

logo

NTDB id 1068318 ACFVP7 RS15285 WP 396582430.1
NTDB id 354 BSU 27620 NP 390640.1
consensus

MLASKMRWEIQRPDQDKVKSLEQLHITPLVASLLVKRGFDTAESARLFLHTKDADFYPFEMKGMKEAADRIKQAISSQQ
MLASKMRWEIQRPDQDKVKSLEQLHITPLVASLLVKRGFDTAESARLFLHTKDADFYPFEMKGMKEAADRIKQAISSQQ
MLASKMRWEIQRPDQDKVKSLEQLHITPLVASLLVKRGFDTAESARLFLHTKDADFYPFEMKGMKEAADRIKQAISSQQ
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80
80

logo

NTDB id 1068318 ACFVP7 RS15285 WP 396582430.1
NTDB id 354 BSU 27620 NP 390640.1
consensus

EKIMYGDYDADGVTSTSVMLHTLQKLSAQVDFYIPDRFKEGYGPNQAFRSIKERGFSLIITVDTGIAAVHEAKVAKEL
EKIMYGDYDADGVTSTSVMLHTLQKLSAQVDFYIPDRFKEGYGPNQAFRSIKERGFSLIITVDTGIAAVHEAKVAKEL
EKIMYGDYDADGVTSTSVMLHTLQKLSAQVDFYIPDRFKEGYGPNQAFRSIKERGFSLIITVDTGIAAVHEAKVAKEL
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160
160

logo

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consensus

GLDVIITDHEPGPEL PDVRAIVHPKQPGCTYPFKELAGVGVAFKLAHALLGELPDELLDLAAIGTIADLVPLHDENRLI
GLDVIITDHEPGPEL PDVRAIVHPKQPGCTYPFKELAGVGVAFKLAHALLGELPDELLDLAAIGTIADLVPLHDENRLI
GLDVIITDHEPGPEL PDVRAIVHPKQPGCTYPFKELAGVGVAFKLAHALLGELPDELLDLAAIGTIADLVPLHDENRLI
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240
240

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consensus

ATLGLERLRRTNRLGLKELIKLSSGGDIGEANEETVGFQLAPRLNAVGRIEQADPAVHLLMSEDSFEAEELAAEIDQLNKE
ATLGLERLRRTNRLGLKELIKLSSGGDIGEANEETVGFQLAPRLNAVGRIEQADPAVHLLMSEDSFEAEELAAEIDQLNKE
ATLGLERLRRTNRLGLKELIKLSSGGDIGEANEETVGFQLAPRLNAVGRIEQADPAVHLLMSEDSFEAEELAAEIDQLNKE
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320
320

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consensus

RQKMSKMTDEAIEVVEQQGLDQTAIVVAKAGWNPVGVIVASKLVDLRFYRPAIVLGI DEEKGI AKGSARSIRGFNLFES
RQKMSKMTDEAIEVVEQQGLDQTAIVVAKAGWNPVGVIVASKLVDLRFYRPAIVLGI DEEKGI AKGSARSIRGFNLFES
RQKMSKMTDEAIEVVEQQGLDQTAIVVAKAGWNPVGVIVASKLVDLRFYRPAIVLGI DEEKGI AKGSARSIRGFNLFES
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400
400

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consensus

LSECRDILPHFGGHPMAAGMTLKAEDVPDLRSRLNEIADNTL TEEDFIPVQEVLDVCGVEDITVESIAEMNMLSPFGMLN
LSECRDILPHFGGHPMAAGMTLKAEDVPDLRSRLNEIADNTL TEEDFIPVQEVLDVCGVEDITVESIAEMNMLSPFGMLN
LSECRDILPHFGGHPMAAGMTLKAEDVPDLRSRLNEIADNTL TEEDFIPVQEVLDVCGVEDITVESIAEMNMLSPFGMLN
!!

480
480

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consensus

PKPHVLVENAVLEDVRKIGANKTHVKMTIRNESSQLDCVGFNKGELQEGIVPGSRISIVGEMSIENWNNRKKPQLMIKDA
PKPHVLVENAVLEDVRKIGANKTHVKMTIRNESSQLDCVGFNKGELQEGIVPGSRISIVGEMSIENWNNRKKPQLMIKDA
PKPHVLVENAVLEDVRKIGANKTHVKMTIRNESSQLDCVGFNKGELQEGIVPGSRISIVGEMSIENWNNRKKPQLMIKDA
!!

560
560

