

Additional file 1

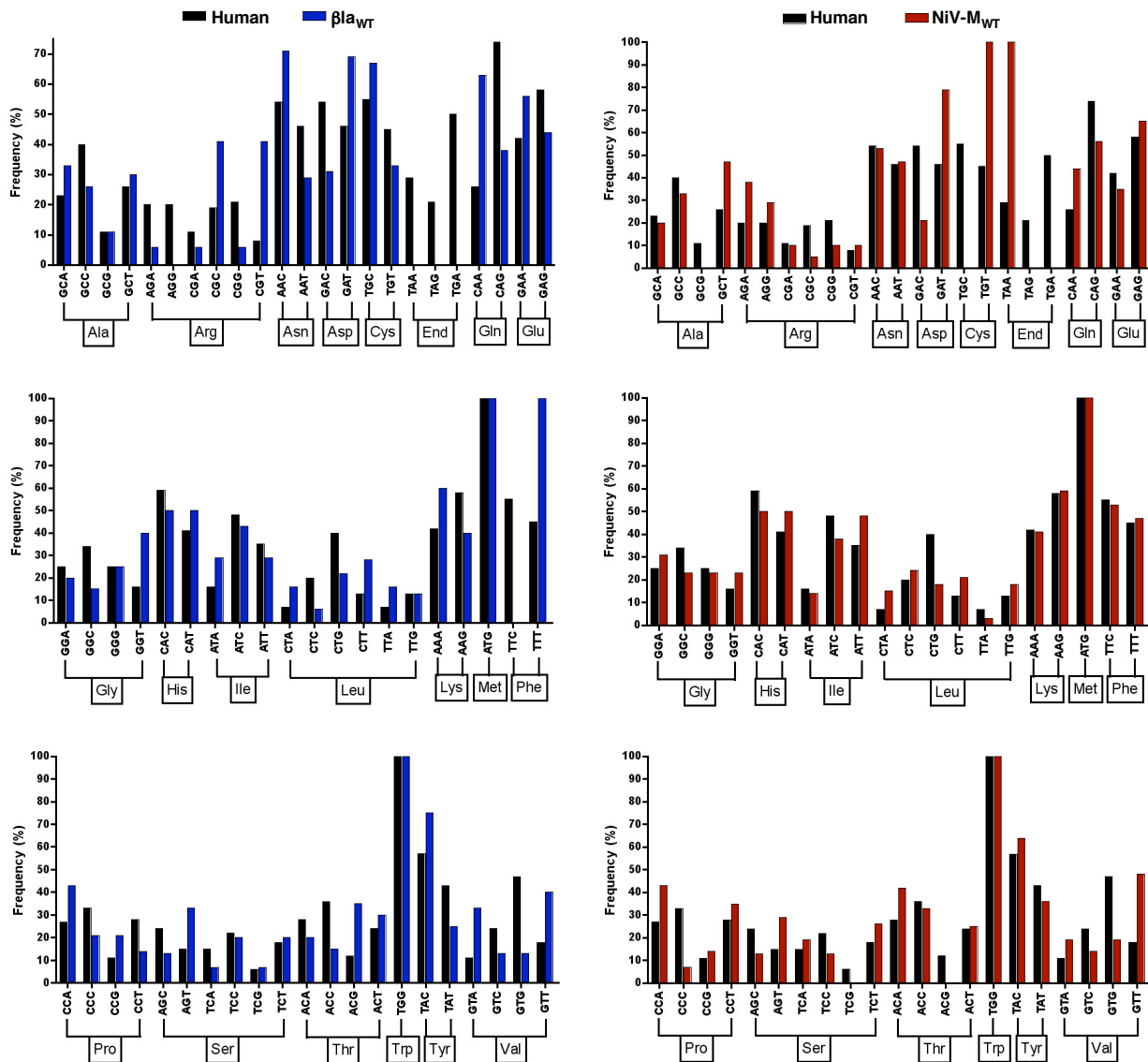


Table 1. Codon usage comparisons between wild-type Nipah matrix (henipavirus), beta-lactamase (bacteria) and average Homo sapiens genes. Note the skewing towards more rarely used mammalian codons. Overall, codon usage for amino acids cumulatively demonstrates a pattern of rare mammalian codon usage.