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

humanDHCR7	-MAAKSQPNIPKAKSLDGVTNDRTASQGQWGRAWEVDWFSLASVIFLLLFAPFIVYYF	57
mouseDHCR7	-MASKSQHNAPKVKSPNGKAGSQGQWGRAWEVDWFSLASIIFLLLFAPFIVYYF	
zebrafishDHCR7	MMASDRVRKRHKGSANGAQTVEKEPSKEPAQWGRAWEVDWFSLSGVILLLCFAPFLVSFF	60
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humanDHCR7	IMACDQYSCALTGPVVDIVTGHARLSDIWAKTPPITRKAAQLYTLWVTFQVLLYTSLPDF	117
mouseDHCR7	IMACDQYSCSLTAPALDIATGHASLADIWAKTPPVTAKAAQLYALWVSFQVLLYSWLPDF	
zebrafishDHCR7	IMACDQYQCSISHPLLDLYNGDATLFTIWNRAPSFTWAAAKIYAIWVTFQVVLYMCVPDF	
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humanDHCR7	CHKFLPGYVGGIOEGAVTPAGVVNKYOINGLOAWLLTHLLWFANAHLLSWFSPTIIFDNW	177
mouseDHCR7	CHRFLPGYVGGVOEGAITPAGVVNKYEVNGLOAWLITHILWFVNAYLLSWFSPTIIFDNW	
zebrafishDHCR7	LHKILPGYVGGVQDGARTPAGLINKYEVNGLQCWLITHVLWVLNAQHFHWFSPTIIIDNW	180
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humanDHCR7	IPLLWCANILGYAVSTFAMVKGYFFPTSARDCKFTGNFFYNYMMGIEFNPRIGKWFDFKL	237
mouseDHCR7	IPLLWCANILGYAVSTFAMIKGYLFPTSAEDCKFTGNFFYNYMMGIEFNPRIGKWFDFKL	233
zebrafishDHCR7	IPLLWCTNILGYAVSTFAFIKAYLFPTNPEDCKFTGNMFYNYMMGIEFNPRIGKWFDFKL	240
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humanDHCR7	FFNGRPGIVAWTLINLSFAAKQRELHSHVTNAMVLVNVLQAIYVIDFFWNETWYLKTIDI	297
mouseDHCR7	FFNGRPGIVAWTLINLSFAAKQQELYGHVTNSMILVNVLQAIYVLDFFWNETWYLKTIDI	
zebrafishDHCR7	FFNGRPGIVAWTLINLSYAAKQQELYGYVTNSMILVNVLQAVYVVDFFWNEAWYLKTIDI	300
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humanDHCR7	CHDHFGWYLGWGDCVWLPYLYTLQGLYLVYHPVQLSTPHAVGVLLLGLVGYYIFRVANHQ	357
mouseDHCR7	CHDHFGWYLGWGDCVWLPYLYTLQGLYLVYHPVQLSTPNALGILLLGLVGYYIFRMTNHQ	353
zebrafishDHCR7	CHDHFGWYLGWGDCVWLPFLYTLQGLYLVYNPIQLSTPHAAGVLILGLVGYYIFRVTNHQ	360
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humanDHCR7	KDLFRRTDGRCLIWGRKPKVIECSYTSADGQRHHSKLLVSGFWGVARHFNYVGDLMGSLA	417
mouseDHCR7	KDLFRRTDGRCLIWGKKPKAIECSYTSADGLKHHSKLLVSGFWGVARHFNYTGDLMGSLA	413
zebrafishDHCR7	KDLFRRTEGNCSIWGKKPTFIECSYQSADGAIHKSKLMTSGFWGVARHMNYTGDLMGSLA	420
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humanDHCR7	YCLACGGGHLLPYFYIIYMAILLTHRCLRDEHRCASKYGRDWERYTAAVPYRLLPGIF 47	'5
mouseDHCR7	YCLACGGGHLLPYFYIIYMTILLTHRCLRDEHRCANKYGRDWERYTAAVPYRLLPGIF 47	1
zebrafishDHCR7	YCLACGGNHLLPYFYIIYMTILLVHRCIRDEHRCSNKYGKDWERYTAAVSYRLLPNIF 47	8
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