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Axmi296 : MNVSTELVPSYRNVLAHPIRLDSFFDPFVSVENELKDCDDDEGKTGYMPLQQNFNLAWN : 60
Cry21Aa2 : MNVPTILVPSYRNVLAHPIRLDSFFDPFVETFKLLKGADEEFGKTGYMDPLKCHLQIAGD : 60

Cons : MxNxTxLYPSYxNVLAHPIRLDSFFDPFVxxFxxLKxxWxxFGKTGYMxPLxQxxxiAWx

Axmi296 : AYKTEKVDLALTKASLSLIGLIPGAGPVVPYINMFVNYIWEHLFG-ASGVDPYEEFFKM : 119
Cry21Aa2 : TSQNCITVYLALTKASLSLIGLIPGADAVVPFINMFVDFIFKILFGSGSQNAQAGTEEL : 120

Cons : xxxxxGxVDxLALTKASxSxIGLIPGAGxVVPxINMFVxxIxPxLFGxxSxxxxxxAxFFxx

Axmi296 : LMKAEELVDTQKIAEFTLNNLANSILGCTYLVDFQOKTGLATCOGEKCAILLTACTPEP : 179
Cry21Aa2 : LIEKVELVDELIRNFTLNLLNVLDEMCTALSHFQNDVGLATCOGEQCEILMLIQ-TPTA : 179

Cons : LxxxVxELVDxxFxxFTLNNLxNxLxGMQTxLxxFQxxxQxAICQGExPxLxLDxxTPxx

Axmi296 : CIPTPERDKDVTIRSEDDAMKYIDACLINFKNPMOLSNDFPKDFQANFVLLTLEMYTTAAT : 239
Cry21Aa2 : CIPTTDHILISVRESEKDRRTITETALDFHKNPMLSTNDNTDFDNSDTVLLTLEMYTTAAT : 239

Cons : CIPTxxxxLxxVxxSFxKDAxxxxLxxxxLPxFKNPMxxxxNDxxxxDFxxxxVLLTLEMYTTAAT

Axmi296 : LNLVIFQCYIQERERWKDVAYLQGMMDQKANKSRRIKEYSNTVYTYKCYLESINPSNK : 299
Cry21Aa2 : LNLIIHQCYIQERERWKSUNYLESEFINCLEVDLQRIQDYSTTVSTTFEERKPLINPSNK : 299

Cons : LNLxLxQCYIQxERERWxVxYDxxxxxxQxKxxxxQxRIxxYSxTVxTTxxKxxPxLNPSNK

Axmi296 : NSVNTYNRYVRSMTLQSLIFAPLTPTLDEVNYPQNTGSHYDQTRIIFSDVAGPWEGNENI : 359
Cry21Aa2 : ESVNRYNRYVRSMTLQSLIIAPLTPTLDEVNYPNSVDIQIDQTRIIFSDVAGPWEGNDNI : 359

Cons : xSVNxYNRYVRSMTLQSLDxAAxWPTLDxVNYPxNxxxxxxDQIRLxFSVAGPWEGNxNI

Axmi296 : AANITNVLKPVSTPTCYEESDDLRFQYPRMELQSMQFHGQYVSKSVTHCYSDGLKLN : 419
Cry21Aa2 : TSNITNVLKPVSTPTCYEESDDLRFQYPRMELQSMQFHGQYVSKSVTHCYSDGLKLN : 419

Cons : xxxNIIxVLxPxTxIGxxEESDLRxFxYPRxELQSMQFHGQYVxSKSVxHCYSDGLKLN

Axmi296 : KNKTITAGVSTIDESNQNNKHNYGHLINPITDININSONSQYLDLNSVIVN-GQKLAGC : 478
Cry21Aa2 : KNKTITAGVSNIDESNQNNKHNYGVLINSPITDININSONSQYLDLNSVIVNCGQKLAGC : 479

Cons : KNKTITAGVSxIDESNQNNKHNYGPyINxPITDINxNSQNSQYLDLNSVxVNxGQKxAGC

Axmi296 : SEISGASNISNNDALENQKINVTIYVQSNNTPEKHADNYRKWGYMSSHIPYDLVPENIIGH : 538
Cry21Aa2 : SELSSNGNISNNDALENQKINVTIYVQSNNDPEKHADTYRKWGYMSSHIPYDLVPENIIGH : 539

Cons : SPxxxxxxxSNNxALPNQKINVTIYxVQSNxxPEKHADxYRKWGYMSSHIPYDLVPENxIGD

Axmi296 : IDPNTKQBPANIIRGEPAEKSSNGVLQLIKEPLNGANAVQISSGENIVLPITNITTKQYDI : 598
Cry21Aa2 : IDPNTKQBSLLIRGEPAERGYGDSIAYVSEPLNGANAVKITTSYQVLKMEVITNCTTKQYRI : 599

Cons : IDPxTKQPxxxxKGFPAEKxxxxxxxxxxEPLNGANAVxLxSxxxLxxxxTNxTTQKYxI

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FIGURA 2