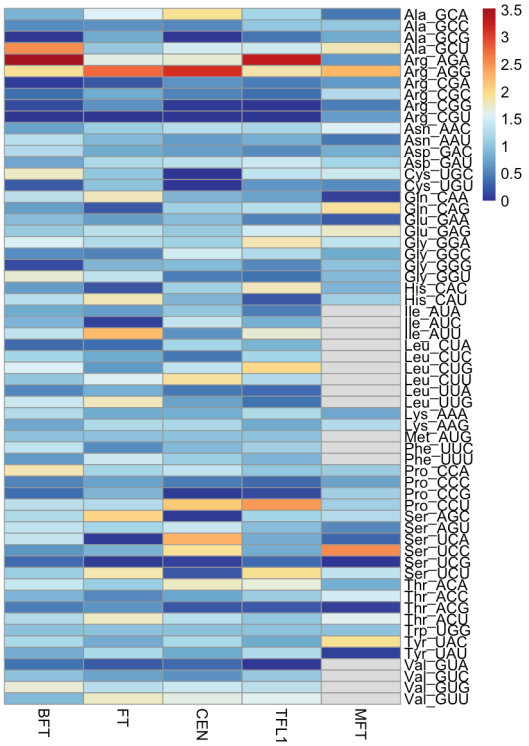Figure S6. Heatmap of average RSCU scores estimated for different codons in five *Rosaceae PEBP* gene lineages. Average codon usage below 1.5 are colored with blue, while codons with RSCU values larger than 2.0 are colored from yellow to red.