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SPIT2 HUMAN   MAQLCGLRRSRAFLALLGSLLLSGVLAADRERSIHDFCLVSKVVGRCRASMPRWWYNVTDGSCQLFVYGGCDGNSNNYLIKEECLKKCATVTENATGDLATSRNAADSSV   110
SPIT2 MACFA   MAQLCGLRRGRALLALLGSLLLSGVLAADRERSIHDFCLVSKVVGRCRASMPRWWYNVTDGSCQLFVYGGCDGNSNNYMSKEECLKKCAAVTENATGDLATSRNAADSSV   110
SPIT2 MOUSE   MAQLCELRRGRALLALVASLLLSGAQVASRELDVHESCGVSKVVGKCRASIPRWWYNITDGSCQPFVYGGCEGNGNNYQSKEECLDKCAGVTENTTDDMARNRNGADSSV   110
consensus     !!!!!*!!!*!!*!!!**!!!!!***!*!!***!**!*!!!!!!*!!!!*!!!!!!*!!!!!!*!!!!!!*!!*!!!  *!!!!!*!!!  !!!!*!*!*!**!!*!!!!!

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SPIT2 HUMAN   PSAPRRQDSEDHSSDMFNYEEYCTANAVTGPCRASFPRWYFDVERNSCNNFIYGGCRGNKNSYRSEEACMLRCFRQQENPPLPLGSKVVVLAGLFVMVLILFLGASMVYL   220
SPIT2 MACFA   PSVPRRQDSDDHSSDMFNYEEYCAAKAVTGPCRASFPRWYFDVERNSCNNFIYGGCRGNKNSYRSEEACMLRCFRQQENPSLPLGSKVVVLAGLFVMVLILFLGASMVYL   220
SPIT2 MOUSE   LSVPRKQSAEDLSAEIFNYEEYCVPKAVTGPCRAAFPRWYYDTEKNSCISFIYGGCRGNKNSYLSQEACMQHCSGKQMHPFLTPGLKAVILVGLFLMVLILLLGTSMVCL   220
consensus     *!*!!*!***!*!***!!!!!!! **!!!!!!!!!*!!!!!*!*!*!!!**!!!!!!!!!!!!!!!*!*!!!!**!***!**! !**!*!*!*!*!!!*!!!!!*!!*!!!*!

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SPIT2 HUMAN   IRVARRNQERALRTVWSSGDDKEQLVKNTYVL   252
SPIT2 MACFA   IRVARRNQERALRTVWSSGDDKEQLVKNTYVL   252
SPIT2 MOUSE   IRVVRRKQERALRTVWSTADDKEQLVKNTCVL   252
consensus     !!!*!!*!!!!!!!!!!!**!!!!!!!!!!!*!!
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