



S7 Fig. Time-dependent predictor poultry production of effective population dynamics of H9N2 influenza viruses in 12 countries/regions in Asia estimated by 526 sequences. The 50% prior mass was specified on no predictors being included in the GLM. Parameters and figure elements here are the same as in S3 Fig. National poultry production is a strongly supported driver to virus genetic diversity through time in mainland China.