

logo

NTDB id 1060126 ACF3NW RS10545 WP 392423323.1
NTDB id 1133 OK783 RS06970 WP 010951252.1
consensus



57
80

logo

NTDB id 1060126 ACF3NW RS10545 WP 392423323.1
NTDB id 1133 OK783 RS06970 WP 010951252.1
consensus



57
160

logo

NTDB id 1060126 ACF3NW RS10545 WP 392423323.1
NTDB id 1133 OK783 RS06970 WP 010951252.1
consensus



57
240

logo

NTDB id 1060126 ACF3NW RS10545 WP 392423323.1
NTDB id 1133 OK783 RS06970 WP 010951252.1
consensus



57
320

logo

NTDB id 1060126 ACF3NW RS10545 WP 392423323.1
NTDB id 1133 OK783 RS06970 WP 010951252.1
consensus



57
400

logo

NTDB id 1060126 ACF3NW RS10545 WP 392423323.1
NTDB id 1133 OK783 RS06970 WP 010951252.1
consensus



57
480

logo

NTDB id 1060126 ACF3NW RS10545 WP 392423323.1
NTDB id 1133 OK783 RS06970 WP 010951252.1
consensus



57
560

logo

NTDB id 1060126 ACF3NW RS10545 WP 392423323.1
NTDB id 1133 OK783 RS06970 WP 010951252.1
consensus

AVQNLENEWVKWQEESTLAGFFGQLPQQTIVIRHIRRFLDSESEAGFLRGGITFCSMVPMRSLPFKVICLLGLNDGDFPRN
AVQNLENEWVKWQEESTLAGFFGQLPQQTIVIRHIRRFLDSESEAGFLRGGITFCSMVPMRSLPFKVICLLGLNDGDFPRN

57
640

logo

NTDB id 1060126 ACF3NW RS10545 WP 392423323.1
NTDB id 1133 OK783 RS06970 WP 010951252.1
consensus

TKAAVFDLVAKHPAKGDRARRDDDRYLFLEALISAREILYLSYIGRDIRKDEELASSSLLGELIDTVAAMTGTNSRQLAQ
TKAAVFDLVAKHPAKGDRARRDDDRYLFLEALISAREILYLSYIGRDIRKDEELASSSLLGELIDTVAAMTGTNSRQLAQ

57
720

logo

NTDB id 1060126 ACF3NW RS10545 WP 392423323.1
NTDB id 1133 OK783 RS06970 WP 010951252.1
consensus

NWIEQHPLQAFSRRYFQEGGRSDGIFGTRTDYAAALGQTPEPPQPFDDQPVENAEPVAEIQGDEFIRFWRNPVKVWLQQL
NWIEQHPLQAFSRRYFQEGGRSDGIFGTRTDYAAALGQTPEPPQPFDDQPVENAEPVAEIQGDEFIRFWRNPVKVWLQQL

57
800

logo

NTDB id 1060126 ACF3NW RS10545 WP 392423323.1
NTDB id 1133 OK783 RS06970 WP 010951252.1
consensus

AWSEPHIGEAWEPAEPEPQHAEQIAETIYGARREGQDFSQTAARIAAESLLPSGELGKLWQSFQTAAKQIDTAVLNSP
AWSEPHIGEAWEPAEPEPQHAEQIAETIYGARREGQDFSQTAARIAAESLLPSGELGKLWQSFQTAAKQIDTAVLNSP

57
880

logo

NTDB id 1060126 ACF3NW RS10545 WP 392423323.1
NTDB id 1133 OK783 RS06970 WP 010951252.1
consensus

KLPPFPYAIPSDGQILKGS LGNLYRCGQVFYAYGKPNAPQRVAFLEHLIFCAVMPSEAE TRQTFIVQSGETEVLAEIAQ
KLPPFPYAIPSDGQILKGS LGNLYRCGQVFYAYGKPNAPQRVAFLEHLIFCAVMPSEAE TRQTFIVQSGETEVLAEIAQ

57
960

logo

NTDB id 1060126 ACF3NW RS10545 WP 392423323.1
NTDB id 1133 OK783 RS06970 WP 010951252.1
consensus

DRALQLLSEWMAFFNIGQNRPLPFFAKTSLAAAEFAQKQDWEAALKKAQTAYYGSKVSKGQKDYTEVALVFGNASQNPL
DRALQLLSEWMAFFNIGQNRPLPFFAKTSLAAAEFAQKQDWEAALKKAQTAYYGSKVSKGQKDYTEVALVFGNASQNPL

57
1040

logo

NTDB id 1060126 ACF3NW RS10545 WP 392423323.1
NTDB id 1133 OK783 RS06970 WP 010951252.1
consensus

EQPLFENLARLLADTLAAAEKREGTGAA
EQPLFENLARLLADTLAAAEKREGTGAA

57
1068

 non conserved
 similar
 $\geq 50\%$ conserved