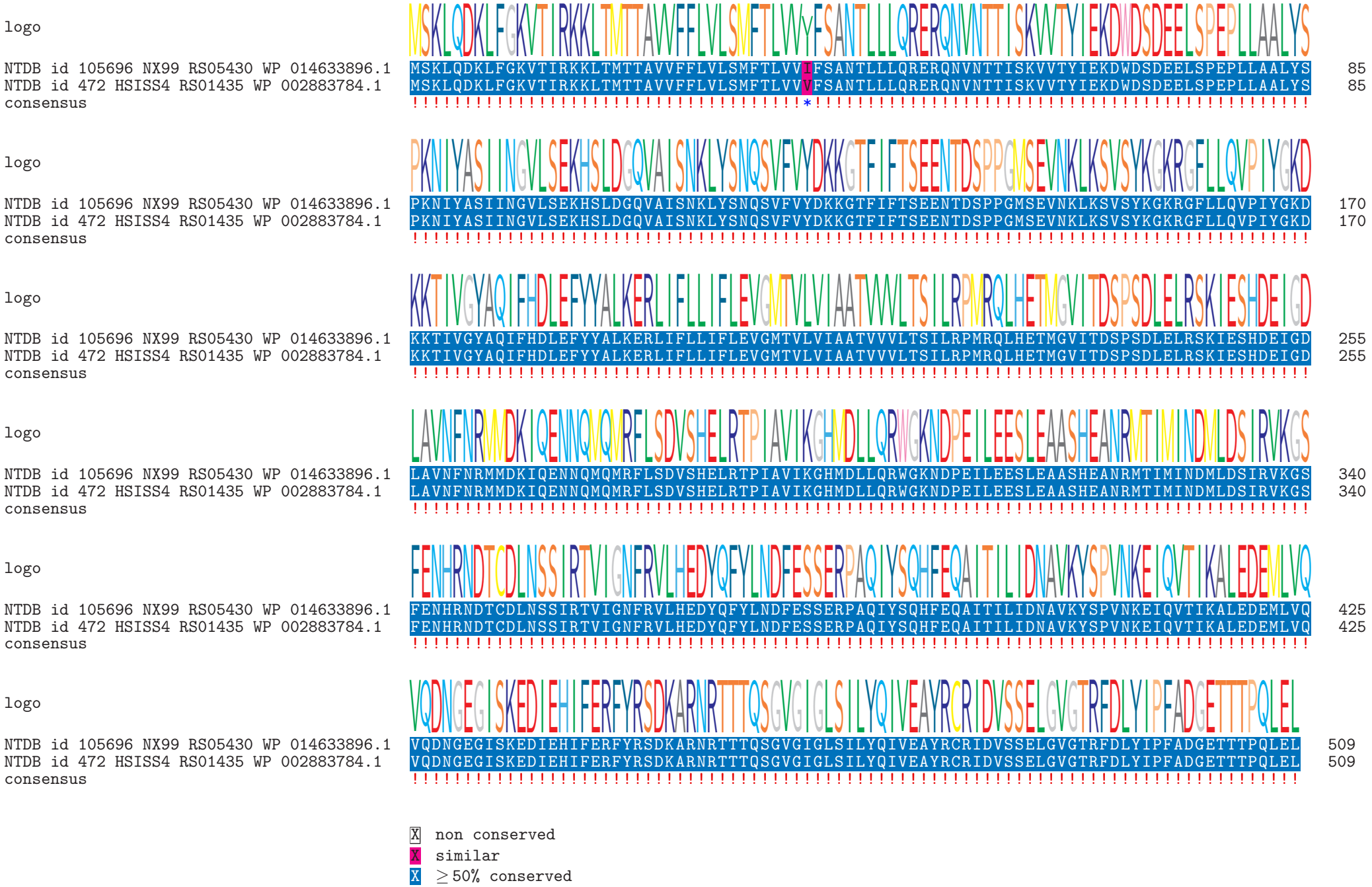

logo
NTDB id 105696 NX99 RS05430 WP 014633896.1 MSKLQDKLFGKVTIRKKLTMTTAVVFFLVLSMFTLVVIFSANTLLLQRERQNVNTTISKVVTYIEKDWDSDEELSPEPLLAALYS 85
NTDB id 472 HSISS4 RS01435 WP 002883784.1 MSKLQDKLFGKVTIRKKLTMTTAVVFFLVLSMFTLVVVFSANTLLLQRERQNVNTTISKVVTYIEKDWDSDEELSPEPLLAALYS 85
consensus
logo
NTDB id 105696 NX99 RS05430 WP 014633896.1 PKNIYASIINGVLSEKHSLDGQVAISNKLYSNQSVFVYDKKGTFIFTSEENTDSPPGMSEVNKLKSVSYKGKRGFLLQVPIYGKD 170
NTDB id 472 HSISS4 RS01435 WP 002883784.1 PKNIYASIINGVLSEKHSLDGQVAISNKLYSNQSVFVYDKKGTFIFTSEENTDSPPGMSEVNKLKSVSYKGKRGFLLQVPIYGKD 170
consensus
logo
NTDB id 105696 NX99 RS05430 WP 014633896.1 KKTIVGYAQIFHDLEFYYALKERLIFLLIFLEVGMTVLVIAATVVVLTSILRPMRQLHETMGVITDSPSDLELRSKIESHDEIGD 255
NTDB id 472 HSISS4 RS01435 WP 002883784.1 KKTIVGYAQIFHDLEFYYALKERLIFLLIFLEVGMTVLVIAATVVVLTSILRPMRQLHETMGVITDSPSDLELRSKIESHDEIGD 255
consensus
logo
NTDB id 105696 NX99 RS05430 WP 014633896.1 LAVNFNRMMDKIQENNQMQMRFLSDVSHELRTPIAVIKGHMDLLQRWGKNDPEILEESLEAASHEANRMTIMINDMLDSIRVKGS 340
NTDB id 472 HSISS4 RS01435 WP 002883784.1 LAVNFNRMMDKIQENNQMQMRFLSDVSHELRTPIAVIKGHMDLLQRWGKNDPEILEESLEAASHEANRMTIMINDMLDSIRVKGS 340
consensus
logo
NTDB id 105696 NX99 RS05430 WP 014633896.1 FENHRNDTCDLNSSIRTVIGNFRVLHEDYQFYLNDFESSERPAQIYSQHFEQAITILIDNAVKYSPVNKEIQVTIKALEDEMLVQ 425
NTDB id 472 HSISS4 RS01435 WP 002883784.1 FENHRNDTCDLNSSIRTVIGNFRVLHEDYQFYLNDFESSERPAQIYSQHFEQAITILIDNAVKYSPVNKEIQVTIKALEDEMLVQ 425
consensus
logo
NTDB id 105696 NX99 RS05430 WP 014633896.1 VQDNGEGISKEDIEHIFERFYRSDKARNRTTTQSGVGIGLSILYQIVEAYRCRIDVSSELGVGTRFDLYIPFADGETTTPQLEL 509
NTDB id 472 HSISS4 RS01435 WP 002883784.1 VQDNGEGISKEDIEHIFERFYRSDKARNRTTTQSGVGIGLSILYQIVEAYRCRIDVSSELGVGTRFDLYIPFADGETTTPQLEL 509
consensus
X non conserved
X similar
X ≥ 50% conserved