humanEBP MTTNAGPL-HPYWPQHLRLDNFVPNDRPTWHILAGLFSVTGVLVVTTWLLSG-RAAVVPL 58 mouseEBP MTTNTVPL-HPYWPRHLKLDNFVPNDLPTSHILVGLFSISGGLIVITWLLSS-RASVVPL 58 zebrafishEBP MPMEAEAVNHPYWPRNLSIPNYVANDRSMSEILVFLFSVSGILLFATWSLTGCKVSGSRL 60 \*. :: .: \*\*\*\*\*: \* : \*:\*.\*\* . .\*\*. \*\*\*: \* \*:. \*\* \*:. ::: humanEBP GTWRRLSLCWFAVCGFIHLVIEGWFVLYYEDLLGDQAFLSQLWKEYAKGDSRYILGDNFT 118 mouseEBP GAGRRLALCWFAVCTFIHLVIEGWFSLYNGILLEDQAFLSQLWKEYSKGDSRYILSDSFV 118 zebrafishEBP SAARRLALCWFTVCGFIHGVIEGWFSLYYAIIPEDOSFLSOLWKEYSKGDSRYAIADNFT 120 \*\*\*\*\*\*\*\*\*\*\*\*\*\* \*\*\*\*\*\*\*\*\*\*\*\* humanEBP VCMETITACLWGPLSLWVVIAFLRQHPLRFILQLVVSVGQIYGDVLYFLTEHRDGFQHGE 178 mouseEBP VCMETVTACLWGPLSLWVVIAFLRQQPFRFVLQLVVSMGQIYGDVLYFLTELHEGLQHGE 178 zebrafishEBP VSMETVTAYMWGPFSVWTVLAFLYNRPYRFVLQLIVSLGQLYGAVLYFFTEHRDGYIHSE 180 humanEBP LGHPLYFWFYFVFMNALWLVLPGVLVLDAVKHLTHAQSTLDAKATKAKSKKN 230

white-space: pre-wrap; ">CLUSTAL 2.1 multiple sequence alignment

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mouseEBP zebrafishEBP </body></html>