A multiple sequence alignment of PE-PGRS family protein PE_PGRS14 from various Mycobacterium tuberculosis strains, showing a sequence logo at the top and a consensus sequence at the bottom.

Sequence logo (consensus): GGNGGAGGAASGVGNGGVGGAGGAGGALVAIGGAGGAGGAATTGTGGAGGAGSNALGLFLGLGGSGGQGGDSAMGSGGAGG

Accession	Description
KBS91563.1	PE-PGRS family protein PE_PGRS14, partial [Mycobacterium tuberculosis UT0117]
KBM14237.1	PE-PGRS family protein PE_PGRS14, partial [Mycobacterium tuberculosis UT0115]
KEA82016.1	PE-PGRS family protein PE_PGRS14 [Mycobacterium tuberculosis TKK_05SA_0024]
KBG18801.1	PE-PGRS family protein PE_PGRS14 [Mycobacterium tuberculosis]
KBD66208.1	PE-PGRS family protein PE_PGRS14 [Mycobacterium tuberculosis M2249]
KAR10586.1	PE-PGRS family protein PE_PGRS14 [Mycobacterium tuberculosis MD5758]
KBV42412.1	PE-PGRS family protein PE_PGRS14 [Mycobacterium tuberculosis NRITLD38]
KAN19269.1	PE-PGRS family protein PE_PGRS14 [Mycobacterium tuberculosis 2228BH]
KAT38388.1	PE-PGRS family protein PE_PGRS14 [Mycobacterium tuberculosis TKK_04_0059]
KEA70737.1	PE-PGRS family protein PE_PGRS14 [Mycobacterium tuberculosis 2104HD]
KEB05078.1	PE-PGRS family protein PE_PGRS14 [Mycobacterium tuberculosis 2995AB]
KAN96224.1	PE-PGRS family protein PE_PGRS14 [Mycobacterium tuberculosis Mr 4387]
KBK18923.1	PE-PGRS family protein PE_PGRS14 [Mycobacterium tuberculosis str. Erdman = ATCC 35801]
KBJ31846.1	PE-PGRS family protein PE_PGRS14 [Mycobacterium tuberculosis str. Haarlem]
KBH36450.1	PE-PGRS family protein PE_PGRS14 [Mycobacterium tuberculosis MAL010136]
KBS62290.1	PE-PGRS family protein PE_PGRS14 [Mycobacterium tuberculosis XTB13-253]
KAN12228.1	PE-PGRS family protein PE_PGRS14 [Mycobacterium tuberculosis 3067CJ]
KBU10916.1	PE-PGRS family protein PE_PGRS14 [Mycobacterium tuberculosis TKK_04_0155]
KBF55063.1	PE-PGRS family protein PE_PGRS14 [Mycobacterium tuberculosis T85]
KBN02133.1	PE-PGRS family protein PE_PGRS14 [Mycobacterium tuberculosis KZN 4207]
AIB47412.1	PE-PGRS family protein PE_PGRS14 [Mycobacterium tuberculosis K]
KCB14797.1	PE-PGRS family protein PE_PGRS14 [Mycobacterium tuberculosis TKK-01-0092]
KBH24576.1	PE-PGRS family protein PE_PGRS14 [Mycobacterium tuberculosis variant africanum MAL010131]
KBH18713.1	PE-PGRS family protein PE_PGRS14 [Mycobacterium tuberculosis variant africanum MAL010129]
KCE57703.1	PE-PGRS family protein PE_PGRS14 [Mycobacterium tuberculosis TKK_04_0007]
KAL00947.1	PE-PGRS family protein PE_PGRS14 [Mycobacterium tuberculosis SK-B]
KCH60069.1	PE-PGRS family protein PE_PGRS14 [Mycobacterium tuberculosis UG-D]
KBN24787.1	PE-PGRS family protein PE_PGRS14 [Mycobacterium tuberculosis 44503]
KBM74503.1	PE-PGRS family protein PE_PGRS14 [Mycobacterium tuberculosis KT-0042]
KCL98484.1	PE-PGRS family protein PE_PGRS14 [Mycobacterium tuberculosis TB_RSA180]
KCR07884.1	PE-PGRS family protein PE_PGRS14 [Mycobacterium tuberculosis BTB12-002]
KAM43940.1	PE-PGRS family protein PE_PGRS14 [Mycobacterium tuberculosis Kc 32216]
KAQ34373.1	PE-PGRS family protein PE_PGRS14 [Mycobacterium tuberculosis MD16257]
KBC87776.1	PE-PGRS family protein PE_PGRS14 [Mycobacterium tuberculosis M2248]
KBJ42541.1	PE-PGRS family protein PE_PGRS14 [Mycobacterium tuberculosis OFXR-1]
KBF55769.1	PE-PGRS family protein PE_PGRS14 [Mycobacterium tuberculosis EAS054]
KBF18853.1	PE-PGRS family protein PE_PGRS14 [Mycobacterium tuberculosis H2398]
KAM38315.1	PE-PGRS family protein PE_PGRS14 [Mycobacterium tuberculosis 2196MM]
KAN74715.1	PE-PGRS family protein PE_PGRS14 [Mycobacterium tuberculosis Wt 21231]

consensus

GGNGGAGGAASGVGNGGVGGAGGAGGALVAIGGAGGAGGAATTGTGGAGGAGSNALGLFLGLGGSGGQGGDSAMGSGGAGG