

**Figure S4. Percentage ORFs incorrect when AGEs have a consensus start codon coordinate prediction.**

The error-rates for consensus start codon coordinate prediction for four AGEs (Table 1) and of various combinations of these four AGEs are plotted for the four reference organisms. Error-rates were calculated as discussed in materials and methods. A: BASys; B: ISGA; C: RAST and D: xBASE.

