

logo

NTDB id 416611 E3S75 RS00450 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

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NTDB id 416611 E3S75 RS00450 WP 000153535.1
NTDB id 4 SA RS09755 WP 000153535.1
NTDB id 83 BSU 35490 NP 391429.1
consensus

[illegible]

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consensus

ELYEMLTEREMEILLIAKGYSNQEIASASHITIKTVKTHVSNILSKLEVQDRTQAVIYAFQHNLIQ
 RPLHLIRRCVQMLDKRGESLFISEIKTVKNHVSNIQLQVWNRDRTQAVIYAFQHNLIQ
 ELYEMLTEREMEILLIAKGYSNQEIASASHITIKTVKTHVSNILSKLEVQDRTQAVIYAFQHNLIQ
 ELYEMLTEREMEILLIAKGYSNQEIASASHITIKTVKTHVSNILSKLEVQDRTQAVIYAFQHNLIQ
 RPLHLITRRECEVLQMLADGKSNRGIGESLFISEIKTVKNHVSNIQLQVWNRDRTQAVIYAFQHNLIQ

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