

logo

NTDB id 127726 IX87 RS16075 WP 000156642.1
NTDB id 1072 ABD1 RS01805 WP 000156662.1
consensus

85
85

logo

NTDB id 127726 IX87 RS16075 WP 000156642.1
NTDB id 1072 ABD1 RS01805 WP 000156662.1
consensus

170
170

logo

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consensus

GVETATLKALQKKGLVDCTLEPHDFSP_APVQL_AQMPLTLNEDQKKATQ_{HV}LN_AQHQQYQAFLLDGLTGS_GKT_EVY_LHIM_HEV_LKQ_G
GVETATLKALQKKGLVDCTLEPHDFSP_APVQL_AQMPLTLNEDQKKATQ_{HV}LN_AQHQQYQAFLLDGLTGS_GKT_EVY_LHIM_HEV_LKQ_G
GVETATLKALQKKGLVDCTLEPHDFSP_SPVQL_AQMPLTLNEDQKKATQ_{HV}LN_AQHQQYQAFLLDGLTGS_GKT_EVY_LHIM_HEV_LKQ_G
* *

255
255

logo

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KQVLVLVPEIGLTPQTISRFSRFNCDIALLHSGLNDSKRLQAWQQAQTGKASIIIGTRSAIYTPLPRLGLIILDEEHDLSYKQQ
KQVLVLVPEIGLTPQTISRFSRFNCDIALLHSGLNDSKRLQAWQQAQTGKASIIIGTRSAIYTPLPRLGLIILDEEHDLSYKQQ
KQVLVLVPEIGLTPQTISRFSRFNCDIALLHSGLNDSKRLQAWQQAQTGKASIIIGTRSAIYTPLPRLGLIILDEEHDLSYKQQ

340
340

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EGFR YHARDVALYRGHLQGCPVLLGSATPSIDSYHLVETGKLTALQLNQRAGHALLPKMHLIDLKIVKKQHGISQPLIEQIKNTL

425
425

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ARKEQVLIFLNRRGYAPVLVCESGWQANCPHCDAHFTLHTQPYSYLCHHCGTVHRLPDHCPECQQKSLKTLGAGTAKVEEHLQ
ARKEQVLIFLNRRGYAPVLVCESGWQANCPHCDAHFTLHTQPYSYLCHHCGTVHRLPDHCPECQQKSLKTLGAGTAKVEEHLQ
ARKEQVLIFLNRRGYAPVLVCESGWQANCPHCDAHFTLHTQPYSYLCHHCGTVHRLPDHCPECQQKSLKTLGAGTAKVEEHLQ

510
510

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ELFPDHDVIRVDRDSTSRVGSWQKIYDRIQQNKPSILLGTQMLAKGHHFPHTLVAILDIDAGLLSVDIRAPERTAQLIVQVAGR
ELFPDHDVIRVDRDSTSRVGSWQKIYDRIQQNKPSILLGTQMLAKGHHFPHTLVAILDIDAGLLSVDIRAPERTAQLIVQVAGR
ELFPDHDVIRVDRDSTSRVGSWQKIYDRIQQNKPSILLGTQMLAKGHHFPHTLVAILDIDAGLLSVDIRAPERTAQLIVQVAGR

595
595

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680
680

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consensus



742
742

-  non conserved
-  similar
-  ≥ 50% conserved