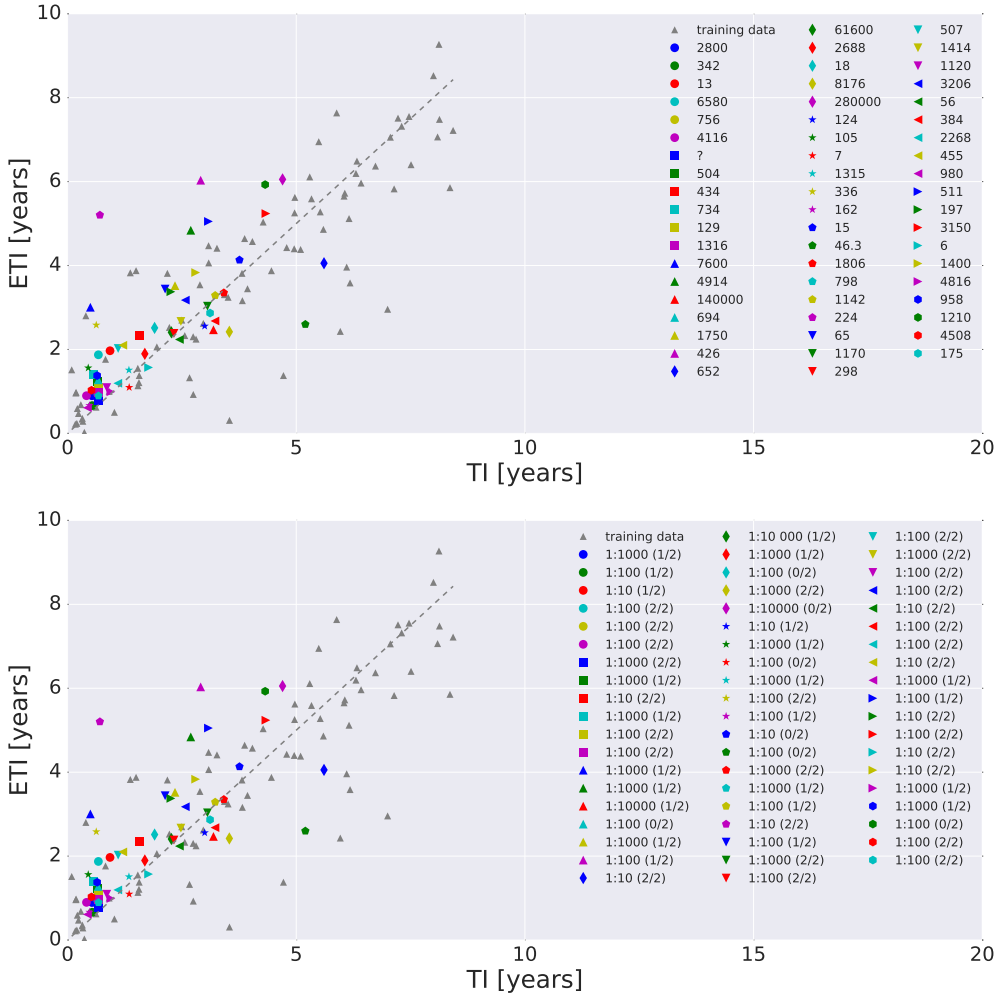Fig S12. Estimated time of infection (ETI) versus actual time of infection (TI). Displayed for the training and the validation data sets. Legend shows the number of templates (top) and the dilutions (bottom) for the validation dataset. (Genetic region: 3rd codon positions in *pol*, diversity measure: average pairwise distance, $x_c = 0.003$.)