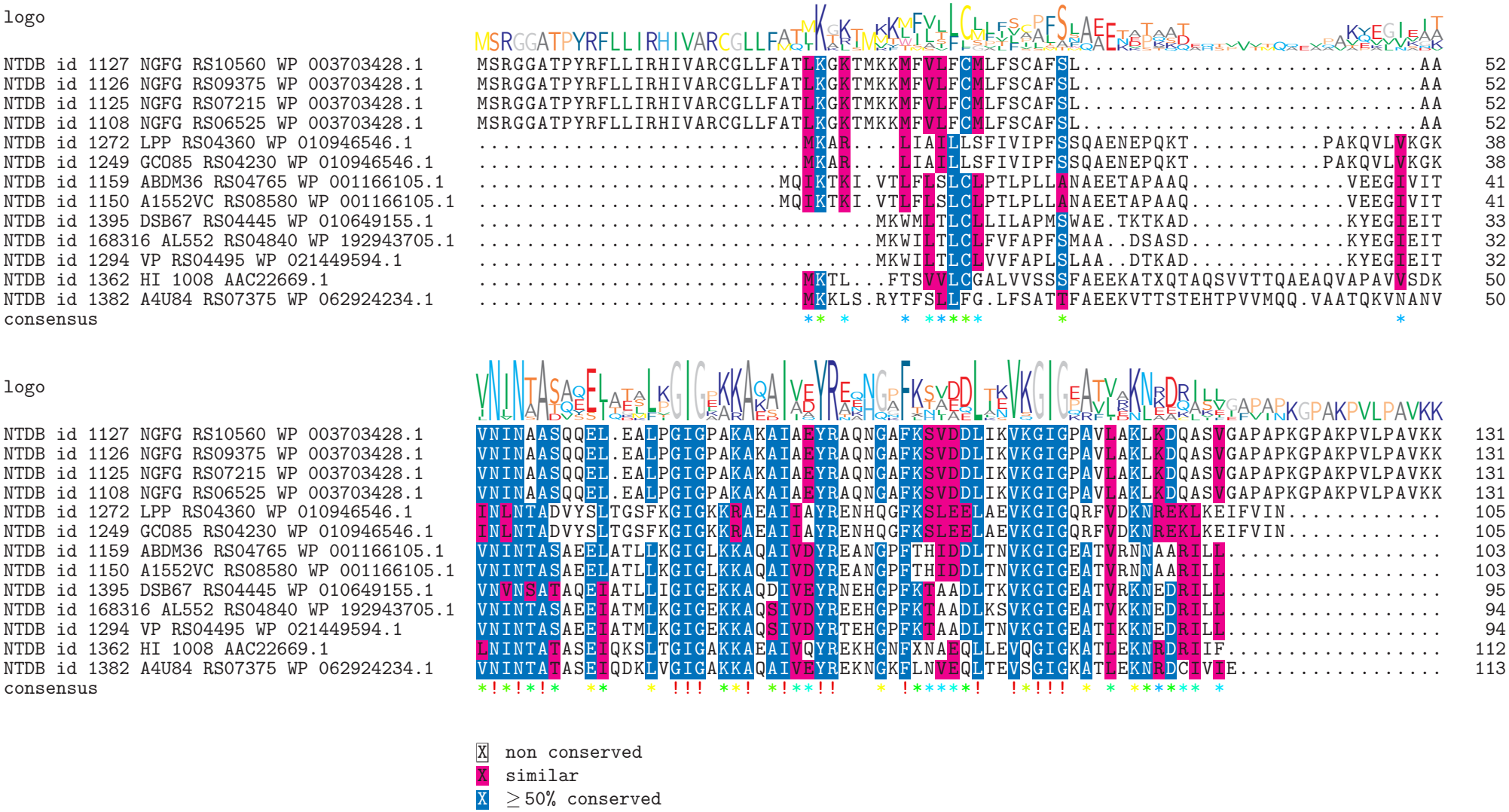
logo
NTDB id 1127 NGFG RS10560 WP 003703428.1 MSRGGATPYRFLLIRHIVARCGLLFATLKGKTMKKMFVLFCMLFSCAFSL.....AA 52
NTDB id 1126 NGFG RS09375 WP 003703428.1 MSRGGATPYRFLLIRHIVARCGLLFATLKGKTMKKMFVLFCMLFSCAFSL.....AA 52
NTDB id 1125 NGFG RS07215 WP 003703428.1 MSRGGATPYRFLLIRHIVARCGLLFATLKGKTMKKMFVLFCMLFSCAFSL.....AA 52
NTDB id 1108 NGFG RS06525 WP 003703428.1 MSRGGATPYRFLLIRHIVARCGLLFATLKGKTMKKMFVLFCMLFSCAFSL.....AA 52
NTDB id 1272 LPP RS04360 WP 010946546.1 .....MKAR....LIAILLSFIVIPFSSQAENEPQKT.....PAKQVLVKGK 38
NTDB id 1249 GCO85 RS04230 WP 010946546.1 .....MKAR....LIAILLSFIVIPFSSQAENEPQKT.....PAKQVLVKGK 38
NTDB id 1159 ABDM36 RS04765 WP 001166105.1 .....MQIKTKI.VTLFLSLCLPTLPLLANAEETAPAAQ.....VEEGIVIT 41
NTDB id 1150 A1552VC RS08580 WP 001166105.1 .....MQIKTKI.VTLFLSLCLPTLPLLANAEETAPAAQ.....VEEGIVIT 41
NTDB id 1395 DSB67 RS04445 WP 010649155.1 .....MKWMLTLCLLILAPMSWAE.TKTKAD.....KYEGIEIT 33
NTDB id 168316 AL552 RS04840 WP 192943705.1 .....MKWILTLCLFVFAPFSMAA..DSASD.....KYEGIEIT 32
NTDB id 1294 VP RS04495 WP 021449594.1 .....MKWILTLCLVVFAPLSLAA..DTKAD.....KYEGIEIT 32
NTDB id 1362 HI 1008 AAC22669.1 .....MKTL...FTSVVLCGALVVSSSFAEEKATXQTAQSVVTTQAEAQVAPAVVSDK 50
NTDB id 1382 A4U84 RS07375 WP 062924234.1 .....MKKLS.RYTFSLLFG.LFSATTFAEEKVTTSTEHTPVVMQQ.VAATQKVNANV 50
consensus
logo
NTDB id 1127 NGFG RS10560 WP 003703428.1 VNINAASQQEL.EALPGIGPAKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK 131
NTDB id 1126 NGFG RS09375 WP 003703428.1 VNINAASQQEL.EALPGIGPAKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK 131
NTDB id 1125 NGFG RS07215 WP 003703428.1 VNINAASQQEL.EALPGIGPAKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK 131
NTDB id 1108 NGFG RS06525 WP 003703428.1 VNINAASQQEL.EALPGIGPAKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK 131
NTDB id 1272 LPP RS04360 WP 010946546.1 INLNTADVYSLTGSFKGIGKKRAEAIIAYRENHQGFKSLEELAEVKGIGQRFVDKNREKLKEIFVIN..... 105
NTDB id 1249 GCO85 RS04230 WP 010946546.1 INLNTADVYSLTGSFKGIGKKRAEAIIAYRENHQGFKSLEELAEVKGIGQRFVDKNREKLKEIFVIN..... 105
NTDB id 1159 ABDM36 RS04765 WP 001166105.1 VNINTASAEELATLLKGIGLKKAQAIVDYREANGPFTHIDDLTNVKGIGEATVRNNAARILL..... 103
NTDB id 1150 A1552VC RS08580 WP 001166105.1 VNINTASAEELATLLKGIGLKKAQAIVDYREANGPFTHIDDLTNVKGIGEATVRNNAARILL..... 103
NTDB id 1395 DSB67 RS04445 WP 010649155.1 VNVNSATAQEIATLLIGIGEKKAQDIVEYRNEHGPFKTAADLTKVKGIGEATVRKNEDRILL..... 95
NTDB id 168316 AL552 RS04840 WP 192943705.1 VNINTASAEEIATMLKGIGEKKAQSIVDYREEHGPFKTAADLKSVKGIGEATVKKNEDRILL..... 94
NTDB id 1294 VP RS04495 WP 021449594.1 VNINTASAEEIATMLKGIGEKKAQSIVDYRTEHGPFKTAADLTNVKGIGEATIKKNEDRILL..... 94
NTDB id 1362 HI 1008 AAC22669.1 LNINTATASEIQKSLTGIGAKKAEAIVQYREKHGNFXNAEQLLEVQGIGKATLEKNRDRIIF..... 112
NTDB id 1382 A4U84 RS07375 WP 062924234.1 VNINTATASEIQDKLVGIGAKKAQAIVEYREKNGKFLNVEQLTEVSGIGKATLEKNRDCIVIE..... 113
consensus
X non conserved
X similar
X ≥ 50% conserved