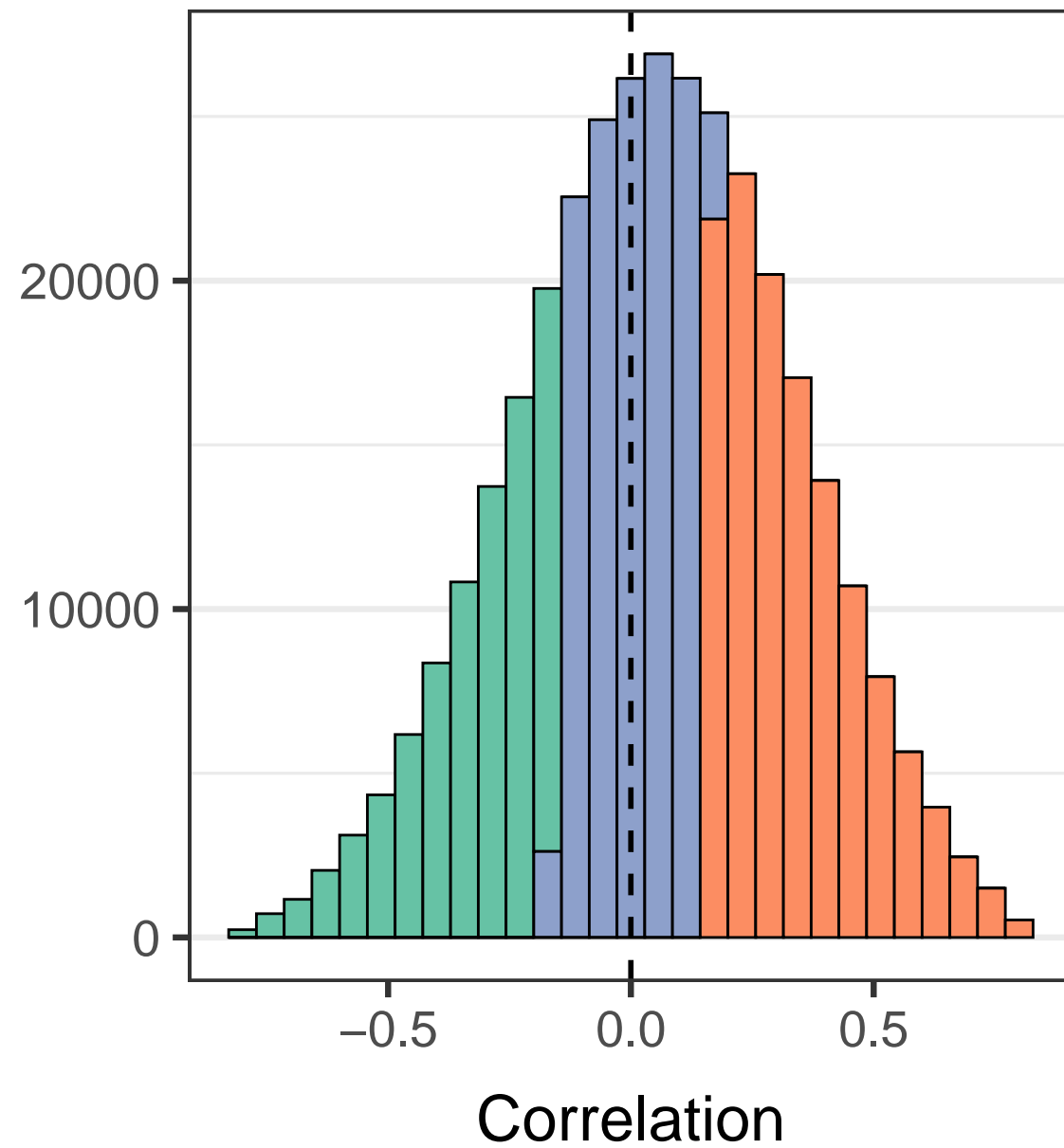


Number of genus pairs

Group    negative    neutral    positive



Model parameters

