

logo

NTDB id 1115 OK783 RS00485 WP 003698035.1
 NTDB id 1100 NMB RS09455 WP 002225652.1
 NTDB id 1092 CAD91899.1 1..2286()
 NTDB id 1009 ACIAD RS15175 WP 004923730.1
 NTDB id 1039 HON27 RS01460 WP 031993779.1
 NTDB id 1052 ABD1 RS15850 WP 001017033.1
 NTDB id 1186 GTF74 RS01320 WP 000788426.1
 NTDB id 1175 A1552VC RS12105 WP 000788426.1
 NTDB id 1408 DSB67 RS13980 WP 010643500.1
 NTDB id 1257 GC085 RS04855 WP 163621000.1
 NTDB id 1284 ACE17W RS12370 WP 010946666.1
 NTDB id 1009993 BIKONL RS07395 WP 010955626.1
 NTDB id 1193 PAKAF RS26665 WP 003161249.1
 consensus

MNTKLT...KTIISGLFVATAAFQTASAGNITDIKVVSSLPNKQKIVKVSFDKEIV.NPTGFTVTSSPARTIALDFEQ 70
 MNTKLT...KTIISGLFVATAAFQTASAGNITDIKVVSSLPNKQKIVKVSFDKEIV.NPTGFTVTSSPARTIALDFEQ 70
 MNTKLT...KTIISGLFVATAAFQTASAGNITDIKVVSSLPNKQKIVKVSFDKEIV.NPTGFTVTSSPARTIALDFEQ 70
 MKQIIRQFSIGTVAYAIMQV.....ASAQVAITNVVPMNIPGQGTEIRVMFNGLPP.QPQAYQLENPSRLILDFDK 70
 MNHVFRQFSMGAVAIAIMQA.....ASAQVSMTNIVPMQIAGQGTEIRVMFNGLPP.QPQAYQLENPSRLILDFDK 70
 MNHVFRQFSMGAVAIAIMQA.....ASAQVSMTNIVPMQIAGQGTEIRVMFNGLPP.QPQAYQLENPSRLILDFDK 70
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 0
 0
M....RKIIAIIIL.ISASLSLAYSKDNSLISVKVLP.LPENRVRIDFQFSKPLKQLPASFITQKPPRLVLDLFIS 69
M....RKIIAIIIL.ISASLSMAYSKDNSLISVKVLP.LPENRVRIDFQFSKPLKQLPASFITQKPPRLVLDLFIS 69
 0
 MNSGLS....RLGIALIIAA.MFA....PALLAADLEKLDVAA.LPGDRVELKLFQFDEPVAA.PRGYTIEQPARTIALDLPG 69
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 consensus

TGISMDQ..QVLEYADPLLSKISAAQNSSRARLVNLNKPQQYNTVEVRGNKVVIFINESDDTVSAPARPAVKAAPAAPAK 148
 TGISMDQ..QVLEYADPLLSKISAAQNSSRARLVNLNKPQQYNTVEVRGNKVVIFINESDDTVSAPARPAVKAAPAAPAK 148
 TGISMDQ..QVLEYADPLLSKISAAQNSSRARLVNLNKPQQYNTVEVRGNKVVIFINESDDTVSAPARPAVKAAPAAPAK 148
 AGQKLAKS..LIPVTTNEASSIDVSSDDKRRVNVNLDKAGAFTRVEGCVNFILKINNIQSTAT.....P..... 133
 AQQGLKQS..KISVATNEASSVDVTSDDQRSRLTVNLDKAGAFTRVEGNTFILKINSAQTSNK.....P..... 133
 AQQGLKQS..KISVATNEASSVDVTSDDQRSRLTVNLDKAGAFTRVEGNTFILKINSAQTSNK.....P..... 133
MKNGLKT..YVAQTWLTLLWVGLA.....L.....C.. 23
MKNGLKT..YVAQTWLTLLWVGLA.....L.....C.. 23
MKMRHGLTLTPRKLTLGLFSLWVGLA.....L.....A.. 27
 TDMQIPSDQKTKTIQLGSLHSYNIVAVGDRVRAILELDRSVSYSGSTAGSVSLVINGKSNLDL.F.....E.. 134
 TDMQIPSDQKTKTIQLGSLHSYNIVAVGDRVRAILELDRSVSYSGSTAGSVSLVINGKSNLDL.F.....E.. 134
 0
 VQNK...GTKNRELSVGNTRSVTVVEAKDRRLIINLTTLSYTTTRVEGNLFFVVGNSPAGASVASTAPVKASPAPA.. 145
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QQAAPSTKSAVSVSEPFPAKQQAAPFTESVSVSAPFSPAKQQA...AASAKQQA...AASAKQTNI 182
 QQAAPSTKSAVSVSEPFPAKQQAAPFTESVSVSAPFSPAKQQA...AASAKQQA...AASAKQTNI 228
 QQAAPSTKSAVSVSEPFPAKQQAAPFTESVSVSAPFSPAKQQA...AASAKQQA...AASAKQTNI 220
V.....ATSAPAPQQGTIGNI 148
L.....PVVS.AQPQGVSN 147
L.....PVVS.AQPQGVSN 147
AS..SMV.....FSAESATANQLENI 42
AS..SMV.....FSAESATANQLENI 42
P...F.....VAAEEVQSNSLKSI 43
NSKEVFI.....TNQVVNAKHEIKRI 155
NSKEVFI.....TNQVVNAKHEIKRI 155
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SYAQPIK.....PKPYVPAGRAIRNI 166
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logo

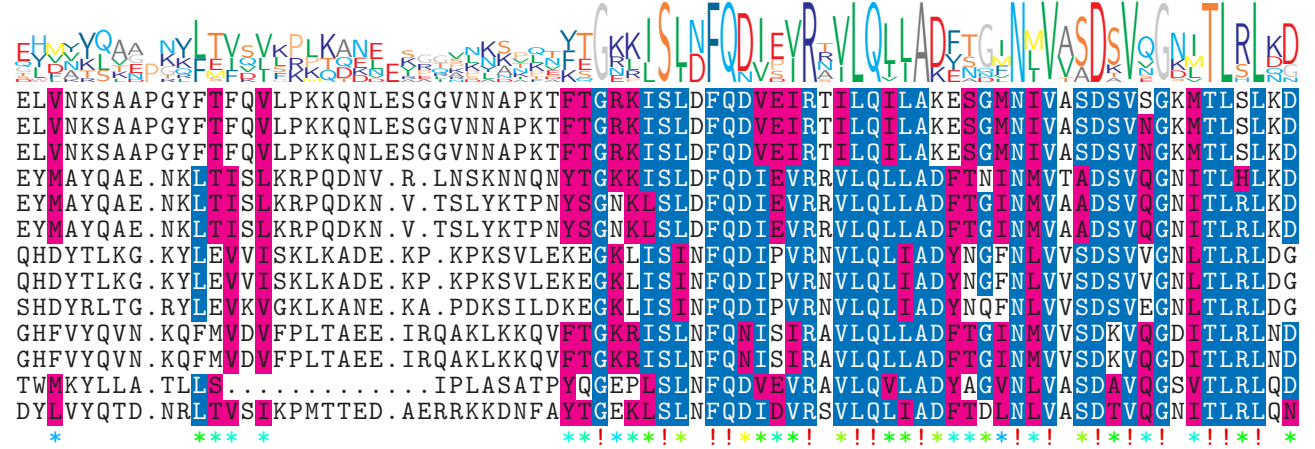
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235
2
246

logo

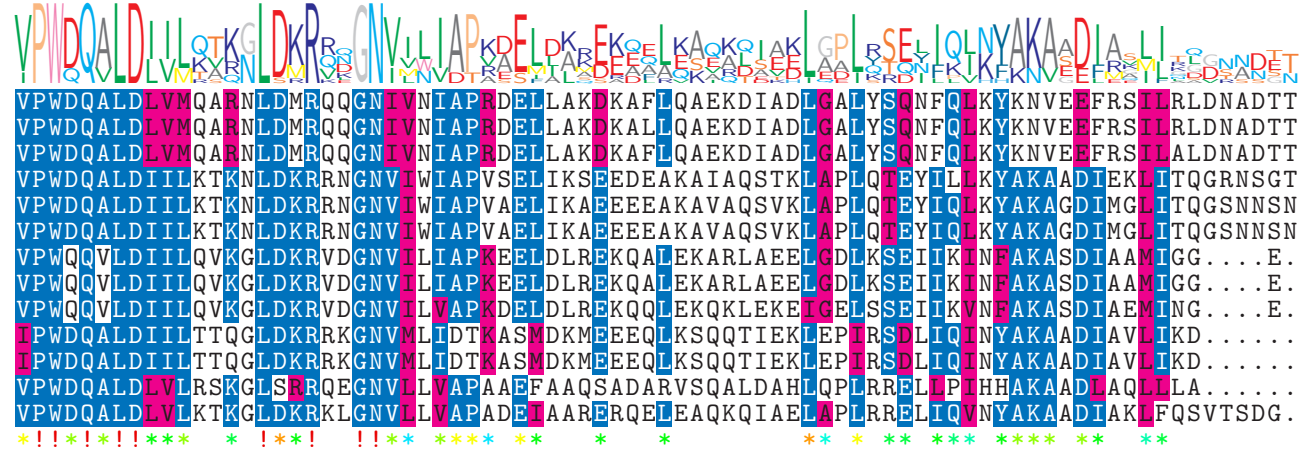
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342
388
380
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304
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199
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200
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422
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