Supplementary Figure 2. DualBrother recombination detection using the 489bp *ompA* fragment alignment of 26 Chinese *C. suis* strains. The top plot shows marginal posterior probabilities of the four most probable tree topologies, where break and change points of topologies are indicative of recombination. The last two plots show 95% Bayesian confidence interval (shaded in green) of the Kappa transition/transversion ratio, and average divergence Mu.