

logo

NTDB id 49456 SALIVA RS02160 WP 014633984.1
NTDB id 480 HSISS4 RS02090 WP 002890133.1
consensus

MSDNRAHLEEKYTWDLTTIFATDADWETEYESTVQDLKKASAYAGHL LDSAKNLLGATELYMSLMRRLEKIYVYASMKND
MSDNRAHLEEKYTWDLTTIFATDADWETEYESTVQDLKKASAYAGHL LDSAKNLLGATELYMSLMRRLEKIYVYASMKND
MSDNRTHLEEKYTWDLTTIFATDADWETEYESTVQDLKKASAYAGHL LDSAKNLLGATELYMSLMRRLEKIYVYASMKND
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80
80

logo

NTDB id 49456 SALIVA RS02160 WP 014633984.1
NTDB id 480 HSISS4 RS02090 WP 002890133.1
consensus

QDTTVGLYQEYQAKASNLYSQLSEAFAYFEPEFMALDDKKLAEFKEQEPGLGLYDHYFERLLANKDHVLSQEAPELLAAA
QDTTVGLYQEYQAKASNLYSQLSEAFAYFEPEFMALDDKKLAEFKEQEPGLGLYDHYFERLLANKDHVLSQEAPELLAAA
QDTTVGLYQEYQAKASNLYSQLSEAFAYFEPEFMALDDKKLAEFKEQEPGLGLYDHYFERLLANKDHVLSQEAPELLAAA
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160
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logo

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consensus

GDIFNGPTDTFNVLDNADILFPWVSDGQGDVVELTHGNFITLMESKDRDIRKGAYEAMYGTYEQFQHTYAQTLQGVVKKVH
GDIFNGPTDTFNVLDNADILFPWVSDGQGDVVELTHGNFITLMESKDRDIRKGAYEAMYGTYEQFQHTYAQTLQGVVKKVH
GDIFNGPTDTFNVLDNADILFPWVSDGQGDVVELTHGNFITLMESKDRDIRKGAYEAMYGTYEQFQHTYAQTLQGVVKKVH
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consensus

NYQAKVRHYN SARHAALANFIPESVYDSLLESVNKHLPLLRHYLDLRKKVGLDELKMYDVYTP LSETETALTYEESLK
NYQAKVRHYN SARHAALANFIPESVYDSLLESVNKHLPLLRHYLDLRKKVGLDELKMYDVYTP LSETETALTYEESLK
NYQAKVRHYN SARHAALANFIPESVYDSLLESVNKHLPLLRHYLDLRKKVGLDELKMYDVYTP LSETETALTYEESLK
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320
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KAEVLAIFGEEYSKGVHAAFTERWIDVHPNKGKRS GAYSGGAYDTN AFMLLNWQD TLDNLF TLVHETGHS LHSTFTRQT
KAEVLAIFGEEYSKGVHAAFTERWIDVHPNKGKRS GAYSGGAYDTN AFMLLNWQD TLDNLF TLVHETGHS LHSTFTRQT
KAEVLAIFGEEYSKGVHAAFTERWIDVHPNKGKRS GAYSGGAYDTN AFMLLNWQD TLDNLF TLVHETGHS LHSTFTRQT
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400
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consensus

QPYVYGDYPIFLAEI ASTT NENIL TETLLKEVKDDKTRFAILNHYLDGFKGT VFRQTQFAEFEHA IHEADASGQILTADF
QPYVYGDYPIFLAEI ASTT NENIL TETLLKEVKDDKTRFAILNHYLDGFKGT VFRQTQFAEFEHA IHEADASGQILTADF
QPYVYGDYPIFLAEI ASTT NENIL TETLLKEVKDDKTRFAILNHYLDGFKGT VFRQTQFAEFEHA IHEADASGQILTADF
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480
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consensus

MNKLYADLN EKYNNLKAEDN YEIQFEWERIPHFYMN YVYQYATGFAAASYLAEKIVHGTEEDKEAYLTYLKAGSSDYPL
MNKLYADLN EKYNNLKAEDN YEIQFEWERIPHFYMN YVYQYATGFAAASYLAEKIVHGTEEDKEAYLTYLKAGSSDYPL
MNKLYADLN EKYNNLKAEDN YEIQFEWERIPHFYMN YVYQYATGFAAASYLAEKIVHGTEEDKEAYLTYLKAGSSDYPL
!!

560
560

