKGLQNTMSVRLPPIITKFAAEEA 469
Bos_taurus        GTLQSFSTVHEKFNKSLGHAIFPSHLTPGLINKKRVKRKGLQNTMSVRLPPIITKFAAEEA 405
Latimeria_chalumnae GTLQSFSTVHERFNKSLGH-----GSINKKSKKKGLQYDT-MKLPPITEFASETA 392
Rhinatrema_bivittatum GTLQSFSTVHEKFNKSLGH-----GSINKKSKKKGLQYDT-MKLPPITAFASEVA 390
Xenopus_tropicalis GTLQSFSTVHEDRYSKSLGH-----GSVDKAKSKKVGLRCLSLRLPPIITAFASEEA 394
.....430.....440.....450.....460.....470.....480

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:::***::***: .:**** : :*: : *.* . . **.***::****.
Pavo_muticus      HQNDWDGIVACHQGYITCTTWNQKTSMGHKLKRPPEFSKNKPIDIYATAVDITTCGNFA 401
Pavo_cristatus    HQNDWDGIVACHQGYITCTTWNQKTSMGHKLKRPPEFSKNKPIDIYATAVDITTCGNFA 401
Gallus_gallus     HQNDWDGIIACHQGYITCTTWNQKTSMGHKLKRPPEFSKNKPIDIYATAVDITTCGNFA 450
Anas_platyrhynchos HQSDWDGIVACHQGYITCTTWNQKTSMGHKLKRPPEFSKNKPIDIYATAVDITTCGNFA 457
Chelonia_mydas    RQNDWDGIIACHQGYITCTTWNQKTSMGHKLKPKAFSKNKPLDVYATVVDITTCGNFA 461
Mus_musculus      RQSDWDGIIACHQKRSCTSTWNYQSTIGAYFLKPRGVKTSN--LTATAVDITTCGNFA 457
Rattus_norvegicus RQSDWDGVIACHQKLSCTSTWNYQSTIGAYFLKPKNVKANS---ATATAVDITTCGNFA 453
Homo_sapiens      RESDWDGIIACHQKLSCTSTWNYQSTIGAYFLKPKELKKDD---ITATAVDITTCGNFA 453
Pan_troglodytes   RESDWDGIIACHQKLSCTSTWNYQSTIGAYFLKPKELKTDD---ITATAVDITTCGNFA 509
Macaca_mulatta    RESDWDGIIACHQKLSCTSTWNYQSTIGAYFLKPKEMKKDD---ITATAVDITTCGNFA 482
Callithrix_jacchus RESDWDGIIACHQKLSCTSTWNYQSTIGAYFLKPKELKKDD---ITATAVDITTCGNFA 510
Heterocephalus_glaber RESDWDGIIACHQKLSCTSTWNYQSTIGAYFLKPKELKRDD---ITATAVDITTCGNFA 453
Canis_lupus_familiaris RESDWDGIIACHQKLSCTSTWNYQSTIGAYFLKPKELKKDD---ITATAVDITTCGNFA 449
Equus_caballus    RESDWDGIIACHQKLSCTSTWNYQSTIGAYFLKPKELKKDD---VTATAVDITTCGNFA 526
Bos_taurus        RESDWDGIVACHQKLSCTSTWNYQSTIGAYFLKPKELKTD---ITATAVDITTCGNFA 462
Latimeria_chalumnae RQSDWDGIIACHRGYLIATTWNYHRSMSGVHKLKPKRFHKDRINVTATVVDITTCGNFA 452
Rhinatrema_bivittatum HQSDWDGIIACHQGYVSCCTTWNQKTSMGHKLKPKDRFKKNKPLHVYATAVDISSCGNFV 450
Xenopus_tropicalis RQSDWDGIVACHQQLSCTTWNVYKCTMGHKLKPPDLKNSKTLEIYATASDITTCGNFT 454
.....490.....500.....510.....520.....530.....540

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Pavo_muticus      VIGMSTGQVDVYNMQSGIHRGRYGRERAHEGSIRGVAVDGLNQLTITAGSDGLIKFWKFK 461
Pavo_cristatus    VIGMSTGQVDVYNMQSGIHRGRYGRERAHEGSIRGVAVDGLNQLTITAGSDGLIKFWKFK 461
Gallus_gallus     VIGMSTGQVDVYNMQSGIHRGRYGREKAHEGSIRGVAVDGLNQLTITAGSDGLIKFWKFK 510
Anas_platyrhynchos VIGMSTGRVDVYNMQSGLHRGCGYKERAHEGSIRGLAVDGLNQLTITAGSEGLIKFWKFK 517
Chelonia_mydas    VIGLSTGHVDVYNMQSGIHRGHYGKEKAHEGAIRGVAVDGLNQLTITAGSEGLIKFWKFK 521
Mus_musculus      IIGLSSGAVDVYNMQSGIHRGNFGDDKAHTGSVRGVAVDGLNQLVVTAGSERLLKFWNFK 517
Rattus_norvegicus IIGLSSGAVDVYNMQSGIHRGSFSGDEKAHAGSVRGVAVDGLNQLAITAGSERLLKFWNFK 513
Homo_sapiens      VIGLSSGTVDVYNMQSGIHRGSFSGKDAQHKGSVRGVAVDGLNQLTIVTTGSEGLLKFWNFK 513
Pan_troglodytes   VIGLSSGTVDVYNMQSGIHRGSFSGKDAQHKGSVRGVAVDGLNQLTIVTTGSEGLLKFWNFK 569
Macaca_mulatta    VIGLSSGTVDVYNMQSGIHRGSFSGKDAQHKGSVRGVAVDGLNQLTIVTTGSEGLLKFWNFK 542
Callithrix_jacchus VIGLSSGTVDVYNMQSGIHRGSFSGKDAQHKGSIIRGVSDGLNQLTIVTTGSEGLLKFWNFK 570
Heterocephalus_glaber VIGLSSGTVDVYNMQSGIHRGSFSGKDAQHKGSVRGVAVDGLNQLTITTTGSEGLLKFWNFK 513
Canis_lupus_familiaris VIGLSSGTVDVYNMQSGIHRGSFSGEDQAHKGSVRGVAVDGLNQLTITAGSEGLLKFWNFK 509
Equus_caballus    VIGLSSGTVDVYNMQSGIHRGSFSGKDAQHKGSVRGVAVDGLNQLTIVTTGSEGLLKFWNFK 586
Bos_taurus        VIGLSSGAVDVYNMQSGIHRGSFSGRDAQHKGSVRVFAVDGLNQLTITAGSEGLVLFWNFK 522
Latimeria_chalumnae VVGIISSGHVDVYNMQSGLHRGHYGKEKAHEGPIRGVAVDGLNQLTITTTGGDGIIFWKFK 512
Rhinatrema_bivittatum VIGLSTGEVDRYNIQSGIHRGHYGAEKAHEGAVRGVAVDGLNELTITAGSDRQIKFWKFK 510
Xenopus_tropicalis VIGLNSGSVDVYNIQSGIHRGHYGKEKAHGGPIRGVVVDGLNQVTITAGSDKLIKFWKFK 514
.....550.....560.....570.....580.....590.....600

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Pavo_muticus      AKDLVHSTDLSSSPSGILLHRD SGILGIAFDDFSISVLDVETWKIVRKFSGHHGRINDMT 521
Pavo_cristatus    AKDLVHSTDLSSSPSGILLHRD SGILGIAFDDFSISVLDVETWKIVRKFSGHHGRINDMT 521
Gallus_gallus     AKDLVHSTDLSSSPSGILLHRD SGILGIAFDDFSISVLDIETWKIVRKFSGHHGRINDMT 570
Anas_platyrhynchos TKDLVHSTDLSSSPSGILLHRD SGILGIAFDDFSISVFDIETRKIVRKFSGHHGRINDMT 577
Chelonia_mydas    TMDLVYSANLSSSPSAMLHRD SGILGIAFDDFSISVLDVETRRIVRKFSGHHGRINDMT 581
Mus_musculus      SKVLIHSLGLDSSPNMMLLHRD SGILGLAMDDFSIAVLDIETRKIVREFSGHHGRINDMT 577
Rattus_norvegicus SKVLVHSLGLDSSPNMMLLHRD SGILGLALDDFSITVLDIETRKIVREFSGHHGRINDMT 573
Homo_sapiens      NKILIHSVSLSSSPNIMLLHRD SGILGLALDDFSISVLDIETRKIVREFSGHHGRINDMT 573
Pan_troglodytes   NKILIHSVSLSSSPNIMLLHRD SGILGLALDDFSISVLDIETRKIVREFSGHHGRINDMT 629
Macaca_mulatta    NKILIHSVSLSSSPNIMLLHRD SGILGLALDDFSISVLDIETRKIVREFSGHHGRINDMT 602
Callithrix_jacchus SKSLIHSVGLSSSPNIMLLHRD SGILGLALDDFSISVLDIETRKIVREFSGHHGRINDMT 630
Heterocephalus_glaber SKGLIHSVSLDSSPNMMLLHRD SGILGLALDDFSISVLDIETRKIVREFSGHHGRINDMT 573
Canis_lupus_familiaris NKILIHSMSLSSSPNIMLLHRD SGILGLALDDFSICVLDIETRKIVREFSGHHGRINDMT 569
Equus_caballus    SKILIQMSLSLSSPNIMLLHRD SGILGLALDDFSISVLDIETRKIVREFSGHHGRINDMT 646
Bos_taurus        NKVLIHSVSLDSSPNMMLLHRD SGILGLALDDFSISVLDIETRKIVREFSGHHGRINDMT 582
Latimeria_chalumnae TKELVHTTIELNTSPAIMLLHRES SMLAIASDDFSICVLDIETRKIVRKFSGHHGRINDMT 572
Rhinatrema_bivittatum TFLVHTTDLSSPPNVLLQRD SGMLGIAFDDFSICVLDIETRKIVRKFSGHHGRINDMT 570
Xenopus_tropicalis TKELIQTTSLVPPPTAVLLHRES SAMLVAALDDFTLNIFDIETRRIVRREFSGHHGRINDMT 574
.....610.....620.....630.....640.....650.....660

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*****:***:*** :*. *****:*. *.:*** :.***:*** *****:***:***
Pavo_muticus      FSPDGRWLITSSMDCTIKTWDLPSGCLIDCFLLDSAAVSLTMSPTGDFLASAHVDDLGIY 581
Pavo_cristatus    FSPDGRWLITSSMDCTIKTWDLPSGCLIDCFLLDSAAVSLTMSPTGDFLASAHVDDLGIY 581
Gallus_gallus     FSPDGRWLITSSMDCTIKTWDLPSGCLIDCFLLDSAAVSLTMSPTGDFLASAHVDDLGIY 630
Anas_platyrhynchos FSPDGRWLITSSMDCTIKTWDLPSGCLIDCFLLDSAAVSLTMSPTGDFLASAHVDDLGIY 637
Chelonia_mydas    FSPDGRWLITASMDCTIKTWDLPSGCLIDCFLLDSAAISISMSPTGDFLASSHVDDLGIY 641
Mus_musculus      FSPDGRWLISAAMDCSVRTWDLPSGCLIDCFLLDSAPLNVMTMSPTGDFLATSHVDHLGIY 637
Rattus_norvegicus FSPDGRWLISAAMDCSVRTWDLPSGCLIDCFLLDSAPLNVMTMSPTGDFLATSHVDHLGIY 633
Homo_sapiens      FSPDGRWLISAAMDCSIRTWDLPSGCLIDCFLLDSAPLNVMSPTGDFLATSHVDHLGIY 633
Pan_troglodytes   FSPDGRWLISAAMDCSIRTWDLPSGCLIDCFLLDSAPLNVMSPTGDFLATSHVDHLGIY 689
Macaca_mulatta    FSPDGRWLISAAMDCSIRTWDLPSGCLIDCFLLDSAPLNVMSPTGDFLATSHVDHLGIY 662
Callithrix_jacchus FSPDGRWLISAAMDCSVRTWDLPSGCLIDCFLLDSAPLNVMSPTGDFLATSHVDHLGIY 690
Heterocephalus_glaber FSPDGRWLISAAMDCSIRTWDLPSGCLVDCFLLD SAPLNVMTMSPTGDFLATSHVDHLGIY 633
Canis_lupus_familiaris FSPDGRWLISASMDCSVRTWDLPSGCLIDSFLVDSAPLNVMTMSPTGDFLATSHVDHLGIY 629
Equus_caballus    FSPDGRWLISASMDCSIRTWDLPSGCLIDSFLVDSAPLNVMTMSPTGDFLATSHVDHLGIY 706
Bos_taurus        FSPDGRWLISASMDCSIRTWDLPSGCLIDSFLVDSAPLNVMTMSPTGDFLATSHVDHLGIY 642
Latimeria_chalumnae FSPDGRWLVTASMDCTLRTWDLPSGCLIDCFLLDSAAVSITMSPTGDFLASSHVDDLGIY 632
Rhinatrema_bivittatum FSPDGRWLITSSMDCTIKVWDLPSGCLVDCFLLD SAALSITMSPTGDFLASSHVDDLGIY 630
Xenopus_tropicalis FSPDGRWLLTASMDCTIKVWDLPSGCLIDCFLLDSAAVSLTMSPTGDFLATAHVDDLGIY 634
.....670.....680.....690.....700.....710.....720

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CLUSTAL 2.1 MULTIPLE SEQUENCE ALIGNMENT

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****  :*:::*:**.*: *   : *** .      :      ::::*.** **
LWSNRSLSYSLVSLRPLPADYEPSVITLPGTCLPLQDVDAEG--EDKC-DEMIEYDSPEQLG  639
LWSNRSLSYSLVSLRPLPADYEPSVITLPGTCLPLQDVDAEG--EDKC-DEMIEYDSPEQLG  639
LWSNRSLSYSLVSLRPLPADYEPSVITLPGTCLPLQDVDAEG--EDKC-DEMIEYDSPEQLG  688
LWSNRSLSYSLVSLRPLPADYEPSMVLPGTCLPLEDVDAEG--EETC-DEMIEYDSPEQLG  695
LWSNRSLSYSLVSLRPLSADYEPSVIMLPGTCAQDVDTITGD--EETS--NEMIDYNSPAQLE  699
LWSNISLSYSVVSLRPLPPDYVPSIVMLPGTCQTQGMEDLEEKTEPS--DEMIEYESPEQLS  696
LWSNISLSYSVVSLRPLPPDYVPSIVMLPGTCQTQGLEALEEQIEPS--DEMIEYESPEQLN  692
LWSNISLSYSVVSLRPLPADYVPSIVMLPGTCQTQDVEVSEETVEPS--DELIEYDSPEQLN  692
LWSNISLSYSVVSLRPLPADYVPSIVMLPGTCQTQDVEVSEETVEPS--DEMIEYDSPEQLN  748
LWSNISLSYSVVSLRPLPADYVPSVVMLPGTCQTQDVEVSEETVEPS--DEMIEYDSPEQLN  721
LWSNISLSYSVVSLRPLPTDYVPSVVMLPGTCQTQDVEVSEETVEPS--DEMIEYDSPEQLD  749
LWSNISLSYSVVSLRPLPTDYVPSVVMLPGICQTQDVEVSEENAEP--DEMIEYSPEQLS  692
LWSNISLSYSVVSLRPLPTDYVPLVVMLPGTCQNEDEVFSEETIEPS--DEMIEYDSPEQLN  688
LWSNVLSYSVVSLRPLPTDYVPSVVMLPGTCQIQDVEFSEETIEPS--DEMIEYDSPEQLN  765
LWSNISLSYSVVSLRPLPTDYVPSVVMLPGTCQTQDVELSEETVEPS--EEMIEYDSPEQLN  701
LWSNSTLSYSLVSLHPLPADYEPSVVTLPGTCPALDVGATEVEEPS--DEMIEYDSPEQLG  690
LWSNKSLSYSLVSLRPLPADYEPATVMLPGTCVPQVEEPTTEDEAETR--DEMIEYESPEQLE  689
LWSNKSLSLSLISLRPLPADFEPTVVMLPGGADEMEVEETVVEENTNEDIMEYESPEQLG  694
.....730.....740.....750.....760.....770.....780

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***.**.:*****:***:***:***:*****:*****:***:*.:.:
EQLVTLSSLPE$SRWKNLLSLDVIKKKNKPREPPKVPKSAPFFFIPTVPGLIPRYAAPEQEN 699
EQLVTLSSLPE$SRWKNLLSLDVIKKKNKPREPPKVPKSAPFFFIPTVPGLIPRYAAPEQEN 699
EQLVTLSSLPE$SRWKNLLSLDVIKKKNKPREPPKVPT$APFFFIPTVPGLIPRYAAPEQDN 748
EQLVTLSSLPE$SRWKNLLNLDIIKKKNKPREPPKVPT$APFFFIPTVPGLIPRYAAPEQEN 755
EQLVTLSSLPE$SRWKNLLNLDVIKKKNKPREPPKVPKSAPFFFIPTVPGLIPRYAAAEQEN 759
EQLVTLSSLPE$SRWKNLLNLDVIKKKNKPKPEPPKVPQ$APFFFIPTVPGLVPRFAVPEPSS 756
EQLVTLSSLPE$SRWKNLLNLDVIKKKNKPKPEPPKVPQ$APFFFIPTVPGLVPRFAVPEPTS 752
EQLVTLSSLPE$SRWKNLLNLDVIKKKNKPKPEPPKVPKSAPFFFIPTIPGLVPRYAAPEQNN 752
EQLVTLSSLPE$SRWKNLLNLDVIKKKNKPKPEPPKVPKSAPFFFIPTIPGLVPRYAAPEQNN 808
EQLVTLSSLPE$SRWKNLLNLDVIKKKNKPKPEPPKVPKSAPFFFIPTIPGLVPRYAAPEQNN 781
EQLVTLSSLPE$SRWKNLLNLDVIKKKNKPREPPKVPR$APFFFIPTIPGLVPRYAAPEQNN 809
EQLVTLSSLPE$SRWKNLLNLDVIKKKNKPKPEPPKVPKSAPFFFIPTVPGLVPRYAAPEQNN 752
EQLVTLSSLPE$SRWKNLLNLDVIKKKNKPKPEPPKVPKSAPFFFIPTVPGLVPRYAVPEQNN 748
EQLVTLSSLPE$SRWKNLLNLDVIKKKNKPKPEPPKVPKSAPFFFIPTVPGLVPRYAVPEQNN 825
EQLVTLSSLPE$SRWKNLLNLDVIKKKNKPKPEPPKVPKSAPFFFIPTIPGLVPRYAAPEQNN 761
EQLVTLSSLPE$SRWKNLLNLDVIKKRNKPKPEPPKVPKSAPFFFIPTVPGLVPRFAVPEGEQA 750
AHLVTMSTLPSRWKNLLNLDVIKKRNKPKPEPPKVPKTAPFFFIPTVPGLVPHFAS-LDQA 748
EKLVTLSLLPE$SRWKNLLNLDIIKQRNKPEPPKVPKSAPFFFIPTLGLVPRFAS-DAQG 753
.....790.....800.....810.....820.....830.....840

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.....850.....860.....870.....880.....890.....900		
D-TQSKVVNIGVLAQKSDFYVHLEELALSTNEYTTPLNLLKDLGPSNIETELRSLAPEGGG	758	
D-TQSKVVNIGVLAQKSDFYVHLEELALSTNEYTAPLNLLKDLGPSNIETELRSLAPEGGG	758	
D-TQSKVVNIGVLAQKSDFYVHLEELALSTNEYTAPLNLLKDLGPSNIETELRSLAPEGGG	807	
D-AQSKVVNIGVLAQKSDFYIHLEELALSTNKYAVPLNVLKDLGPSNIETELRSLAPEGGG	814	
D-GHPKVVNLGVLAQKSDFYVQLEELALSSNEYKSPSLXKMGKPSIETELRGLAPDAGG	818	
DPQQSKVVNLGILAQKSNFYLKLEEGLLNNQYEAALNLLKELGPSGIETELRNLSPPDGG	816	
DPQQSKVVNLGILAQKSNFYLKLEEGLLNNQYEGALNLLKELGPSGIETELRNLSPPDGG	812	
DPQQSKVVNLGVLAQKSDFCLKLEEGLVNNKYDTALNLLKESGPSGIETELRSLSPDCGG	812	
DPQQSKVVNLGVLAQKSDFCLKLEEGLVNNKYDTALNLLKESGPSGIETELRSLSPDCGG	868	
DPQQSKVVNLGVLAQKSDFCLKLEEGLVNNKYDTALNLLKESGPSGIETELRSLSPDCGG	841	
DPQQSKVVNLGVLAQKSDFCLKLEEGLVNNKYDMALNLLKELGPSGIETELRSLSPDCGG	869	
NPQQSKVVNLGILAQKSDFCLKLEEGLLNNKYETALNLLKELGPSGIETELRNLSPPDGG	812	
DPHQSKVVNLGVLAQKSDFCLKLEEGLVNNKYEAALNLLKELGPSGIETELRSLSPDCGG	808	
DPQQSKVVNLGILAQKSDFCLKLEEGLVNNKYEAALNLLKELGPSGIETELRSLSPDCGG	885	
DPQQSKVVNLGILAQKSDFCLKLEEGLVNNKYEAALNLLKELGPSGIETELRSLSPDCGG	821	
NTEQSKVLNLGVLAQKSNFYMQLEDALLHEDYNSPVKLLKEMGPSIDTELRLAPEGGG	810	
DGPQSKVVNLGILAQKSNFCLQLEELALSNQDYDSAVKMLKEMGPSAVETELRGLAPEGGG	808	
DTSQSKIVNLGVLEQSAFFVQLEELALTRQNSVPLKTIKEMGPSVDTELRLAPENGG	813	

CLUSTAL 2.1 MULTIPLE SEQUENCE ALIGNMENT

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*: * ** :* *: : *: *****:*****: : .:* *:** : .:~:
SLEVMLSFLKMGIMMLNKKYNFELAQAYLALFLKLHLKIISSSEPSLLEEISRLSTQVEET 818
SLEVMLSFLKMGIMMLNKKYNFELAQAYLALFLKLHLKIISSSEPSLLEEISRLSTQVEET 818
SLEVMLSFLRMIGIMMLNKKYNFEIAQAYLALFLKLHLKIISSSEPSLLEEISVLKSQLEET 867
SVEVMLSFLRMIGIMMLNKKYNFELAQAYLALFLKLHLKIISSSEPSLLEEISRLSTQLEET 874
SVEVMKSFLRLIGTMLNTKCDFFEXAQAYLALFLKLHLKILSSSEPVILLEEIAKVSTQLEET 878
SVEVMRSFLSMIGIMMLDRKDRDFELAQAYLALFLKLHLRLTLPSEPALLEELVKLSSQVEKD 876
SIEAMRSFLSMIGVMLDRKDRDFELAQAYLALFLKLHLRMLPSEPVLLEEIVKLSSQVEED 872
SIEVMQSFLKMGIMMLDRKDRDFELAQAYLALFLKLHLKMLPSEPVLLEEITNLSSQVEEN 872
SIEVMQSFLKMGIMMLDRKDRDFELAQAYLALFLKLHLKMLPSEPVLLEEITNLSSQVEEN 928
SVEVMQSFLKMGIMMLDRKDRDFELAQAYLALFLKLHLKMLPSEPVLLEEITNLSSQVEEN 901
SIEVMQSFLKMGIMMLDRKDRDFELAQAYLALFLKLHLKMLPSEPILLEEITNLSSQVEEN 929
SIEVMQSFLKMIEMMLERKDRDFELAQAYLALFLKLHLKMLPSEPVLLEEMTKLSSQVEEN 872
SVDIMQSFLKMGIMLDRKDRDFELAQAYLALFLKLHLKMLPSEPVLLEEITKLSSQVEEN 868
SVEVMQSFLKMGIMMLDRKDRDFELAQAYLALFLKLHLKMLPSEPVLLEEITKLSSQVEEN 945
SVGIMKSFLKMGITMLDRKDRDFELAQAYLALFLKLHLKTLPSSEPVLLEEMTKLSSQLEEN 881
SLEVMSFLRMIGSMHTRKDRDFELAQAYLALFLKLHLKVIVEDPTLLEETSKVSTQLEET 870
SIEVMLSFLRLIGSMLKTKRDFELAQAYLALFLKLHLKLLPSEPTLVEEMSTLSGQLEET 868
SVEVMQSFLKMIQSMLDTKKDFEVAQAYLALFLKLHLKFISSEPILLEEINAVLSQLEET 873
.....910.....920.....930.....940.....950.....960

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* ::: *****: :* *:***:*
WIHLRSLFNQSLCVLTYMK$SALL      841
WIHLRSLFNQSLCVLTYMK$SALL      841
WIHLRSLFNQSLCVLTYMK$SALL      890
WIHLRTLFNQSLCVLTYMK$SALL      897
WIHLQTLFNQSLCVLNYMK$SALL      901
WTHLQSLFNQSMCVLNYIK$SAFL      899
WTHLQSLFNQSMCVLNYIK$SALL      895
WTHLQSLFNQSMCILNYLK$SALL      895
WTHLQSLFNQSMCILNYLK$SALL      951
WTHLQSLFNQSMCILNYLK$SALL      924
WIHLQSLFNQSMCILNYLK$SALL      952
WIHLQSLFNQSMCILNYIK$SALL      895
WIHLQSLFNQSMCILNYIK$SALL      891
WIHLQSLFNQSLCVLNYIK$SALL      968
WIHLQLLFNQSLCVLNYIK$SALL      904
WAHMQTLFNQSMCLLGYMK$SALL      893
WLHLQTLFNQSLNLVSYMK$SALL      891
WLNLQTLFNQSLCVLNYMK$SALL      896
.....970.....980...
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