

Sequence logo showing amino acid conservation across two sequences. The top sequence is LEKELVQRPGSLHIRRQLSLLNRASISTLHSFCLQVLKKYYMIDLDPGFRMADQTEGELTGDEVLDLFEDEYAKGNQA. The bottom sequence is LEKELVQRPGSLHIRRQLSLLNRASISTLHSFCLQVLKKYYYIDLDPGFRIMADQTEGELTGDEVLDLFEDEYAKGEKA. Asterisks mark positions 9, 10, 16, 17, 23, 24, 28, 29, 35, 36, 42, 43, 49, 50, 56, 57, 63, 64, 70, 71, 77, 78, 84, 85, 91, 92, 98, 99, 105, 106, 112, 113, 119, 120, 126, 127, 133, 134, 140, 141, 147, 148, 154, 155, 161, 162, 168, 169, 175, 176, 182, 183, 189, 190, 196, 197, 203, 204, 210, 211, 217, 218, 224, 225, 231, 232, 238, 239, 245, 246, 252, 253, 259, 260, 266, 267, 273, 274, 280, 281, 287, 288, 294, 295, 301, 302, 308, 309, 315, 316, 322, 323, 329, 330, 336, 337, 343, 344, 350, 351, 357, 358, 364, 365, 371, 372, 378, 379, 385, 386, 392, 393, 399, 400.

FFE_AL_ADRYTTDRHDL_DLD_DL_DQ_DLV_KR_RVVEYSRSH_PDP_DPEAW_LQ_QSFV_RRL_LYDV_VTE_ESK_KMEEL_LPFYQV_VRED_DEA_AMA_ALF_FGAK_KQ_QKL_LERA_A

FFE_AL_ADRYTTDRHDL_DLD_DL_DQ_DLV_KR_RVVEYSRSH_PDP_DPEAW_LQ_QSFV_RRL_LYDV_VTE_ESK_KMEEL_LPFYQV_VRED_DEA_AMA_ALF_FGAK_KQ_QKL_LERA_A

320
 320

logo

NTDB id 206498 BSF20 RS01595 WP 003155147.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



640
640

logo

NTDB id 206498 BSF20 RS01595 WP 003155147.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



720
720

logo

NTDB id 206498 BSF20 RS01595 WP 003155147.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



800
800

logo

NTDB id 206498 BSF20 RS01595 WP 003155147.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



880
880

logo

NTDB id 206498 BSF20 RS01595 WP 003155147.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



960
958

logo

NTDB id 206498 BSF20 RS01595 WP 003155147.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



1040
1038

logo

NTDB id 206498 BSF20 RS01595 WP 003155147.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



1120
1118

logo

NTDB id 206498 BSF20 RS01595 WP 003155147.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



logo

NTDB id 206498 BSF20 RS01595 WP 003155147.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



- non conserved
- similar
- $\geq 50\%$ conserved