S15 Fig. Bayesian Skyline Plots fo the whole HV sequences and for major lineages within HV.
X axis: time in years ago. Y axis: effective population size per generation time. The run for the whole haplogroup HV*(xH, V) is performed with the with and without a partition in coding and non-coding and a relaxed clock, the other runs for single lineages are performed with a strick clock and no partition (see Methods for details).