

hAAK1	46	VTVDDEVLAEGGFAIVFLVRTSNGMKCAL	KRMFV-NNEHDLQVCKREIQIMRDL	SGHKNI	VG	IDSSINN	VSSGDV	WEVLI	124	
SEL-5	47	VTIEKQIAEGGFAIVYVASDRKNNKFAL	KRQFTKDNEKQLEACCREHSFLKQC	IGHKN	IVEFV	DSYTN	CLGNG-	IWECML	125	
		::: :***::: .. * **** *. :***:::.* ** :::: ***** ::* * :..* :** ::								
hAAK1	125	LMDFCRGGQVVNLMNQRLQT--GFTENE	VLQIFCDTCEAVARLHQCKTPII	HRDL	KVEN	ILL--	HDRGH	YVLCDFGS	ATN	200
SEL-5	126	LTEYHQ-KNVLQLMNERISQNQYL	TNDEILSIFTDLCEAVSFIHNR	PQPII	HRDL	KVEN	VLISSHKPPHY	VLCDFGS	ATT	204
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hAAK1	201	KFQNPQTEGVNAVEDEIKKYTTLSY	RAPEMVNLYSGKIITTKADI	WALGCL	LYKLCY	FTLP	FGESQVA	ICDGNFT	IPDNS	280
SEL-5	205	QILSVEKYGVEYVKSEVERNTTMCY	RSPERMIDFYSGLEIGLKSDI	WALGV	LLYRLC	FFCVP	FEESPLA	IQSVNYQ	FPSVP	284
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hAAK1	281	RYSQDMHCLIRYMLEPDPDKRPDI	YQVSYFSFKLL	315						
SEL-5	285	NIPDEIKVLIYMLLDIDVNR	RPSIYQTSVLA	FE--	317					
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