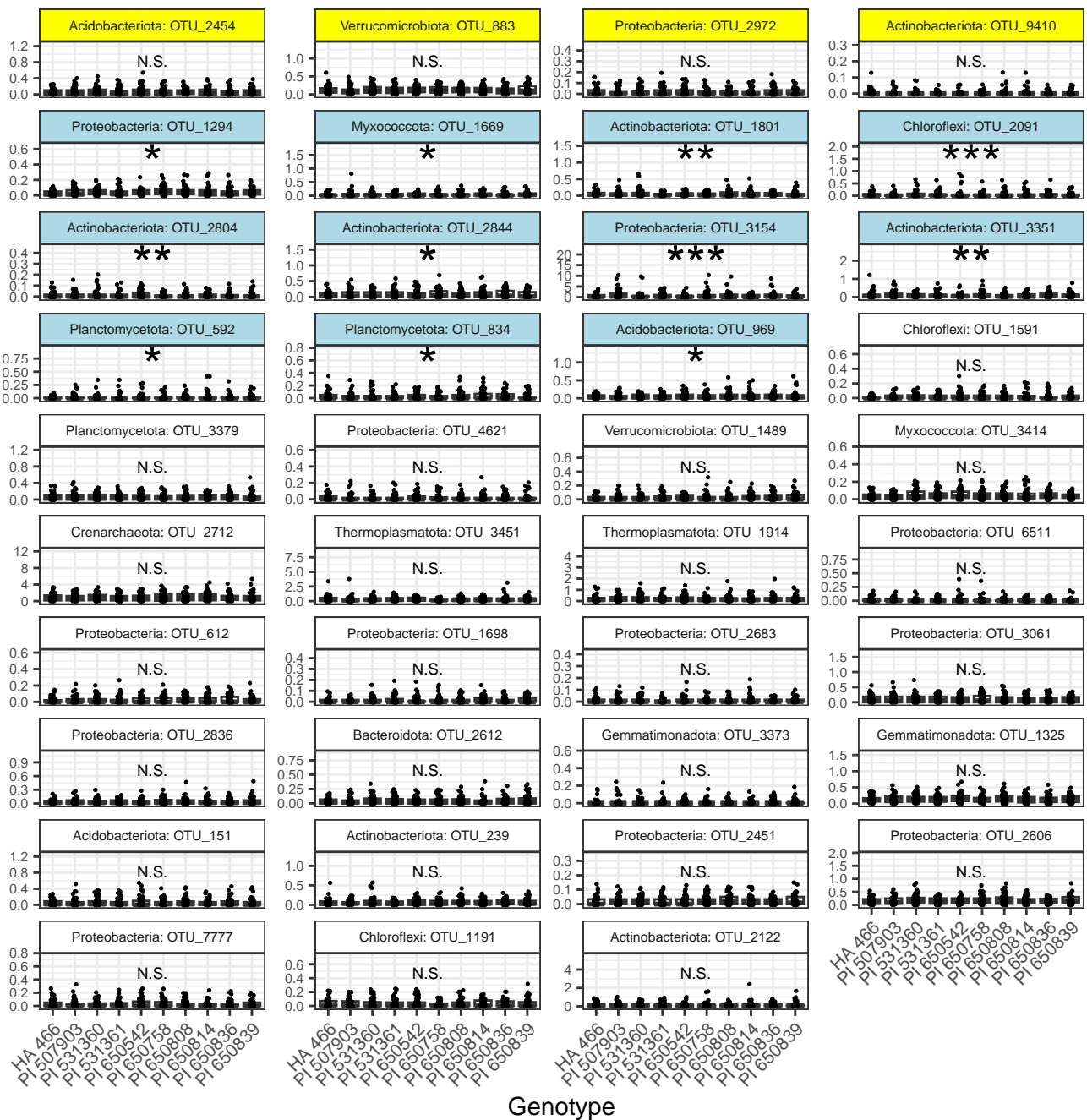


% Relative abundance



Genotype