

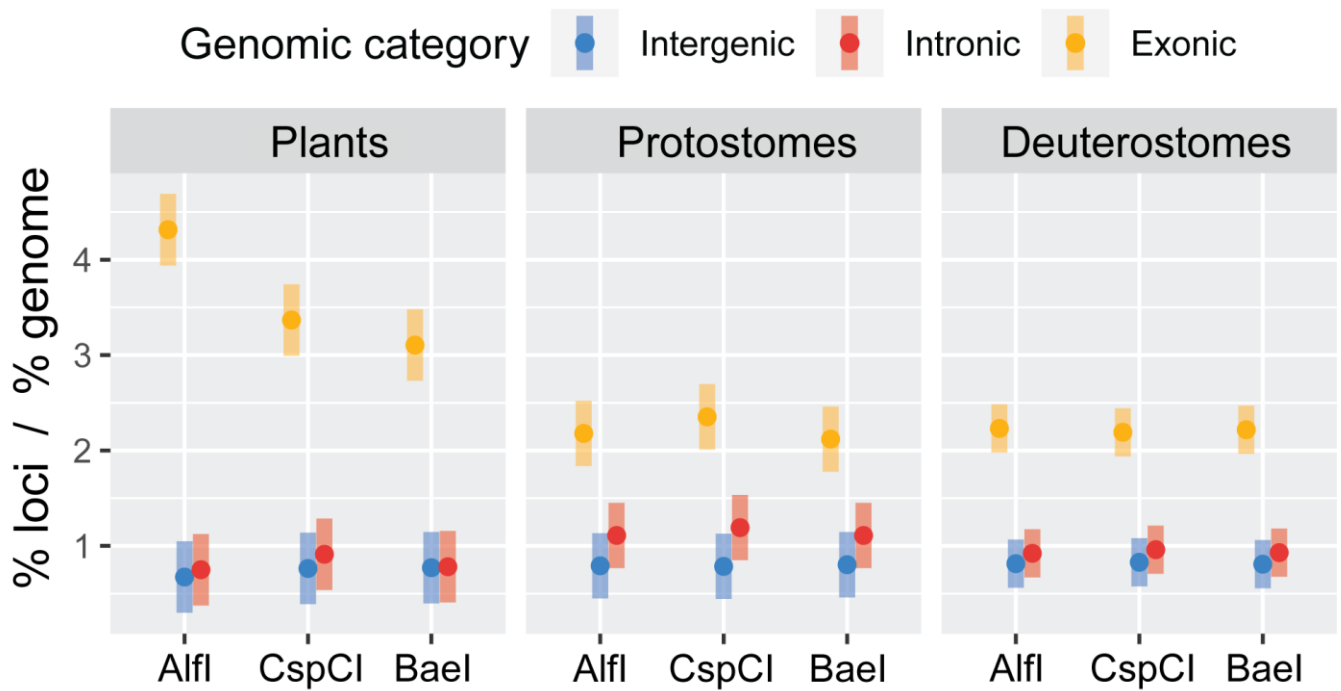


Figure S3: Predicted values of the ratio between the percentage of unique loci in a genomic category and the percentage of genome in the same genomic category with the GLMM provided in Table S13. Mean values are marked with a dot and their 95% confidence intervals are represented with lines.