




[illegible]

Accession	Type	Protein	Sequence	Position
Acr00336	type VI-B	FHNKHEVAANTNNAVP.....AENLFDYAYYYLEVN	GKNAQAKNFQSLYFFFRNMFTLGI	VSIVILIALVYSTITSVGKDVLLSEIVFKIAFFAVTIG
Acr00334	type VI-B	FHNKHEVAANTNNAVP.....AENLFDYAYYYLEVN	GKNA	
Acr00335	type VI-B	FHNKHEVAANTNNAVP.....AENLFDYAYYYLEVN	GKNAQAKNFQSLYFFFRNMFTLGI	VSIVILIALVYSTITSVGKDVLLSEIVLKIaffAVTIG
Acr00337	type VI-B	RTNKG...ISVTENIS.....NGSVFDTAYYYLEAQ	GKISQAKNFQSIYFLFRNIVTSL	FLVLPVSVIFLLASFFMN.DCSLS..EKIITIIIGTFVLGG
Acr00339	type VI-B	RTNKG...ISVTENIS.....NGSVFDTAYYYLEAQ	GKISQAKNFQSIYFLFRNIVTSL	FLVLPVSVIFLLASFFMN.DCSLS..EKIITIIIGTFVLGG
Acr00338	type VI-B	RANKG...ISATENIS.....NGSVFDTAYYYLEAQ	GKISQAKNFQSIYFLFRNIVTSL	FLVLPVSVIFLLASFFMN.DCKLS..GKIITIVIGTLVIGG
Acr00323	type VI-B	SKIHN...INSIDEIS.....KLNIFEAVITT		
Acr00333	type VI-B	IAHRKHTE.KPVNENE.....ENNLFDAAYYYLEIND	KIAA AKSFQSLYFLFRNFFTITL	FFLPISVITILILYTQNYCCCLQ..RTALTAFILLIITAF
Acr00332	type VI-B	NGTKPYCTVEQYDAMDASAQKEIEDAQDIFYDHMYR	LDCKGKLEGAKAFQSYYL	CFLHSFLGLLIFGVYLLICILLSYFMDVLLADT..WQVAFFLMNL
Acr00331	type VI-B	NGTKPYCTVEQYDAMDASAQKEIEDAQDIFYDHMYR	LDCKGKLEGAKAFQSYYL	CFLHSFLGLLIFGVYLLICILLSYFMDVLLADT..WQVAFFLMNL
Acr00325	type VI-B	GLGKIYLSSEEEFCQSEE..KEKIRMLQSDFYDRM	WYRLDFRGKLGSAKSFQCYFFFRHSFWGLV	LISLILSYKLLAYIPACDMEDIGWREYSDIAVPIMILSA
Acr00329	type VI-B	GLGKIYLSSEEEFCQSEE..KEKIRMLQSDFYDRM	WYRLDFRGKLGSAKSFQCYFFFRHSFWGLV	LISLILSYKLLAYIPACDMEDIGWREYSDIAVPIMILSA
Acr00328	type VI-B	GLGKIYLSSEEEFCQSEE..KEKIRMLQSDFYDRM	WYRLDFRGKLGSAKSFQCYFFFRHSFWGLV	LISLILSYKLLAYIPACDMEDIGWREYSDIAVPIMILSA
Acr00326	type VI-B	GLGKIYLSSEEEFCQSEE..KEKIRMLQSDFYDRM	WYRLDFRGKLGSAKSFQCYFFFRHSFWGLV	LISLILSYKLLAYIPACDMEDIGWREYSDIAVPIMILSA
Acr00327	type VI-B	GLGKIYLSSEEEFCQSEE..KEKIRMLQSDFYDRM	WYRLDFRGKLGSAKSFQCYFFFRHSFWGLV	LISLILSYKLLAYIPACDMEDIGWREYSDIAVPIMILSA
Acr00330	type VI-B	GLGKIYLSSEEEFCQSEE..KEKIRMLQSDFYDRM	WYRLDFRGKLGSAKSFQCYFFFRHSFWGLV	LISLILSYKLLAYIPACDMEDIGWREYSDIAVPIMILSV
Acr00008	type I-D	VSVSFYFVKEDNIYKLV.....QVTFDKIFFGKLYEL	KRQLSQKKQ	
Acr00119	type I-F	HGYS		
Acr00014	type I-D	IESIFN.....VERIFVDLIQLIRYPR		
Acr00317	type VI-A	LSKDK		
Acr00324	type VI-B	HWYFRQEQGREYILALS.....ENLFNKG	YGVFMPGPAKENLYHFRGMIYKLLQDSKSN	NDNNEVLLKNQSMVQKKDKQAAPNKKSV
consensus		*	** ** *	** * *

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			LEVELAQLFRKMKMYWTFYASLIEQENSNL	
Acr00336	type	VI-B	I A V P V A N W L R K K M I I T V F G C Y Y A D R V H Q T N K . . . .	226
Acr00334	type	VI-B	. . . . .	137
Acr00335	type	VI-B	I A V P V A N W L R K K M I I T V F G C Y Y A D R V H Q T N K . . . .	226
Acr00337	type	VI-B	L S S V I A Q W F R V K M T D R I F G L Y Y A E L T H N K K . . . . .	219
Acr00339	type	VI-B	L S S V I A Q W F R V K M T D R I F G L Y Y A E L T H N K K . . . . .	201
Acr00338	type	VI-B	I S S V I A Q W F R V K M T D R I F G L Y Y A E L T H H K K . . . . .	219
Acr00323	type	VI-B	. . . . .	124
Acr00333	type	VI-B	I L V Y I G R W L R E K T V E K V I W S Y Y V A I T H E E N N N K T K	229
Acr00332	type	VI-B	L F M R L A R W F R Q R M V L K M Y W A F Y E S L I E . . . . .	235
Acr00331	type	VI-B	L F M R L A R W F R Q R M V L K M Y W A F Y E S L I E . . . . .	235
Acr00325	type	VI-B	L F V F L A Q W F R I K M V E K M Y W T F Y I S L I E Q E N S N L . .	238
Acr00329	type	VI-B	L F V F L A Q W F R I K M V E K M Y W T F Y I S L I E Q E N S N L . .	238
Acr00328	type	VI-B	L F V F L A Q W F R I K M V E K M Y W T F Y I S L I E Q E N S N L . .	238
Acr00326	type	VI-B	L F V F L A Q W F R I K M V E K M Y W T F Y I S L I E Q E N S N L . .	238
Acr00327	type	VI-B	L F V F L A Q W F R I K M V E K M Y W T F Y I S L I E Q E N S N L . .	238
Acr00330	type	VI-B	L F V F L A Q W F R I K M V E K M Y W T F Y I S L I E Q E N S N L . .	238
Acr00008	type	I-D	. . . . .	126
Acr00119	type	I-F	. . . . .	69
Acr00014	type	I-D	. . . . .	107
Acr00317	type	VI-A	. . . . .	77
Acr00324	type	VI-B	N M L Q K S E I D L D . . . . .	211
consensus			* * * * * * * *	

-  non conserved
-  similar
-  ≥ 50% conserved