

logo

NTDB id 81184 OCUAC RS15980 WP 072689176.1
NTDB id 1072 ABD1 RS01805 WP 000156662.1
consensus

85

85

logo

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NTDB id 1072 ABD1 RS01805 WP 000156662.1
consensus

LDEQVL_NLLTWSAQYYQFP|GEVMTALPALLRQGKPM_DVL_FHL_WKI|T_PCDN_AEALLKRS_IKQQDAYQILKLHPAGTTENILNLS
 LDEQVL_NLLTWSAQYYQFP|GEVMTALPALLRQGKPM_DVL_FHL_WKI|T_PCDN_AEALLKRS_IKQQDAYQILKLHPAGTTENILNLS
 LDEQVL_SLLTWSAQYYQFP|GEVMTALPALLRQGKPM_DVL_FHL_WKI|T_PCDN_VEALLKRS_GKQQDAYQILKLHPAGTTENILNLS

170

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GVETATLKALQKKGLVDCTLEPHDFSP_APVQLAQMP_LTLNEDQKKATQ_HY_LNAQHQQYQAFLLDGLTGS_GKT_EVY_LHIMHEV_LKQ_G
GVETATLKALQKKGLVDCTLEPHDFSP_APVQLAQMP_LTLNEDQKKATQ_HY_LNAQHQQYQAFLLDGLTGS_GKT_EVY_LHIMHEV_LKQ_G
GVETATLKALQKKGLVDCTLEPHDFSP_SPVQLAQMP_LTLNEDQKKATQ_YY_LNAQHQQYQAFLLDGLTGS_GKT_EVY_LHIMHEV_LKQ_G

255

255

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KQVLVLVPEIGLTPQTSRFKSRFNCDIALHSGLNDSKRLQAWQQAQTGKASIIIGTRSAIYTPLPRLGLIILDEEHDLSYKQQ
KQVLVLVPEIGLTPQTSRFKSRFNCDIALHSGLNDSKRLQAWQQAQTGKASIIIGTRSAIYTPLPRLGLIILDEEHDLSYKQQ
KQVLVLVPEIGLTPQTSRFKSRFNCDIALHSGLNDSKRLQAWQQAQTGKASIIIGTRSAIYTPLPRLGLIILDEEHDLSYKQQ

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EGFR_{YHARDVALYRGHLQGCPVLLGSATPSIDSYHLVETGKLTALQLNQRAGHALLPKMHLIDLKIVKKQHGISQPLIEQIKNTL}

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ARKEQVLIFLNRRGYAPVLVCECGWQANCPHCDAHFTLHTQPYSYLCHHCGTVHRLPDHCPECQQKSLKTLGAGTAKVEEHLQ
ARKEQVLIFLNRRGYAPVLVCECGWQANCPHCDAHFTLHTQPYSYLCHHCGTVHRLPDHCPECQQKSLKTLGAGTAKVEEHLQ
ARKEQVLIFLNRRGYAPVLVCECGWQANCPHCDAHFTLHTQPYSYLCHHCGTVHRLPDHCPECQQKSLKTLGAGTAKVEEHLQ

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ELFPDHDVIRVDRDST_SNRVGSWQKIYDRIQQNKPSILLGTQMLAKGHHFPHVTLVAILDIDAGLLSVDIRAPERTAQLIVQVAGR
ELFPDHDVIRVDRDST_NNRVGSWQKIYDRIQQNKPSILLGTQMLAKGHHFPHVTLVAILDIDAGLLSVDIRAPERTAQLIVQVAGR
ELFPDHDVIRVDRDST_SNRVGSWQKIYDRIQQNKPSILLGTQMLAKGHHFPHVTLVAILDIDAGLLSVDIRAPERTAQLIVQVAGR

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680
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consensus



742
742

- ☐ non conserved
- ☒ similar
- ☒ ≥ 50% conserved