

The figure displays a sequence alignment of various proteins. The top row shows a reference sequence with a blue highlight indicating a specific region. Below it, numerous other protein sequences are aligned, each represented by a colored bar corresponding to its amino acid composition. The x-axis at the bottom indicates the position of each amino acid from 10 to 200.

	810	820	830	840	850	860	870	880	890	900	910		
24582767.f		
17136512.f	FPIKWPAPPVPELVNYTRFSSKSDVAYIQLMEIFTCCKMPYGRLKNDI	EIVTCCKMPYGRLKNDI	EVVERVORCILIEBKP	KPKSCKEIYDVMKLC	WNSHGP	PDPERPAFRVLM	DQLALVA	CTLTD		
24582764.f	FPIKWPAPPVPELVNYTRFSSKSDVAYIQLMEIFTCCKMPYGRLKNDI	EIVTCCKMPYGRLKNDI	EVVERVORCILIEBKP	KPKSCKEIYDVMKLC	WNSHGP	PDPERPAFRVLM	DQLALVA	CTLTD		
17136510.f	FPIKWPAPPVPELVNYTRFSSKSDVAYIQLMEIFTCCKMPYGRLKNDI	EIVTCCKMPYGRLKNDI	EVVERVORCILIEBKP	KPKSCKEIYDVMKLC	WNSHGP	PDPERPAFRVLM	DQLALVA	CTLTD		
31222210.i	FPIKWPAPPVPELVNYTRFSSKSDVAYIQLMEVTCCKMPYGRLKNDI	EVTCCKMPYGRLKNDI	EVVERVORCILIEBKP	KACKEIYDVMKKM	SHP	PDPERPAFRVLM	DQLALVA	CTLTD		
34880171.m	FPVKWCPPPEVFNYSRFSKSDVSPQVLMEIFTCGRMFPEKNTYV	EVTCGRMFPEKNTYV	TMTRCRRLP	ASLAKSYLYBVM	MRCW	MDP	PERPGRCS	EDLLR	I	DECEET	FGR		
34877298.m	FPVKWCPPPEVFNYSRFSKSDVSPQVLMEIFTCGRMFPEKNTYV	EVTCGRMFPEKNTYV	TMTRCRRLP	ASLAKSYLYBVM	MRCW	MDP	PERPGRCS	EDLLR	I	DECEET	FGR		
7305569.m	FPVKWCPPPEVFNYSRFSKSDVSPQVLMEIFTCGRMFPEKNTYV	EVTCGRMFPEKNTYV	TMTRCRRLP	LATKYL	BVM	MRCW	MDP	PERPGRCS	EDLLR	I	DECEET	FGR	
TEC.h	FPVKWCPPPEVFNYSRFSKSDVSPQVLMEVTCSEKIDPYE	NRSN	EVVEDIST	ERLYKPLASCHYQIN	HCK	KEP	PDPERPAFRVLM	DSLL	QE	DECEET	FGR		
6754386.m	FPVKWCPPPEVFNYSRFSKSDVSPQVLMEVTCSEKIDPYE	NRSN	EVVEDIST	ERLYKPLASCHYQIN	HCK	KEP	PDPERPAFRVLM	DSLL	QE	DECEET	FGR		
34871240.m	FPVKWCPPPEVFNYSRFSKSDVSPQVLMEVTCSEKIDPYE	NRSN	EVVEDIST	ERLYKPLASPHVQVM	NHC	WCK	KDP	PDPERPAFRVLM	DSLL	QE	DECEET	FGR	
ITK.h	FPVKWCPPPEVFNYSRFSKSDVSPQVLMEVTCSEKIDPYE	NRSN	EVVEDIST	ERLYKPLASPHVQVM	NHC	WCK	KDP	PDPERPAFRVLM	DSLL	QE	DECEET	FGR	
18858911.d	FPVRMSPPPEVHFKGKYSSKSDVSPQVLMEVTCSEKIDPYE	DRN	SEVVEINTNA	CHRLKPLRCPQS	Y	BLMQWS	WCK	KDP	PERPGRCS	FAILL	LEACLO	DEEP	
34880173.m	FPVKWCPPPEVHNKYSSKSDVSPQVLMEVTCSEKIDPYE	DRN	SEVVEINTNA	CHRLKPLRCPQS	Y	BLMQWS	WCK	KDP	PERPGRCS	FAILL	LEACLO	DEEP	
34877399.m	FPVKWCPPPEVHNKYSSKSDVSPQVLMEVTCSEKIDPYE	DRN	SEVVEINTNA	CHRLKPLRCPQS	Y	BLMQWS	WCK	KDP	PERPGRCS	FAILL	LEACLO	DEEP	
7305601.m	FPVKWCPPPEVHNKYSSKSDVSPQVLMEVTCSEKIDPYE	DRN	SEVVEINTNA	CHRLKPLRCPQS	Y	BLMQWS	WCK	KDP	PERPGRCS	FAILL	LEACLO	DEEP	
38079278.m	FPVKWCPPPEVHNKYSSKSDVSPQVLMEVTCSEKIDPYE	DRN	SEVVEINTNA	CHRLKPLRCPQS	Y	BLMQWS	WCK	KDP	PERPGRCS	FAILL	LEACLO	DEEP	
TKX.h	FPVKWCPPPEVHNKYSSKSDVSPQVLMEVTCSEKIDPYE	DRN	SEVVEINTNA	CHRLKPLRCPQS	Y	BLMQWS	WCK	KDP	PERPGRCS	FAILL	LEACLO	DEEP	
6753196.m	FPVKWCPPPEVHNKYSSKSDVSPQVLMEVTCSEKIDPYE	DRN	SEVVEINTNA	CHRLKPLRCPQS	Y	BLMQWS	WCK	KDP	PERPGRCS	FAILL	LEACLO	DEEP	
BMx.h	FPVKWCPPPEVHNKYSSKSDVSPQVLMEVTCSEKIDPYE	DRN	SEVVEINTNA	CHRLKPLRCPQS	Y	BLMQWS	WCK	KDP	PERPGRCS	FAILL	LEACLO	DEEP	
34880098.m	FPVKWCPPPEVHNKYSSKSDVSPQVLMEVTCSEKIDPYE	DRN	SEVVEINTNA	CHRLKPLRCPQS	Y	BLMQWS	WCK	KDP	PERPGRCS	FAILL	LEACLO	DEEP	
34881511.m	FPVKWCPPPEVHNKYSSKSDVSPQVLMEVTCSEKIDPYE	DRN	SEVVEINTNA	CHRLKPLRCPQS	Y	BLMQWS	WCK	KDP	PERPGRCS	FAILL	LEACLO	DEEP	
7709994.m	FPVRMSPPPEVIMMSKFSSKSDIAFQVLMEIFSK	Y	SE	TCRFLRPL	LA	MT	YI	RVM	SCWHS	PERPGRCS	FAILL	LEACLO	DEEP
BTk.h	FPVRMSPPPEVIMMSKFSSKSDIAFQVLMEIFSK	Y	SE	TCRFLRPL	LA	MT	YI	RVM	SCWHS	PERPGRCS	FAILL	LEACLO	DEEP
1k2p_BTK	FPVRMSPPPEVIMMSKFSSKSDIAFQVLMEIFSK	Y	SE	TCRFLRPL	LA	MT	YI	RVM	SCWHS	PERPGRCS	FAILL	LEACLO	DEEP
1gpc_A	FPPIKWTAPAEIYNGTHTKSDVSPQVLMEIFTC	SE	TCRFLRPL	LA	SE	YI	QML	RCM	PERPGRCS	FAILL	LEACLO	DEEP	
2src	FPPIKWTAPAEIYNGTHTKSDVSPQVLMEIFTC	SE	TCRFLRPL	LA	SE	YI	QML	RCM	PERPGRCS	FAILL	LEACLO	DEEP	