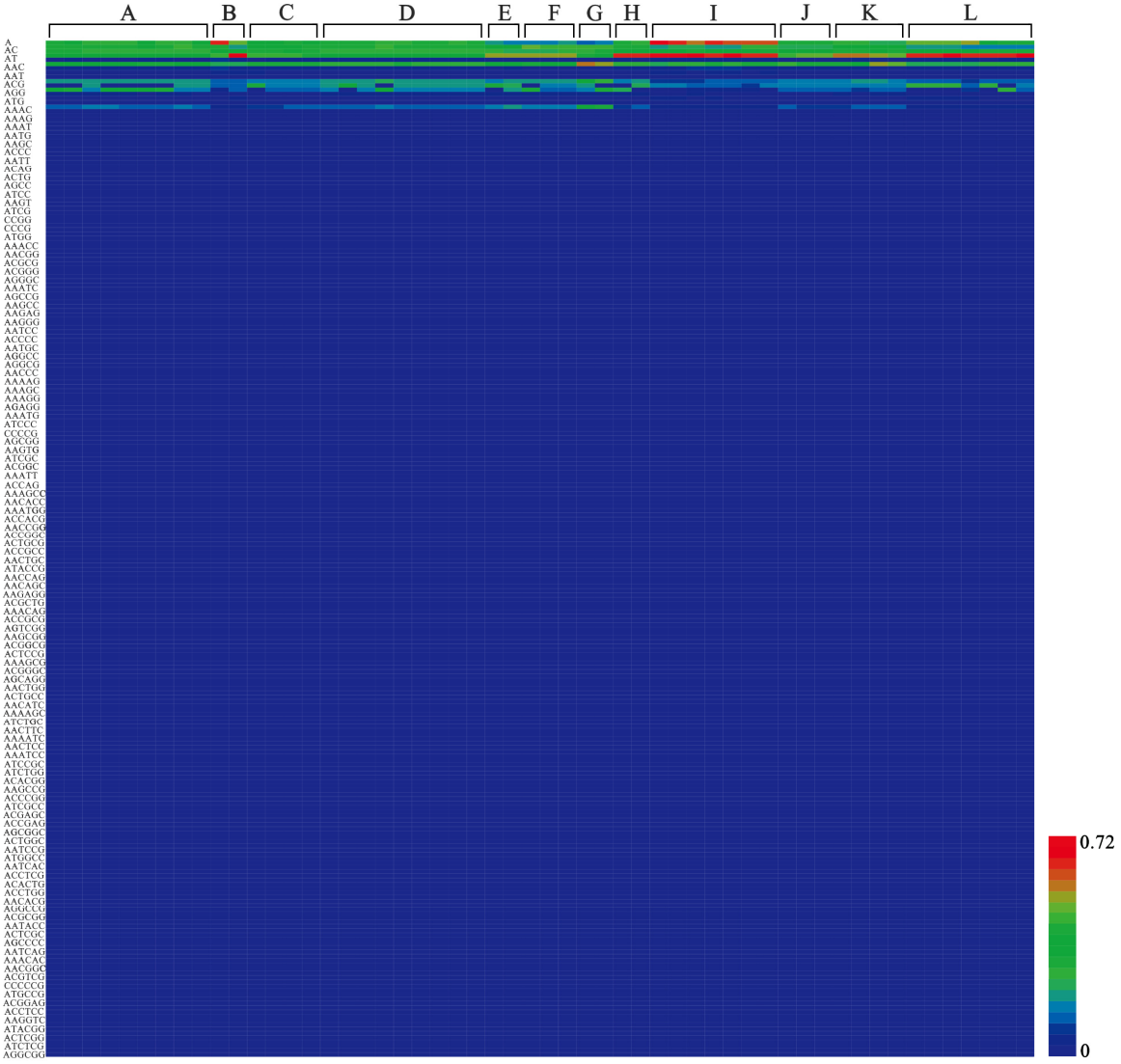

Figure S1. Relative abundance of standard motifs identified in genomes of 54 *Parasynechococcus*-like strains.