

logo

NTDB id 176031 A3772 RS05815 WP 003244988.1
NTDB id 120 BSU 10620 NP 388943.2
consensus

MGAEFLVGRSGSGKTKLIINSIQDELRRAPFGKPIIFLVPDQMTFLMEYELAKTPDMGGMIRAQVFSFSRLAWRVLQHTG
MGAEFLVGRSGSGKTKLIINSIQDELRRAPFGKPIIFLVPDQMTFLMEYELAKTPDMGGMIRAQVFSFSRLAWRVLQHTG
!!

80
80

logo

NTDB id 176031 A3772 RS05815 WP 003244988.1
NTDB id 120 BSU 10620 NP 388943.2
consensus

GMSRPFLTSTGVQMLLRKLIIEEHKQEFKVVYQKASDKSGFTAQVERMLTEFKRYCLEPEDIRRMESGTASEYRGERVLS
GMSRPFLTSTGVQMLLRKLIIEEHKQEFKVVYQKASDKSGFTAQVERMLTEFKRYCLEPEDIRRMESGTASEYRGERVLS
GMSRPFLTSTGVQMLLRKLIIEEHKQEFKVVYQKASDKSGFTAQVERMLTEFKRYCLEPEDIRRMESGTASEYRGERVLS
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160
160

logo

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consensus

KLHDLSILYQQMEKSLADQYLHSEDYLTLLAEHIPLAEDIKGAHIYVDGFYQFTPQEFRVLEQLMVHAEHITFSLTADKP
KLHDLSILYQQMEKSLADQYLHSEDYLTLLAEHIPLAEDIKGAHIYVDGFYQFTPQEFRVLEQLMVHAEHITFSLTADKP
KLHDLSILYQQMEKSLADQYLHSEDYLTLLAEHIPLAEDIKGAHIYVDGFYQFTPQEFRVLEQLMVHAEHITFSLTADKP
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240
240

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consensus

SYEREPHELELFRMTGKTYRRLHQAQKELNLDITYKELSGTERHTKTPELAHLEAQYEARPAIPYAEKQEALTMQAANR
SYEREPHELELFRMTGKTYRRLHQAQKELNLDITYKELSGTERHTKTPELAHLEAQYEARPAIPYAEKQEALTMQAANR
SYEREPHELELFRMTGKTYRRLHQAQKELNLDITYKELSGTERHTKTPELAHLEAQYEARPAIPYAEKQEALTMQAANR
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320
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RAELEGIAREIHALVREKGYRYKDVAIILARQPEDYKDMVKEVFADYEIPYFIDGKASMLNHPLIEFIRSSLDVVKGNWRY
RAELEGIAREIHALVREKGYRYKDVAIILARQPEDYKDMVKEVFADYEIPYFIDGKASMLNHPLIEFIRSSLDVVKGNWRY
RAELEGIAREIHALVREKGYRYKDVAIILARQPEDYKDMVKEVFADYEIPYFIDGKASMLNHPLIEFIRSSLDVVKGNWRY
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400
400

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consensus

EAVFRCVKTELLFPLNEPKAKVREQVDQLENYCIAYGIKGRWTKGDRFYRRFVSLDDDDFAQTDQEIEEMNMLNDRDW
EAVFRCVKTELLFPLNEPKAKVREQVDQLENYCIAYGIKGRWTKGDRFYRRFVSLDDDDFAQTDQEIEEMNMLNDRDW
EAVFRCVKTELLFPLNEPKAKVREQVDQLENYCIAYGIKGRWTKGDRFYRRFVSLDDDDFAQTDQEIEEMNMLNDRDW
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480
480

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IVPPLFQLQKRMKKAKTVQEKAEALRYRLEETDVPLKLDQERQRAEDDGRITIEAQQHQQAWDAVIQLLEEFVEMMGDDEI
IVPPLFQLQKRMKKAKTVQEKAEALRYRLEETDVPLKLDQERQRAEDDGRITIEAQQHQQAWDAVIQLLEEFVEMMGDDEI
IVPPLFQLQKRMKKAKTVQEKAEALRYRLEETDVPLKLDQERQRAEDDGRITIEAQQHQQAWDAVIQLLEEFVEMMGDDEI
!!

560
560

