

Additional File 16.
Co-localization network diagrams for individual ruminant enzyme classes.

Numbers in parentheses indicate relative gene frequencies per 1 million proteins in 391 ruminant metagenomes. Maximum intervening distance between nodes is 25 genes. CAZy and SulfAtlas classes lacking ruminant MAG co-localizations within the 25 gene limit have been omitted, but remaining classes are presented in the same order as main Figures 7 and 8,

