

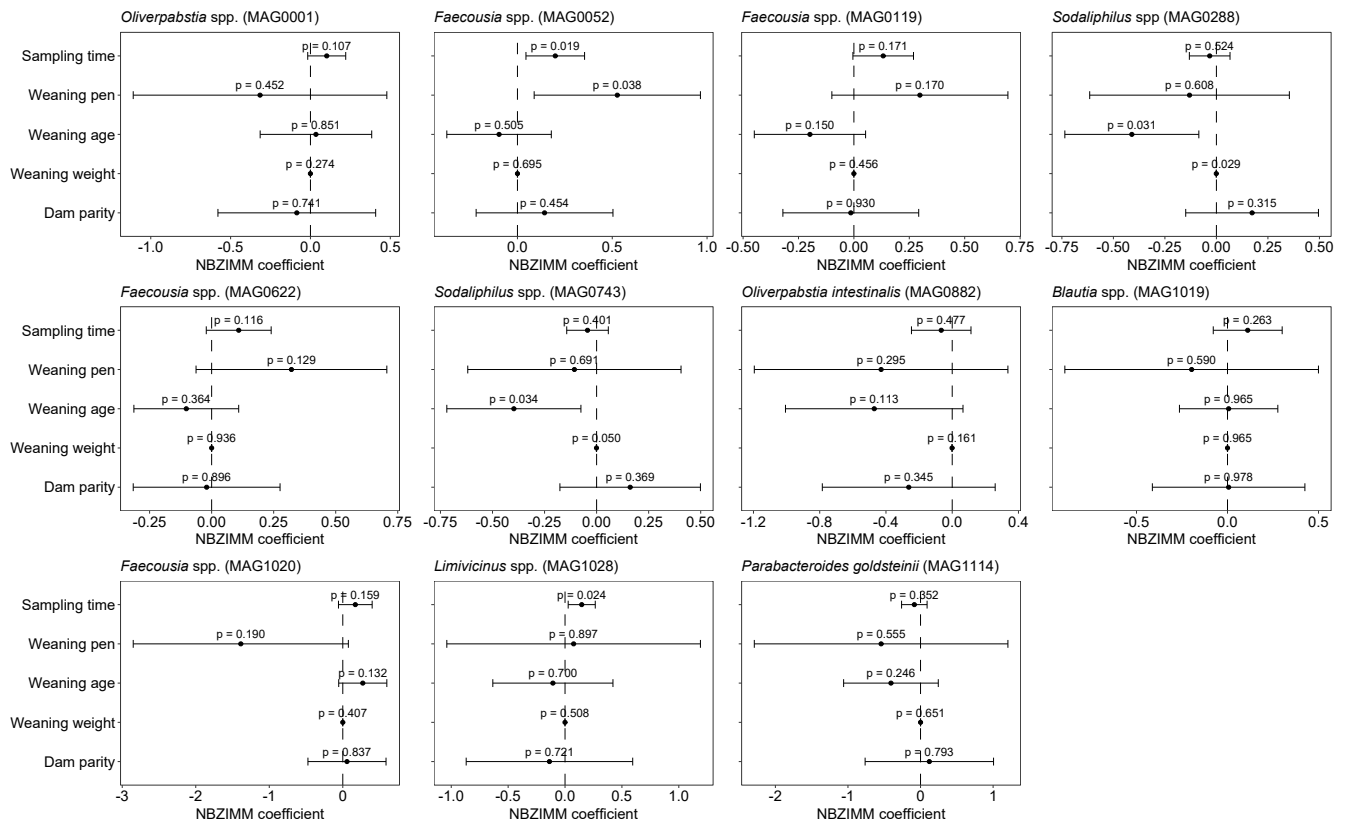


Figure S8. Negative binomial and zero-inflated mixed model (NBZIMM) results on the relative abundance of metagenome-assembled genomes (MAGs) from farm A, showing 95% confidence intervals for all of the fixed effects (coefficients) included. Statistical associations (p-values) are indicated next to each coefficient.