

logo

NTDB id 966008 NST10 RS00505 WP 000153535.1
NTDB id 4 SA RS09755 WP 000153535.1
NTDB id 83 BSU 35490 NP 391429.1
consensus

Row	Sequence	Score
1	.MTIKVLFVDDHEMVRIGISSYLSTQSDIEVVGEGASGKEATAKAHELKPDLLIMDLLMEDMDGVEATTQIKKDLDPQIKV	79
2	.MTIKVLFVDDHEMVRIGISSYLSTQSDIEVVGEGASGKEATAKAHELKPDLLIMDLLMEDMDGVEATTQIKKDLDPQIKV	79
3	MTKNIVIIIDHQLFREGVKRIIDFEPTFEVVAEGDDGEAARIVEHYHPDVIIMDINMPNVNGVEATKQLVELYPESKV	80

logo

NTDB id 966008 NST10 RS00505 WP 000153535.1
NTDB id 4 SA RS09755 WP 000153535.1
NTDB id 83 BSU 35490 NP 391429.1
consensus

[illegible]

logo

NTDB id 966008 NST10 RS00505 WP 000153535.1
NTDB id 4 SA RS09755 WP 000153535.1
NTDB id 83 BSU 35490 NP 391429.1
consensus

Sequence	Position
ELYEMLTEREMEILLIAKGYSNQEIASASHITIKTVKTHVSNILSKLEVQDRTQAVIYAFQHNLIQ..	209
ELYEMLTEREMEILLIAKGYSNQEIASASHITIKTVKTHVSNILSKLEVQDRTQAVIYAFQHNLIQ..	209
RPLHLITRRECEVLQMLADGKSNRGIGESLFISEIKTVKNHVSNIQKNNVNDRTQAVVVAIKNGWVEMR	229

- X non conserved
- X similar
- X > 50% conserved