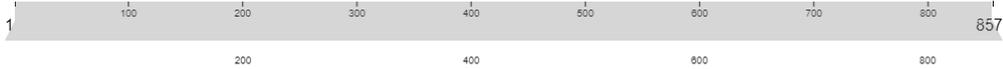
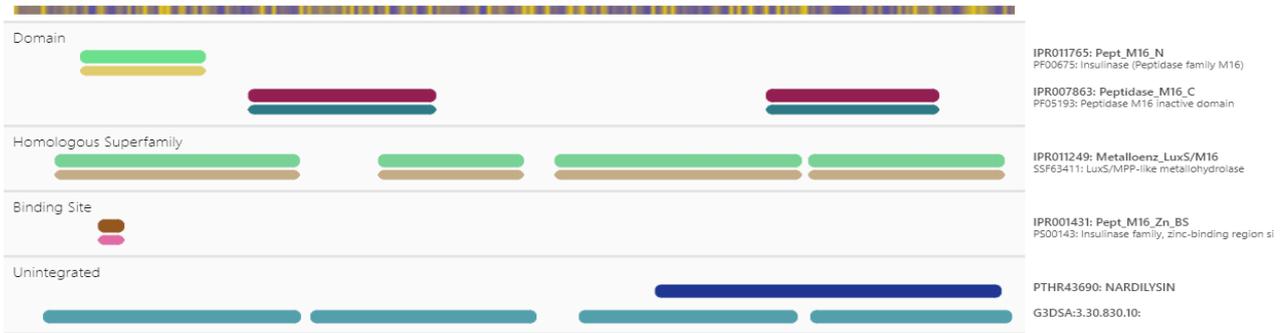
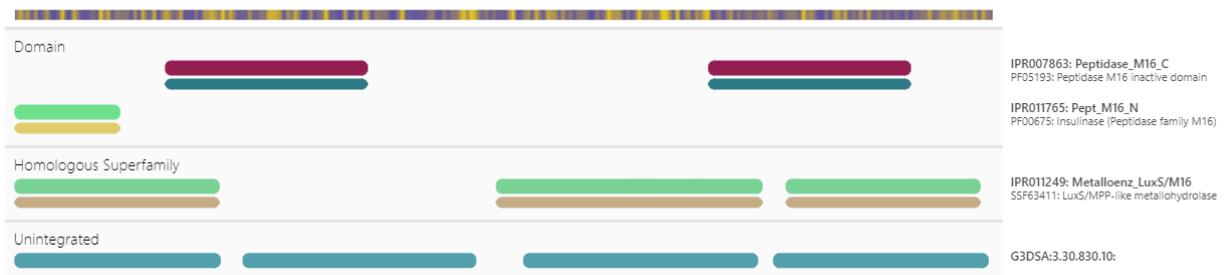




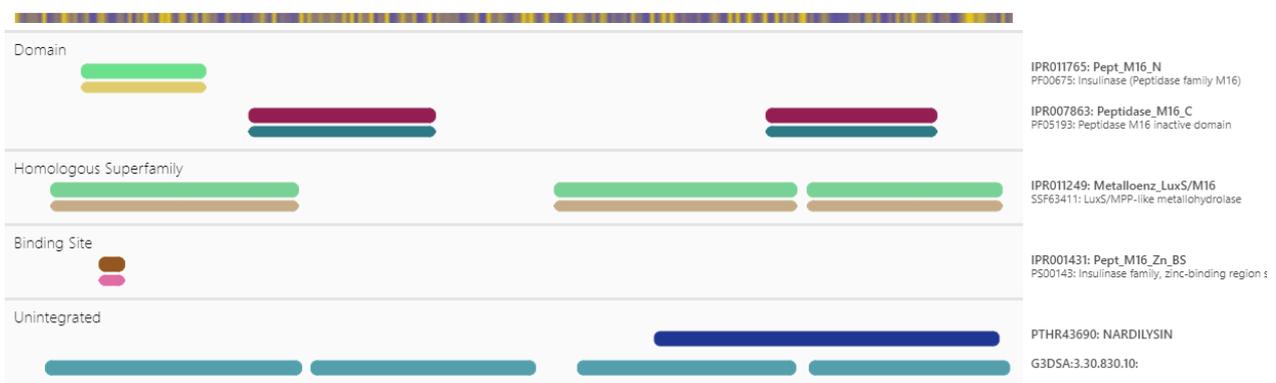
## WSM597

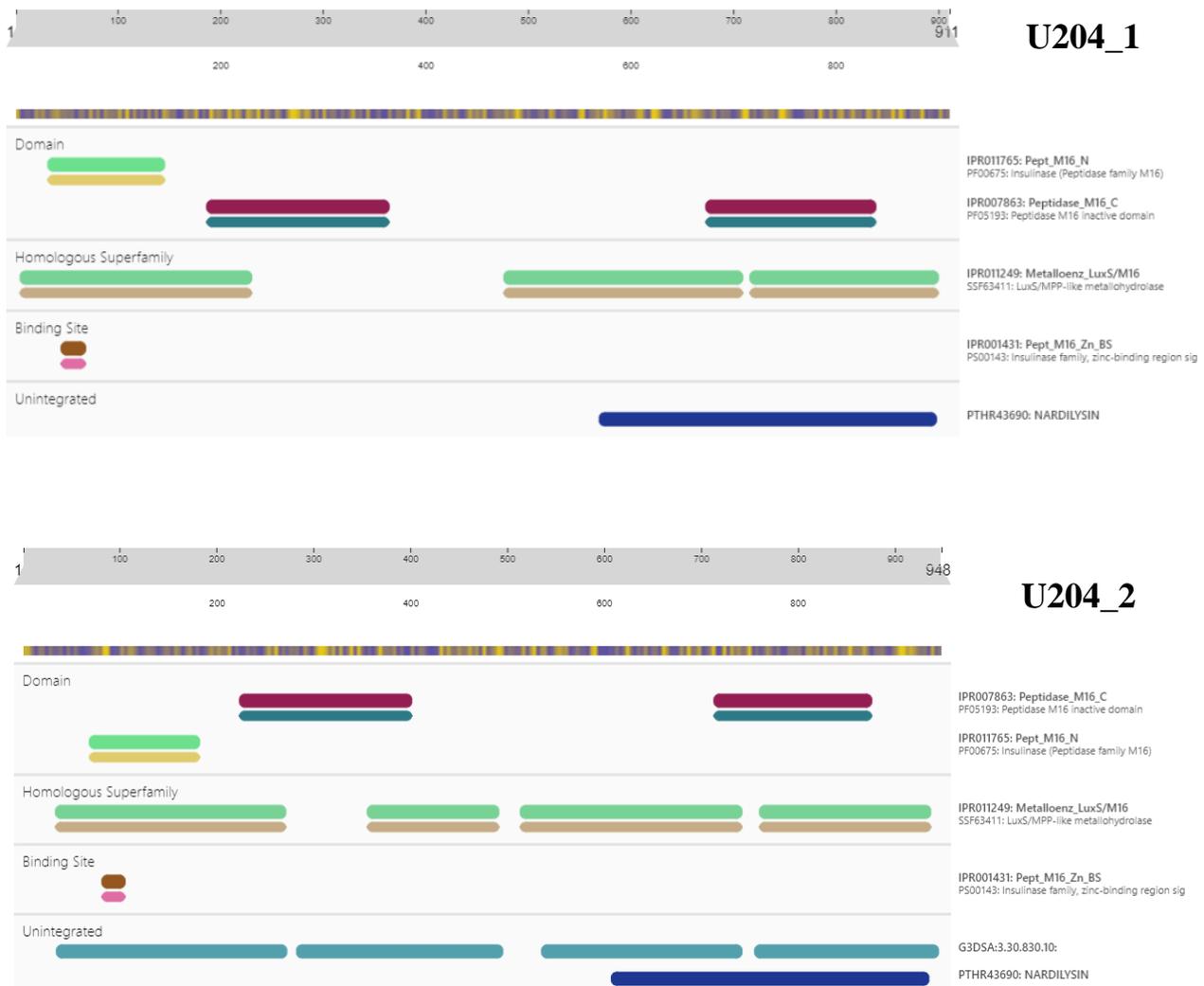


## U268\_1



## U268\_2





**Figure S5.** Graphical view of The InterPro search and protein classification of M16 peptidase sequences found in the genome of strains U268, U204 and WSM 597 using InterPro database. The InterPro entry was the aminoacidic sequence for the putative M16 peptidases. In the graphical view, matches to the InterPro databases and the relevant positions of these matches on the query sequence are displayed.