

logo

NTDB id 86841 K6K01 RS05735 WP 003244988.1
NTDB id 120 BSU 10620 NP 388943.2
consensus

MGAEFLVGRSGSGKTKLIINSIQDELRRAPFGKPIIFLVPDQMTFLMEYELAKTPDMMGGMIRAQVFSFSRLAWRVLQHTGGMSR
MGAEFLVGRSGSGKTKLIINSIQDELRRAPFGKPIIFLVPDQMTFLMEYELAKTPDMMGGMIRAQVFSFSRLAWRVLQHTGGMSR
MGAEFLVGRSGSGKTKLIINSIQDELRRAPFGKPIIFLVPDQMTFLMEYELAKTPDMMGGMIRAQVFSFSRLAWRVLQHTGGMSR

85
85

logo

NTDB id 86841 K6K01 RS05735 WP 003244988.1
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consensus

FLTSTGVQMLLRKLLIEEHKQEFKVYQKASDKSGFTAQVERMLTEFKRYCLEPEDIRRMESGTASEYRGERVLSEKLDLSILY
FLTSTGVQMLLRKLLIEEHKQEFKVYQKASDKSGFTAQVERMLTEFKRYCLEPEDIRRMESGTASEYRGERVLSEKLDLSILY
FLTSTGVQMLLRKLLIEEHKQEFKVYQKASDKSGFTAQVERMLTEFKRYCLEPEDIRRMESGTASEYRGERVLSEKLDLSILY

170
170

logo

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consensus

QMEKSLADQYLHSEDYLTLLAEHIPLAEDIKGAHIYVDGFYQFTPQEFRVLEQLMVHAEHITFSLTADKPSYEREPHELELFMR
QMEKSLADQYLHSEDYLTLLAEHIPLAEDIKGAHIYVDGFYQFTPQEFRVLEQLMVHAEHITFSLTADKPSYEREPHELELFMR
QMEKSLADQYLHSEDYLTLLAEHIPLAEDIKGAHIYVDGFYQFTPQEFRVLEQLMVHAEHITFSLTADKPSYEREPHELELFMR

255
255

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G K T Y Y R L H Q K A K E L N L D I T Y K E L S G T E R H T K T P E L A H L E A Q Y E A R P A I P Y A E K Q E A L T V M Q A A N R R A E L E G I A R E I H A L V R E K G
 G K T Y Y R L H Q K A K E L N L D I T Y K E L S G T E R H T K T P E L A H L E A Q Y E A R P A I P Y A E K Q E A L T V M Q A A N R R A E L E G I A R E I H A L V R E K G
 G K T Y Y R L H Q K A K E L N L D I T Y K E L S G T E R H T K T P E L A H L E A Q Y E A R P A I P Y A E K Q E A L T V M Q A A N R R A E L E G I A R E I H A L V R E K G

340
340

logo

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RYKDVAILARQPEDYKDMVKEVFADYEIPYFIDGKASMLNHPLIEFIRSSLDVLKGNWRYEAVFRCVKTELLFPLNEPKAKVRE
RYKDVAILARQPEDYKDMVKEVFADYEIPYFIDGKASMLNHPLIEFIRSSLDVLKGNWRYEAVFRCVKTELLFPLNEPKAKVRE
RYKDVAILARQPEDYKDMVKEVFADYEIPYFIDGKASMLNHPLIEFIRSSLDVLKGNWRYEAVFRCVKTELLFPLNEPKAKVRE

425
425

logo

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VDQLENYCIAYGIKGDRWTKGDRFQYRRFVSLDDDFAQTDQEIEVENMLNDTRDNIWVPLFQLQKRMKKAKTVQEKA EALYRYLI
VDQLENYCIAYGIKGDRWTKGDRFQYRRFVSLDDDFAQTDQEIEVENMLNDTRDNIWVPLFQLQKRMKKAKTVQEKA EALYRYLI
VDQLENYCIAYGIKGDRWTKGDRFQYRRFVSLDDDFAQTDQEIEVENMLNDTRDNIWVPLFQLQKRMKKAKTVQEKA EALYRYLI

510
510

logo

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ETDVP LKLDQERQRAEDDGR I EAQQHQQA WDAVIQLLEEFVEMMGDDEISLDLFQQMIEAGAESLTFSLIPPALDQVFVGNMD
ETDVP LKLDQERQRAEDDGR I EAQQHQQA WDAVIQLLEEFVEMMGDDEISLDLFQQMIEAGAESLTFSLIPPALDQVFVGNMD
ETDVP LKLDQERQRAEDDGR I EAQQHQQA WDAVIQLLEEFVEMMGDDEISLDLFQQMIEAGAESLTFSLIPPALDQVFVGNMD

595
595

logo

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consensus



680
680

logo

NTDB id 86841 K6K01 RS05735 WP 003244988.1
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consensus



765
765

logo

NTDB id 86841 K6K01 RS05735 WP 003244988.1
NTDB id 120 BSU 10620 NP 388943.2
consensus



850
850

logo

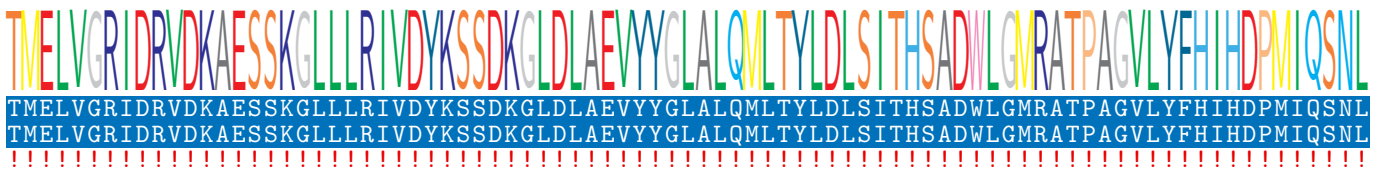
NTDB id 86841 K6K01 RS05735 WP 003244988.1
NTDB id 120 BSU 10620 NP 388943.2
consensus



935
935

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consensus



1020
1020

logo

NTDB id 86841 K6K01 RS05735 WP 003244988.1
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consensus



1105
1105

logo

NTDB id 86841 K6K01 RS05735 WP 003244988.1
NTDB id 120 BSU 10620 NP 388943.2
consensus



1166
1166

 non conserved
 similar
 $\geq 50\%$ conserved