

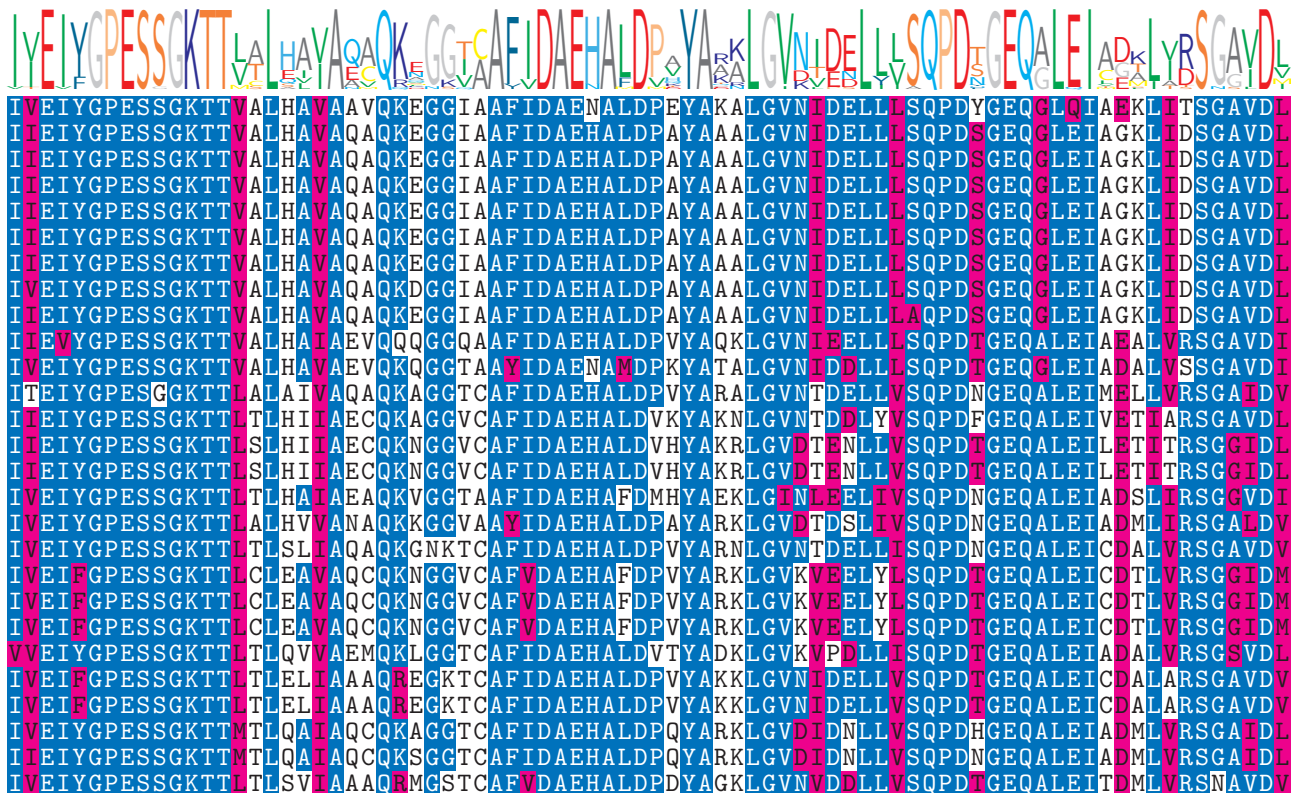
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NTDB id 269 KZH43 RS08665 WP 001085462.1	. M A K K P K L E E I S K K F G A E R E K A L N D A L K L I E K D F G K G S I M R L G E R . A E Q K V Q V M S S G S L A L D I A L G S G G Y P K G R	73
NTDB id 228 SPD RS09265 WP 001085462.1	. M A K K P K L E E I S K K F G A E R E K A L N D A L K L I E K D F G K G S I M R L G E R . A E Q K V Q V M S S G S L A L D I A L G S G G Y P K G R	73
NTDB id 194 SPR RS08825 WP 001085462.1	. M A K K P K L E E I S K K F G A E R E K A L N D A L K L I E K D F G K G S I M R L G E R . A E Q K V Q V M S S G S L A L D I A L G S G G Y P K G R	73
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NTDB id 527 SMSK321 RS11065 WP 001085454.1	. M A K K P K L D E I S K K F G A E R E K A L N D A L K L I E K D F G K G S I M R L G E R . A E Q K V Q V M S S G S L A L D I A L G S G G Y P K G R	73
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NTDB id 115 BSU 16940 NP 389576.2 M S D R Q A A L D M A L K Q I E K Q F G K G S I M K L G E K . T D T R I S T V P S G S L A L D T A L G I G G Y P R G R	58
NTDB id 625 LCA RS02525 WP 011374196.1 M A K D E R Q A A L D A A L K K I E K N F G K G S I M R M G E K . V D T Q V S T V S S G S L A L D E A L G V G G Y P R G R	60
NTDB id 1313 DR RS12030 WP 010888966.1	M S K D A T K E I S A P T D A K E R S K A I E T A M S Q I E K A F G K G S I M K L G A E . S K L D V Q V V S T G S L S L D L A L G V G G I P R G R	72
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NTDB id 232043 BB017W439 RS06600 WP 015438912.1	M A L E T K P A Q D P A T E . . T K H E L D P K R K A A L D T A L A Q V E K S F G K G S A M R L G D Q . P V Q N V E V I P T G S L A L D M A L G I G G L P K G R	77
NTDB id 1389 A4U84 RS07560 WP 021112712.1	M A . E N K K S S K N N T V . V K Q V E P E Q K K Q A L E A A L A Q I E K Q F G K G S I M K L G Q T . E Q L D I E S V S T G S L G L D L A L G I G G L P M G R	76
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consensus	* * * * * K A L A A A L K Q I E K F G K G S I M R L G E A N Q K V S V S S G S L A L D I A L G G G Y P K G R	

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NTDB id 1427 RS RS02760 WP 011000510.1	I	V	I	D	S	V	A	A	L	V	P	K	A	E	I	E	G	E	M	G	D	A	L	P	G	L	Q	A	R	L	M	S	Q	A	L	R	K	L	T	G	T	I	K	R	T	N	C	L	V	I	F	I	N	Q	L	R	M	K	I	G	V	M	.	F	G	S	P	E	T	T	T	P	G	G	N	A	L	K	F	Y	A	227	
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NTDB id	Accession	Sequence	Position
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269	KZH43_RS08665_WP_001085462.1	GDELEIEIEE	388
228	SPD_RS09265_WP_001085462.1	GDELEIEIEE	388
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consensus

-  non conserved
-  similar
-  ≥ 50% conserved