CLUSTAL 2.1 MULTIPLE SEQUENCE ALIGNMENT

File: /media/morpheus/disk1/fst/pep_msa/SLC**D5Ac5** II pre Feb 1 14:32:58 2022 Page 1 of 3

Xenopus_tropicalis Chelonia_mydas Mus_musculus Rattus_norvegicus Homo_sapiens Pan_troglodytes Macaca_mulatta Callithrix_jacchus Bos_taurus Equus_caballus Canis_lupus_familiaris Felis_catus Heterocephalus_glaber Pavo_muticus Pavo_cristatus Gallus_gallus Anas_platyrhynchos		32 32 44 60
Xenopus_tropicalis Chelonia_mydas Mus_musculus Rattus_norvegicus Homo_sapiens Pan_troglodytes Macaca_mulatta Callithrix_jacchus Bos_taurus Equus_caballus Canis_lupus_familiaris Felis_catus Heterocephalus_glaber Pavo_muticus Pavo_cristatus Gallus_gallus Anas_platyrhynchos	QTVEILQLFSLMMDSEAHEKRKS- GAVGLGRRWPRRRVPGPAPVSGRERPWSRRGRGQHPYDERKSS	29 49 12 12 12 12 12 12 11 12 12 78 78 91 120
Xenopus_tropicalis Chelonia_mydas Mus_musculus Rattus_norvegicus Homo_sapiens Pan_troglodytes Macaca_mulatta Callithrix_jacchus Bos_taurus Equus_caballus Canis_lupus_familiaris Felis_catus Heterocephalus_glaber Pavo_muticus Pavo_cristatus Gallus_gallus Anas_platyrhynchos	:::	84 109 65 65 65 65 65 65 137 150 179

CLUSTAL 2.1 MULTIPLE SEQUENCE ALIGNMENT

File: /media/morpheus/disk1/fst/pep_msa/SLC**D5Ac5** II pre Feb 1 14:32:58 2022 Page 2 of 3

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Xenopus_tropicalis	RDAVROLOTDGIRNLYRGILPPLMOKTTTLALMFGLYEDFSSLLLRHTN-SPEVVTRSVA	143
Chelonia_mydas	RDAVCQLQKDGIRNLYRGLLPPLMQKTTTLALMFGLYQDFSSLLQRHTN-APELVTCSVA	168
Mus_musculus	RDAVLOLRKDGFRNLYRGILPPLMOKTTTLALMFGLYEDLSRLLHKHVSSAPEFATRSVA	125
Rattus_norvegicus	RDAILQLRKDGFRNLYRGILPPLMQKTTTLALMFGLYEDLSRLLRKHVSNAPEFATRSVA	125
Homo_sapiens	RDAILQLRRDGFRNLYRGILPPLMQKTTTLALMFGLYEDLSCLLHKHVS-APEFATSGVA	124
Pan troglodytes	RDAILOLRRDGFRNLYRGILPPLMOKTTTLALMFGLYEDLSCLLHKHVS-APEFATSGVA	124
Macaca mulatta	RDAILQLRRDGFRNLYRGILPPLMQKTTTLALMFGLYEDLSCLLHKHVS-APEFATCGVA	124
Callithrix_jacchus	RDAILOLRRDGFRNLYRGILPPLMOKTTTLALMFGLYEDLSCLLHKHVS-APEFATRSVA	124
Bos taurus	RDAILOLRRDGFRNLYRGILPPLMOKTTTLALMFGLYEDLSCLLRKHVS-TPEFATRSMA	124
Equus_caballus	RDAVLQLRRDGFRNLYRGILPPLMQKTTTLALMFGLYEDLSCLLHKHVS-TPEFATRSMA	123
Canis lupus familiaris	RDAVLQLRRDGFRNLYRGILPPLMQKTTTLALMFGLYEDLSCLLRKHIS-TPEFATRSVA	124
Felis_catus	RDAVLOLRRDGFRNLYRGILPPLLOKTTTLALMFGLYEDLSCLLRKHIS-PPEFATRSVA	124
Heterocephalus glaber	RDAVLQLRKDGFRNLYRGILPPLMQKTTTLALMFGLYEDLSCILHKHVS-APEFVTRGAA	124
Pavo muticus	KDAIYQLQKDGIRNLYRGILPPLMQKTTTLALMFGLYEDFSSLLHSHTT-APELLTRSVA	196
Pavo_cristatus	KDAIYQLQKDGIRNLYRGILPPLMQKTTTLALMFGLYEDFSSLLHSHTT-APELLTRSVA	196
Gallus_gallus	KDAIHQLQKDGIRNLYRGILPPLMQKTTTLALMFGLYEDFSSLLHSHTT-APELLTRSMA	209
Anas_platyrhynchos	KDAVHQLQKDGMRNLYRGILPPLMQKTTTLALMFGLYEDFSSLLHSHTT-APELLTRSMA	238
	190200210220230240	
Vananus tuanisalis	*:*****::*:	202
Xenopus_tropicalis	AILAGTTEALLTPFERVQTLLQDYKHHDRFTNTFQAFKVLRPYGIREYYRGLVPILLRNG	203
Chelonia_mydas	AVLAGTTEALLTPFERVQTLLQDYKHHDRFTNTYQAFKVLKDYGIKEYYRGLVPILIRNG	228
Mus_musculus	ALLAGITEAILTPFERVQTLLQDHKHHDKFTNTYQAFRALRCHGIAEYYRGMVPILFRNG	185
Rattus_norvegicus	ALLAGITEAILTPLERVQTLLQDHKHHDKFTNTYQAFRALRCHGVAEYYRGLVPILFRNG	185
Homo_sapiens	AVLAGITEAIFTPLERVQTLLQDHKHHDKFTNTYQAFKALKCHGIGEYYRGLVPILFRNG	184
Pan_troglodytes	AVLAGTTEAIFTPLERVQTLLQDHKHHDKFTNTYQAFKALKCHGIGEYYRGLVPILFRNG	184
Macaca_mulatta	AVLAGTTEAIFTPLERVQTLLQDHKHHDKFTNTYQAFKALKCHGIGEYYRGLVPILFRNG	184
Callithrix_jacchus	AVLAGTTEAIFTPLERVQTLLQDHKHHDKFTNTYQAFKALKCHGIGEYYRGLVPILFRNG	184
Bos_taurus	AVLAGTTEAIFTPLERVQTLLQDHRHHDKFTNTYQAFKALKCHGIREYYRGLVPVLFRNG	184
Equus_caballus	AVLAGTTEAIFTPLERVQTLLQDHKHHDKFTNTYHAFKALKCHGMREYYRGLVPILFRNG	183
Canis_lupus_familiaris		184
Felis_catus	AVLAGTTEAIFTPLERVQTLLQDHRHHDKFTNTYQAFKALRCHGIREYYRGLVPVLFRNG	184
Heterocephalus_glaber	AILAGTTEAIFTPLERVQTLLQDHKHHDKFTNTYQAFRVLKCHGIREYYRGLVPILFRNG	184
Pavo_muticus	AVLAGTTEAVLTPFERVQTLLQDYKHHDKFTNTYQAFKVLKVYGMREYYRGLVPILLRNG	256
Pavo_cristatus	AVLAGTTEAVLTPFERVQTLLQDYKHHDKFTNTYQAFKVLKVYGMREYYRGLVPILLRNG	256
Gallus_gallus	AVLAGTTEAVLTPFERVQTLLQDYKHHDKFTNTYQAFKVLRVYGMREYYRGLVPILLRNG	269 298
Anas_platyrhynchos	AVLAGTTEAVLTPFERVQTLLQDYKHHDKFTNTYQAFKVLKVYGMREYYRGLVPILLRNG 250260270280290300	298
	250260270200290300	
	.* :****.**: ** *.: * :*:******:**:* **:***:*	
Xenopus_tropicalis	PSNALFFGLRSPIKQCLPEAKTYSANLINDFICGGLLGAMLGFLFFPINVVKARMQSQIG	263
Chelonia_mydas	PSNVLFFGLRGPIKQCLPEATSYSAHLVNDFICGGLLGAMLGFLFFPMNVVKARMQSQIG	288
Mus_musculus	FGNVLFFGLRGPIKESLPTATTYSAHLVNDFICGGVLGAVLGFLSFPINVVKARIQSQIG	245
Rattus_norvegicus	F <mark>SN</mark> ILFFGLRGPIKEHLPTATTNSEHLVNDFICGGVLGAMLGFLSFPVNVVKARIQSQIG	245
Homo_sapiens	L <mark>SN</mark> VLFF <mark>GLRGPIKEHLPTATTHSAHLVNDFICGGLLG</mark> AML <mark>G</mark> FLFFPINVVKTRIQSQIG	244
Pan_troglodytes	L <mark>SN</mark> VLFF <mark>G</mark> LRGPIKEHLPTATTHSAHLVNDFICGGLLGAMLGFLFFPINVVKTRIQSQIG	244
Macaca_mulatta	L <mark>SN</mark> VLFF <mark>G</mark> LRGPIKEHLPTATTHSAHLVNDFICGGLLGAMLGFLFFPINVVKTRIQSQIG	244
Callithrix_jacchus	LSNVFFFGLRGPIKEHLPTATTHSAHLVNDFICGGLLGAMLGFLFFPINVVKTRIQSQIG	244
Bos_taurus	FSNVLFFGLRGPIKEHLPTATTHSAHLVNDFICGGLLGAMLGILFFPVNVVKTRMQSQIG	244
Equus_caballus	FSNVLFFGLRGPIKEHLPTATTHSAHLVNDFICGGLLGAVLGILFFPVNVVKTRMQSQIG	243
Canis_lupus_familiaris		244
Felis_catus_	FSNVLFFGLRGPIKEHLPTATTHSAHLVNDFICGGLLGAMLGILFFPVNVVKTRMQSQIG	244
Heterocephalus_glaber	FSNVLFFGLRGPIKEHLPTATTHSAHLVNDFICGGVLGAMLGFLFFPVNVVKTRIQSQIG	244
Pavo_muticus	PSNVLFFGLRGPIKQCLPEATSYSTHLVNDFICGGLLGAMLGFLFFPMNVVKARMQAQIG	316
Pavo_cristatus	PSNVLFFGLRGPIKQCLPEATSYSTHLVNDFICGGLLGAMLGFLFFPMNVVKARMQAQIG	316
Gallus_gallus	PSNVLFFGLRGPIKQCLPEATSYSTHLVNDFICGGLLGAMLGFLFFPMNVVKARMQAQIG	329
Anas_platyrhynchos	PSNVLFFGLRGPIKQCLPEATSYSTHLVNDFICGGLLGAMLGFLFFPVNVVKARMQAQIG	358
	310320330340350360	

CLUSTAL 2.1 MULTIPLE SEQUENCE ALIGNMENT

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Xenopus_tropicalis	GE	'IS	FGF	VL	II IW	TERI	GKLI	HLF	'RGAH	L <mark>N</mark> YH	RSII	SWG	IIN	ATYE	LLL	V L	316
Chelonia_mydas	GE	'QS	FPF	VFN	I <mark>K</mark> IW	LERI	RKLI	RLF	'RGAH	L <mark>N</mark> YH	RSII	SWG	IIN	ATYE	FLL	LL	341
Mus_musculus	GPE	LS	LPN	IVFI	TIW	IERI	RKLI	NLF	'RGAH	L <mark>N</mark> YH	R <mark>S</mark> LI	SWG	IIN	ATYE	FLL	CIV	298
Rattus_norvegicus	GP E	QS	LPN	IVF	T <mark>IW</mark>	IERI	RKLI	N <mark>LF</mark>	'RGAH	L <mark>N</mark> YH	R <mark>S</mark> LI	SWG	IIN	ATYE	FLL	C IV	298
Homo_sapiens	GE	QS	FPF	VFÇ	KIW	LERI	RKLI	N <mark>LF</mark>	'RGAH	L <mark>N</mark> YH	R <mark>S</mark> LI	SWG	IIN	ATYE	FLL	C VI	297
Pan_troglodytes	GE	QS	FPF	VFÇ	KIW	LERI	RKLI	N <mark>LF</mark>	'RGAH	L <mark>N</mark> YH	RS <mark>L</mark> I	SWG	IIN	ATYE	FLL	C VI	297
Macaca_mulatta	GE	QS	FPF	VFÇ	KIW	LERI	RKLI	N <mark>LF</mark>	'RGAH	L <mark>N</mark> YH	RS <mark>L</mark> I	SWG	IIN	ATYE	FLL	C VI	297
Callithrix_jacchus	GE	'QS	FPF	VFÇ	KIW	LERI	RKLI	N <mark>LF</mark>	'RGAH	L <mark>N</mark> YH	R <mark>S</mark> LI	SWG	IIN	ATYE	FLL	V I	297
Bos_taurus	GE	'QS	FPF	VFÇ	KIW	LERI	RKLI	' <mark>N</mark> LF	'RGAH	L <mark>N</mark> YH	R <mark>S</mark> LI	SWG	IIN	ATYE	FLL	TI	297
Equus_caballus	GE	'QS	FPF	VFÇ	KIW	LERI	RKLI	' <mark>N</mark> LF	'RGAH	L <mark>N</mark> YH	R <mark>S</mark> LI	SWG	IIN	ATYE	FLL	TI	296
Canis_lupus_familiaris	GE	QS	FPF	VFÇ	KIW	LERI	RKLI	' <mark>N</mark> LF	'RGAH	L <mark>N</mark> YH	RS <mark>L</mark> I	SWG	IIN	ATYE	FLL	CII	297
Felis_catus	GE	QS	SPE	VFÇ	KIW	LERI	RKLI	' <mark>N</mark> LF	'RGAH	L <mark>N</mark> YH	R <mark>S</mark> LI	SWG	IIN	ATYE	FLL	KMI	297
<pre>Heterocephalus_glaber</pre>	GE	'QS	FSF	VFI	TIW	LERI	RKLI	' <mark>N</mark> LF	'RGAH	L <mark>N</mark> YH	R <mark>S</mark> LI	SWG	IIN	ATYE	FLL	TI	297
Pavo_muticus	GE	'QS	FS1	'VF'\	KIW	LERI	RKLI	N <mark>LF</mark>	'RGAH	L <mark>N</mark> YH	RSVI	SWG	IIN	ATYE	FLL	LL	369
Pavo_cristatus	GE	'QS	FS1	'VF'\	KIW	LERI	RKLI	N <mark>LF</mark>	'RGAH	L <mark>N</mark> YH	RSVI	SWG	IIN	ATYE	FLL	LL	369
Gallus_gallus	GE	'QS	FS1	'VF'\	KIW	LERI	RKLI	N <mark>LF</mark>	'RGAH	L <mark>N</mark> YH	RSVI	SWG	IIN	ATYE	FLL	LL	382
Anas_platyrhynchos	GE	QS	FS1	'VF'	KIW	LERI	RKLI	N <mark>LF</mark>	'RGAH	L <mark>N</mark> YH	RSVI	SWG	IIN	ATYE	FLL	L L	411
			3	70.		3	380		39	0		400			410		