

## CLUSTAL 2.1 MULTIPLE SEQUENCE ALIGNMENT

**File: /media/morpheus/disk1/fst/pep\_msa/NUD142.p Tue Feb 1 14:11:34 2022**

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Mus_musculus	-----MSSVKRNP	PKKEMISELHSS	19
Rattus_norvegicus	-----MSSVKRNP	KREIISELHSS	19
Canis_lupus_familiaris	-----MAQFVLYHFQ	SILRQHSIREMSSIK-SPKQEIISQFHYS	38
Bos_taurus	-----MSSVKRSLN	QEIISQFHYS	19
Equus_caballus	-----MSSIKGSPKQEIISQFHCS		19
Homo_sapiens	-----MSSVKRSLKQEIIVTQFHCS		19
Pan_troglodytes	-----MEEMSSVKRSPKQEIIVTQFHCS		22
Macaca_mulatta	-----MSSVKRSPKQEIIVTQFHCS		19
Callithrix_jacchus	-----MSSVKRSPKQEIIVTQFHCS		19
Heterocephalus_glaber	-----MSSVKGSPKREIIYEFHSA		19
Pavo_muticus	-----MTNFEKNFROEMVSQLHNF		19
Pavo_cristatus	-----MTNFEKNFROEMVSQLHNF		19
Gallus_gallus	-----MTNFEKNFHOEMVSQLHNF		19
Anas_platyrhynchos	-----MTAFEKNLHOEMISQLHNF		19
Chelonia_mydas	-----MFVSLPPAPTS	SKITVTAARRGLSSSAETMTTTERNPMQOMISQLHSF	47
Xenopus_tropicalis	-----MQ-NPRKEMVSQLHNL		15
Rhinatrema_bivittatum	MRVSWALRCTWLPAGSARLTLCRPTQATGVAVLSRAFTGVNNMSEMORITPKQEMVSQLHNS		60
	1.....10.....20.....30.....40.....50.....60		

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Mus_musculus	AAEGNVAKLAGILSHSPSLNNETSENGWTALMYAARNGHDPVVOFLLEKGCDRSLVNKAR	79
Rattus_norvegicus	AAEGDVAKLAGILSHSPSLNNETSENGWTALMYAARNGHDPDAVOFLLEKGCDRSSVNKSR	79
Canis_lupus_familiaris	AAEGDIARLTVILSHSPSLNNETSENGWTALMYAARNGHDPVVOFLLDKGCDRSIVNKS	98
Bos_taurus	AAEGDIAKLTAILSHSPSLNNETSENGWSALMYAARNGHDPVVOFLLEKGCDRSIVNKS	79
Equus_caballus	AAEGDIARLTGILSLSPSLNNETSENGWTALMYAARNGHDPHVVOFLLEKGCDRSIVNKS	79
Homo_sapiens	AAEGDIAKLTGILSHSPSLNNETSENGWTALMYAARNGHPEIVVOFLLEKGCDRSIVNKS	79
Pan_troglodytes	AAEGDIAKLTGILSHSPSLNNETSENGWTALMYAARNGHPEIVVOFLLEKGCDRSIVNKS	82
Macaca_mulatta	AAEGDIAKLTGILSHSPSLNNETSENGWTALMYAARNGHPEIVVOFLLEKGCDRSIVNKS	79
Callithrix_jacchus	AAEGDIAKLTGILSHSPSLNNETSENGWTALMYAARNGHPEIVVOFLLEKGCDRSLVNKS	79
Heterocephalus_glaber	AAEGDMAKLTGLSHSPSLNDETSSENGWTALMYAARNGHSDVAOFLLEKGCDRSIVNKS	79
Pavo_muticus	AAVGDAARLKALLSRSPSLINATAGNGWTALMYAARNGHFDVVOILLEGGCDRSIINKS	79
Pavo_cristatus	AAVGDAARLKALLSRSPSLINATAGNGWTALMYAARNGHFDVVOILLEGGCDRSIINKS	79
Gallus_gallus	AAVGDAARLKALLSRSPSLINATAGNGWTALMYAARNGHFDVVRILLEGGCDRSIINKS	79
Anas_platyrhynchos	AAVGDVAKLKALLSHSPSLINAAADNGWTALMYGARNGHFDVVRILLEGGCDRSIVNKS	79
Chelonia_mydas	AATGDKIRLTALCCHSPSLINEAAENGWTALMYAARNGHFEIVVOILLEKGCDRSIVNKS	107
Xenopus_tropicalis	CALGDTTKLHTLLSHSSSIINERSEHGWSALMFGARNGHFDVVNMLEKGCDRTLVNKS	75
Rhinatrema_bivittatum	AASGDKMQLSALLGHSLSLIDEATEKGWTALMYGARNGHFSVVOMLLEKGCDRSLVNKTG	120
	.....70.....80.....90.....100.....110.....120	

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Mus_musculus	QTALDIAAFWGYR	GHIANLLANAKGGK	PPWFLTN	-EV	DECENYFSRTLLDR	RS	DKR	NSDW	138
Rattus_norvegicus	QTALDIAVFWGYKH	IANLLANAKGGK	PPWFLTS	-DVD	GCENYFSRTLLDR	RS	DKRT	NSDW	138
Canis_lupus_familiaris	QTALDIAKFWGYKH	IANLLANAKSGMK	PPWFLSN	-EVE	ECENYFSRTLLDR	K	SEKR	NSDW	157
Bos_taurus	QTALDIAKFWGYKH	IANLLANAKGGK	PPWFLTN	-EVE	ECENYFSKTL	LLDRK	SEKR	NSDW	138
Equus_caballus	QTALDIAKFWGYKH	IANLLANAKGGK	PPWFLTN	-DVE	ECENYFSRTLLDR	K	SEKR	NSDW	138
Homo_sapiens	QTALDIAVFWGYKH	IANLLATAKGGK	PPWFLTN	-EVE	ECENYFSRTLLDR	K	SEKR	NSDW	138
Pan_troglodytes	QTALDIAVFWGYKH	IANLLATAKGGK	PPWFLTN	-EVE	ECENYFSKTL	LLDRK	SEKR	NSDW	141
Macaca_mulatta	QTALDIAVFWGYKH	IANLLATAKGGK	PPWFLTN	-EVE	ECENYFSKTL	LLDRK	SEKR	NADW	138
Callithrix_jacchus	QTALDIAAFWGYKH	IADLLTPAKGGK	HPWFLTN	-EVE	ECENYFSKTL	LLDRK	SEKR	NSDW	138
Heterocephalus_glaber	QTALDIAVFWGYKH	TANLLANAKGGK	PPWFLTN	-EVE	ECENYFSRTLLDR	K	SEKR	NSDW	138
Pavo_muticus	QTALDIAKFWGYKH	IANLLANAKGGQ	KPGFLPT	-EV	KEYSNYFGT	TLLDRR	SDKR	ISDW	138
Pavo_cristatus	QTALDIAKFWGYKH	IANLLANAKGGQ	KPGFLPT	-EV	KEYSNYFGT	TLLDRR	SDKR	ISDW	138
Gallus_gallus	QTALDIAKFWGYKH	IANLLANAKGGQ	KPGFLPT	-EV	KEYSNYFGT	TLLDRR	SDKR	ISDW	138
Anas_platyrhynchos	QTALDIAKFWGYKH	IANLLANVKGQ	KPNFLST	-EV	KERENYFSMT	LLDRR	SDKR	TSNW	138
Chelonia_mydas	QTALDIAKFWGYKH	IANLLANVKGQ	KPFFLPN	-DA	EYENYFSRTLLDR	R	SDKR	TSKW	166
Xenopus_tropicalis	QTALDIAKFWGHK	HANLLTNTRGG	SKPHFLN	-A	KEEHNYSIT	ILLDRK	SDKR	TDMNW	134
Rhinatrema_bivittatum	QTALDIAKFWGHK	HIVDLLSSKGG	PKPWFPLD	GAA	EGPENYFGS	LLDRK	SEKR	TANW	180
	.....	130	140	150	160	170	180		

## CLUSTAL 2.1 MULTIPLE SEQUENCE ALIGNMENT

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Mus_musculus	LQAKESHPTTVYLLFSDLNPLVTLGGNKESQOPEVRLCOLNYPDVKGYLAAPEKITLVF	198
Rattus_norvegicus	LQAKESHPTTVYILFSDLNPLVTLGGNKESQOPEVRLCOLNYADIKDYLAQPEKITLVF	198
Canis_lupus_familiaris	LLAKESHPATVYILFSDLNPLVTLGGNKETFOQPEVRLCOLNYTDVKDYLSQPEKITTLIF	217
Bos_taurus	LLAKESHPATVYILFSDLNPLVTLGGNKESFOQPEVRLCOLNYTDIKDYLAQPEKITTLIF	198
Equus_caballus	LLAKESHPATVYILFSDLNPLVTLGGNKESQOPEVRLCOLNYTDIKDYLAQPEKITTLIF	198
Homo_sapiens	LLAKESHPATVFILFSDLNPLVTLGGNKESFOQPEVRLCOLNYTDIKDYLAQPEKITTLIF	198
Pan_troglodytes	LLAKESHPATVFILFSDLNPLVTLGGNKESFOQPEVRLCOLNYTDIKDYLAQPEKITTLIF	201
Macaca_mulatta	LLAKESHPATVFILFSDLNPLVTLGGNKESFOQPEVRLCOLNYKDIKDYLAQPEKITTLIF	198
Callithrix_jacchus	LLAKESHPATVFILFSDLNPLVTLGGNKESFOQPEVRLCOLNYTDIKDYLAQPEKITTLIF	198
Heterocephalus_glaber	LLAKESHPTVYILFSDLNPLVTLGGNKESFOQPEVRLCOLNYTDVKDYLAQPEKITTLIF	198
Pavo_muticus	LSKKQNHHPATVYILFSDLSPVLVTLGGAEKSQOPEVRLCRLHKKDVQQCMSQTEEVTLIF	198
Pavo_cristatus	LSKKQNHHPATVYILFSDLSPVLVTLGGAEKSQOPEVRLCRLHKKDVQQCMNQTTEVTLIF	198
Gallus_gallus	LSKKQSHPATVYILFSDLSPVLVTLGGAEKSQOPEVRLCRLHKKDVEQYMIQTTEFTLIF	198
Anas_platyrhynchos	LSRKQSHPATVYILFSDLNPLVTLGMGTERSOPEVKLCRLYHKDVERYSQTEEVTLIF	198
Chelonia_mydas	LNTKQNHHPATVYILFSDLNPLVTLGGEEGDSQOPEVRLCRLCHKDVKEYMSQTEEVTLIF	226
Xenopus_tropicalis	LKSKQTQASTVYILFSDLNPLVTLVHVGGRDTPEEPEIKLCRLQSDVKEYLLNPEDVTLIF	194
Rhinatrema_bivittatum	LKTKQSQAATVYVILFSLKNPLVTLSTREEENSQPKVKLCRLHDEVKEFLSQPEDVFLIF	240
	.....190.....200.....210.....220.....230.....240	

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Mus_musculus	LGVELEMRKGS--PAQAGGVP-EEEEGLVAVFALGIEPGAAEEFKORHENCYFLHPPMP			255
Rattus_norvegicus	LGVELEMRKGS--HAHAGGVP-EGEEDGLVAVFVLGIEPGAAEEFKORHENCYFLHPPMP			255
Canis_lupus_familiaris	LGVELEMKKES--FNYAGEVP-REE-DGLVAVFALGVDSVAEEFKORHENCYFLHPPMP			273
Bos_taurus	LGVELEMKKEF--FNYAGEIS-KEEEDGLVAVFALGIDTVAAEEFKORHENCYFLHPPMP			255
Equus_caballus	LGVELEMKKEL--LNYGVEVP-GGE-DGLVAVFALADPVAEEFKORNENCYFLHPPMP			254
Homo_sapiens	LGVELEIKDKL--LNYAGEVP-REEEDGLVAVFALGIDPIAAEEFKORHENCYFLHPPMP			255
Pan_troglodytes	LGVELEIKDKL--LNYAGEVP-REEEDGLVAVFALGIDPIAAEEFKORHENCYFLHPPMP			258
Macaca_mulatta	LGVELEMKDKL--LNYAGEVP-REEEDGLVAVFALGIDPIAAEEFKORHENCYFLHPPMP			255
Callithrix_jacchus	LGVELEMKKEL--LNFAGEVP-REEEDGLVAVFALGIDPIAAEEFKORHENCYFLHPPMP			255
Heterocephalus_glaber	LGVELEMNKKL--HDHAREVP-REEEDQLVAVFALGIDLIAAEFKORHENCYFLHPPMP			255
Pavo_muticus	LGVDLQLHMSL-LAAINGKVLQEDDEDGLVAVFALSINPTSAAERFKOKHEDCYFLHPPMP			257
Pavo_cristatus	LGVDLQLHMSL-LAAINGKVLQEDDEDGLVAVFALSINPTSAAERFKOKHEDCYFLHPPMP			257
Gallus_gallus	LGVDLQFHMLTVAAHNGKVLQEDDEDGLVAVFALSINPTSAAERFKOKHEDCYFLHPPMP			258
Anas_platyrhynchos	LGVELQFNKNF-MAARNEKVL EEEEDDGLVAVFALSIDPTSAAEKFKOKHEDCYFLHPPMP			257
Chelonia_mydas	LGVELQL-----ATLSGGVLKEDEEDGLVAVFALSINATFAEQFKOKHEDCYFLHPPMP			280
Xenopus_tropicalis	LGVEKQWKT---VHSSDPKRDSGGEDDGLIAWFGLNADKVSTEQFEKKHEGSCYFLQPPMP			251
Rhinatrema_bivittatum	LGVELPLNRNL-LEPTEGSRMEEQEDGLVAVFALNVDSAEQFSQKHDSYFLQPPMP			299
	.....250.....260.....270.....280.....290.....300			

	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48	49	50	51	52	53	54	55	56	57	58	59	60	61	62	63	64	65	66	67	68	69	70	71	72	73	74	75	76	77	78	79	80	81	82	83	84	85	86	87	88	89	90	91	92	93	94	95	96	97	98	99	100	101	102	103	104	105	106	107	108	109	110	111	112	113	114	115	116	117	118	119	120	121	122	123	124	125	126	127	128	129	130	131	132	133	134	135	136	137	138	139	140	141	142	143	144	145	146	147	148	149	150	151	152	153	154	155	156	157	158	159	160	161	162	163	164	165	166	167	168	169	170	171	172	173	174	175	176	177	178	179	180	181	182	183	184	185	186	187	188	189	190	191	192	193	194	195	196	197	198	199	200	201	202	203	204	205	206	207	208	209	210	211	212	213	214	215	216	217	218	219	220	221	222	223	224	225	226	227	228	229	230	231	232	233	234	235	236	237	238	239	240	241	242	243	244	245	246	247	248	249	250	251	252	253	254	255	256	257	258	259	260	261	262	263	264	265	266	267	268	269	270	271	272	273	274	275	276	277	278	279	280	281	282	283	284	285	286	287	288	289	290	291	292	293	294	295	296	297	298	299	300	301	302	303	304	305	306	307	308	309	310	311	312	313	314	315	316	317	318	319	320	321	322	323	324	325	326	327	328	329	330	331	332	333	334	335	336	337	338	339	340	341	342	343	344	345	346	347	348	349	350	351	352	353	354	355	356	357	358	359	360
Mus_musculus	ALL	L	K	E	K	E	A	G	V	V	A	Q	A	R	S	V	L	A	W	H	S	R	Y	K	F	C	P	T	C	G	S	A	T	K	I	E	E	G	G	Y	K	R	V	C	V	R	E	T	C	P	S	L	Q	G	V	H	N	315																																																																																																																																																																																																																																																																																																														
Rattus_norvegicus	ALL	L	K	E	K	E	A	G	V	V	A	Q	A	R	S	V	L	A	W	H	S	R	Y	K	F	C	P	T	C	G	S	T	T	K	I	E	E	G	G	Y	K	R	V	C	V	R	E	N	C	P	S	L	H	G	V	H	N	315																																																																																																																																																																																																																																																																																																														
Canis_lupus_familiaris	ALL	L	K	E	K	E	A	G	V	V	A	Q	A	R	S	V	L	A	W	H	S	R	Y	K	F	C	P	T	C	G	S	A	T	K	I	E	E	G	G	Y	K	R	V	C	L	K	E	D	C	P	S	L	H	G	V	H																																																																																																																																																																																																																																																																																																																

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* .*****  *** *:**** :*****:* **.***:*****:**
TSYPRVDPVVIMQVIHPDGTKCLLGRQKRFPFGMFTCLAGFIEPG----- 360
TSYPRVDPVVIMQVIHPDGTKCLLGRQKRFPFGMFTCLAGFIEPG----- 360
TSYPRVDPVVIMQVIHPDGTKCLLGRQKRFPFGMFTCLAGFIEPG----- 378
TSYPRVDPVVIMQVIHPDGTKCLLGRQKRFPFGMFTCLAGFIEPG----- 360
TSYPRVDPVVIMQVIHPDGTKCLLGRQKRFPFGMFTCLAGFIEPG----- 359
TSYPRVDPVVIMQVIHPDGTKCLLGRQKRFPFGMFTCLAGFIEPG----- 360
TSYPRVDPVVIMQVIHPDGTKCLLGRQKRFPFGMFTCLAGFIEPG----- 363
TSYPRVDPVVIMQVIHPDGTKRCLLGRQKRFPFGMFTCLAGFIEPG----- 360
TSYPRVDPVVIMQVIHPDGTKCLLGRQKRFPFGMFTCLAGFIEPG----- 360
TSYPRVDPVVIMQVIHPDGNHCLLGRQKRFPFGMFTCLAGFVEPG----- 362
TSYPRVDPVVIMQVIHPDGNHCLLGRQKRFPFGMFTCLAGFVEPG----- 362
TSYPRVDPVVIMQVIHPDGNHCLLGRQKRFPFGMFTCLAGFVEPG----- 363
TSYPRVDPVVIMQVIHPDGNHCLLGRQKRFPFGMFTCLAGFVEPG----- 362
TSYPRVDPVAVIMQVHPDGNQCLLGRQKRFPFGMFTCLAGFVEPVRCKGGS LKGRGMCQR 400
TSYPRVDPVVIMLVHPDGNHCLLGRKKIFPAGMFSCLAGFIEPG----- 356
TCYPRVDLSVIMLVHPDGNQCLLGRKKQFPAGMFSCLAGFIEPG----- 404
.....370.....380.....390.....400.....410.....420

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[illegible]

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WFTREQVVDVLTGKGQQAFFVPPSRAIAHQLIKHWGMNPNL      462
WFTREQVVDVLTGKGQQAFFVPPSRAIAHQLIKHWGMNPNSL      462
WFTREQVVDVLTGKGQQAFFVPPSRAIAHQLLKHWIGMNPNL      480
WFTREQVVDVLTGKGQQAFFVPPSRAIAHQLIKHWIGMNPNL      462
WFTREQVVDVLTGKGQQAFFVPPSRAIAHQLIKHWIGMNPNL      461
WFTREQVLDVLTGKGQQAFFVPPSRAIAHQLIKHWIRINPNL      462
WFTREQVLDVLTGKGQQAFFVPPSRAIAHQLIKHWIRINPNL      465
WFTREQVLDVLTGKGQQAFFVPPSRAIAHQLIKHWIRINPNL      462
WFTREQVLDVLTGKGQQAFFVPPSRAIAHQLIKHWIRINPNL      462
WFTREQVVDVLTGKGQQAFFVPPSRAIAHQLIKHWIRMNPNL      462
WFTREQVVDVLIKGNRSFFVPPSRAIAHQLIKHWIGMNPANL      464
WFTREQVVDVLIKGNRSFFVPPSRAIAHQLIKHWIGMNPANL      464
WFTREQVVDVLIKGNRSFFVPPSRAIAHQLIKHWIGMNPANL      465
WFTREQVVEVLIKGNRSFFVPPSQAIAHQLIKHWIGMNPANL      464
WFTREQIPALFILLN--SSLSEKRS-----      484
WFTREQVVDVAVIKGNHHAITVPPROAIAYOLIKHWIGMNPANL      458
WFTREQVLEVVMK--KKRSFFVPPROAIAHQLIKHWIRLNANL      505
.....490.....500.....510.....520..
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