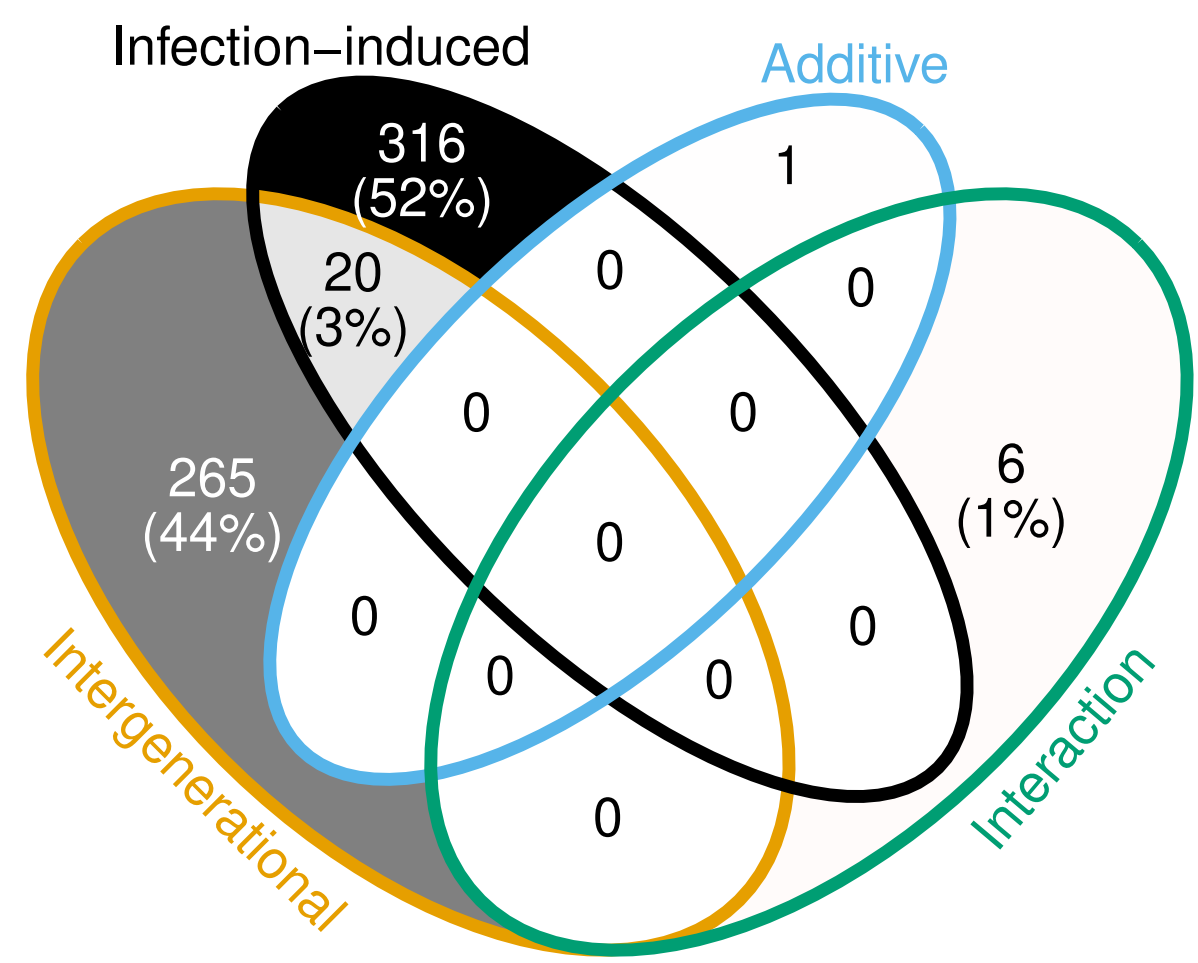
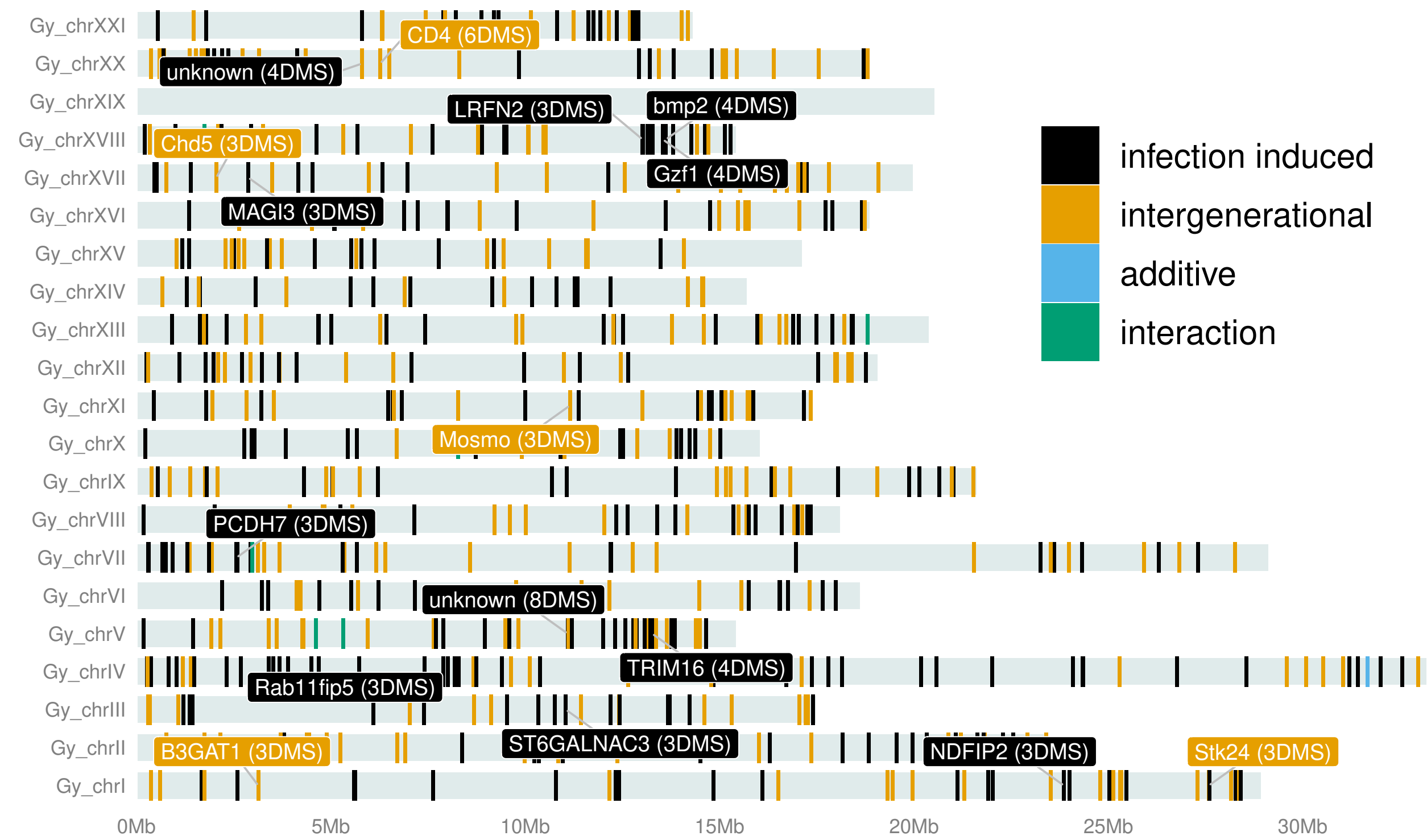


A. Number of genes containing DMS associated with each category



B. Genomic positions of DMS coloured by category



C. Gene ontology enrichment (simplified with GO slim) for infection-induced and intergenerational DMS

