

sp Q62651 ECH1_RAT	Delta(3,5)-Delta(2,4)-dieno	Ech1	15	49.5	36.171	327	0	126.01	50.51387432	49.30408983	56.55001174	47.52025781	56.76611379	51.71772861
tr D4A3V2 D4A3V2_RAT	NADH dehydrogenase [ub	Ndufa6	8	52.3	15.224	130	0	323.31	170.0101938	173.8747795	150.6414742	180.9849019	158.7143007	153.6881347
tr D3ZTN2 D3ZTN2_RAT	COX assembly mitochondr	Cmc1	3	28.3	12.572	106	0	7.7271	1.804210291	1.232281416	1.109732235	1.036946793	1.376225022	1.704518963
sp P06685 AT1A1_RAT	Sodium/potassium-transpo	Atp1a1	31	31.1	113.05	1023	0	323.31	79.40331496	98.27254264	109.5233404	86.58941717	84.75009294	114.6234901
tr F1LRV9 F1LRV9_RAT	Myosin heavy chain 4	Myh1	4	2.4	223.28	1943	0	323.31	899.0947833	850.0994485	957.5631234	928.5457778	738.3477712	1048.903485
sp P35280 RAB8A_RAT	Ras-related protein Rab-8	Rab8a	2	7.2	23.668	207	0	15.551	3.793909053	3.36569885	2.674325248	3.311218014	3.885291566	2.679357101
sp Q9Z270 VAPA_RAT	Vesicle-associated membr	Vapa	11	43.4	27.841	249	0	82.418	41.55905083	45.07259556	49.50641597	41.78366413	48.43293889	46.18568727
sp Q64428 ECHA_RAT	Trifunctional enzyme subu	Hadha	62	76	82.664	763	0	323.31	855.7455705	1033.515204	984.1824269	958.8861171	815.6825681	1091.574879
tr D4A2V6 D4A2V6_RAT	Junctional sarcoplasmic re	Jsrp1	9	41.1	36.751	338	0	106.99	20.61646414	29.34602724	31.33531102	25.33418332	24.22779462	31.43459302
sp Q5BK63 NDUA9_RAT	NADH dehydrogenase [ub	Ndufa9	32	72.4	42.559	377	0	323.31	359.632896	377.8285582	329.928429	387.7949273	326.8015356	354.3859209
tr M0R735 M0R735_RAT	Heterogeneous nuclear rib	Syncrip	3	5.9	62.672	562	0.00055127	3.9222	0.782744292	0.845854627	0.745431973	0.788012671	0.924026785	0.655991392
tr A0A0H2UHK5 A0A0H2U	Leucyl-cystinyl aminopepti	Lnpep	7	9.6	103.19	900	0	27.327	5.093030773	4.683222195	4.585655271	4.474005546	5.346293645	4.523878873
sp P67779 PHB_RAT	Prohibitin	Phb	21	80.5	29.82	272	0	323.31	138.2057193	149.9606424	132.794649	141.0586877	130.3743873	149.1873187
sp Q5HZE0 MCATL_RAT	Mitochondrial basic amino	Slc25a29	2	6.9	32.634	306	0.0010672	3.4163	0.741953887	0.832478505	0.631545262	0.705771239	0.764988908	0.738022066
tr A0A0G2K4U8 A0A0G2K	Syntaxin-12 (Syntaxin-13)	Stx12	2	8.8	31.106	272	0	13.417	0.815005338	0.846496286	0.72750282	0.748355219	0.7816531	0.861159515
tr Q2I6B2 Q2I6B2_RAT	V-V-type proton ATPase 116	Atp6v0a1	6	7.9	95.584	832	0	20.869	2.408088907	3.219252524	2.698474719	2.511320053	2.351862267	3.479547323
sp P68136 ACTS_RAT	Actin, alpha skeletal muscl	Acta1	2	5.8	42.051	377	0	323.31	841.0449694	578.5789778	1030.3317	869.2985011	839.4654449	745.4590801
tr F8WG17 F8WG17_RAT	Troponin I, fast skeletal m	Tnni2	10	49.5	21.311	182	0	281.27	48.45980398	36.19549019	60.34166167	52.4947595	57.07351152	35.18053176
tr D3ZYX8 D3ZYX8_RAT	Cytochrome c oxidase sub	Cox7a2l	5	35.8	13.273	120	0	58.235	20.50357557	24.30554906	20.99494653	23.04193485	20.32492207	22.47156847
sp Q561S0 NDUAA_RAT	NADH dehydrogenase [ub	Ndufa10	3	13	40.493	355	0	323.31	369.8631096	370.7357584	360.3393825	401.3226574	359.649949	340.3246808
sp P20788 UCRI_RAT	Cytochrome b-c1 complex	Uqcrcf1	19	62.4	29.445	274	0	323.31	599.7644553	686.5257637	537.4172096	659.3648541	576.7752109	587.859401
tr A0A0A6YYM0 A0A0A6Y	Long-chain-fatty-acid-CoA	Acsf6	31	47.6	80.995	722	0	323.31	104.3391468	79.55584341	68.8808781	81.88308107	96.98128675	73.86316571