logo

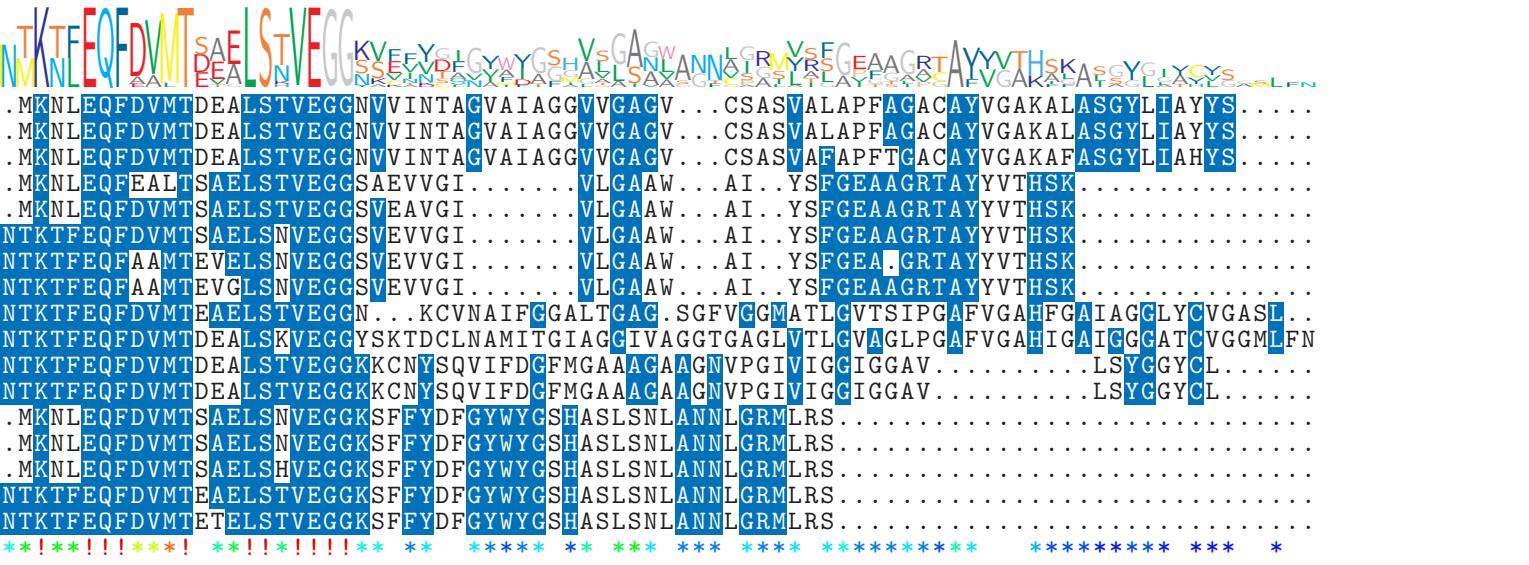| Label	Sequence
BKANDP 06005	..MKNLEQFDVMTDEALSTVEGGNVVINTAGVAIAGGVVGAGV...CSASVALAPFAGACAYVGAKALASGYLIAYYS.....
GINLMA 04435	..MKNLEQFDVMTDEALSTVEGGNVVINTAGVAIAGGVVGAGV...CSASVALAPFAGACAYVGAKALASGYLIAYYS.....
ECDHJF 06265	..MKNLEQFDVMTDEALSTVEGGNVVINTAGVAIAGGVVGAGV...CSASVAFAPFTCACAYVGAKAFASGYLIAHYS.....
NOCGMI 06630	..MKNLEQFEALTSAELSTVEGGSAEVVGI.......VLGAAW...AI.YSFGEAAGRTAYYVTHSK.............
DDIDDG 02255	..MKNLEQFDVMTSAELSTVEGGSVEAVGI.......VLGAAW...AI.YSFGEAAGRTAYYVTHSK.............
ECDHJF 06275	MNTKTFEQFDVMTSAELSNVEGGSVEVVGI.......VLGAAW...AI.YSFGEAAGRTAYYVTHSK.............
BKANDP 05995	MNTKTFEQFAAMTEVELSNVEGGSVEVVGI.......VLGAAW...AI.YSFGEA.GRTAYYVTHSK.............
GINLMA 04445	MNTKTFEQFAAMTEVGLSNVEGGSVEVVGI.......VLGAAW...AI.YSFGEAAGRTAYYVTHSK.............
EGPFLP 00955	MNTKTFEQFDVMTEAELSTVEGGN...KCVNAIFGGALTGAG.SGFVGGMATLGVTSIPGAFVGAHFGAIAGGLYCVGASL..
JHGBMJ 09345	MNTKTFEQFDVMTDEALSKVEGGYSKTDCLNAMITGIAGGIVAGGTGAGLVTLGVAGLPGAFVGAHIGAIGGCATCVGGMLFN
KALHLD 05610	MNTKTFEQFDVMTDEALSTVEGGKKCNYSQVIFDGFMGAAAGAAGNVPGIVIGGIGGAV..........LSYGGYCL......
MOCIPG 03375	MNTKTFEQFDVMTDEALSTVEGGKKCNYSQVIFDGFMGAAAGAAGNVPGIVIGGIGGAV..........LSYGGYCL......
HDEPEA 06125	..MKNLEQFDVMTSAELSNVEGGKSFFYDFGYWYGSHASLSNLANNLGRMLRS...............
NDNJMB 09570	..MKNLEQFDVMTSAELSNVEGGKSFFYDFGYWYGSHASLSNLANNLGRMLRS...............
KCLKHH 07005	..MKNLEQFDVMTSAELSHVEGGKSFFYDFGYWYGSHASLSNLANNLGRMLRS...............
IKMHFE 06975	MNTKTFEQFDVMTEAELSNVEGGKSFFYDFGYWYGSHASLSNLANNLGRMLRS...............
OLMONE 04600	MNTKTFEQFDVMTETELSTVEGGKSFFYDFGYWYGSHASLSNLANNLGRMLRS...............
consensus	***!**!!!***! **!!*!!!!** ** ***** ** *** *** **** ********** ******** *** *

X non-conserved
X ≥ 10% conserved