

logo

NTDB id 394504 FPL20 RS12350 WP 003230267.1  
NTDB id 352 BSU 24240 NP 390304.2  
consensus

MLAELS|KNFAI|EELTVSFERGLTVLTGETGAGKSI|IDAISLLVGGRSSEFVRYGEAKAELEGLFLL|ESGHPVLGVC  
MLAELSIKNFAIIEELTVSFERGLTVLTGETGAGKSIIDAISLLVGGRSSEFVRYGEAKAELEGLFLL|ESGHPVLGVC  
MLAELSIKNFAIIEELTVSFERGLTVLTGETGAGKSIIDAISLLVGGRSSEFVRYGEAKAELEGLFLL|ESGHPVLGVC  
!!

80  
80

logo

NTDB id 394504 FPL20 RS12350 WP 003230267.1  
NTDB id 352 BSU 24240 NP 390304.2  
consensus

AEQGI|DVSDEM|VRRD|STSGKSV|CRVNGKLV|T|IASLRE|IGRLLLD|IHGQHDNQL|MEDENHLQL|LDKFAGAEVESALK  
AEQGIDVSD|EMIVRRD|ISTSGKSV|CRVNGKLV|T|IASLRE|IGRLLLD|IHGQHDNQL|MEDENHLQL|LDKFAGAEVESALK  
AEQGIDVSD|EMIVRRD|ISTSGKSV|CRVNGKLV|T|IASLRE|IGRLLLD|IHGQHDNQL|MEDENHLQL|LDKFAGAEVESALK  
!!

160  
160

logo

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consensus

TYQEGYQRYV|KLLK|KQLSESE|QEMAHRLD|L|IQFLEE|ESAKLE|LNEDEQLQEERQQ|SNFEK|YESLQ|NAYNALRSE  
TYQEGYQRYV|KLLK|KQLSESE|QEMAHRLD|L|IQFLEE|ESAKLE|LNEDEQLQEERQQ|SNFEK|YESLQ|NAYNALRSE  
TYQEGYQRYV|KLLK|KQLSESE|QEMAHRLD|L|IQFLEE|ESAKLE|LNEDEQLQEERQQ|SNFEK|YESLQ|NAYNALRSE  
!!

240  
240

logo

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consensus

QGGLD|WVGMASAQ|LED|ISD|INEPL|KKMSESV|SNSYYL|LEDATF|QMRNML|DELEFD|PERLNY|IETRLNE|IKQLKRKYGATV  
QGGLDWVGMASAQLEDISDINEPLKKMSESVSNSYYLLEDATFQMRNMLDELEFDPERLNYIETRLNEIKQLKRKYGATV  
QGGLDWVGMASAQLEDISDINEPLKKMSESVSNSYYLLEDATFQMRNMLDELEFDPERLNYIETRLNEIKQLKRKYGATV  
!!

320  
320

logo

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consensus

ED|ILEYASKI|EEE|DQI|ENRDSHL|QSLK|KELDSV|GKDVAVEAANV|SQIRKTWAKK|LADE|IHRELK|SLYMEKSTFDTEFKV  
EDILEYASKI|EEE|DQI|ENRDSHL|QSLK|KELDSV|GKDVAVEAANV|SQIRKTWAKK|LADE|IHRELK|SLYMEKSTFDTEFKV  
EDILEYASKI|EEE|DQI|ENRDSHL|QSLK|KELDSV|GKDVAVEAANV|SQIRKTWAKK|LADE|IHRELK|SLYMEKSTFDTEFKV  
!!

400  
400

logo

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consensus

RTASRNEE|APLVNGQP|VQLTEQGI|DLVKFL|ISTNTGE|PLKSLSKV|ASGGELSRV|MLA|IKSIFSSQ|QDVTSI|IFDEVD|TGV  
RTASRNEE|APLVNGQP|VQLTEQGI|DLVKFL|ISTNTGE|PLKSLSKV|ASGGELSRV|MLA|IKSIFSSQ|QDVTSI|IFDEVD|TGV  
RTASRNEE|APLVNGQP|VQLTEQGI|DLVKFL|ISTNTGE|PLKSLSKV|ASGGELSRV|MLA|IKSIFSSQ|QDVTSI|IFDEVD|TGV  
!!

480  
480

logo

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consensus

SGRVAQAI|AEK|HKVS|IGSQV|LCITHL|PQVAAM|ADTHLY|IAKELK|DGR|TTTRV|KPLSKQEK|VAE|IGRMI|AGVEV|TDLTKR  
SGRVAQAI|AEK|HKVS|IGSQV|LCITHL|PQVAAM|ADTHLY|IAKELK|DGR|TTTRV|KPLSKQEK|VAE|IGRMI|AGVEV|TDLTKR  
SGRVAQAI|AEK|HKVS|IGSQV|LCITHL|PQVAAM|ADTHLY|IAKELK|DGR|TTTRV|KPLSKQEK|VAE|IGRMI|AGVEV|TDLTKR  
!!

560  
560

logo

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consensus

HAKELLKQADQVKTG  
HAKELLKQADQVKTG  
HAKELLKQADQVKTG  
!!!!!!!!!!!!!!!!

576  
576

non conserved  
 similar  
  $\geq 50\%$  conserved