Results colour-coded for secondary structure

The current colourscheme of the alignment is for **secondary structure type**.

The 3-state (H, E C) secondary structure for each sequence is represented by a colour. If a sequence in the alignment has no colours assigned, this means that either there is no DSSP information available (if this was requested), or that no prediction was possible for that sequence (if this was requested).

The colour assignments are:

HELIX (H) STRAND (E) You have selected to perform secondary structure prediction using DSSP (Kabsch and Sander, 1983) and PSIPRED (Jones, 1999).

		10	20	30	40	50
(PRED)	gi_13633932_sp_					GDEEEKWAEK
	gi_118573879_sp			LLGWKQ		GDE <mark>EEKWAE</mark> K
(PRED)	gi_13633915_sp_	MNVTSL	FSFTS PAVKR	LLGWKQ		GDE <mark>EEKWAEK</mark>
(PRED)	gi_341942042_sp	MNVTSL	FSFTS PAVKR	LL <mark>GWKQ</mark>		GDE <mark>EEKWAEK</mark>
(PRED)	gi_13959533_sp_	MTSMASL	FSFTS PAVKR	LL <mark>GWKQ</mark>		G D E E E K W A E K
	gi_13959566_sp_		F S F T S P A V K R	LLGWKQ		GDE <mark>EEKWAEK</mark>
	gi_13959559_sp_		F S F T S P A V K R	LLGWKQ		GDEEEKWAEK
	gi_82121811_sp_		FSFTSPAVKR	LLGWKQ		GDEEEKWAEK
	gi_21264062_sp_		FSFTSPAVKR	LLGWKQ		GDEEEKWAEK
	gi_13633872_sp_		FSFTSPAVKR	LLGWKQ		GDEEKWAEK
	gi_13959539_sp_: gi_21264049_sp_		FSFTSPAVKR FSFTSPAVKR	LLGWKQ		GDEEEKWAEK GDEEEKWAEK
	gi_21204049_sp_ gi_66774168_sp_:		FSFTSPAVKR	LLGWKQ		GDEEEKWAEK
	gi_13959527_sp_		FSFT SPAVKR	LLGWKQ		GDEEEKWAEK
	gi_51338669_sp_		LPFTPPIVKR	LLGWKK		NGQEEKWCEK
	gi_60414603_sp_		LPFTPPIVKR	LLGWKK		NGQEEKWCEK
	gi_117949830_sp		LPFTPPVVKR	LLGWKKSAGG	SG-GAGGGEQ	NGQEEKWCEK
	gi_13633871_sp_		LPFTP <mark>PVVKR</mark>	LLGWK KSAGG	SG-GAGGEQ	NGQEEKWCEK
(PRED)	gi_13633914_sp_	MSSI	LPFTP <mark>PVVKR</mark>	LLGWK KSAGG	SG-GAGGEQ	NGQEEKWCEK
(PRED)	gi_75041393_sp_	MSSI	LPFTP PVVKR	LLGWK KSAGG	SG-GAGGEQ	NGQEEKWCEK
	gi_110826300_sp		LPFTP PVVKR	LLGWK KSAGG	SG-GAGGEQ	NGQEEKWCEK
(PRED)	gi_21264050_sp_	MSSI	LPFTPPVVKR	LLGWKKSASG	SSGAGGGGEQ	N G Q E E K W C E K
60						
(DDFD)	gi_13633932_sp_			KALS CPGQPS	NCVTIP	
	gi_138573879_sp		KKKGAMEELE	KALS CPGQPS	NCVTIP	
(PRED)	gi_13633915_sp_		KKKGAMEELE	KALSCPGQPS	NCVTIP	
	gi_341942042_sp		KKKGAMEELE	KALSCPGQPS	NCVTIP	
	gi_13959533_sp_		KKKGAMEELE	KALSSPGQPS	KCVTIP	
	gi_13959566_sp_		KKKGA <mark>MEELE</mark>	KALS SPGQPS	KCVTIP	
(PRED)	gi_13959559_sp_	AVDALVKKLK	KKKGA <mark>MEELE</mark>	KALS SPGQPS	KCVTIP	
(PRED)	gi_82121811_sp_	AVDALVKKLK	K K K G A MEELE	KALS SPGQPS	KCVTIP	
	gi_21264062_sp_		KKKGAMEDLE	KALS SPGQPS	KCVTIP	
	gi_13633872_sp_		KKKGAMEELE	KALS CPGQPS	NCVTIP	
	gi_13959539_sp_		KKKGAMDELE	RALSCPGQPS	KCVTIP	
	gi_21264049_sp_		KKKGAMEELE	RALSCPGQPS	NCVTIP KCVTIP	
	gi_66774168_sp_ gi_13959527_sp_		KKKGAMDELE	RALS CPGQPS RALS CPGQPS	KCVTIP	
	gi_13939327_sp_ gi_51338669_sp_			KALTTQNVNT	KCITIP	
	gi_60414603_sp_			KAITTQNINT	KCITIP	
	gi_117949830_sp			KAITTQNCNT	KCVTIPSTCS	EIWGLSTANT
	gi_13633871_sp_			KAITTQNCNT	KCVTIPSTCS	EIWGLSTANT
	gi_13633914_sp_			KAIT TQNCNT	KCVTIPSTCS	EIWGLSTPNT
(PRED)	gi_75041393_sp_	<mark>AVKSLVK</mark> KLK	K-TGRLDELE	KAIT TQNCNT	K <mark>CVTI</mark> PSTC <mark>S</mark>	EI WGLSTPNT
	gi_110826300_sp			KAIT TQNCNT	KCVTIPSTCS	EIWGLSTPNT
(PRED)	gi_21264050_sp_	AVKSLVKKLK	K-TGQLDELE	KAIT TQNRNT	KCVTIP SNCS	EIWGLSTPNT
(PRED)	gi_13633932_sp_					
	gi_118573879_sp			GRLQVSHRKG		
	gi_13633915_sp_			GRLQVSHRKG	LPHVIYCRVW	RWPDLQSHHE
(PRED)	gi_341942042_sp		RSLD	GRLQVSHRKG	LPHVIYCRVW	RWPDLQS <mark>HH</mark> E
(PRED)	gi_13959533_sp_		RSLD	GRLQVSHRKG	LPHVIYCRVW	RWPDLQSHHE
(PRED)	gi_13959566_sp_		RSLD	GRLQVSHRKG	LPHVIYCRVW	RWPDLQSHHE
	gi_13959559_sp_			GRLQVSHRKG	LPHVIYCRVW	RWPDLQSHHE
(PRED)	gi_82121811_sp_		RSLD	GRLQVSHRKG	LPH <mark>VIYCRVW</mark>	RWPDLQSHHE
(PRED)	gi_21264062_sp_		RSLD	GRLQVSHRKG	LPHVIYCRVW	RWPDLQSHHE
(PRED)	gi_13633872_sp_		RSLD	GRLQVSHRKG	LPHVIYCRVW	RWPDLQSHHE
	gi_13959539_sp_			GRLQVSHRKG	LPHVIYCRVW	RWPDLQSHHE
(LKED)	gi_21264049_sp_ gi_66774168_sp_		KSLD	GRLQVSHRKG GRLQVSHRKG	LPHVIYCRVW LPHVIYCRVW	RWPDLQSHHE RWPDLQSHHE
(DDED) (EKFD)	gi_66//4168_sp_ gi_13959527_sp_		K2TD	GRLQVSHRKG GRLQVSHRKG		RWPDLQSHHE RWPDLQSHHE
(PRED)	gi_13939327_sp_ gi_51338669_sp_		RSI.D	GRLOVSHRKG		_ ~
,)	J0_000007_0P_					

```
(PRED) gi_60414603_sp_----- ---- ----RSLD GRLQVSHRKG LPHVIYCRLW RWPDLHSHHE
(PRED) gi_117949830_spVDQWDTTGLY SFSEQTRSLD GRLQVSHRKG LPHVIYCRLW RWPDLHSHHE
(PRED) gi_13633871_sp_VDQWDTTGLY SFSEQTRSLD GRLQVSHRKG LPHVIYCRLW RWPDLHSHHE
(PRED) gi_13633914_sp_IDQWDTTGLY SFSEQTRSLD GRLQVSHRKG LPHVIYCRLW RWPDLHSHHE
(PRED) gi_75041393_sp_IDQWDTTGLY SFSEQTRSLD GRLQVSHRKG LPHVIYCRLW RWPDLHSHHE
(PRED) qi_110826300_spIDQWDTTGLY SFSEQTRSLD GRLQVSHRKG LPHVIYCRLW RWPDLHSHHE
(PRED) gi_21264050_sp_IEQWDTSGLY SYPDQTRSLD GRLQVSHRKG LPHVIYCRLW RWPDLHSHHE
                 (PRED) gi_13633932_sp_LKPLECCEFP FGSKQKEVCI NPYHYKRVES PVLPPVLVPR HSEYNPQHSL
(PRED) gi_118573879_spLKPLECCEFP FGSKQKEVCI NPYHYKRVES PVLPPVLVPR HSEYNPQHSL
(PRED) gi_13633915_sp_LKPLECCEFP FGSKQKEVCI NPYHYKRVES PVLPPVLVPR HSEYNPQHSL
(PRED) gi_341942042_spLKPLECCEFP FGSKQKEVCI NPYHYKRVES PVLPPVLVPR HSEYNPQHSL
(PRED) gi_13959533_sp_LKPLDICEFP FGSKQKEVCI NPYHYKRVES PVLPPVLVPR HNEFNPQHSL
(PRED) gi_13959566_sp_LKPLDICEFP FGSKQKEVCI NPYHYKRVES PVLPPVLVPR HNEFNPQHSL
(PRED) gi 13959559 sp LKPLDICEFP FGSKOKEVCI NPYHYKRVES PVLPPVLVPR HNEFNPOHSL
(PRED) gi_82121811_sp_LKPLDICEFP FGSKQKEVCI NPYHYKRVES PVLPPVLVPR HSEFNPQHSL
(PRED) gi_21264062_sp_LKPLEVCEYP FGSKQKEVCI NPYHYKRVES PVLPPVLVPR HSEFNPQHSL
(PRED) gi_13633872_sp_LKPLECCEFP FGSKQKEVCI NPYHYKRVES PVLPPVLVPR HSEYNPQHSL
(PRED) gi_13959539_sp_LKPLECCEFP FGSKQKEVCI NPYHYRRVET PVLPPVLVPR HSEYNPQLSL
(PRED) gi_21264049_sp_LKALECCEFP FGSKQKDVCI NPYHYKRVDS PVLPPVLVPR NSEFNAKLSM
(PRED) gi_66774168_sp_LKPLECCEFP FGSKQKEVCI NPYHYRRVET PVLPPVLVPR HSEYNPQLSL
(PRED) gi_13959527_sp_LKPLECCEFP FGSKQKEVCI NPYHYRRVET PVLPPVLVPR HSEYNPQLSL
(PRED) gi_51338669_sp_LRAMELCEFA FNMKKDEVCV NPYHYQRVET PVLPPVLVPR HTEIPAEFPP
(PRED) gi_60414603_sp_LRAMEMCEYA FNMKKDEVCV NPYHYQRVET PVLPPVLVPR HTEIPAEFPP
(PRED) gi_117949830_splkAIENCEYA FNLKKDEVCV NPYHYQRVET PVLPPVLVPR HTEILTELPP
(PRED) gi_13633871_sp_LKAIENCEYA FSLKKDEVCV NPYHYQRVET PVLPPVLVPR HTEILTELPP
(PRED) gi_13633914_sp_LKAIENCEYA FNLKKDEVCV NPYHYQRVET PVLPPVLVPR HTEILTELPP
(PRED) gi_75041393_sp_LKAIENCEYA FNLKEDEVCV NPYHYQRVET PVLPPVLVPR HTEILTELPP
(PRED) gi_110826300_splkAIENCEYA FNLKKDEVCV NPYHYORVET PVLPPVLVPR HTEILTELPP
(PRED) gi_21264050_sp_LRAIETCEYA FNLKKDEVCV NPYHYQRVET PVLPPVLVPR HTEILTELPP
                 (PRED) gi_13633932_sp_LAQFRNLG-Q NEPHMPHNAT FPDSFQQPNS HPFPHSPN-- ---SSYPNS
(PRED) gi_118573879_spLAQFRNLG-Q NEPHMPLNAT FPDSFQQPNS HPFPHSPN-- ---SSYPNS
(PRED) gi_13633915_sp_LAQFRNLG-Q NEPHMPLNAT FPDSFQQPNS HPFPHSPN-- ---SSYPNS
(PRED) gi_341942042_spLAQFRNLG-Q NEPHMPLNAT FPDSFQQPNS HPFPHSPN-- ---SSYPNS
(PRED) gi_13959533_sp_LVQFRNLS-H NEPHMPQNAT FPDSFHQPNS TPFPLSPN-- ---SPYPPS
(PRED) gi_13959566_sp_LVQFRNLS-H NEPHMPQNAT FPDSFHQPNN TPFPLSPN-- ----SPYPPS
(PRED) gi_13959559_sp_LVQFRNLS-H NEPHMPQNAT FPDSFHQPNN APFPLSPN-- ----SPYPPS
(PRED) gi_82121811_sp_LVQFRNLS-H NEPHMPHNAT FPDSFQQPNS TPFSISPN-- ----SPYPPS
(PRED) gi_21264062_sp_LVQFRNLS-H NEPHMPLNAT FPESFQQHSG GSSFPISP-- ---NSPYPP
(PRED) gi_13633872_sp_LAQFRNLG-Q NEPHMPLNAT FPDSFQQPHS HPFAQYPN-- ---SSYPNS
(PRED) gi_13959539_sp_LAKFRSASLH SEPLMPHNAT YPDSFQQPPC SALPPSPS-- ----HAFSQS
(PRED) gi_21264049_sp_LPRFRNPLHQ TEPPMPQNAT FPDSFPQQPA NALPFTPNSP TNSYPSSPNS
(PRED) gi_66774168_sp_LAKFRSASLH SEPLMPHNAT YPDSFQQSLC PAP---- ---- ------
(PRED) gi_13959527_sp_LAKFRSASLH SEPLMPHNAT YPDSFQQSLG PAP---- -----
(PRED) gi_51338669_sp_LDDY----- -SHSIPENTN FPAGI-EPQS NI-PETP--- ------
(PRED) gi_60414603_sp_LDDY----- -SHSIPENTN FPAGI-EPQS NYIPETP--- -------
(PRED) gi_117949830_spLDDY----- -THSIPENTN FPAGI-EPQS NYIPETP--- ------
(PRED) gi_13633871_sp_LDDY----- -THSIPENTN FPAGI-EPQS NYIPETP--- ------
(PRED) gi_13633914_sp_LDDY----- -THSIPENTN FPAGI-EPQS NYIPETP--- ------
(PRED) gi_75041393_sp_LDDY----- -THSIPENTN FPAGI-EPQS NYIPETP--- -------
(PRED) gi_110826300_spLDDY----- -THSIPENTN FPAGI-EPQS NYIPETP--- ------
(PRED) gi_21264050_sp_LDDY----- -TNSIPENTN FPTGI-EPPN NYIPETP--- ------
                 (PRED) gi_13633932_sp_PGSSSSTYPH SPAS-SDPGS PFQMPADTPP PAYLPPEDQM THDTSQPMDT
(PRED) gi_118573879_spPGSSSSTYPH SPTS-SDPGS PFQMPADTPP PAYLPPEDPM TQDGSQPMDT
(PRED) gi_13633915_sp_PGSSSSTYPH SPTS-SDPGS PFQMPADTPP PAYLPPEDPM TQDGSQPMDT
(PRED) gi_341942042_spPGGSSSTYPH SPTS-SDPGS PFQMPADTPP PAYLPPEDPM AQDGSQPMDT
(PRED) gi_13959533_sp_PA--SSTYPN SPAS-SGPGS PFQLPADTPP PAYMPPDDQM GQDNSQPMDT
(PRED) gi_13959566_sp_PA--SSTYPN SPAS-SGPGS PFQLPADTPP PAYMPPDDQM GQDNSQPMDT
(PRED) gi_13959559_sp_PA--SSTYPN SPAS-SGPGS PFQLPADTPP PAYMPPDDQM APDNSQPMDT
(PRED) gi_82121811_sp_PA--SSTYPS SPAS-SGPSS PFQLPADTPP PAYMPPDDQM GQDNSQSMDT
(PRED) gi_21264062_sp_SPASSGTYPN SPAS-SGPSS PFQLPADTPP PAYMPPDEQM GQDGSQSMET
(PRED) gi_13633872_sp_PGSSSSTYPH SPTS-SDPGS PFQMPADTPP PAYLPPEDPM AQDGSQPMDT
(PRED) gi_13959539_sp_PC--TASYPH SPGSPSEPES PYQHSVDTPP LPYHATEASE TQSGQPVDAT
(PRED) gi_21264049_sp_GTGSTATFPH SPSS-SDPGS PFQMP-ETPP PAYMPPEEPM TQDCPQPMDT
(PRED) gi_66774168_sp_----- ----- ----PSS PGHVFPQSPC PTSYPHSPGS PSESDSPYQH
(PRED) qi_13959527_sp_ ----- ---- ----PSS PGHVFPQSPC PTSYPQSPGS PSESDSPYQH
(PRED) gi_51338669_sp_----- ----- PPG YLSEDGETSD HQMNHSMDAG SPN-LSPNP-
(PRED) gi_60414603_sp_----- ----- PPG YLSEDGETSD HQMNPSMDAG SPN-LSPNP-
(PRED) gi_117949830_sp----- ---- ----PPG YISEDGETSD QQLNQSMDTG SPAELSPTT-
(PRED) gi_13633871_sp_----- ----- ----PPG YISEDGETSD QQLNQSMDTG SPAELSPTT-
```

```
(PRED) gi_13633914_sp_----- ----- -----PPG YISEDGETSD QQLNQSMDTG SPAELSPTT-
(PRED) gi_75041393_sp_----- ----- PPG YISEDGETSD QQLNQSMDTG SPAELSPTT-
(PRED) gi_110826300_sp----- ----- PPG YISEDGETSD QQLNQSMDTG SPAELSPTT-
(PRED) gi_21264050_sp_----- ---- ----PPG YISEDGEASD QQMNQSMDTG SPAELSPST-
                  (PRED) gi_13633932_sp_N--M-MAPGI HPDIHRGDVQ AVAYEEPKHW CSIVYYELNN RVGEAFHASS
(PRED) gi_118573879_spN--M-MAPSL PSEINRGDVQ AVAYEEPKHW CSIVYYELNN RVGEAFHASS
(PRED) gi_13633915_sp_N--M-MAPPL PSEINRGDVQ AVAYEEPKHW CSIVYYELNN RVGEAFHASS
(PRED) gi_341942042_spN--M-MAPPL PAEISRGDVQ AVAYEEPKHW CSIVYYELNN RVGEAFHASS
(PRED) gi_13959533_sp_S--NNMIPQI MPSISSRDVQ PVAYEEPKHW CSIVYYELNN RVGEAFHASS
(PRED) gi_13959566_sp_S--NNMIPQI MPSISSRDVQ PVAYEEPKHW CSIVYYELNN RVGEAFHASS
(PRED) gi_13959559_sp_S--SNMIPQT MPSISSRDVQ PVAYEEPKHW CSIVYYELNN RVGEAFHASS
(PRED) gi_82121811_sp_S--NTMIPQI MPNISTRDVQ PVAYEEPKHW CSIVYYELNN RVGEAFHASS
(PRED) gi_21264062_sp_G--SSLAPQ- --NMPRGDVQ PVEYQEPSHW CSIVYYELNN RVGEAYHASS
(PRED) qi_13633872_sp_NMTNMTAPTL PAEINRGDVQ AVAYEEPKHW CSIVYYELNN RVGERFHASS
(PRED) gi_13959539_sp_----ADRHV VLSIPNGDFR PVCYEEPQHW CSVAYYELNN RVGETFQASS
(PRED) gi_21264049_sp_N--LLAPNLP LEISNRTDVH PVAYQEPKHW CSIVYYELNN RVGEAFLASS
(PRED) gi_13959527_sp_----- ----- ----SDFR PVCYEEPLHW CSVAYYELNN RVGETFQASS
(PRED) gi_51338669_sp_-----M SPAHNNLDLQ PVTYCEPAFW CSISYYELNQ RVGETFHASQ
(PRED) gi_60414603_sp_-----M SPAHNNLDLQ PVTYCEPAFW CSISYYELNQ RVGETFHASQ
(PRED) gi_117949830_sp-----L SPVNHSLDLQ PVTYSEPAFW CSIAYYELNQ RVGETFHASQ
(PRED) qi_13633871_sp_-----L SPVNHSLDLQ PVTYSEPAFW CSIAYYELNQ RVGETFHASQ
(PRED) gi_13633914_sp_-----L SPVNHSLDLQ PVTYSEPAFW CSIAYYELNQ RVGETFHASQ
(PRED) gi_75041393_sp_-----L SPVNHSLDLQ PVTYSEPAFW CSIAYYELNQ RVGETFHASQ
(PRED) qi_110826300_sp-----L SPVNHSLDLQ PVTYSEPAFW CSIAYYELNQ RVGETFHASQ
(PRED) gi_21264050_sp_-----L SPVNHGMDLQ PVTYSEPAFW CSIAYYELNQ RVGETFHASQ
                  (PRED) gi_13633932_sp_TSILVDGFTD PSNNKNRFCL GLLSNVNRNS TIENTRRHIG KGVHLYYVGG
(PRED) gi_118573879_spTSVLVDGFTD PSNNKNRFCL GLLSNVNRNS TIENTRRHIG KGVHLYYVGG
(PRED) gi_13633915_sp_TSVLVDGFTD PSNNKNRFCL GLLSNVNRNS TIENTRRHIG KGVHLYYVGG
(PRED) gi_341942042_spTSVLVDGFTD PSNNKNRFCL GLLSNVNRNS TIENTRRHIG KGVHLYYVGG
(PRED) gi_13959533_sp_TSVLVDGFTD PANNKSRFCL GLLSNVNRNS TIENTRRHIG KGVHLYYVGG
(PRED) gi_13959566_sp_TSVLVDGFTD PSNNKSRFCL GLLSNVNRNS TIENTRRHIG KGVHLYYVGG
(PRED) gi_13959559_sp_TSVLVDGFTD PSNNKSRFCL GLLSNVNRNS TIENTRRHIG KGVHLYYVGG
(PRED) qi_82121811_sp_TSVLVDGFTD PSNNKNRFCL GLLSNVNRNS TIENTRRHIG KGVHLYYVGG
(PRED) gi_21264062_sp_TSVLVDGFTD PSNNKNRFCL GLLSNVNRNS TIENTRRHIG KGVHLYYVGG
(PRED) gi_13633872_sp_TSVLVDGFTD PSNNKNRFCL GLLSNVNRNS TIENTRRHIG KGVHLYYVGG
(PRED) gi_13959539_sp_RSVLIDGFTD PSNNRNRFCL GLLSNVNRNS TIENTRRHIG KGVHLYYVGG
(PRED) gi_21264049_sp_TSVLVDGFTD PSNNRNRFCL GLLSNVNRNS TIENTRRHIG KGVHLYYVGG
(PRED) gi_66774168_sp_RSVLIDGFTD PSNNRNRFCL GLLSNVNRNS TIENTRRHIG KGVHLYYVGG
(PRED) gi_13959527_sp_RSVLIDGFTD PSNNRNRFCL GLLSNVNRNS TIENTRRHIG KGVHLYYVGG
(PRED) gi_51338669_sp_PSMTVDGFTD PSNS-ERFCL GLLSNVNRNA AVELTRRHIG RGVRLYYIGG
(PRED) gi_60414603_sp_PSMTVDGFTD PSNS-ERFCL GLLSNVNRNA AVELTRRHIG RGVRLYYIGG
(PRED) gi_117949830_spPSLTVDGFTD PSNS-ERFCL GLLSNVNRNA TVEMTRRHIG RGVRLYYIGG
(PRED) gi_13633871_sp_PSLTVDGFTD PSNS-ERFCL GLLSNVNRNA TVEMTRRHIG RGVRLYYIGG
(PRED) qi_13633914_sp_PSLTVDGFTD PSNS-ERFCL GLLSNVNRNA TVEMTRRHIG RGVRLYYIGG
(PRED) gi_75041393_sp_PSLTVDGFTD PSNS-ERFCL GLLSNVNRNA TVEMTRRHIG RGVRLYYIGG
(PRED) gi_110826300_spPSLTVDGFTD PSNS-ERFCL GLLSNVNRNA TVEMTRRHIG RGVRLYYIGG
(PRED) gi_21264050_sp_PSLTVDGFTD PSNS-ERFCL GLLSNVNRNA TVEMTRRHIG RGVRLYYIGG
                  (PRED) gi_13633932_sp_EVYAECLSDS SIFVQSRNCN YHHGFHPTTV CKIPSGCSLK IFNNQEFAQL
(PRED) gi_118573879_sp EVYAECLSDS SIFVQSRNCN YHHGFHPTTV CKIPSGCSLK IFN<mark>NQEFAQL</mark>
(PRED) gi_13633915_sp_EVYAECLSDS SIFVQSRNCN YHHGFHPTTV CKIPSGCSLK IFNNQEFAQL
(PRED) gi_341942042_sp EVYAECLSDS SIFVQSRNCN YHHGFHPTTV CKIPSGCSLK IFNNQEFAQL
(PRED) gi_13959533_sp_<mark>EVYAEC</mark>LSDS S<mark>IFVQ</mark>SRNCN FHHGFHPTT<mark>V CK</mark>IPSSCS<mark>LK IF</mark>N<mark>NQEFAQL</mark>
(PRED) gi_13959566_sp_<mark>EVYAEC</mark>LSDS S<mark>IFVQ</mark>SRNCN FHHGFHPTT<mark>V CK</mark>IPSSCS<mark>LK IFN<mark>NQ</mark></mark>
(PRED) gi_13959559_sp_<mark>EVYAEC</mark>LSDS S<mark>IFVQ</mark>SRNCN FHHGFHPT<mark>TV CK</mark>IPSSCS<mark>LK I</mark>FN<mark>NQEFAQL</mark>
(PRED) gi_82121811_sp_<mark>EVYAEC</mark>LSDS S<mark>IFV</mark>QSRNCN YHHGFHPT<mark>TV CK</mark>IPSGCS<mark>LK I</mark>FN<mark>NQEFAQL</mark>
(PRED) qi 21264062 sp EVYAECLSDT SIFVQSRNCN YHHGFHPTTV CKIPSGCSLK IFNNQEFAQL
(PRED) gi_13633872_sp_<mark>EVYAEC</mark>LSDS S<mark>IFVQ</mark>SRNCN YHHGFHPTT<mark>V CK</mark>IPSGCS<mark>LK I</mark>FN<mark>NQEFAQL</mark>
(PRED) gi_13959539_sp_EVYAECVSDS SIFVQSRNCN YQHGFHPATV CKIPSGCSLK VFNNQLFAQL
(PRED) gi_21264049_sp_EVYAECLSDS SIFVQSRNCN YHHGFHPTTV CKIPSRCSLK IFNNQEFAEL
(PRED) gi_66774168_sp_EVYAECVSDS
                               SIFVQSRNCN YQHGFHPATV CKIPSGCSLK VFNNQLFAQL
(PRED) gi_13959527_sp_E<mark>VYAECV</mark>SDS S<mark>IFVQ</mark>SRNCN YQHGFHPAT<mark>V CKI</mark>PSGC<mark>SLK VF</mark>N<mark>NQLFAQL</mark>
(PRED) gi_51338669_sp_EVFAECLSDS AIFVQSPNCN QRYGWHPATV CKIPPGCNLK IFNNQEFAAL
(PRED) gi_60414603_sp_EVFAECLSDS AIFVQSPNCN QRYGWHPATV CKIPPGCNLK IFNNQEFAAL
(PRED) gi_117949830_spEVFAECLSDS AIFVQSPNCN QRYGWHPATV CKIPPGCNLK IFNNQEFAAL
(PRED) gi_13633871_sp_EVFAECLSDS AIFVQSPNCN QRYGWHPATV CKIPPGCNLK IFNNQEFAAL
(PRED) gi_13633914_sp_EVFAECLSDS AIFVQSPNCN QRYGWHPATV CKIPPGCNLK IFNNQEFAAL
(PRED) gi_75041393_sp_EVFAECLSDS A<mark>IFVQ</mark>SPNCN QRYGWHPAT<mark>V CK</mark>IPPGC<mark>NLK IFNNQEFAAL</mark>
(PRED) gi_110826300_spEVFAECLSDS AIFVQSPNCN QRYGWHPATV CKIPPGCNLK IFNNQGFAAL
```

```
(PRED) gi_21264050_sp_EVFAECLSDS AIFVQSPNCN QRYGWHPATV CKIPPGCNLK IFNNQEFAAL
                  (PRED) gi_13633932_sp_----LAQSVN HGFETVYELT KMCTLRMSFV KGWGAEYHRQ DVTSTPCWIE
(PRED) qi_118573879_sp----LAQSVN HGFETVYELT KMCTIRMSFV KGWGAEYHRQ DVTSTPCWIE
(PRED) gi_13633915_sp_----LAQSVN HGFETVYELT KMCTIRMSFV KGWGAEYHRQ DVTSTPCWIE
(PRED) gi_341942042_sp----LAQSVN HGFETVYELT KMCTIRMSFV KGWGAEYHRQ DVTSTPCWIE
(PRED) gi_13959533_sp_----LAQSVN HGFEAVYELT KMCTIRMSFV KGWGAEYHRQ DVTSTPCWIE
(PRED) gi_13959566_sp_----LAQSVN HGFEAVYELT KMCTIRMSFV KGWGAEYHRQ DVTSTPCWIE
(PRED) gi_13959559_sp_----LAQSVN HGFEAVYELT KMCTIRMSFV KGWGAEYHRQ DVTSTPCWIE
(PRED) gi_82121811_sp_----LAQSVN HGFEAVYELT KMCTIRMSFV KGWGAEYHRQ DVTSTPCWIE
(PRED) gi_21264062_sp_----LAQSVN HGFEAVYELT KMCTIRMSFV KGWGAEYHRQ DVTSTPCWIE
(PRED) gi_13633872_sp_----LAQSVN HGFETEYELT KMCTIRMSFV KGWGAEYHRQ DVTSTPCWIE
(PRED) gi_13959539_sp_----LAQSVH HGFEVVYELT KMCTIRMSFV KGWGAEYHRQ DVTSTPCWIE
(PRED) gi_21264049_sp_----LAQSVN HGFEAVYELT KMCTIRMSFV KGWGAKYHRQ DVTSTPCWIE
(PRED) gi_66774168_sp_----LAQSVH HGFEVVYELT KMCTIRMSFV KGWGAEYHRQ DVTSTPCWIE
(PRED) gi_13959527_sp_LAQLLAQSVH HGFEVVYELT KMCTIRMSFV KGWGAEYHRQ DVTSTPCWIE
(PRED) gi_51338669_sp_----LAQSVN QGFEAVYQLT RMCTIRMSFV KGWGAEYRRQ TVTSTPCWIE
(PRED) gi_60414603_sp_----LAQSVN QGFEAVYQLT RMCTIRMSFV KGWGAEYRRQ TVTSTPCWIE
(PRED) gi_117949830_sp----LAQSVN QGFEAVYQLT RMCTIRMSFV KGWGAEYRRQ TVTSTPCWIE
(PRED) gi_13633871_sp_----LAQSVN QGFEAVYQLT RMCTIRMSFV KGWGAEYRRQ TVTSTPCWIE
(PRED) gi_13633914_sp_----LAQSVN QGFEAVYQLT RMCTIRMSFV KGWGAEYRRQ TVTSTPCWIE
(PRED) gi_75041393_sp_----LAQSVN QGFEAVYQLT RMCTIRMSFV KGWGAEYRRQ TVTSTPCWIE
(PRED) gi_110826300_sp----LAQSVN QGFEAVYQLT RMCTIRMSFV KGWGAEYRRQ TVTSTPCWIE
(PRED) gi_21264050_sp_----LAQSVN QGFEAVYQLT RMCTIRMSFV KGWGAEYRRQ TVTSTPCWIE
                  . . . . . . . . . 510 . . . . . . . . . 520 . . . . . . .
(PRED) gi_13633932_sp_IHLHGPLQWL DKVLTQMGSP HNPISSVS
(PRED) gi_118573879_sp IHLHGPLQWL DKVLTQMGSP HNPISSVS
(PRED) gi_13633915_sp_IHLHGPLQWL DKVLTQMGSP HNPISSVS
(PRED) gi_341942042_sp IHLHGPLQWL DKVLTQMGSP HNPISSVS
(PRED) gi 13959533 sp IHLHGPLQWL DKVLTQMGSP LNPISSVS
(PRED) gi_13959566_sp_IHLHGPLQWL DKVLTQMGSP LNPISSVS
(PRED) gi_13959559_sp_IHLHGPLQWL DKVLTQMGSP LNPISSVS
(PRED) gi_82121811_sp_IHLHGPLQWL DKVLTQMGSP LNPISSVS
(PRED) gi_21264062_sp_VHLHGPLQWL DKVLTQMGSP LNPISSVS
(PRED) gi_13633872_sp_IHLHGPLQWL DKVLTQMGSP HNPISSVS
(PRED) gi_13959539_sp_IHLHGPLQWL DKVLTQMGSP HNPISSVS
(PRED) gi_21264049_sp_IHLHGPLQWL DKVLTQMGSP HNPISSVS
(PRED) gi_66774168_sp_IHLHGPLQWL DKVLTQMGSP HNPISSVS
(PRED) gi_13959527_sp_IHLHGPLQWL DKVLTQMGSP HNPISSVS
(PRED) gi_51338669_sp_LHLNGPLQWL DKVLTQMGSP SIRCSSVS
(PRED) gi_60414603_sp_LHLNGPLQWL DKVLTQMGSP SIRCSSVS
(PRED) gi_117949830_spLHLNGPLQWL DKVLTQMGSP SVRCSSMS
(PRED) gi_13633871_sp_LHLNGPLQWL DKVLTQMGSP SVRCSSMS
(PRED) gi_13633914_sp_LHLNGPLQWL DKVLTQMGSP SVRCSSMS
```

(PRED) gi_75041393_sp_LHLNGPLQWL DKVLTQMGSP SVRCSSMS (PRED) gi_110826300_sp_LHLNGPLQWL DKVLTQMGSP SVRCSSMS (PRED) gi_21264050_sp_LHLNGPLQWL DKVLTQMGSP SVRCSSMS