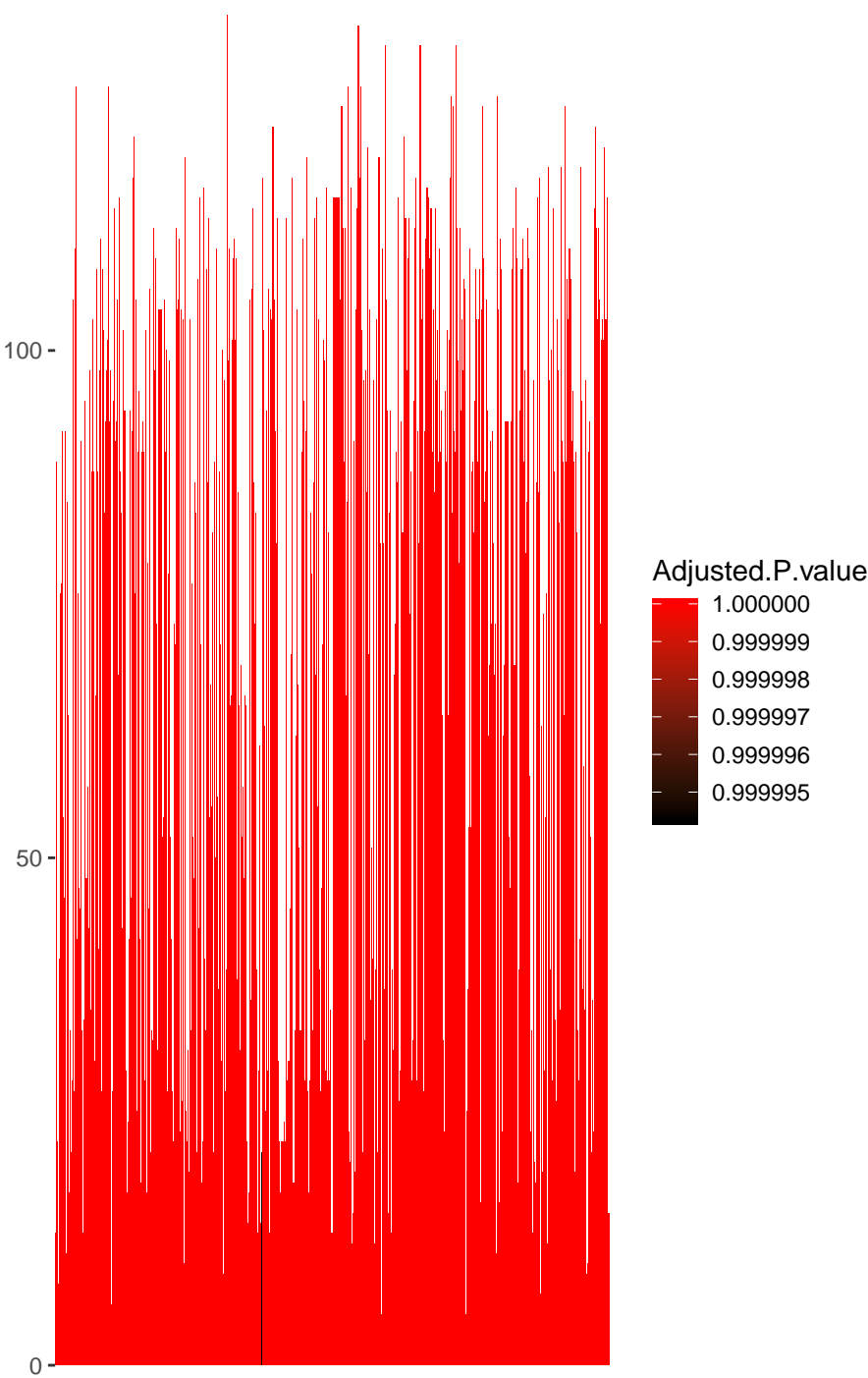


Number of genes overlap



Module names

based on data from enrichR