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

humanDPM1	MASLEVSRSPRRSRRELEVRSPRQNKYSVLLPTYNERENLP-LIVWLLVKSFSESGINYE	59
mouseDPM1	MASTGASRSLAASPRPPQGRSSRQDKYSVLLPTYNERENLP-LIVWLLVKSFSESAINYE	59
zebrafishDPM1	MASRRSNAKSRDKPDKYSVLLPTYNERENLP-LIVWLLVKYFGESGYNYE	49
drosophilaDPM1	MPTNGHKYSILMPTYNEKDNLP-IIIWLIVKYMKASGLEYE	40
celegansDPM1	MTSTPKYSIILPTYNEKENLP-ICIWLIENYLKEVSHE	37
yeastDPM1	MSIEYSVIVPAYHEKLNIKPLTTRLFAGMSPEMAKKTE	38
	:**::*:*: *:	
humanDPM1	TTTTDDCCDDCMDDWAEOTERTYCCDDTTTDDDERRTCTCMAYTUCMRUAMCNYTTMDA	110
mouseDPM1	IIIIDDGSPDGTRDVAEQLEKIYGSDRILLRPREKKLGLGTAYIHGMKHATGNYIIIMDA	
zebrafishDPM1	IIIIDDGSPDGTREVAEQLAEIYGPDRILLRPREKKLGLGTAYIHGIKHATGNYVIIMDA	
	IIVIDDGSPDGTLQIAEQLQKIYGADKILLRPRAEKLGLGTAYIHGIKHATGNFVIIMDA	
drosophilaDPM1	VIVIDDGSPDGTLDVAKDLQKIYGEDKIVLRPRGSKLGLGTAYIHGIKHATGDFIVIIDA	
celegansDPM1	VIIVDDASPDGTQGIAKLLQKEYGDDKILIKPRVGKLGLGTAYSHGLSFARGEFIILMDA	
yeastDPM1	LIFVDDNSQDGSVEEVDALAHQGYNVRIIVRTNERGLSSAVLKGFYEAKGQYLVCMDA **.:** * **:	96
	:*.:** * **: *	
humanDPM1	DLSHHPKFIPEFIRKQKEGNFDIVSGTRYKGNGGVYG-WDLKRKIISRGANFLTQILLRP	178
mouseDPM1	DLSHHPKFIPEFIRKQKEGNFDIVSGTRYKGNGGVYG-WDLKRKIISRGANFITQILLRP	178
zebrafishDPM1	DLSHHPKFIPQFIEKQKEGGYDLVSGTRYRGDGGVYG-WDLRRKLISRGANFVTQVLLRP	168
drosophilaDPM1	DLSHHPKFIPEFIKLQQEGNYDIVSGTRYAGNGGVFG-WDFKRKLISRGANFLSQVLLRP	159
celegansDPM1	DLSHHPKFIPEMIALQHKYKLDIVTGTRYKDGGGVSG-WDLKRKTISKGANFLAQFLLNP	156
yeastDPM1	DLQHPPETVPKLFESLHDHAFTLGTRYAPGVGIDKDWPMYRRVISSTARMMARPLT-I	153
-	**.* *: :*::: :	
humanDPM1	GASDLTGSFRLYRKEVLEKLIEKCVSKGYVFQMEMIVRARQLNYTIGEVPISFVDRV	225
mouseDPM1	GASDLTGSFRLYRKEVLEKLIEKCVSKGYVFQMEMIVRARQMNYTIGEVPISFVDRV	
zebrafishDPM1	GASDLTGSFRLYKKEVLGKLIEKCVSKGIVFQMEMIVRARQLGYTIGEVPISFVDRV	
drosophilaDPM1	NASDLTGSFRLYKKDVLEKCIASCVSKGYVFQMEMLVRARQHGYTIAEVPITFVDRI	
celegansDPM1	GVSDLTGSFRLYKRDILSKLIAESVSKGYVFQMEMMFRAKKSGYRIGEVPISFVDRF	
yeastDPM1	ASDPMSGFFGLQKKYLENCNPRDINSQGFKIALELLAKLPLPRDPRVAIGEVPFTFGVRT	
yeascbrHi	ASDEMSGIFGUÇAKTILEMCMEKDINSQGFKTALIELLLAKTIFLEKDEKVATGEVEFTFGVKT	213
humanDPM1	YGESKLGGNEIVSFLKGLLTLFATT 260	
mouseDPM1	YGESKLGGNEIVSFLKGLLTLFATT 260	
zebrafishDPM1	YGESKLGGNEIVSFLKGLLTLFATT 250	
drosophilaDPM1	YGTSKLGGTEIIQFAKNLLYLFATT 241	
celegansDPM1	FGESKLGSQEIVDYAKGLLYLFAFVW 239	
yeastDPM1	EGESKLSGKVIIQYLQQLKELYVFKFGANNLILFITFWSILFFYVCYQLYHLVF 267	

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