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

humanDHCR7 mouseDHCR7 zebrafishDHCR7		
humanDHCR7 mouseDHCR7 zebrafishDHCR7	**:. : * :: : *: .*********************	113
humanDHCR7 mouseDHCR7 zebrafishDHCR7	**************************************	173
humanDHCR7 mouseDHCR7 zebrafishDHCR7	*::********** IPLLWCANILGYAVSTFAMVKGYFFPTSARDCKFTGNFFYNYMMGIEFNPRIGKWFDFKL IPLLWCANILGYAVSTFAMIKGYLFPTSAEDCKFTGNFFYNYMMGIEFNPRIGKWFDFKL IPLLWCTNILGYAVSTFAFIKAYLFPTNPEDCKFTGNMFYNYMMGIEFNPRIGKWFDFKL	233
humanDHCR7 mouseDHCR7 zebrafishDHCR7	**************************************	293
humanDHCR7 mouseDHCR7	**************************************	357
zebrafishDHCR7	CHDHFGWYLGWGDCVWLPFLYTLQGLYLVYNPIQLSTPHAAGVLILGLVGYYIFRVTNHQ ************************************	417
mouseDHCR7 zebrafishDHCR7 humanDHCR7	KDLFRRTDGRCLIWGKKPKAIECSYTSADGLKHHSKLLVSGFWGVARHFNYTGDLMGSLA KDLFRRTEGNCSIWGKKPTFIECSYQSADGAIHKSKLMTSGFWGVARHMNYTGDLMGSLA *****:.* ***:** ***********************	420
mouseDHCR7 zebrafishDHCR7	YCLACGGGHLLPYFYIIYMTILLTHRCLRDEHRCANKYGRDWERYTAAVPYRLLPGIF 47 YCLACGGNHLLPYFYIIYMTILLVHRCIRDEHRCSNKYGKDWERYTAAVSYRLLPNIF 47 ************************************	

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