<html><head><style type="text/css"></style></head><body>CLUSTAL 2.1 multiple sequence alignment

humanATP7B mouseATP7B zebrafishATP7B drosophilaATP7B celegansATP7B	MPEQERQITAREGASRKILSKLSLPTRAWEPAMKKSFAFDNVGYEGGL MDPRKNLASVGTMPEQERQVTAKE-ASRKILSKLALPGRPWEQSMKQSFAFDNVGYEGGLMNKLSPFSN-LAKFISKSPDNGPVGGE	59
humanATP7B mouseATP7B zebrafishATP7B drosophilaATP7B celegansATP7B	DGLGPSSQVATSTVRILGMTCQSCVKSIEDRISNLKGIISMKVSLEQGSATVKYVPSVVC DSTS-SSPAATDVVNILGMTCHSCVKSIEDRISSLKGIVNIKVSLEQGSATVRYVPSVMNQLCMKDCKPLCHCDPELCVCSLPQNSKDGPTEYKENGSQQEGWIP	108 118 71 28
humanATP7B mouseATP7B zebrafishATP7B drosophilaATP7B celegansATP7B	LQQVCHQIGDMGFEASIAEGKAASWPSRSLPAQEAVVKLRVEGMTCQSCVSSIEGKVRKL LQQICLQIEDMGFEASAAEGKAASWPSRSSPAQEAVVKLRVEGMTCQSCVSSIEGKIRKLTKHAFDNFGYEPDGLKHNLVHQLPSEEGMVKIQVEGMTCQSCVRSIEEQIGRLMPSDERVEATMSTVRLPIVGMTCQSCVRNITEHIGQK -KNIQLLVDFGAPKTDGNVQETMLEIKGMTCNSCVKNIQDVIGAK : : ****:** .* :	178 124 37
humanATP7B mouseATP7B zebrafishATP7B drosophilaATP7B celegansATP7B	QGVVRVKVSLSNQEAVITYQPYLIQPEDLRDHVNDMGFEAAIKSKVAPLSLGPIDIERLQ QGVVRIKVSLSNQEAVITYQPYLIQPEDLRDHICDMGFEAAIKNRTAPLRLGPIDVNKLE EGVIGVQVSLSDKEAILRFNPAKVTPEDMRKRIEDMGFDALILALQGQIQPLSTDWSE SGILGVRVILEENAGYFDYDPRQTDPARIASDIDDMGFECSYPGDAAD PGIHSIQVNLKEENAKCSFDTTKWTAEKVAEAVDDMGFDCKVLKKEPP *: ::* *.:: :: :: ****:	238
humanATP7B mouseATP7B zebrafishATP7B drosophilaATP7B celegansATP7B	STNPKRPLSSANQNFNNSETLGHQGSHVVTLQLRIDGMHCKSCVLNIEENIGQLLGVQSI STNLKKETVSPVQISNHFETLGHQGSYLATLPLRIDGMHCKSCVLNIEGNIGQLPGVQNI VTLGVEGMHCGSCVKNITETLSGMLGVNSV	298
humanATP7B mouseATP7B zebrafishATP7B drosophilaATP7B celegansATP7B	QVSLENKTAQVKYDPSCTSPVALQRAIEALPPGNFKVSLPDGAEGSGTDHRSSSSHSPGS HVSLENKTAQIQYDPSCVTPMFLQTAIEALPPGHFKVSLPDGVEENEPQSGS FVSLEKGSVDLRFDPSLLTLETVKGFLEEIPPGNFRVSIPGWSPPPP	350 255
humanATP7B mouseATP7B zebrafishATP7B drosophilaATP7B celegansATP7B	PPRNQVQGTCSTTLIAIAGMTCASCVHSIEGMISQLEGVQQISVSLAEGTATVLYNPSVI SQRHQEQGPGRTAVLTISGITCASSVQPIEDMLSQRKGVQQTSISLAEGTGAVLYDPSIV SRLNSASTPTQSVTIGIEGMTCNSCVQAIEGMMSQRAGVCSIKVYLQEKKGIVTFDSTVTETPASAWTNIRVVGMTCQSCVRNIEGNIGTKPGIHSIEVQLAAKNARVQYDPAQY -TQMAEKPKIRRAIVSIEGMTCHACVNNIQDTVGSKDGIVKIVVSLEQKQGTVDYNSEKW .:: *:** :.*	410 315 142
humanATP7B mouseATP7B zebrafishATP7B drosophilaATP7B celegansATP7B	SPEELRAAIEDMGFEASVVSESCSTNPLGNHSAGNSMVQTTDGTPTSVQEVAPHTGRL SLDELRTAVEDMGFEVSVNSETFTINPVRNFKSGNSVPQTMGDIAGSVQKMAPDTRGL CPEELRAEIEDMGFEAWLNQDSEICEVSSVSQMPSGLKHLPSQRHP DPAQIAELIDDMGFEASVQEPRSPSQSPSPAPASSPKKRATPTP NGESVAESIDDMGFDCKLITDQEIAAVEPQKASTTKLSISPLKTVDLSD .: ::***: : :	468 361 186

humanATP7B	PANHAPDILAKSPOSTRAVAPOKCFLOIKGMTCASCVSNIERNLOKEAGVLSVLVALMAG	526
mouseATP7B	PTHQGPGHSSETPSSPGATASQKCFVQIKGMTCASCVSNIERSLQRHAGILSVLVALMSG	528
zebrafishATP7B	SKPSPSPITKENADGTGERELRKCFVHVTGMTCASCVSNIERNLVKHEGIKSVLVALMAG	421
drosophilaATP7B	PPPSYAQNGSAVAIPVEQELLTKCFLHIRGMTCASCVAAIEKHCKKIYGLDSILVALLAA	246
celegansATP7B	GKVELQLNGVKYSKEGSSDHLEKCTFAVEGMTCASCVQYIERNISKIEGVHSIVVALIAA	
	** . : ****** ** : * * * * * * * * * * *	
humanATP7B	KAEIKYDPEVIQPLEIAQFIQ-DLGFEAAVMEDYAGSDGNIELTITGMTCASCVHNIE	583
mouseATP7B	KAEVKYDPEIIQSPRIAQLIQ-DLGFEASVMEDNTVSEGDIELIITGMTCASCVHNIE	585
zebrafishATP7B	KAEVKYDPGLLDPAQIVQLIS-HLGFGASVMEEHSVQDGVLDLSVTGMTCASCVHNIE	478
drosophilaATP7B	KAEVKFNANVVTAENIAKSIT-ELGFPTELIDEPDNGEAEVELEIMGMTCASCVNKIE	303
celegansATP7B	KAEVIYDGRVTSSDAIREHMTGELGYKATLLDSMGANPNYSKIRLIIGNLSTESDANRIE	348
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humanATP7B	SKLTRTNGITYASVALATSKALVKFDPEIIGPRDIIKIIEEIGFHASLAQRNPNAHHL	
mouseATP7B	SKLTRTNGITYASVALATSKAHVKFDPEIVGPRDIIKIIEEIGFHASLAQRNPNAHHL	
zebrafishATP7B	SKLLRTKGIQEASVALATNKAHVKFDSDLVGSRDIVRIIEGLGFGVSLIKNEGLNNTL	
drosophilaATP7B	SHVLKIRGVTTASVTLLTKRGKFRYITEETGPRSICEAIEALGFEAKLMTGRDKMAHNYL	
celegansATP7B	SHVLSKSGIDSCNVSIATSMALVEFSPQVIGPRDIINVVESLGFTADLATRDDQMKRL	406
	*:: *:*:: *: *.** : *	
humanATP7B	DHKMEIKQWKKSFLCSLVFGIPVMALMIYMLIPSNEPHQSMVLDHNIIPGLSILNLI	698
mouseATP7B	DHKTEIKQWKKSFLCSLVFGIPVMGLMVYMLIPSNEFHQSMVLDHNIIPGLSVLNLI	
zebrafishATP7B	DHATEIRQWARSFLCSLVFGIPVMGLMVIMLIPSSIPQEIMVLDHNIIPGLSVLNLI DHQEEIRQWKHSFLFSLVFGIPVMGLMIYMMVMDSQHKEHGGSMPADQNILPGLSIINLA	
	EHKEEIRKWRNAFLVSLIFGGPCMVAMIYFMLEMSDKGHANMCCLVPGLSMENLV	
drosophilaATP7B celegansATP7B		
CelegansATP/B	DHSDDVKKWRNTFFIALIFGVPVMIIMIIFHWILRTPMHPDKQTPIFTPALSLDNFL	463
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humanATP7B	FFILCTFVQLLGGWYFYVQAYKSLRHRSANMDVLIVLATSIAYVYSLVILVVAVAEKAER	758
mouseATP7B	FFILCTFVQFLGGWYFYVQAYKSLRHRSANMDVLIVLATTIAYAYSLVILVVAVAEKAEK	760
zebrafishATP7B	FFLLCTPVQFLGGRYFYIQAYRSLRHGVANMDVLIVLATTIAYVYSFTVLIVAMIEGAKQ	
drosophilaATP7B	MFLLSTPVOFFGGFHFYVOSYRAIKHGTTNMDVLISMVTTISYVYSVAVVIAAVLLEONS	478
celegansATP7B	LLCLCTPVQIFGGRYFYVASWKAIKHGNANMDVLIMLSTTIAYTYSIVVLLLAIIFKWPS	523
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humanATP7B	SPVTFFDTPPMLFVFIALGRWLEHLAKSKTSEALAKLMSLQATEATVVTLGEDNLIIREE	818
mouseATP7B	SPVTFFDTPPMLFVFIALGRWLEHVAKSKTSEALAKLMSLQATEATVVTLGEDNLILREE	820
zebrafishATP7B	SPLTFFDTPPMLFVFIALGRWLEHVAKSKTSEALAKLMSLQATDATIVSLGPDNTIIREE	716
drosophilaATP7B	SPLTFFDTPPMLLIFISLGRWLEHIAKGKTSEALSKLLSLKAADALLVEISPDFDIISEK	
celegansATP7B	SPMTFFDVPPMLIVFIALGRMLEHKAKGKTSEALSKLMSLQAKEATLVTMDSEGRLTSEK	583
	:*.***::**:** ** ** ** ** ** ** ** ** ** **	
humanATP7B	OVERNIT VOD CD TVVVVD COVEDVD CVVV ECNIMA DECI TIICEA MDVIIVVD COUVTA CO IN	878
mouseATP7B	QVPMELVQRGDIVKVVPGGKFPVDGKVLEGNTMADESLITGEAMPVTKKPGSTVIAGSIN	880
zebrafishATP7B	QVPMELVQRGDVIKVVPGGKFPVDGKVLEGNTMADESLITGEAMPVTKKPGSIVIAGSIN	776
	QVSVDLVQRGDVVKVAPGGKFPVDGKVIEGTSMADESLITGEPMPVIKKAGSCVIAGSIN	770 598
drosophilaATP7B	VISVDYVQRGDILKVIPGAKVPVDGKVLYGHSSCDESLITGESMPVAKRKGSVVIGGSIN	0,0
celegansATP7B	GINIELVQRNDLIKVVPGAKVPVDGVVVDGKSSVDESFITGESMPVVKKPGSTVIGGSVN : :: ***.*: ** **.*.** *: * * **.**** *: ** **.********	043
humanATP7B	AHGSVLIKATHVGNDTTLAQIVKLVEEAQMSKAPIQQLADRFSGYFVPFIIIMSTLTLVV	938
mouseATP7B	AHGSVLLKATHVGNDTTLAQIVKLVEEAQMSKAPIQQLADRFSGYFVPFIIIISTLTLVV	940
zebrafishATP7B	AHGALLVEATHVGSETTLSQIVKLVEEAQTSKAPIQQLADKLSGYFVPFIVVISILTVTA	836
drosophilaATP7B	QNGVLLVEATHTGENTTLAQIVRLVEEAQTSKAPIQQLADRIAGYFVPFVVVVSSITLIA	658
celegansATP7B	QKGVLIVKATHVGNDSTLSQIVRLVEEAQTNRAPIQQLADKIAGYFVPFVIVLSLFTLGV	703
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humanATP7B	WIVIGFIDFGVVQRYFPNPNKHISQTEVIIRFAFQTSITVLCIACPCSLGLATPTAVMVG	
mouseATP7B	WIVIGFVDFGVVQKYFPSPSKHISQTEVIIRFAFQTSITVLCIACPCSLGLATPTAVMVG	
zebrafishATP7B	WLIIGFLDFDVVSKNFPGYNQNISRTEVIVRFAFQASITVLSIACPCSLGLATPTAVMVG	
drosophilaATP7B	WIIIGFSNPNLVPVAMEH-KMHMDQNTIIVSYAFKCALSVLAIACPCALGLATPTAVMVA	
celegansATP7B	WTYTEYNSARNANT.PPGT.RFEEAT.KTAFEAATTVT.ATACPCST.GT.ATPTAVMVG	171

WIYIEYN--SARNANLPP----GLRFEEALKIAFEAAITVLAIACPCSLGLATPTAVMVG 757

celegansATP7B

humanATP7B	TGVAAQNGILIKGGKPLEMAHKIKTVMFDKTGTITHGVPRVMRVLLLGDVATLPLRKVLA	1058
mouseATP7B	TGVAAQNGULIKGGKPLEMAHKIKTVMFDKTGTITHGVPRVMRFLLLADVATLPLRKVLA	
zebrafishATP7B	TGVGAONGILIKGGEPLEMAHKVGAVMFDKTGTITNGVPKVMKFLLLADVATLFLKKVLA	
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drosophilaATP7B	TGTGAINGVLVKGATALENAHKVKTVVFDKTGTITHGTPMTSKVTLFVTAQVCSLARALT	777
celegansATP7B	TGVGAANGILIKGGEPLESVHKVTTIVFDKTGTITEGRPRVVQIASFVNPSTMSLKLITF	817
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humanATP7B	VVGTAEASSEHPLGVAVTKYCKEELGTETLGYCTDFQAVPGCGIGCKVSNVEGI	1112
mouseATP7B	VVGTAEASSEHPLGVAVTKYCKEELGTETLGYSTDFQAVPGCGISCKVSNVEGI	
zebrafishATP7B	VVGTAEASSEHFLGVAVIKICKEELGIETLGISIDFQAVFGCGISCKVSNVEGI VVGTAEASSEHPLGMAVAKHCKEELGAETLGFCHDFQAVPGCGISCKVSSVEDL	
drosophilaATP7B	IVGAAEQNSEHPIASAIVHFAKDMLNVGATPQAGSFGKSSHFQAVPGCGIRVTVSNYEQT	
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celegansATP7B	LSGATEALSEHPIGNAVAAFAKQLLNEPTWPNTSRFHVSAGHGVTCRIDSIRQS	8/1
	: *::* ****: . *:	
humanATP7B	LAHSERPLSAPASHLNEAGSLP	1134
mouseATP7B	LARSDLTAHPVGVGNPP	1131
zebrafishATP7B	LQNSPKTQETKASAVKQTPLLVTLQGVTTD	
drosophilaATP7B	LRQACNADRIINYENLHRTHPQGSVPVDNGASIEHLLPQRSVRKSMELNNQQLLSDLVLE	897
celegansATP7B	FSSLALSGSTCEIPRLPDGQTITIPGTEVN	
ccicquibilii 1B	:	J 0 1
humanATP7B	AEKDAVPQTFSVLIGNREWLRRNGLTISSDVSDAMTDHEMKGQTAILVAIDGV	1187
mouseATP7B	TGEGAGPQTFSVLIGNREWMRRNGLTISSDISDAMTDHEMKGQTAILVAIDGV	1184
zebrafishATP7B	ESSLVADTDSGSDSPSYSVLIGNRQWMLRNGLEVTADVDNAMSSHETKGQTAILVAIDGV	1100
drosophilaATP7B	PEEELLTDOKIIDSPEILVLIGNREWMERNAIEVPLEISDCMTHEERKGHTAVLCALNGO	957
celegansATP7B	LLQVSSKEVSQPNPDTANIVIGTERMMERHGIPVSEVVKMTLSEEQRKGHISVICAINAE	961
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humanATP7B	LCGMIAIADAVKQEAALAVHTLQSMGVDVVLITGDNRKTARAIATQVGINKVFAEVLPSH	
mouseATP7B	LCGMIAIADAVKPEAALAIYTLKSMGVDVALITGDNRKTARAIATQVGINKVFAEVLPSH	
zebrafishATP7B	LCAMIAVADTVKAESALAVHTLSSMGIEVYMITGDNRRTARAIATQVGIRKVFAEVLPSH	
drosophilaATP7B	LVCMFAVSDMVKPEAHLAVYTLKRMGIDVVLLTGDNKNTAASIAREVGIRTVYAEVLPSH	1017
celegansATP7B	VVAVISIADQVKKEASLAIYTLREMGLRVVLLTGDNSKTAESTAKQVGIDEVFAEVLPNQ	1021
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humanATP7B	KVAKVQELQNKGKKVAMVGDGVNDSPALAQADMGVAIGTGTDVAIEAADVVLIRNDLLDV	1307
mouseATP7B	KVAKVQELQNKGKKVAMVGDGVNDSFALAQADMGVAIGIGIDVAIEAADVVLIRNDLLDV KVAKVQELQNEGKKVAMVGDGVNDSPALAQADVGIAIGTGTDVAIEAADVVLIRNDLLDV	
zebrafishATP7B	KVAKVQELQNEGKKVAMVGDGVNDSPALAQADVGIAIGIGIDVAIEAADVVLIKNDLLDV KVAKVQELQERGLKVAMVGDGVNDSPALAHADLGIAIGTGTDVAIEAADIVLIRNDLLDV	1220
drosophilaATP7B	KVAKIQRIQANGIRVAMVGDGVNDSPALAQADVGITIAAGTDVAAEASDIVLMRNDLLDV	
celegansATP7B	KQQKIKQLKGYKNKVAMVGDGVNDSPALAEANVGIAIAAGSDVAIESAGIVLVRNDLVDV * *:::: :****************************	1081
humanATP7B	VASIHLSKRTVRRIRINLVLALIYNLVGIPIAAGVFMPIGIVLQPWMGSAAMAASSVSVV	1367
mouseATP7B	VASIHLSKRTVRRIRVNLVLALIYNMVGIPIAAGVFMPIGIVLQPWMGSAAMAASSVSVV	1364
zebrafishATP7B	VASIELSKKTVQRIRINFVFALIYNLLGIPFAAGVFMPLGLVLQPWMGSAAMAASSVSVV	1280
drosophilaATP7B	VACLDLSRCTVRRIRYNFFFASMYNLLGIPLASGLFAPYGFTLLPWMASVAMAASSVSVV	
celegansATP7B	VGAIKLSKMTTRRIRLNFLFAIIYNAIGIPIAAGVFRPFGFMLQPWMAAAAMALSSVSVV	
ocicganonii 75	***: *.:** *: ** :** :** :** * * * : * * *** :*** * ***	
humanATP7B	LSSLQLKCYKKPDLERYEAQAHGHMKPLTASQVSVHIGMDDRWRD-SPRATP	1418
mouseATP7B	LSSLQLKCYRKPDLERYEAQAHGRMKPLSASQVSVHIGMDDRRRD-SPRATA	1415
zebrafishATP7B	LSSLLLRLFKKTSVEEYESRAQSHKLSLSPSQVSTHVGLESRRCSP	1326
drosophilaATP7B	CSSLLLKMYRKPTAKTLRTAEYEAQLAAERASGSEDELDKLSLHRGLDDLPEK-GRMPFK	1196
celegansATP7B	SSSLLLKNFRKPTIANLYTTSFKRHQKFLESGSFQVQVHRGLDDSAVFRGAASSK	1196
<u> </u>	*** *: ::*	
l	VIDOVIGUES 0.00 - 0.00	4.65
humanATP7B	WDQVSYVSQVSLSSLTSDKPSRHSAAADDDGDKWSLLLNGRDEEQYI 14	
mouseATP7B	WDQVSYVSQVSLSSLTSDRLSRHGGAAEDGGDKWSLLLSDRDEEQCI 14	162 264

LSDRKRRSRSAS---GSSSFISGSFHPSTTQPYN--TSGRSIV----- 1364

zebrafishATP7B

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drosophilaATP7B RSSTSLISRIFMHDYGNITSPDAKYGEGLLDPEEQYDGRTKIVRSRFHANDSTELQKL 1254 celegansATP7B LSILSSKVGSLLGSTTSIVSSGSSKKQRLLDNVG-SDLEDLIV----- 1238

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