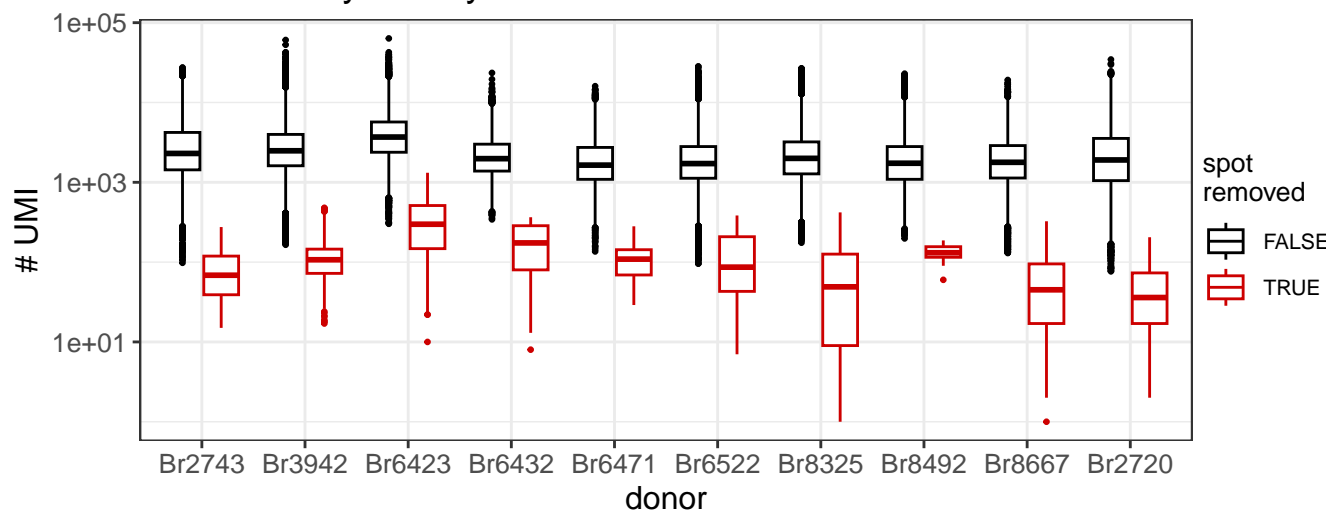
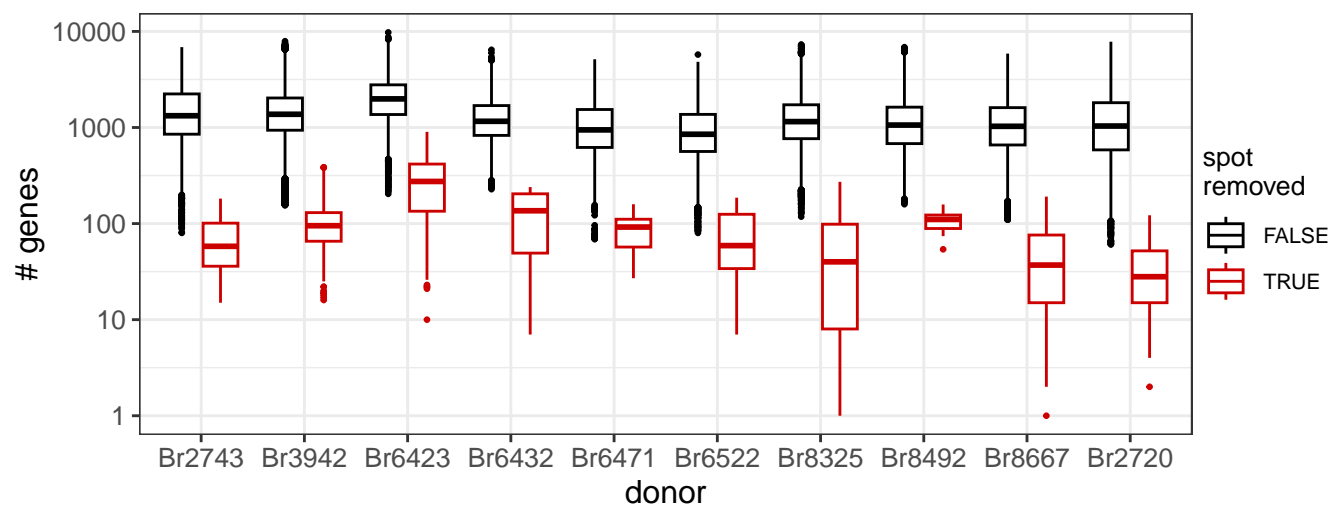


QC summary: library size



QC summary: detected genes



QC summary: spots kept

