

Sequence	Position
MTITTAQKRYDDAMNEFEAITSKELEQTPAFSQDILNDSDYLVITK	46
MTITTAQKRYDDAMNEFEAITSKELEQTPAFSQDILNDSDYLVITK	46
MTITTAQKRYDDAMNEFEAITSKELEQTPAFSQDILNDSDYLVITK	46
MTLTTAQKRYDDAMNEFEAITSKELEQTPAFSQDILNDSDYLVITK	46
MTLTTVQKRYDDAMNEFEAITSKELEQTPAFSQDILNDSDYLVITK	46
MTITTAQKRYDDAMNEFEAITSKELEQTPAFSQDILNDSDYLVITK	46
MTITTAQKRYDDAMNEFEAITSKELEQTPAFSQDILNDSDYLVITK	46
MTITTAQKRYNEAMHFEINMVDDFEESTPDDFAKEVLHDSDYLVITK	46
MTITTAQKRYNEAMHFEINMVDDFEESTPDDFAKEVLHDSDYLVITK	46
MTITTAQKRYNEAMHFEINMVDDFEESTPDDFAKEVLHDSDYLVITK	33
MTITTAQKRYAAMHFEINMVDDFEESTPDDFAKEVLHDSDYLVITK	46
MTITRAQKRYAAMHFEINMVDDFEESTPDDFAKEVLHDSDYLVITK	46
MTLTRAQKRYAAMHFEINMVDDFEESTPDDFAKEVLHDSDYLVITK	46
KMTLTRAQKRYAAMHFEINMVDDFEESTPDDFAKEVLHDSDYLVITK	47
MTLTRAQKRYAAMHFEINMVDDFEESTPDDFAKEVLHDSDYLVITK	46
MTLTRAQKRYAAMHFEINMVDDFEESTPDDFAKEVLHDSDYLVITK	46
MTLTRAQKRYAAMHFEINMVDDFEESTPDDFAKEVLHDSDYLVITK	46
MTLTRAQKRYAAMHFEINMVDDFEESTPDDFAKEVLHDSDYLVITK	46
MTLTRAQKRYAAMHFEINMVDDFEESTPDDFAKEVLHDSDYLVITK	46
MTTARKKRYQAISFEAMTGKDVETPQTADVLNDAEYIAFTK	44
MTNISELIREIKNKDYAVRIEG	21
MSNDRNGIINQIIDYTG	17
MNTQIKVESYVQEEVKVIFSEFEKFLRNKSSSETKNKYNSGFPVTSE	47
MIDFNSIDHAIALIQKDPQANALAGEYAYQMAR	35
MYEIKLNDTLIHQTDDRVNAFVAYRYLLRRGDLPCENIARMYYDGKVIKTDVIDHDSVHSDAQAKVSNNDIKMAISELGNVNFKSLIKKQGYPPFSNGHINSWF	105

[illegible]

logo

logo				
Acr00159	type II-A	...VILKSELN.....	116	
Acr00162	type II-A	...VILKSELN.....	116	
Acr00163	type II-A	...VILKSELN.....	116	
Acr00164	type II-A	...VILKSELN.....	116	
Acr00161	type II-A	...VILKSELN.....	116	
Acr00173	type II-A	...VILKSELN.....	116	
Acr00175	type II-A	...VIVKSELN.....	116	
Acr00176	type II-A	...IILKSELK.....	123	
Acr00174	type II-A	...IILKSELK.....	123	
Acr00169	type II-A	...IILKSELK.....	110	
Acr00167	type II-A	...IILKSELK.....	123	
Acr00166	type II-A	...IILKSELK.....	123	
Acr00160	type II-A	...IILKSELK.....	123	
Acr00165	type II-A	...IILKSELN.....	124	
Acr00168	type II-A	...IILKSELN.....	123	
Acr00170	type II-A	...IILKSEL.....	122	
Acr00177	type II-A	...IILKSELK.....	123	
Acr00171	type II-A	...VILKSELN.....	123	
Acr00172	type II-A	LKAVILKSELN.....	130	
Acr00196	type II-A	...IILKSELN.....	87	
Acr00060	type I-E	...MAVAAEIE.....	65	
Acr00008	type I-D	..LYELKRLSQKKQ.....	126	
Acr00078	type I-F	...IANQSPC.....	86	
Acr00313	type V-A	VPVIGVRDEIERVMTLEEVRNLHK	234	
consensus		*****		

- ⓧ non conserved
- ⓧ similar
- ⓧ ≥ 50% conserved