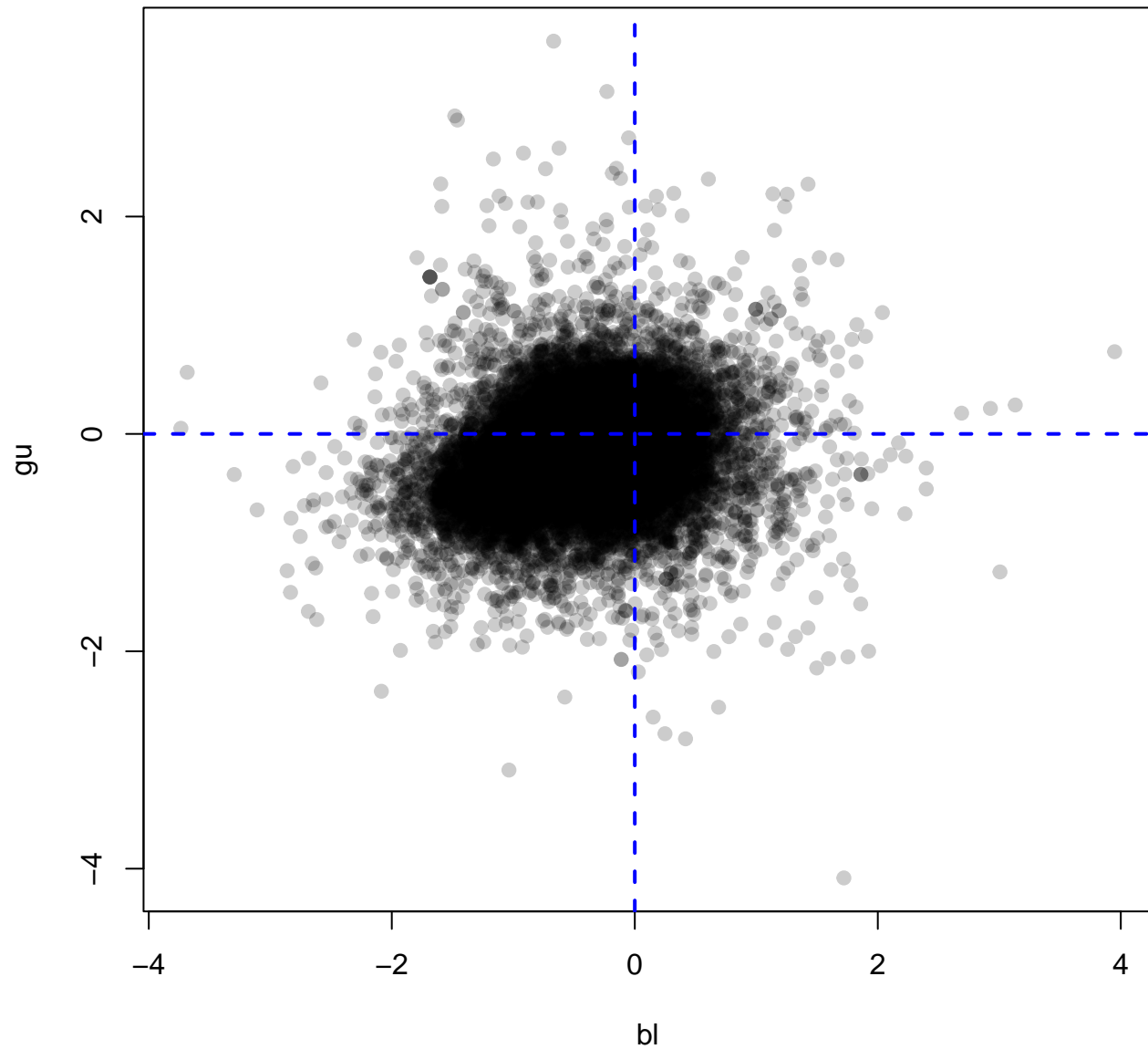
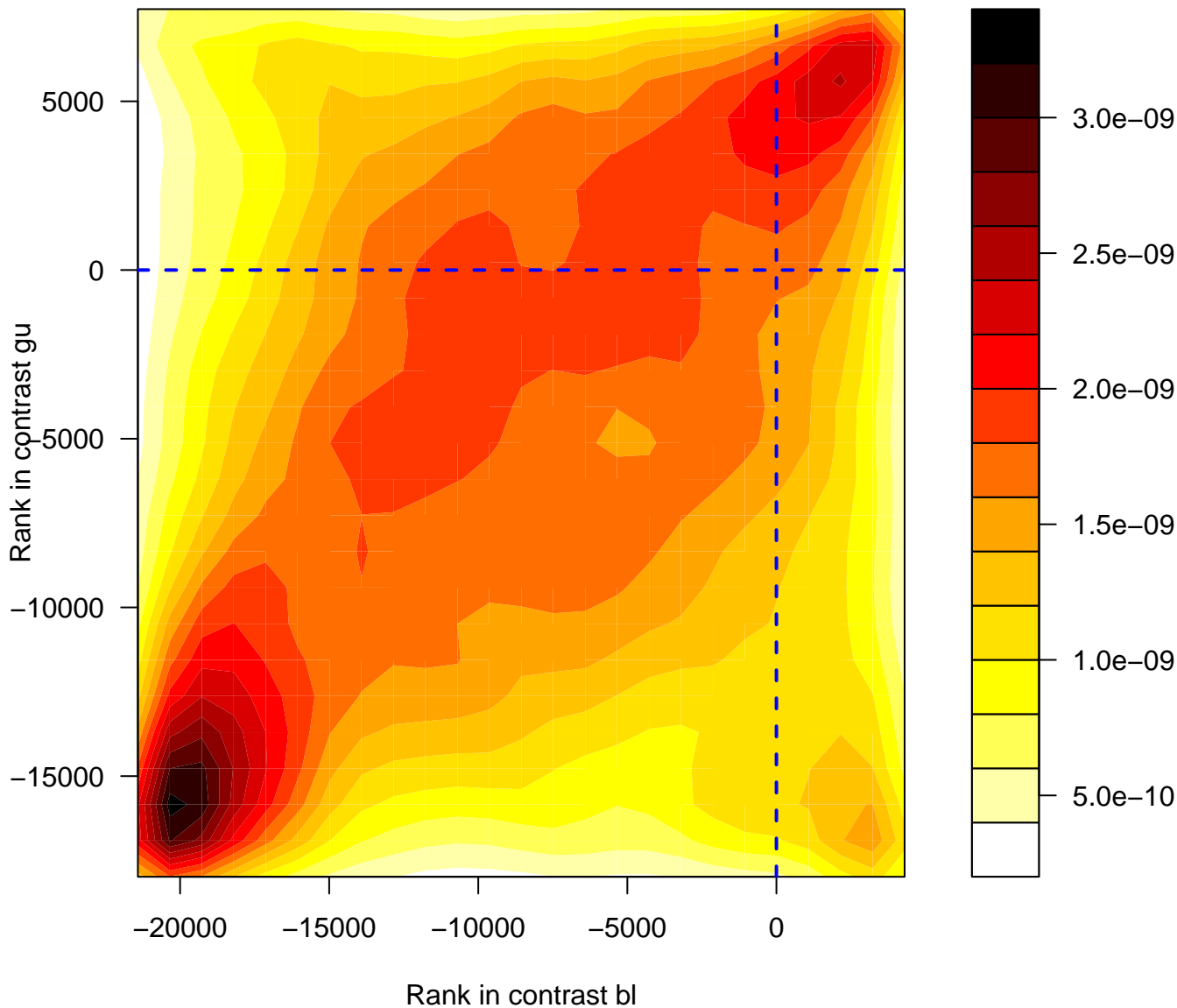


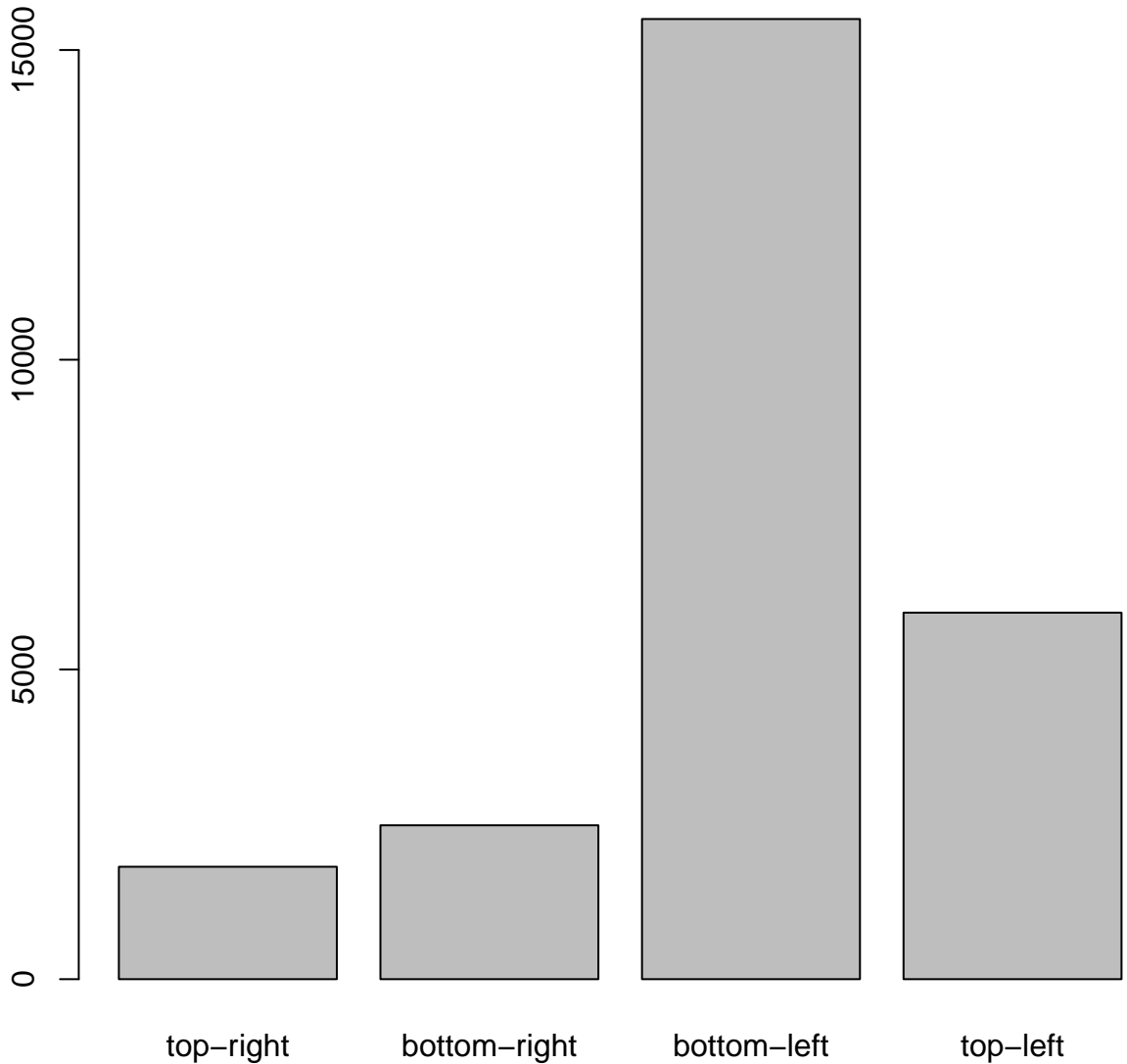
Scatterplot of all genes



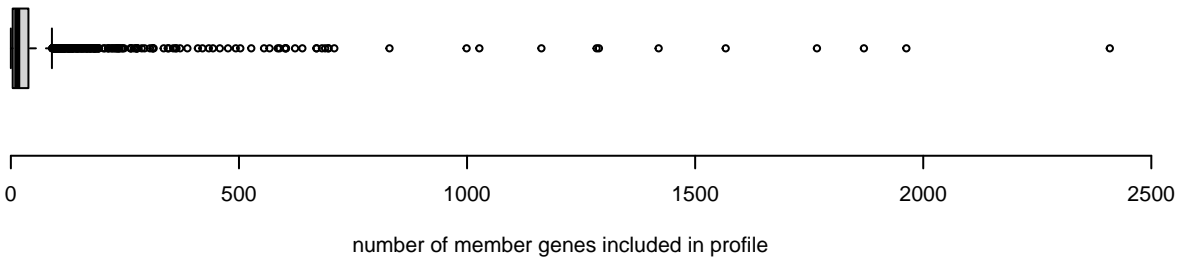
Rank-rank plot of all genes



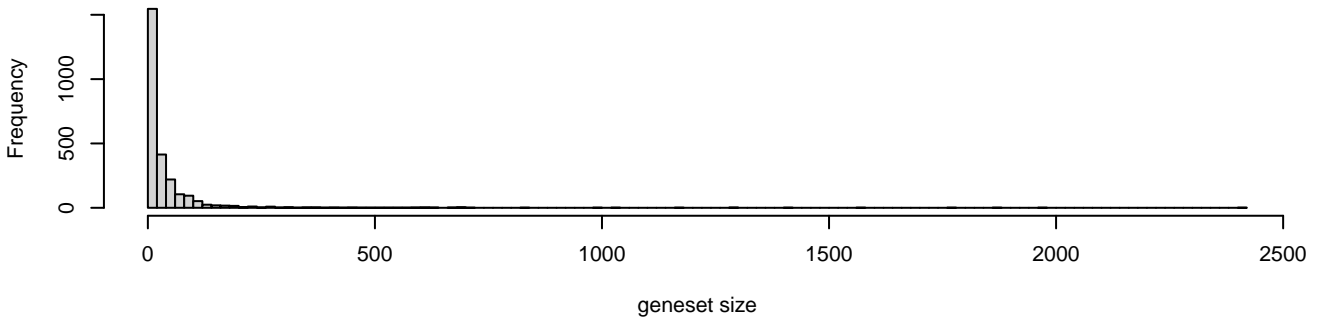
number of genes in each quadrant



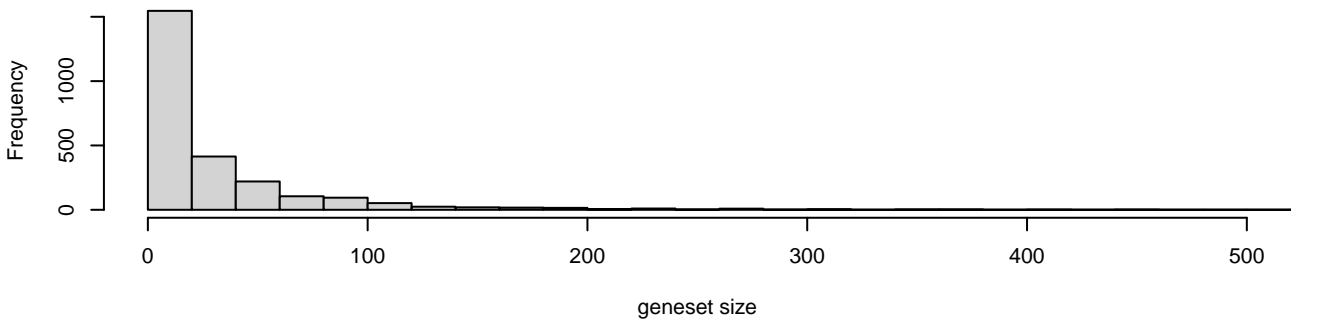
Gene set size



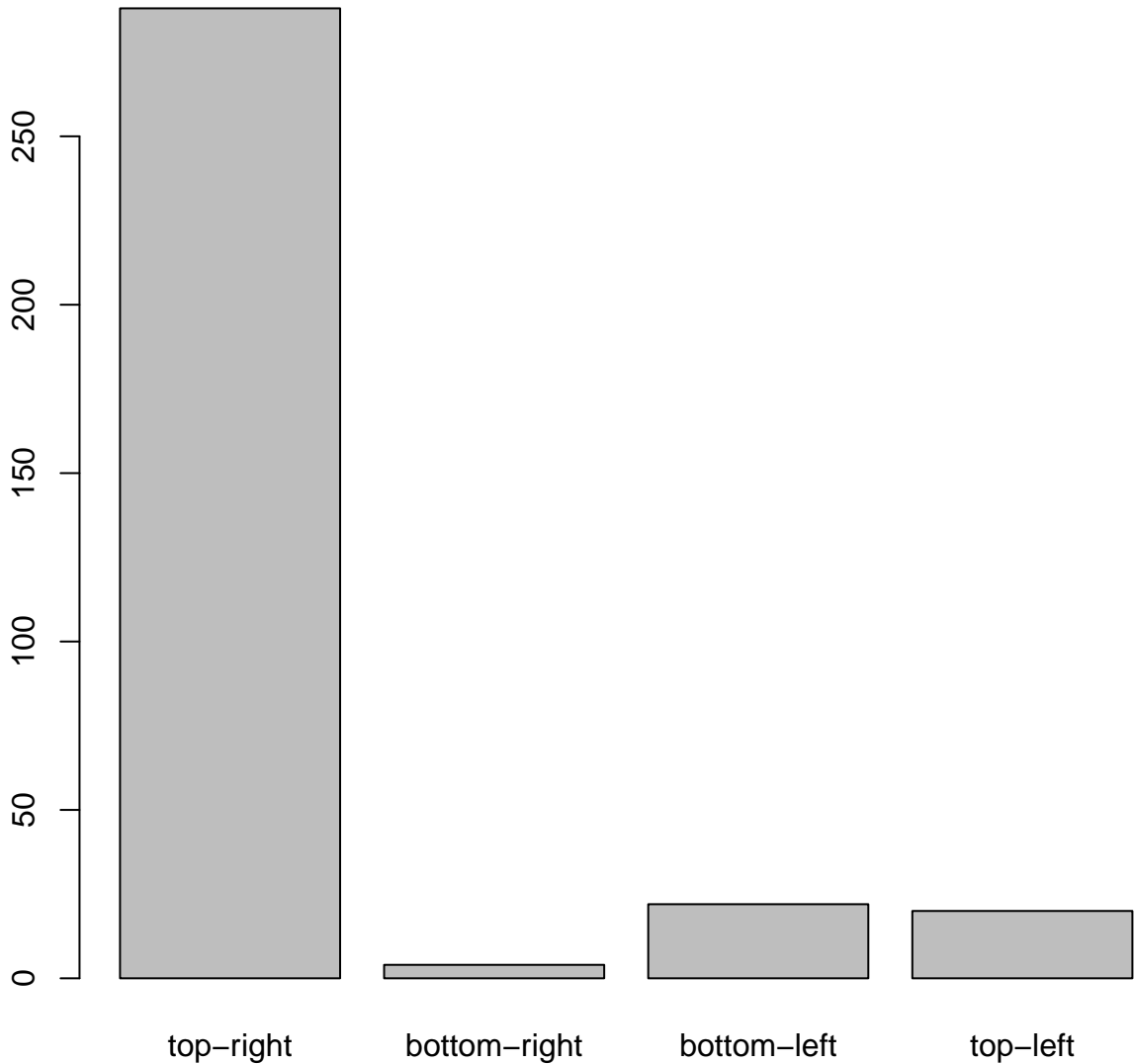
Histogram of geneset size



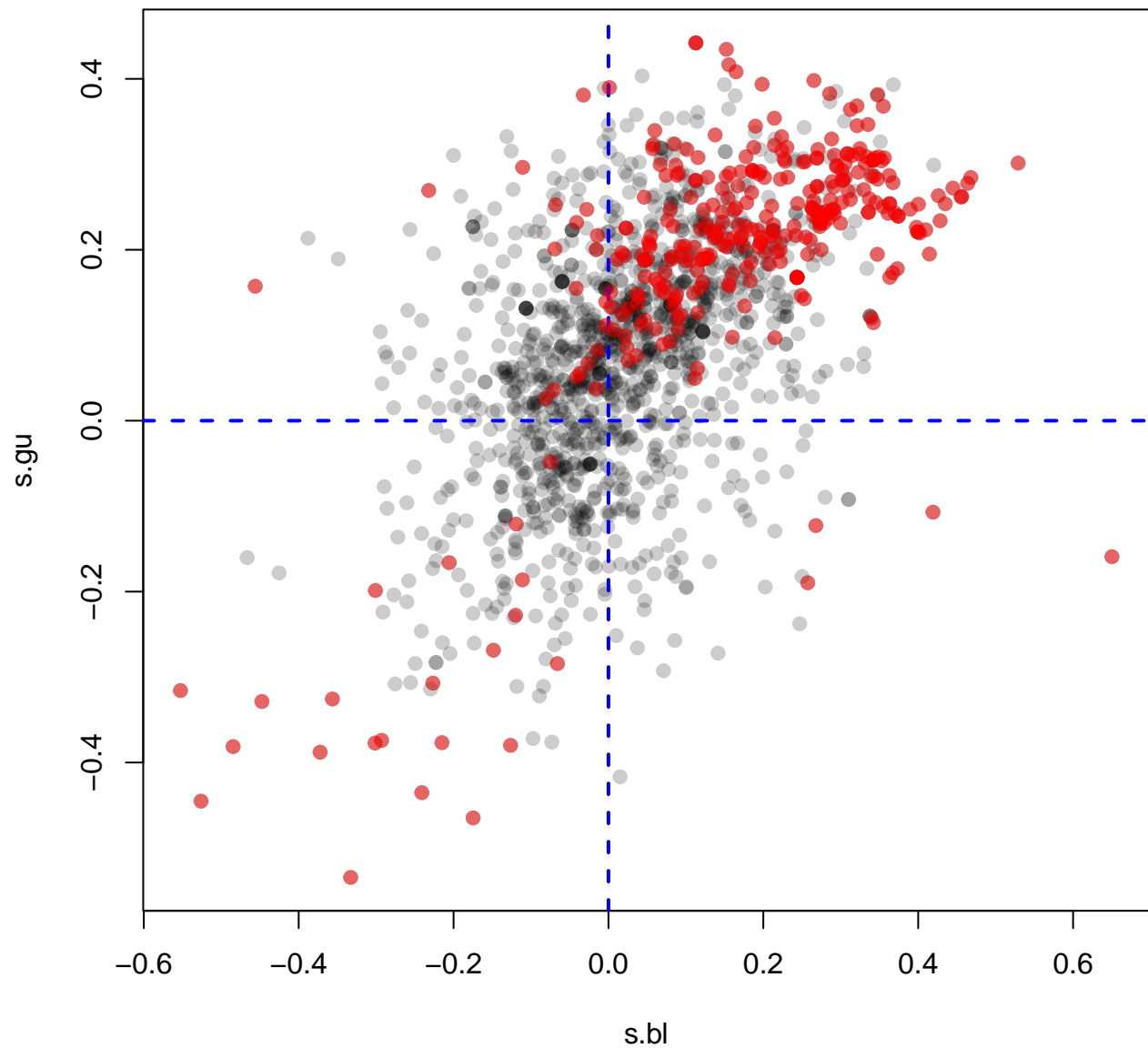
Trimmed histogram of geneset size



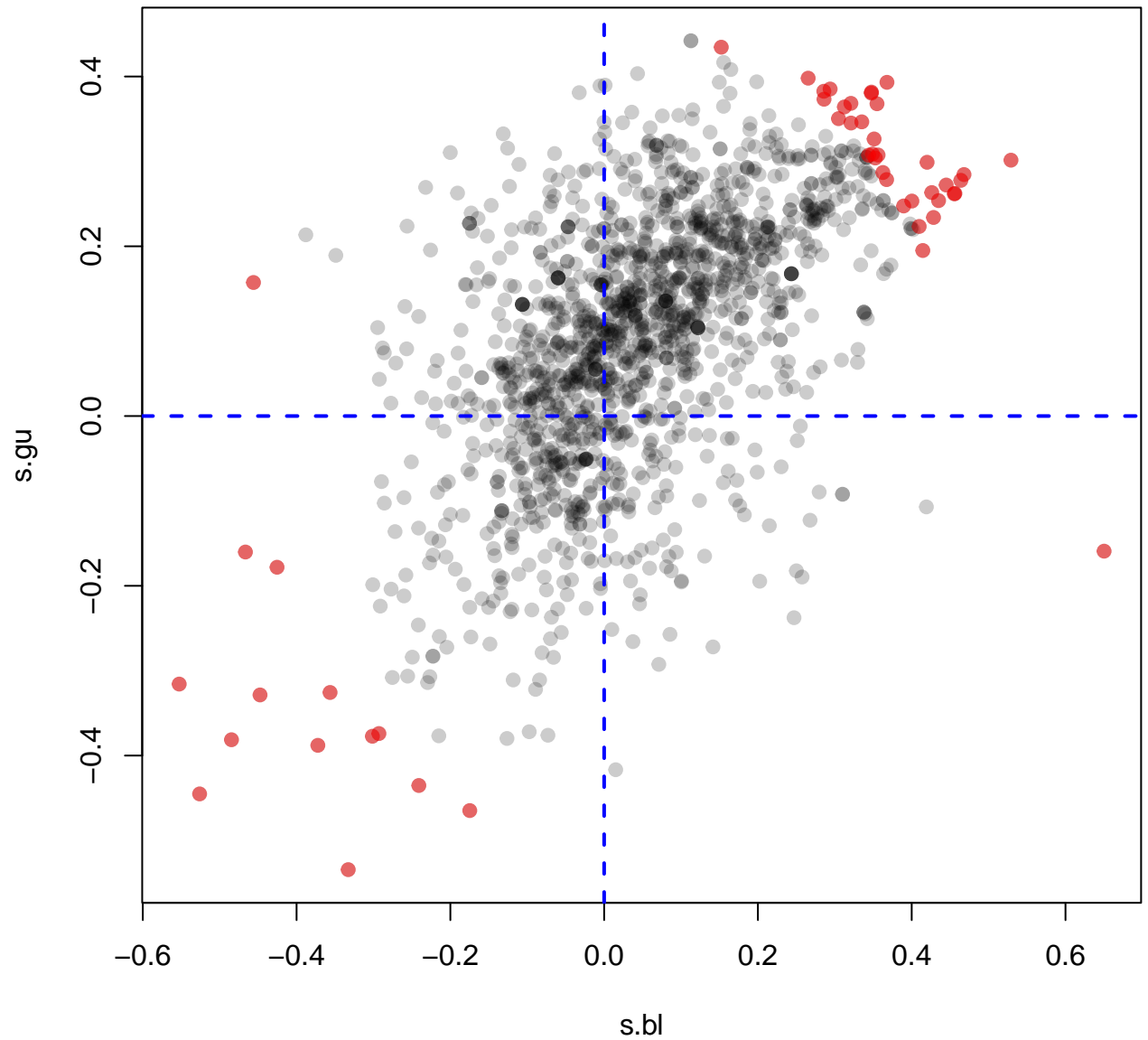
number of genesets FDR<0.05



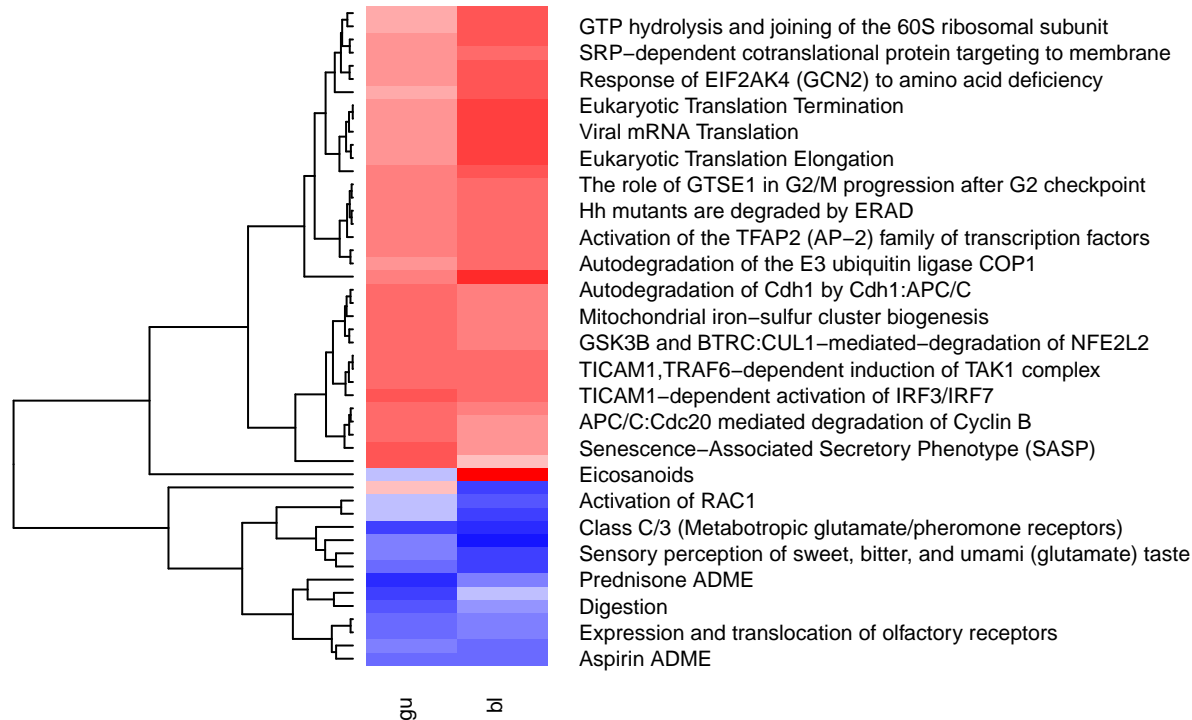
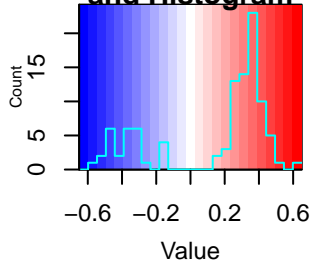
Scatterplot of all gene sets; FDR<0.05 in red



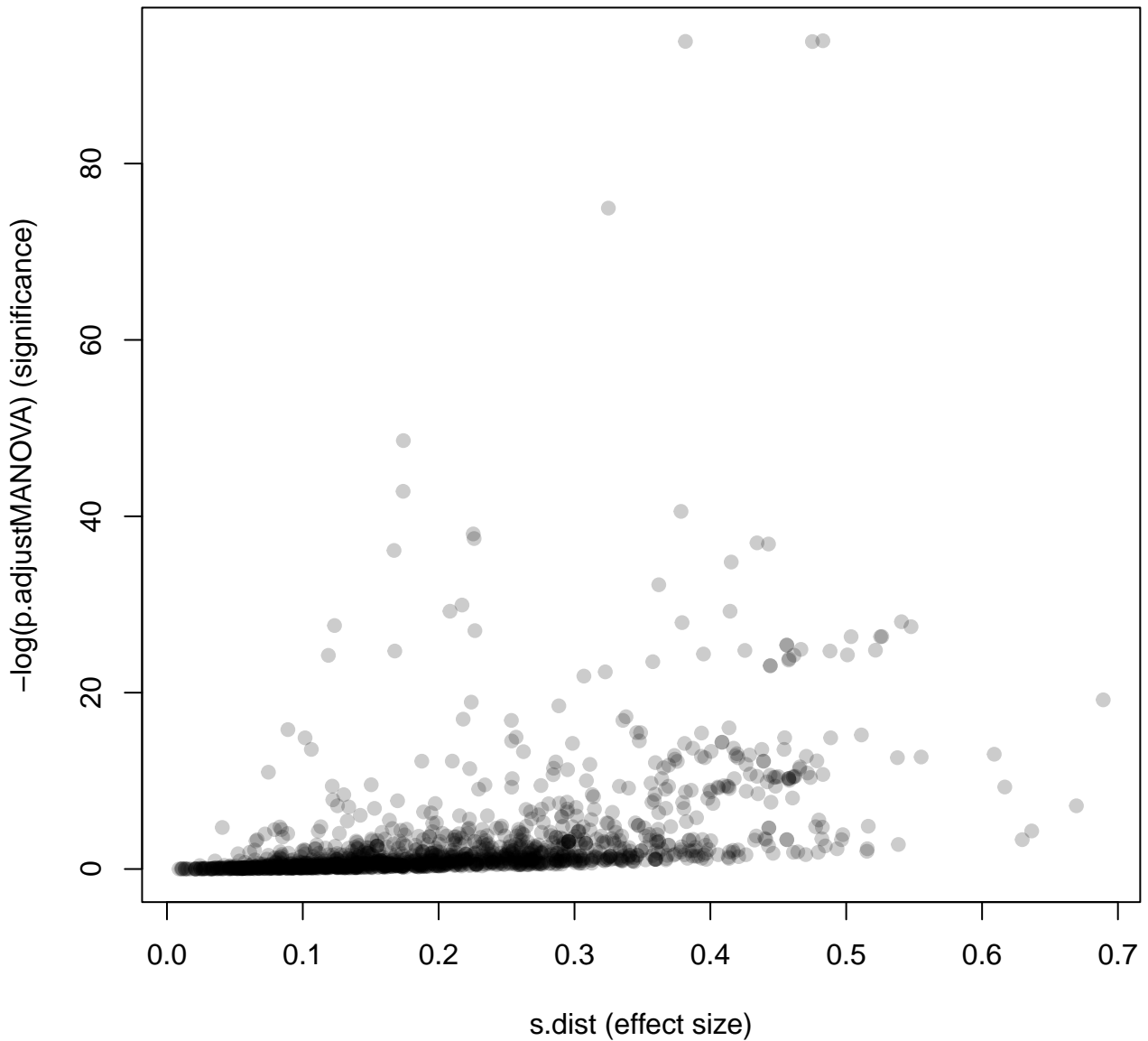
Scatterplot of all gene sets; top 50 in red



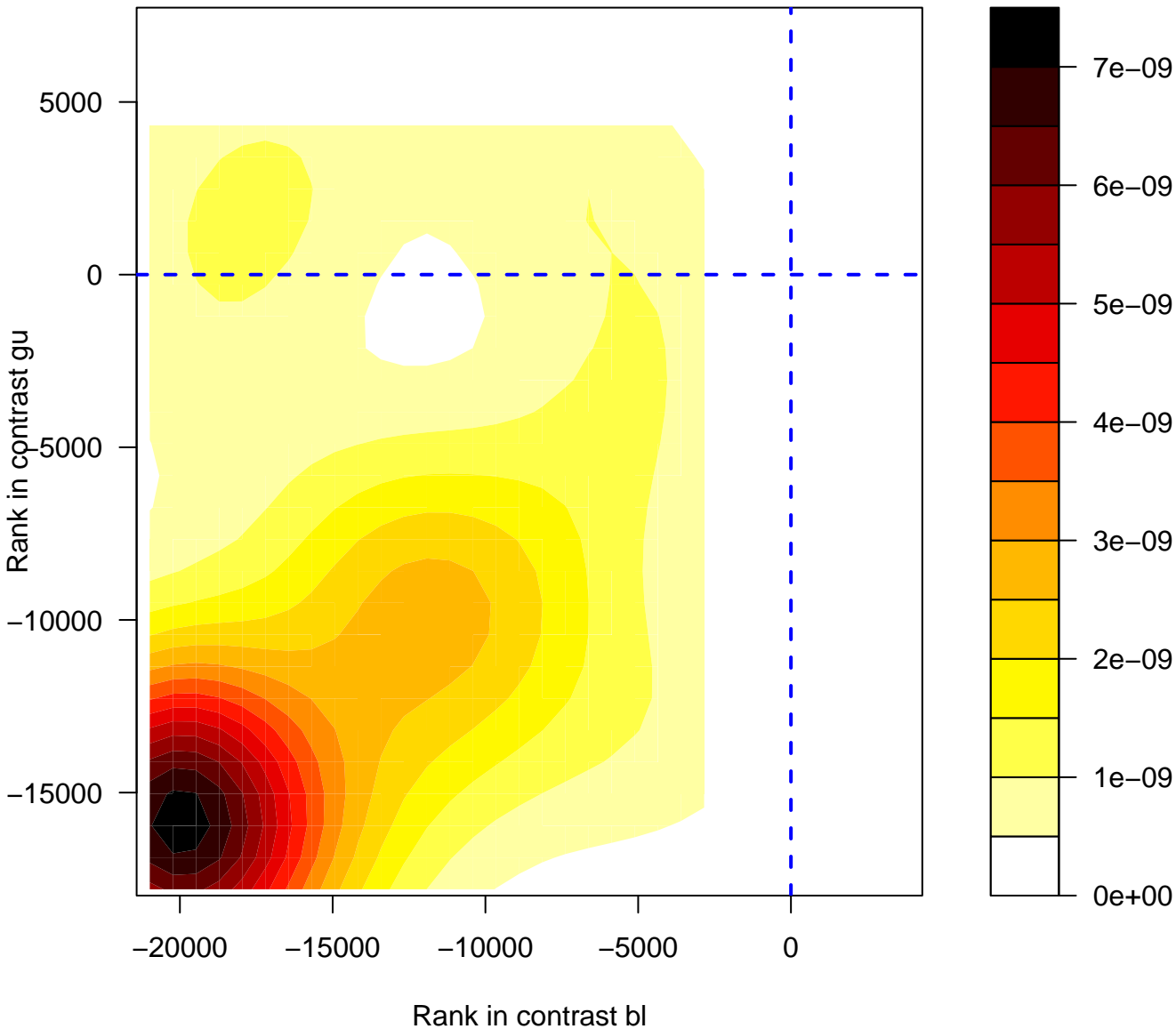
Color Key and Histogram



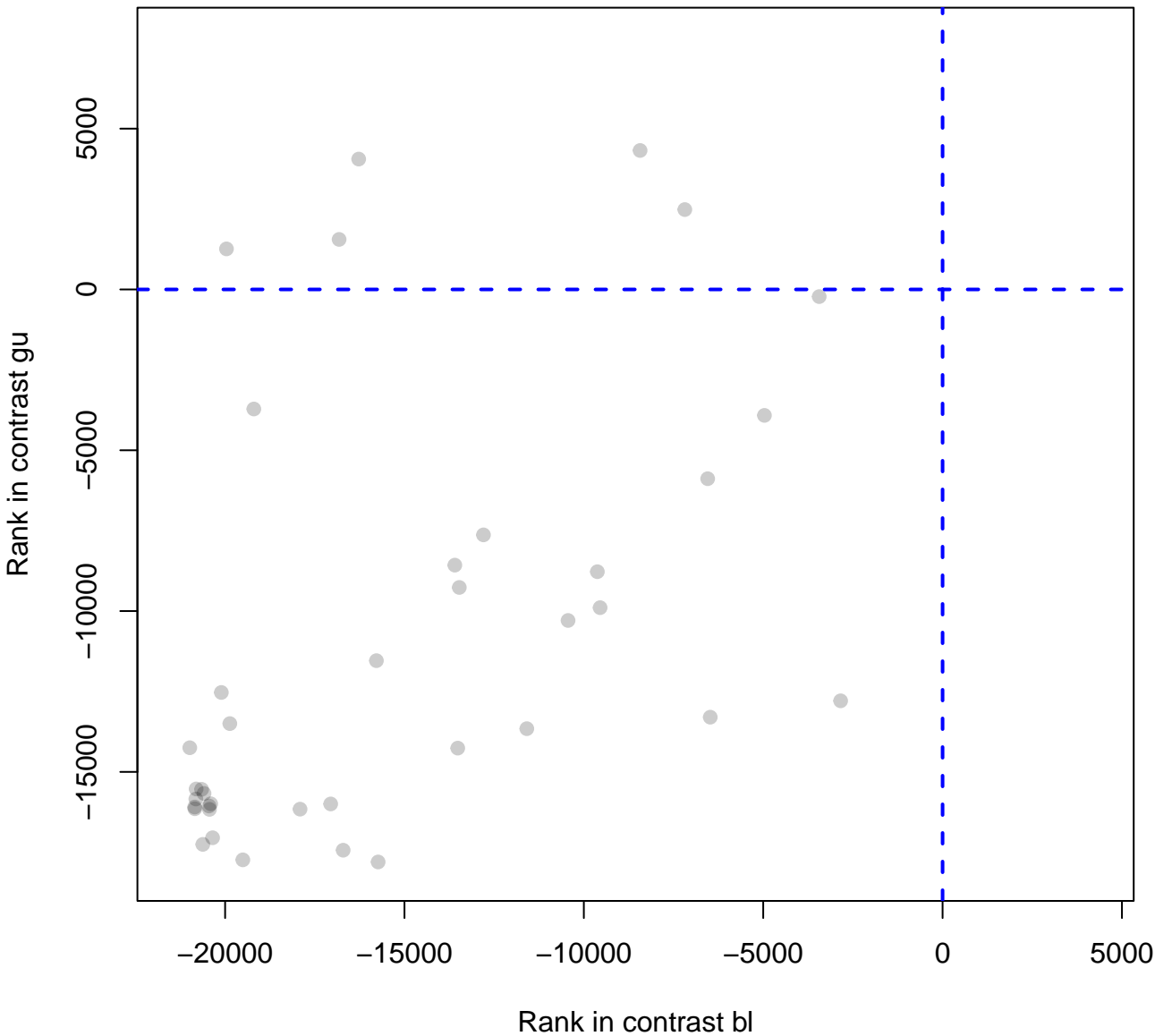
effect size versus statistical significance



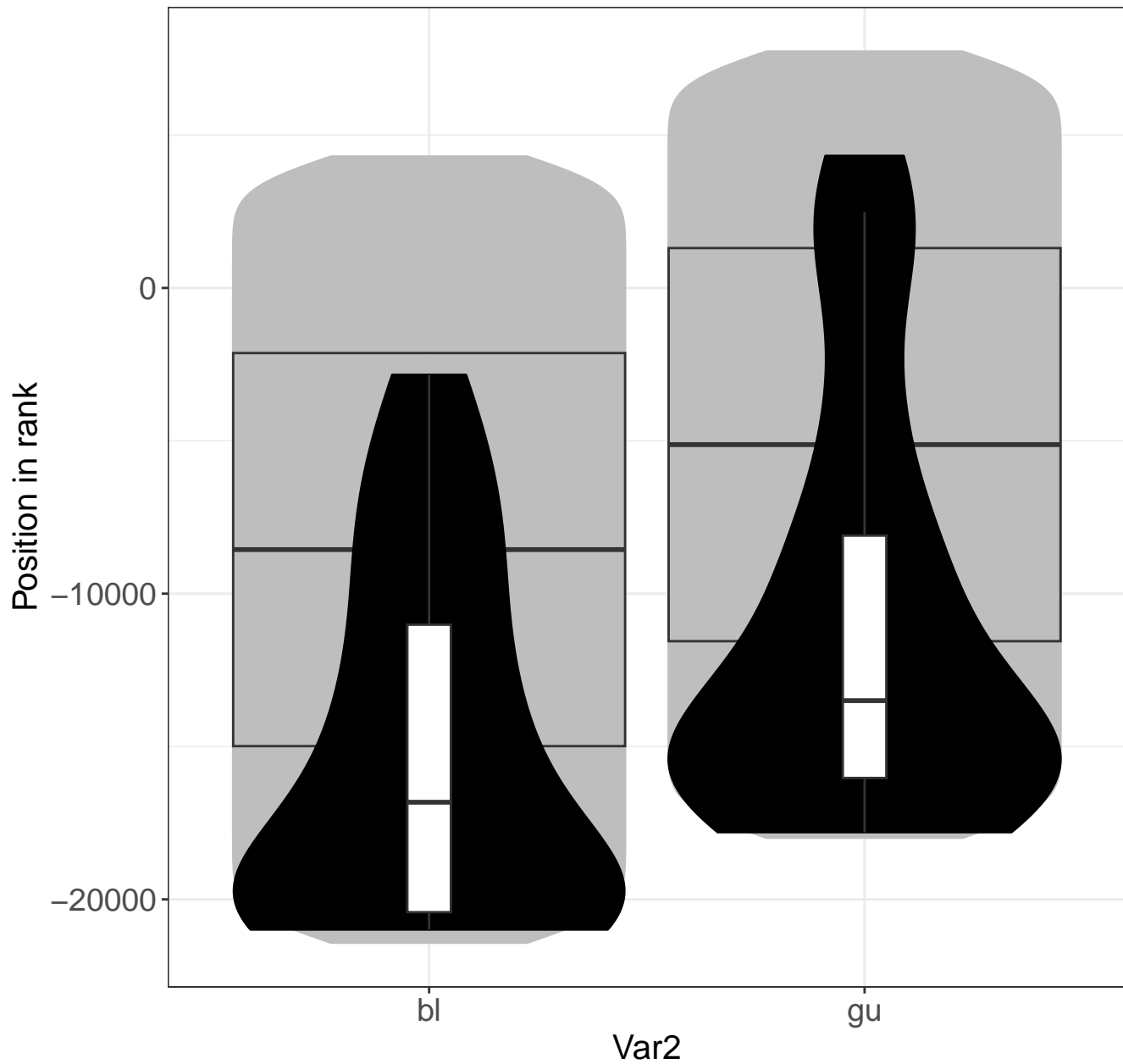
Class C/3 (Metabotropic glutamate/pheromone receptors)



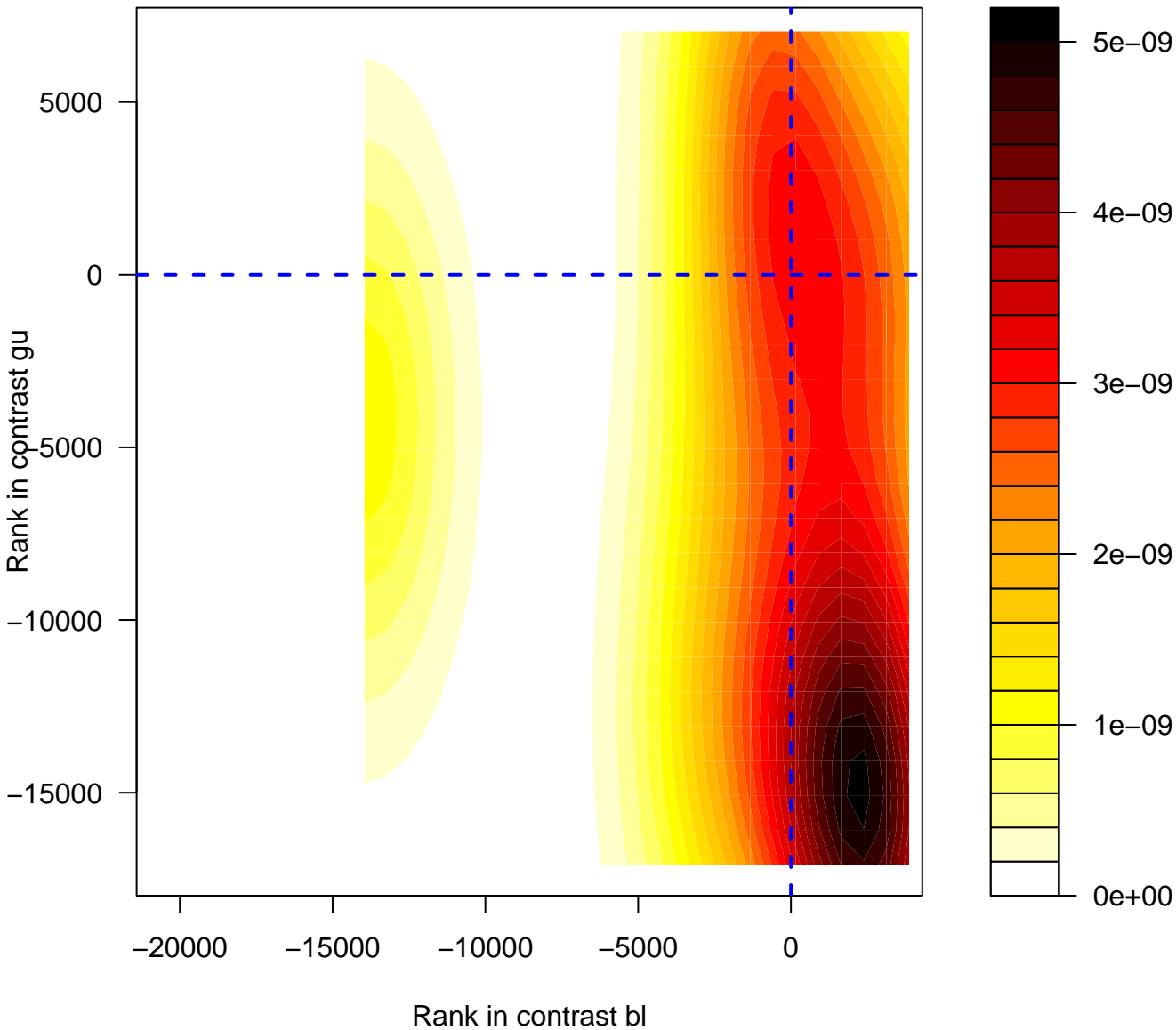
Class C/3 (Metabotropic glutamate/pheromone receptors)



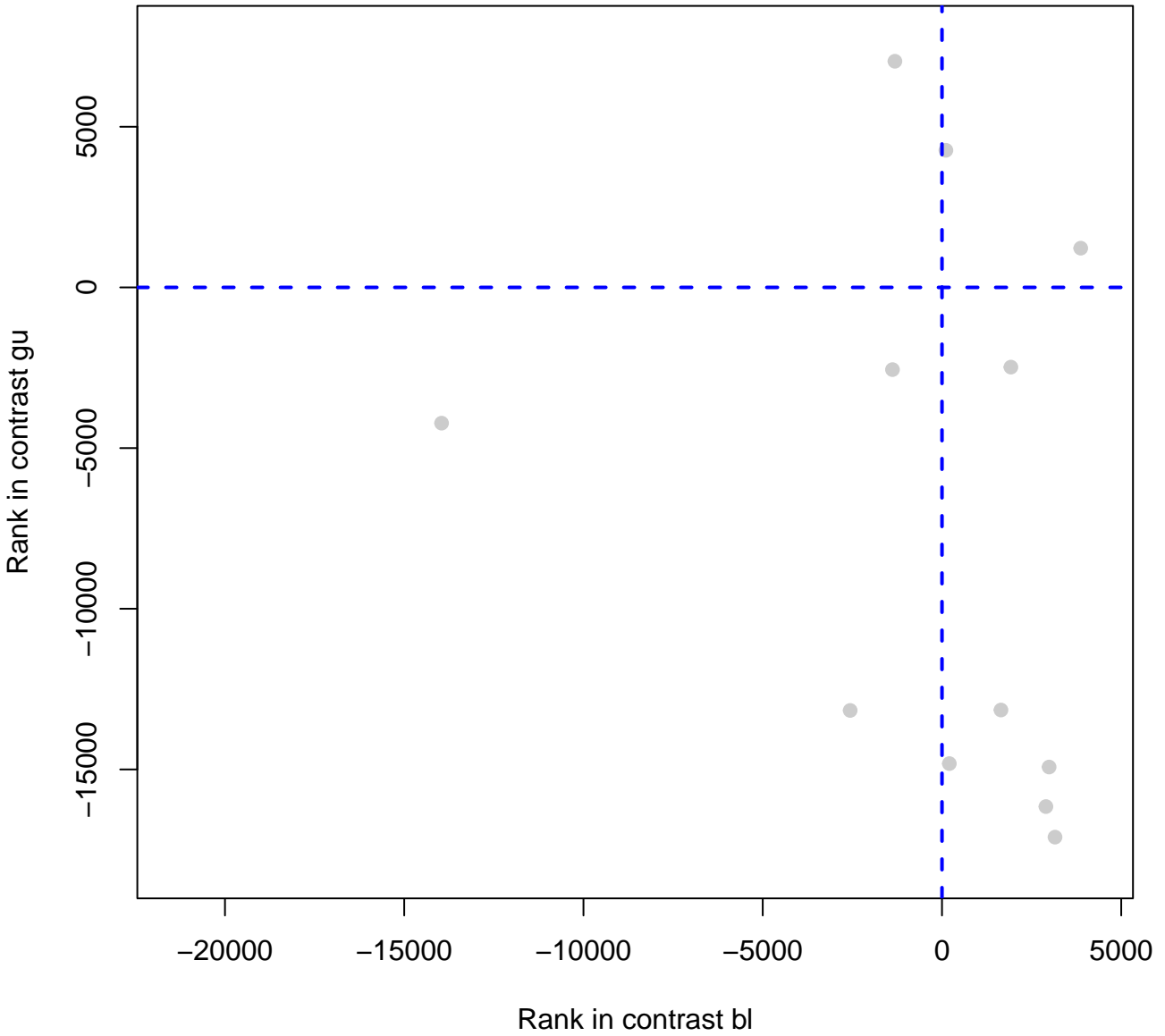
Class C/3 (Metabotropic glutamate/pheromone r



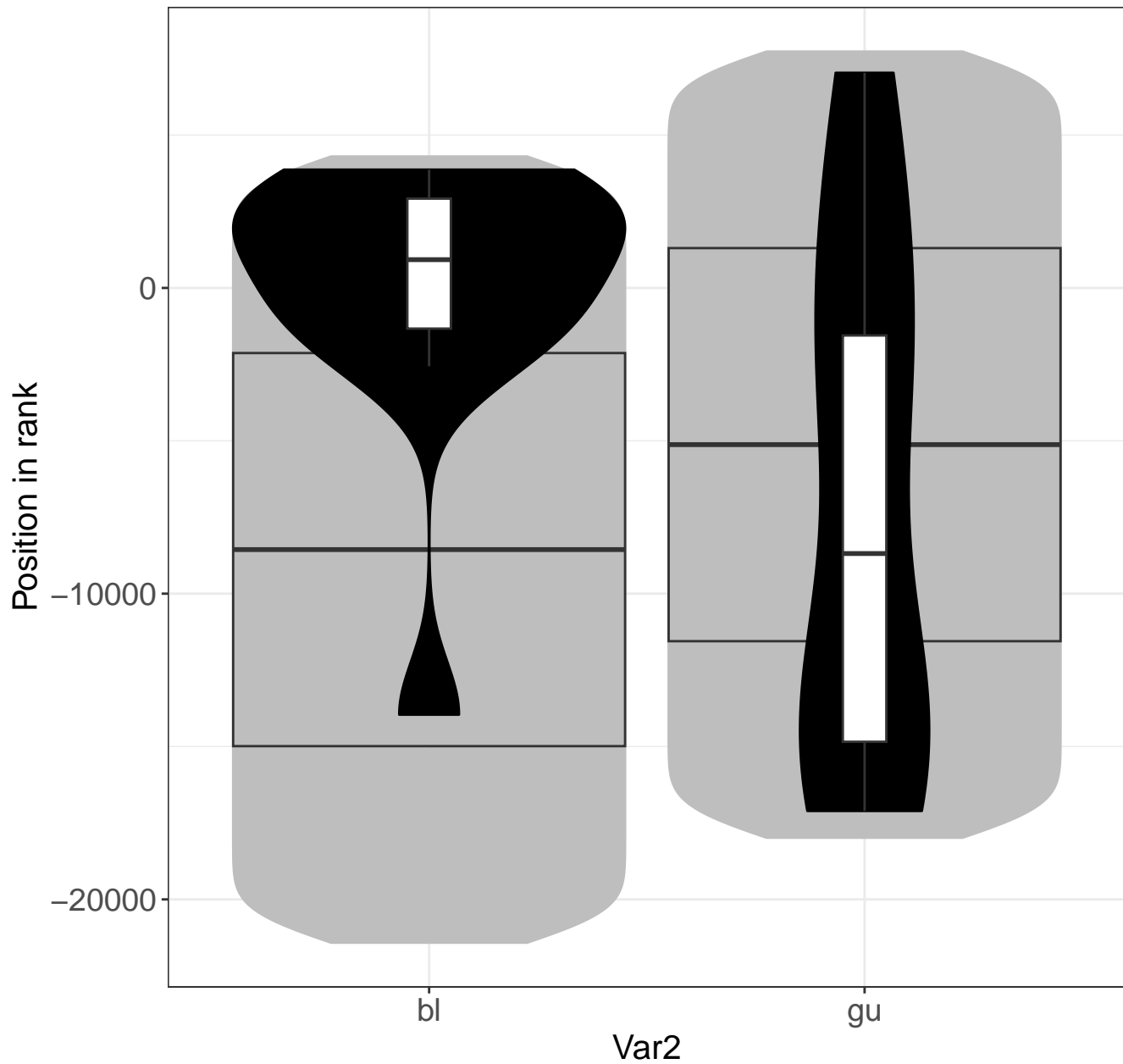
Eicosanoids



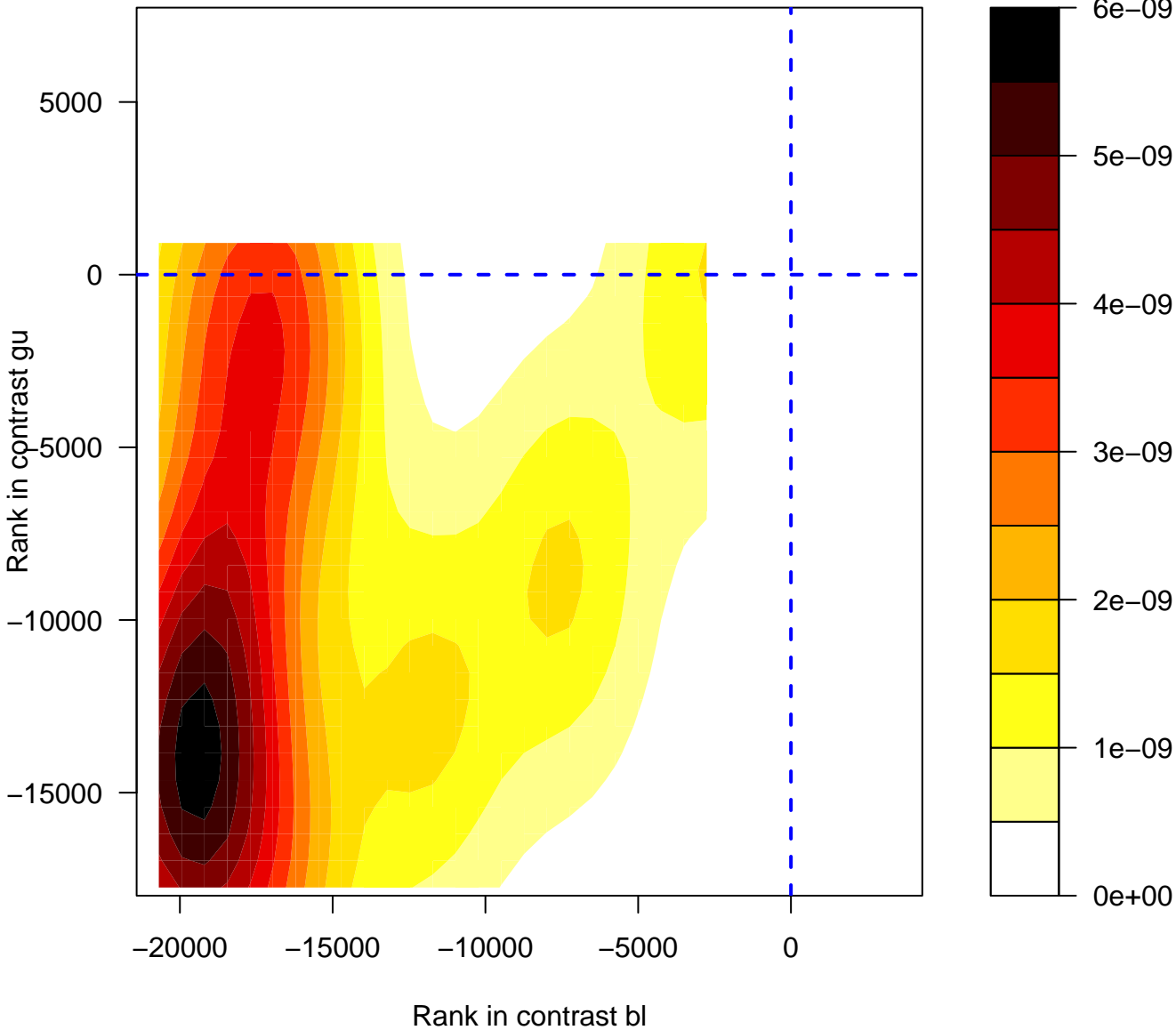
Eicosanoids



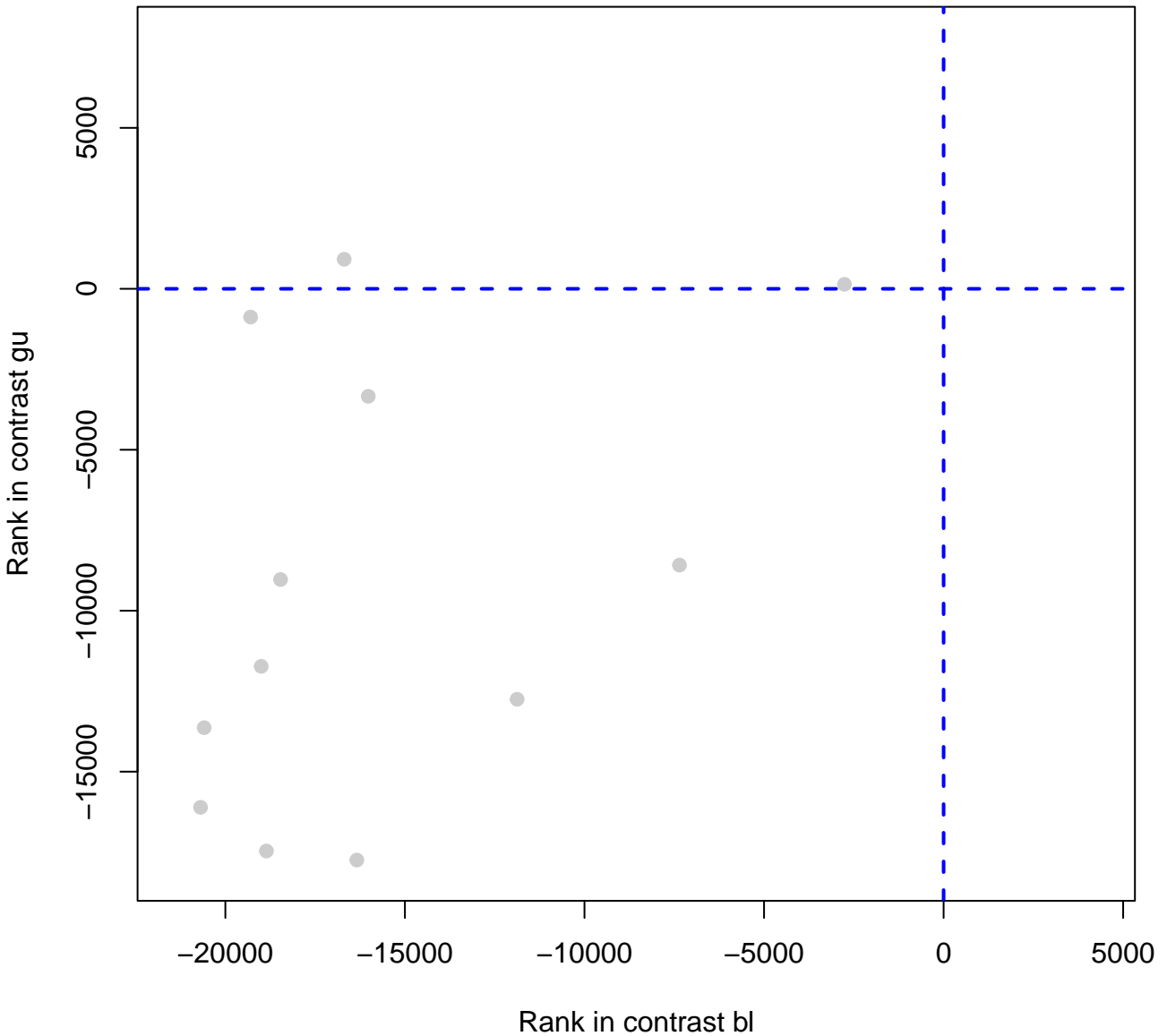
Eicosanoids



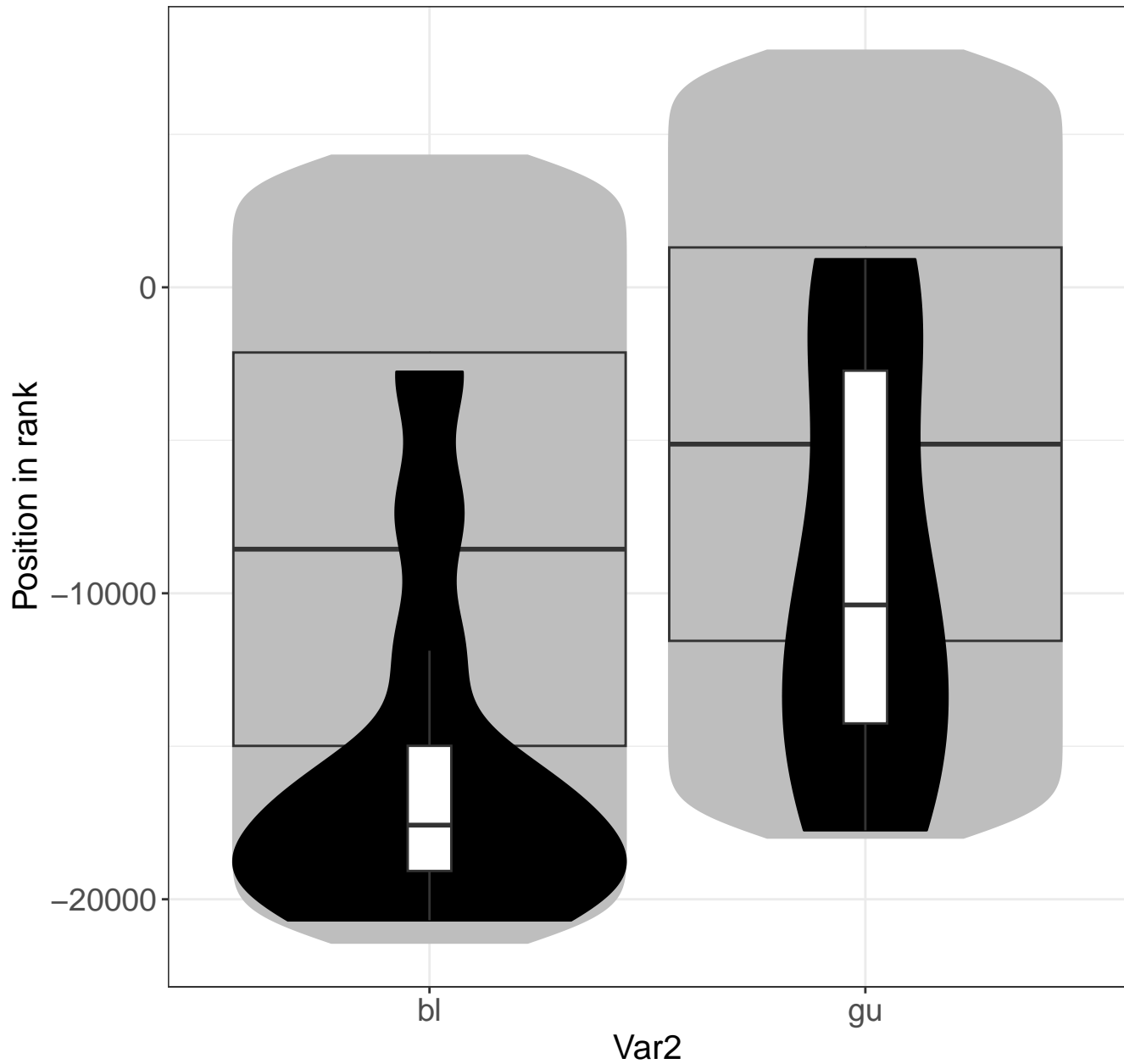
Scavenging of heme from plasma



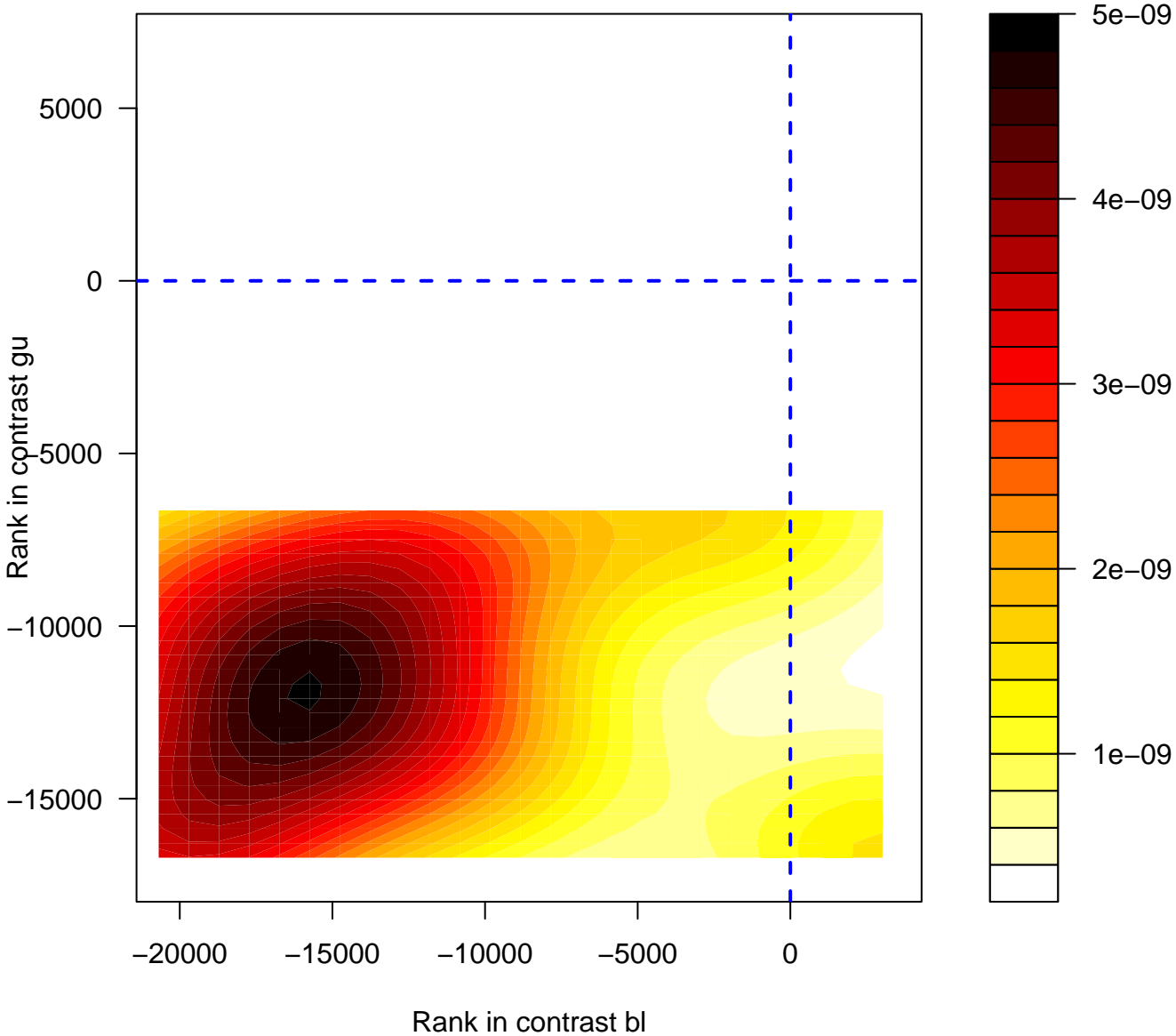
Scavenging of heme from plasma



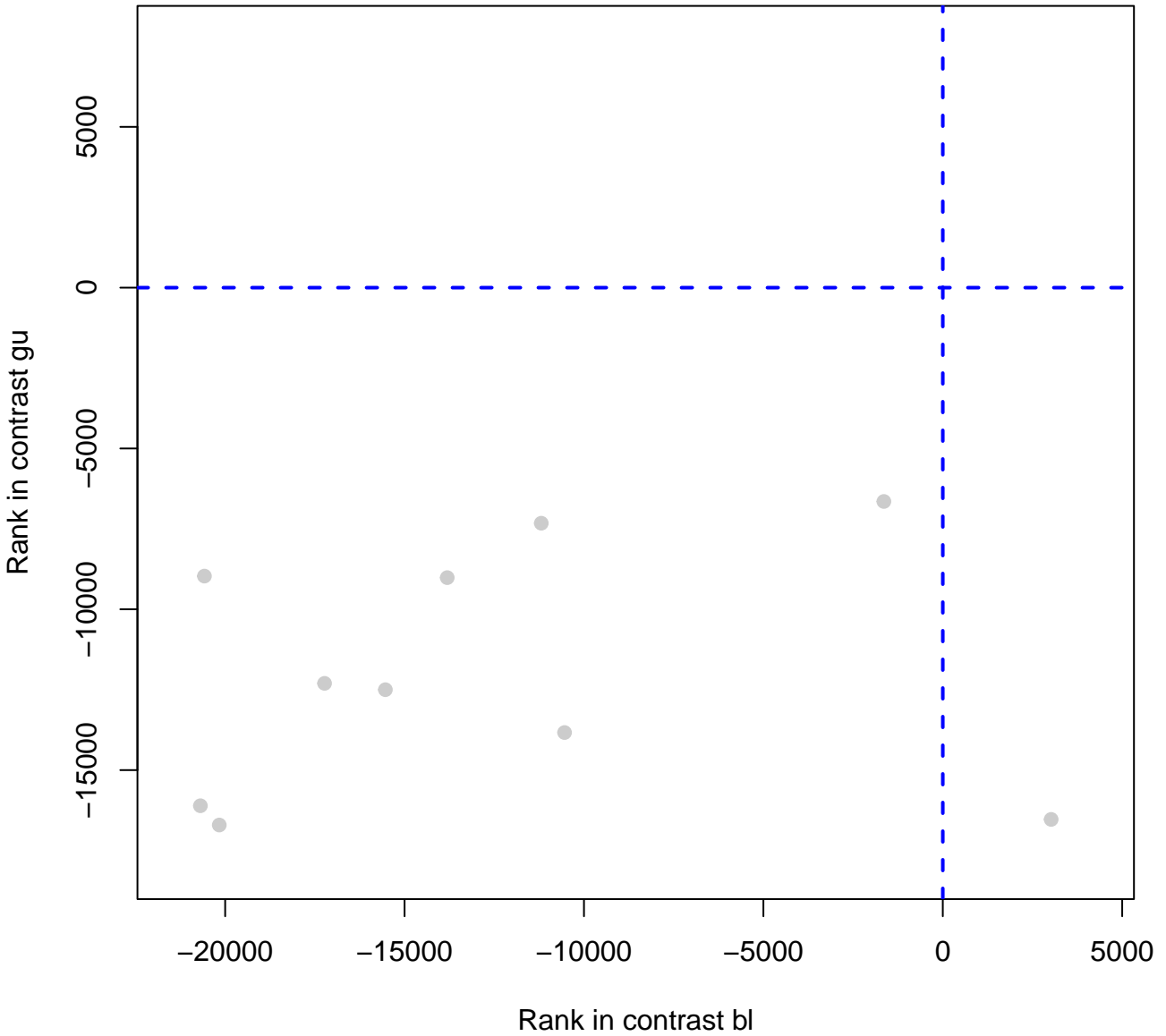
Scavenging of heme from plasma



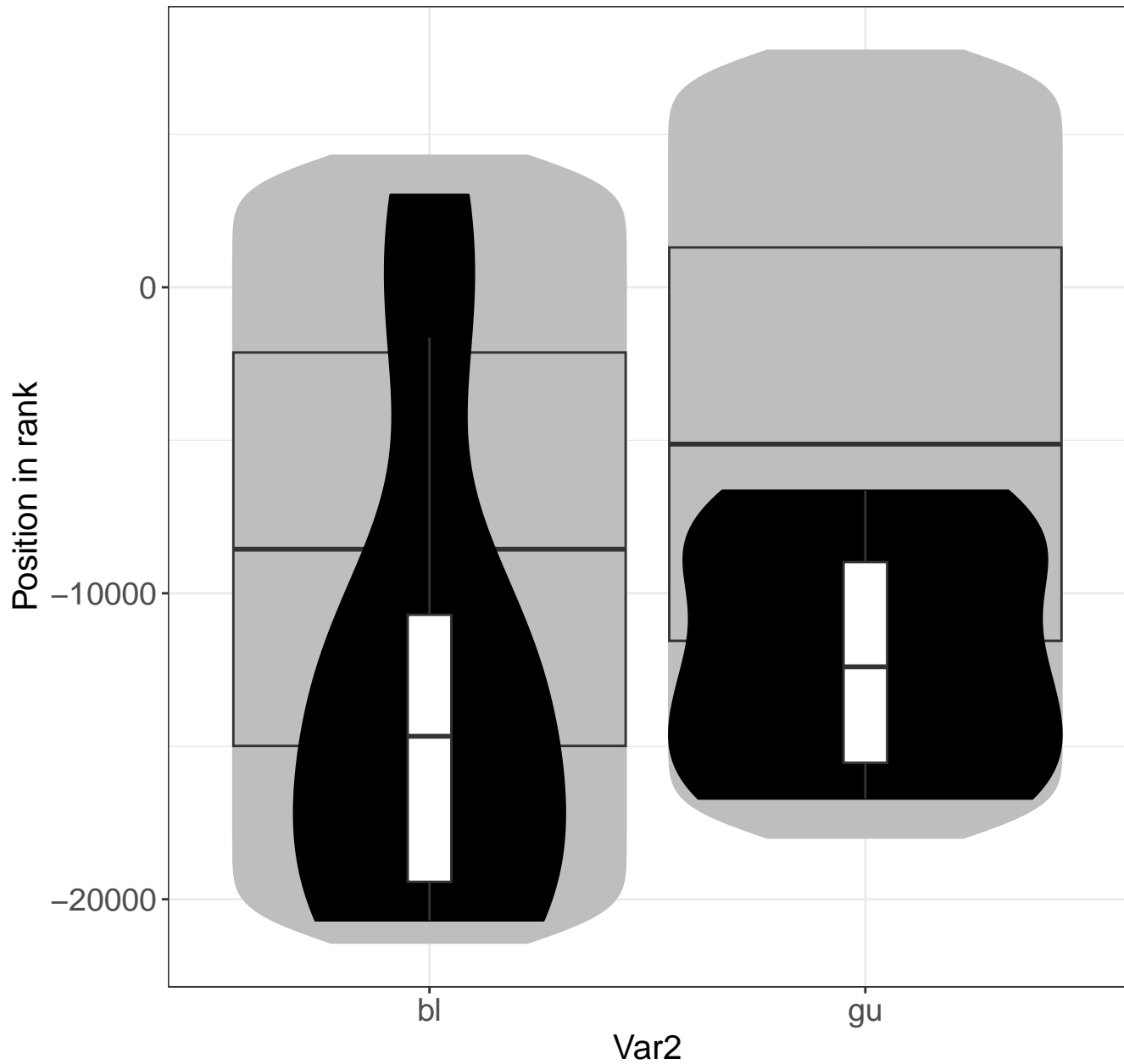
Prednisone ADME



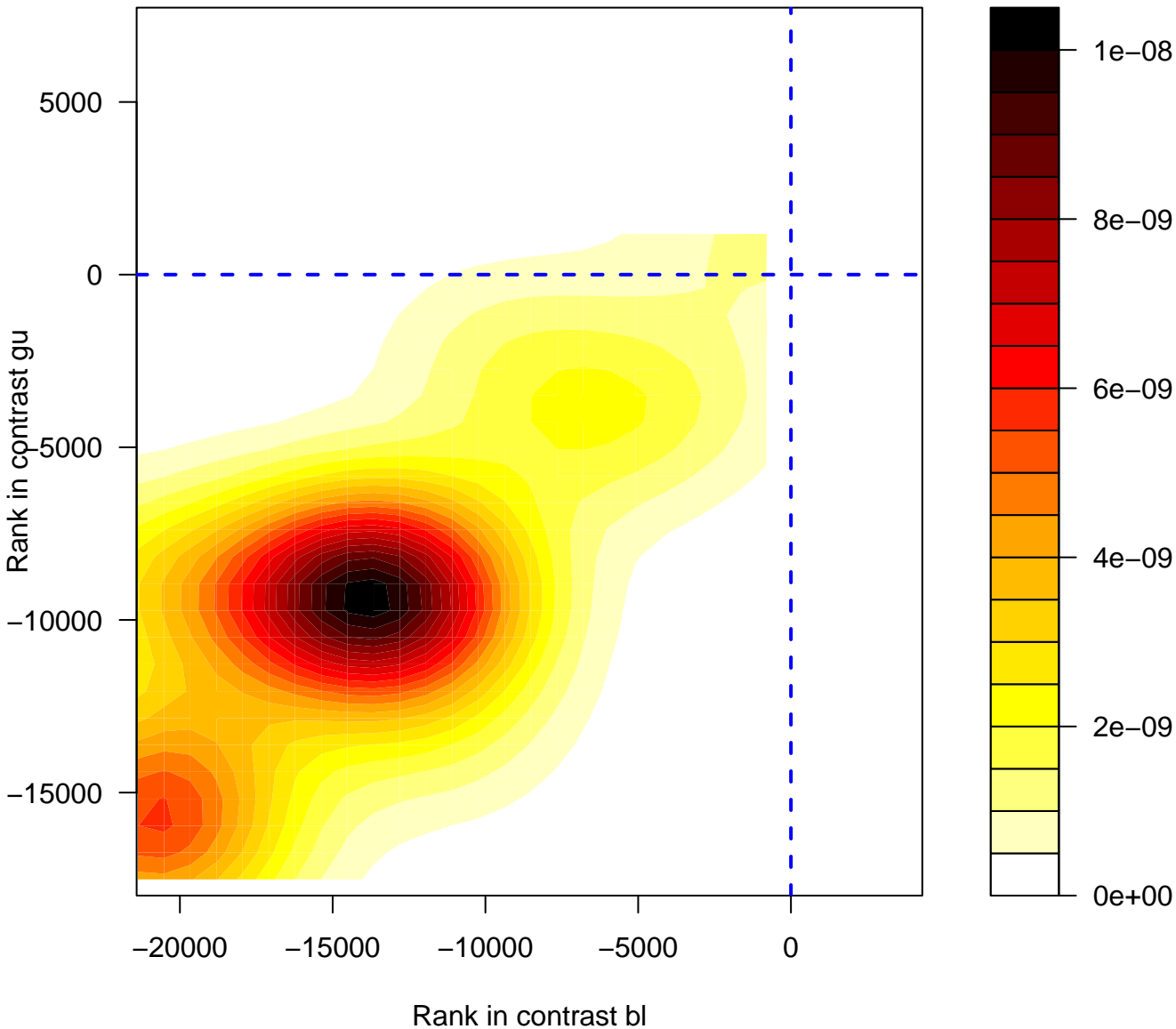
Prednisone ADME



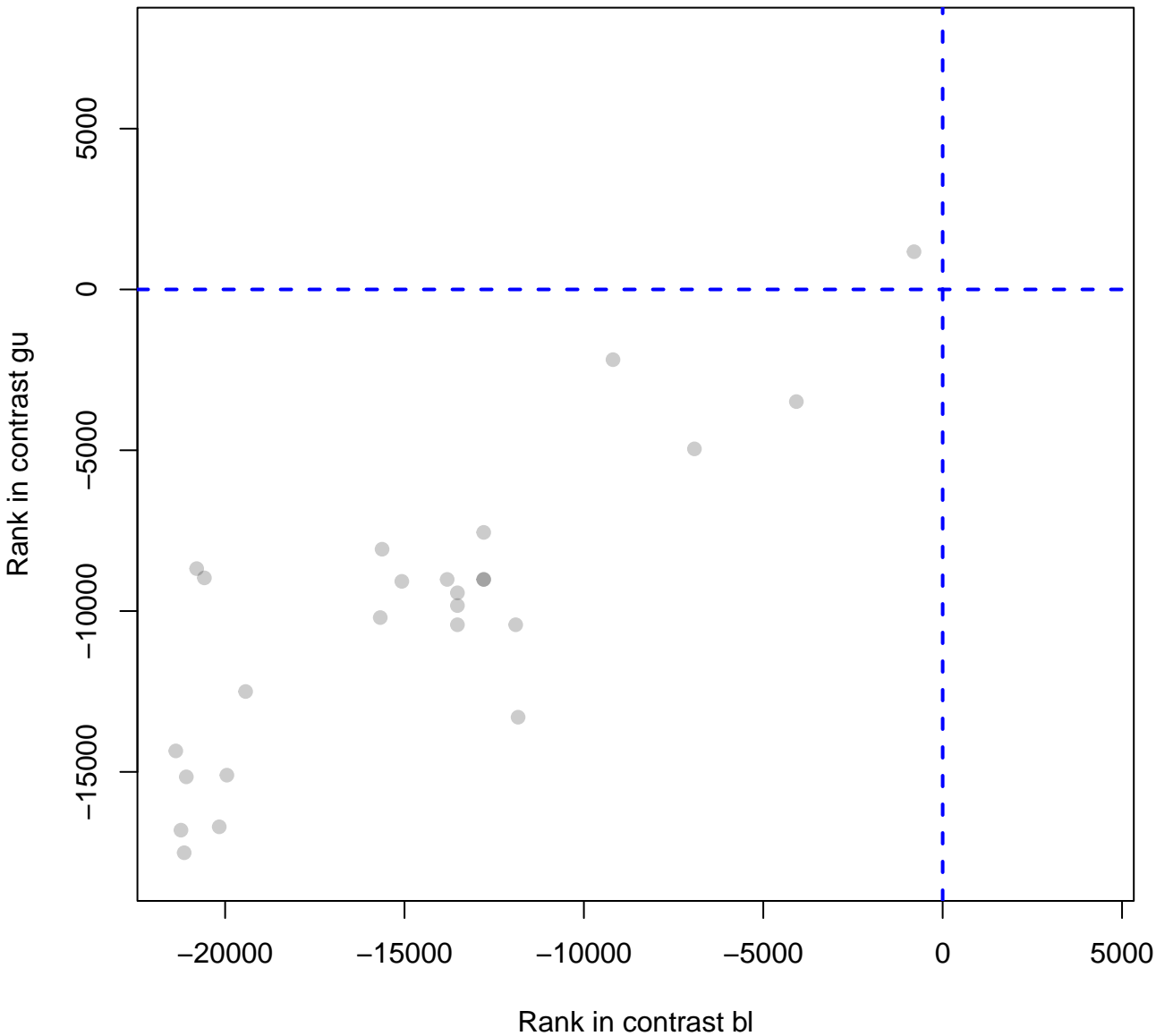
Prednisone ADME



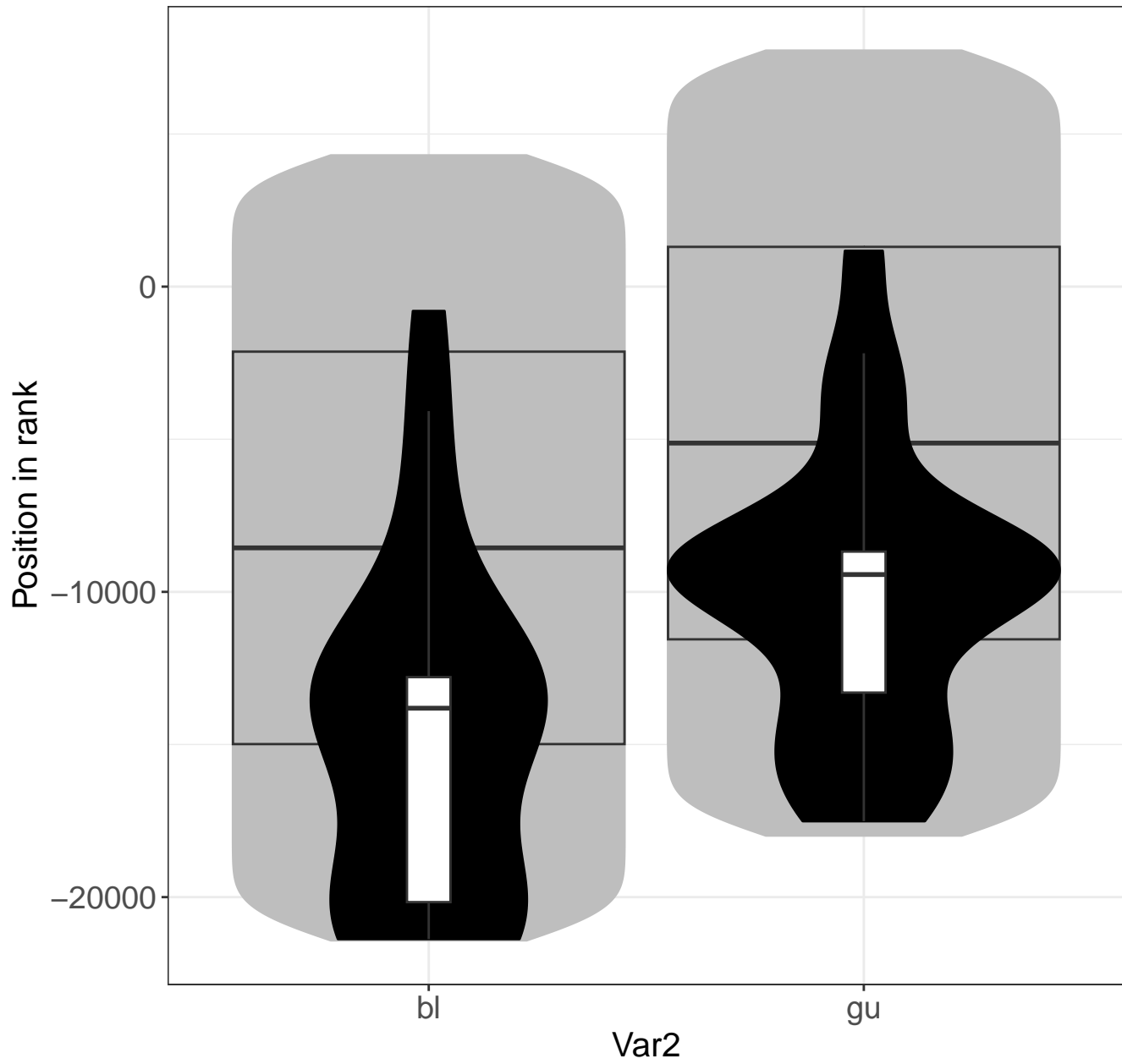
Glucuronidation



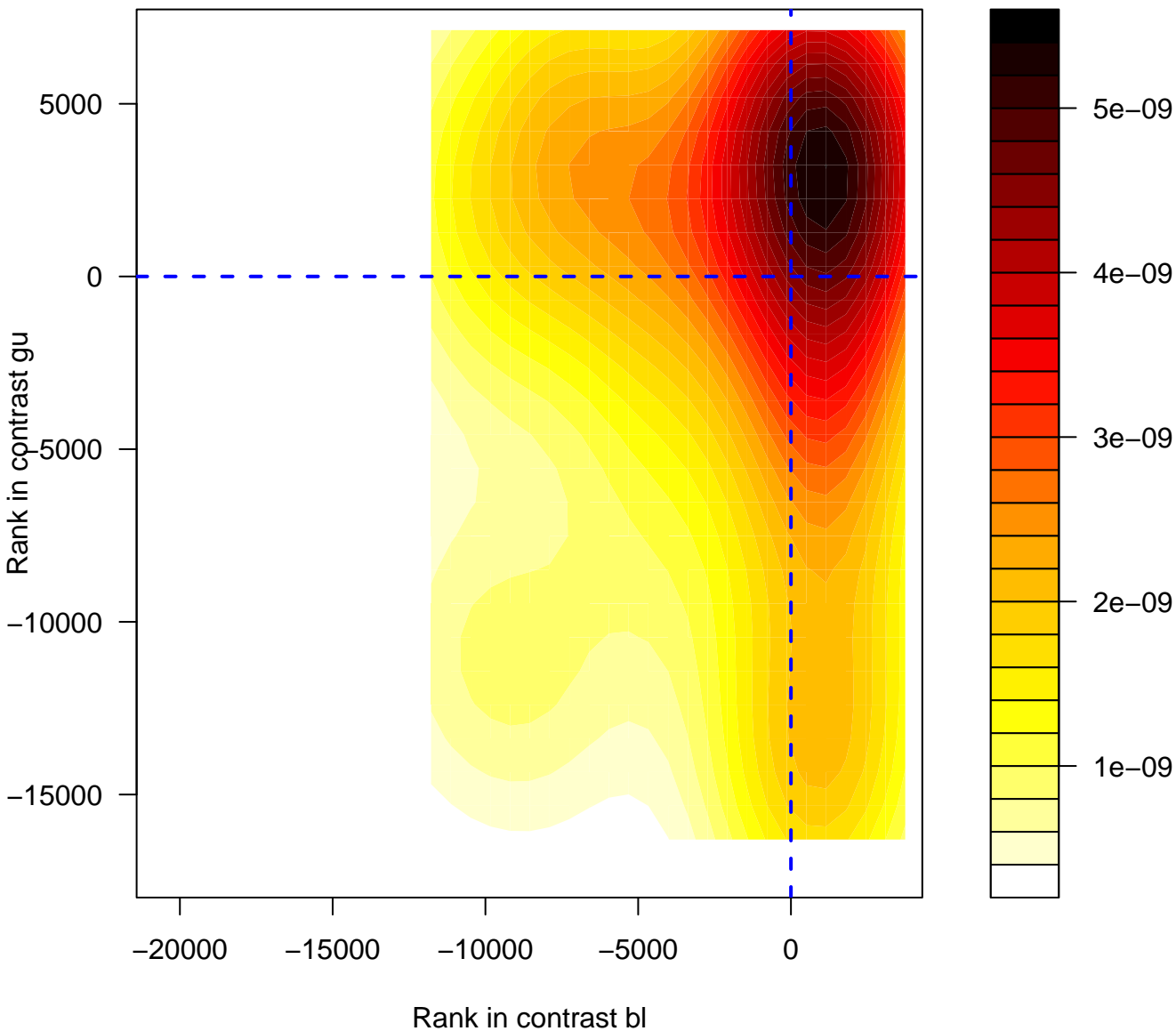
Glucuronidation



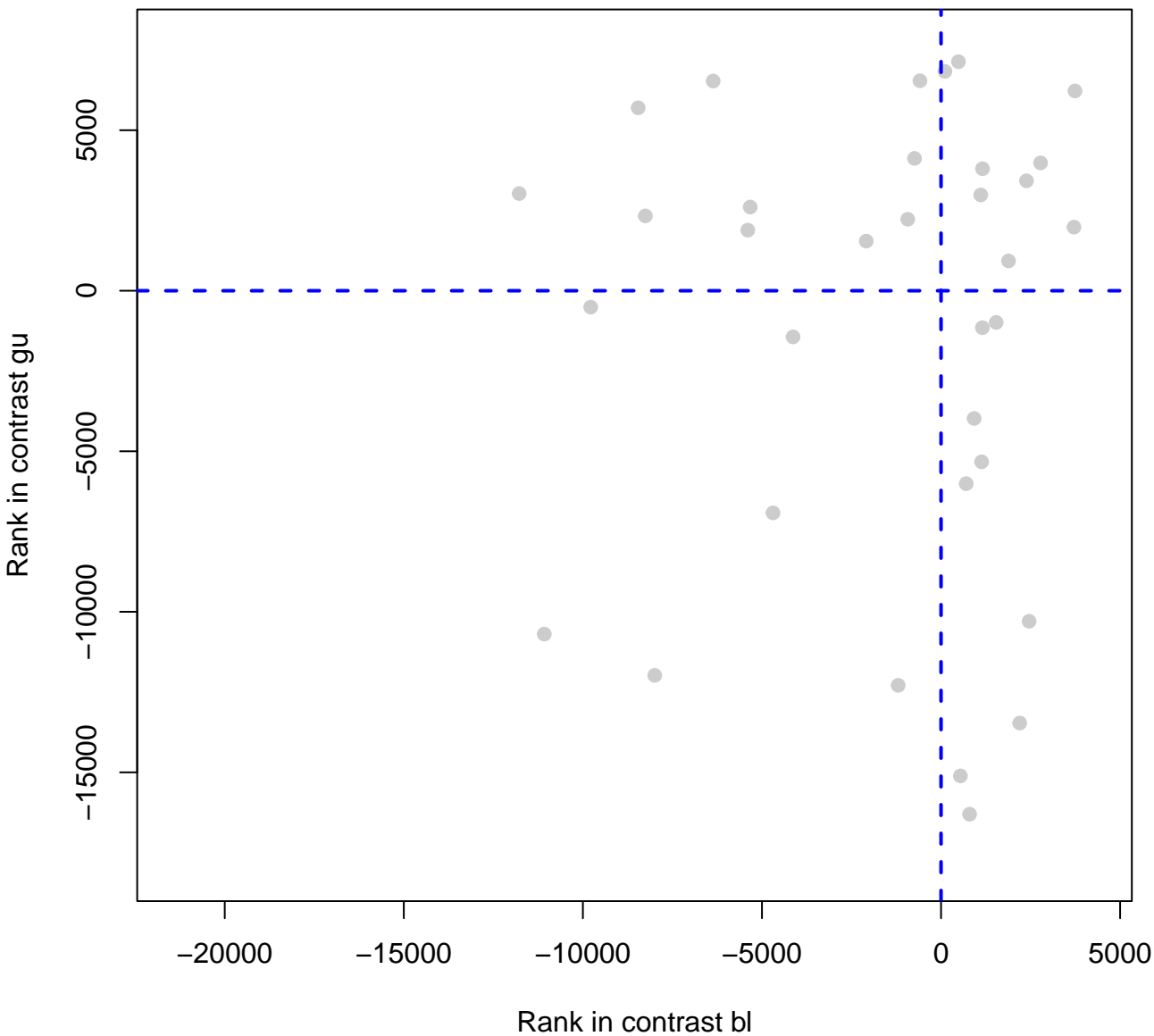
Glucuronidation



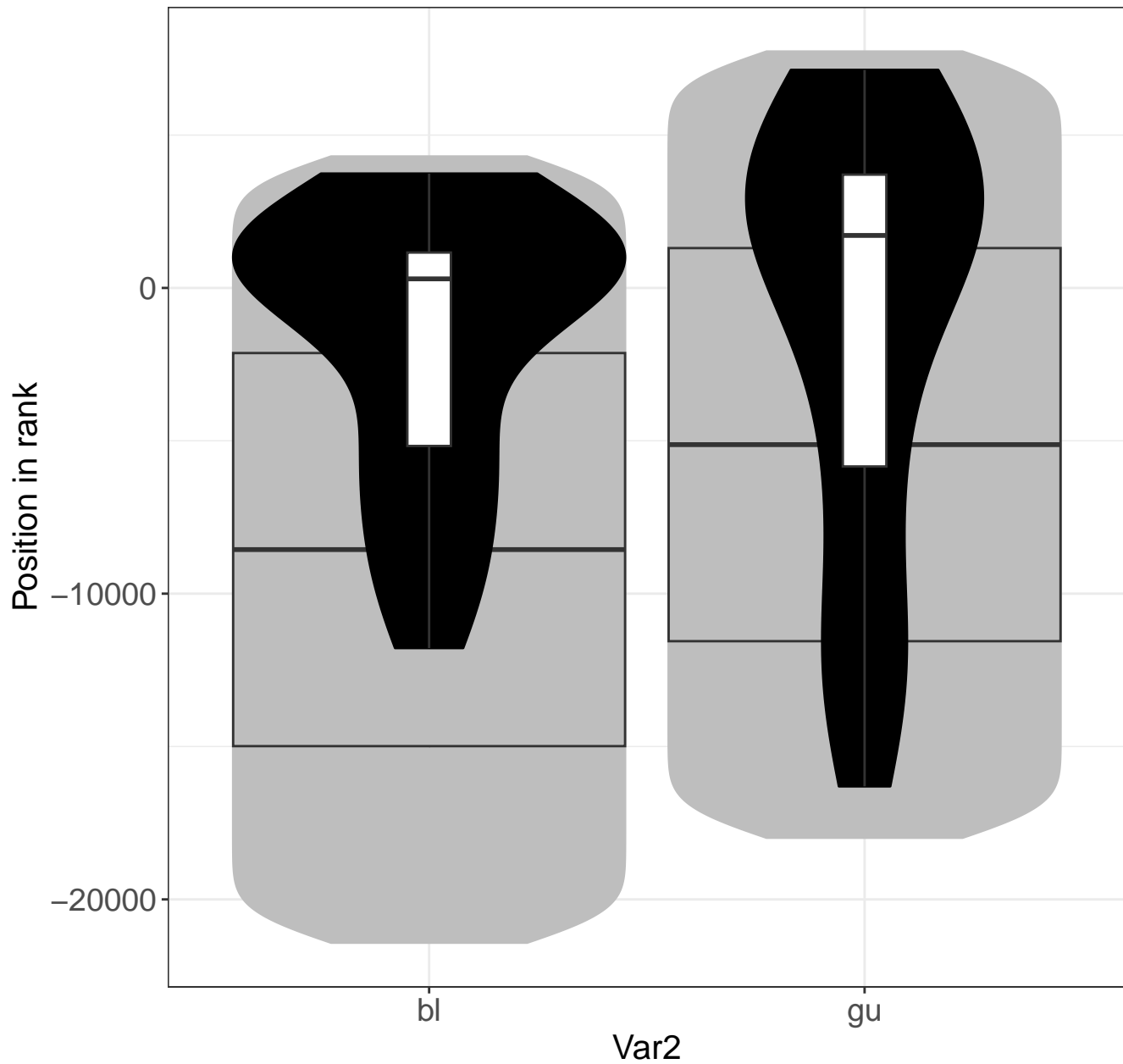
SARS-CoV-1 modulates host translation machinery



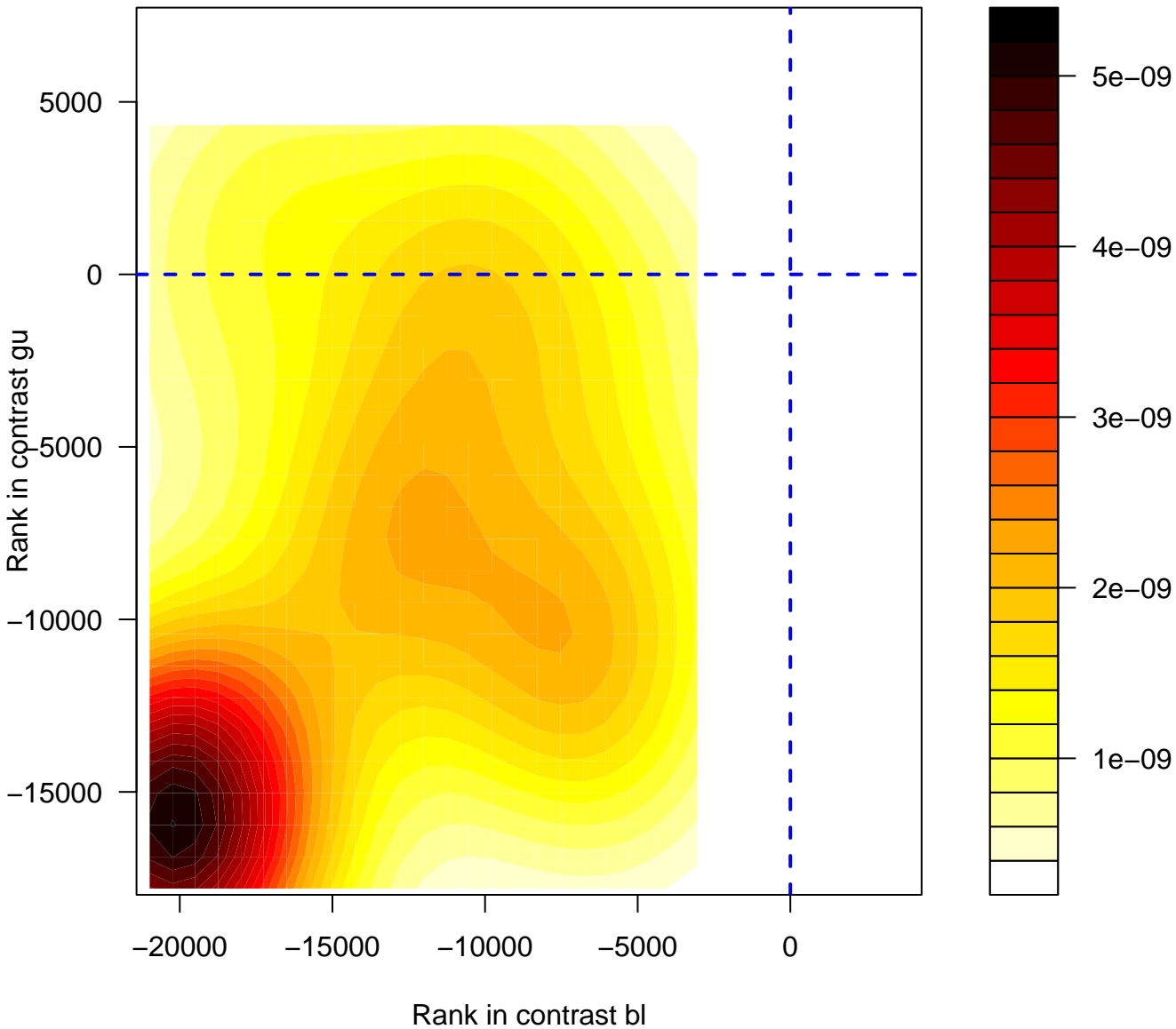
SARS-CoV-1 modulates host translation machinery



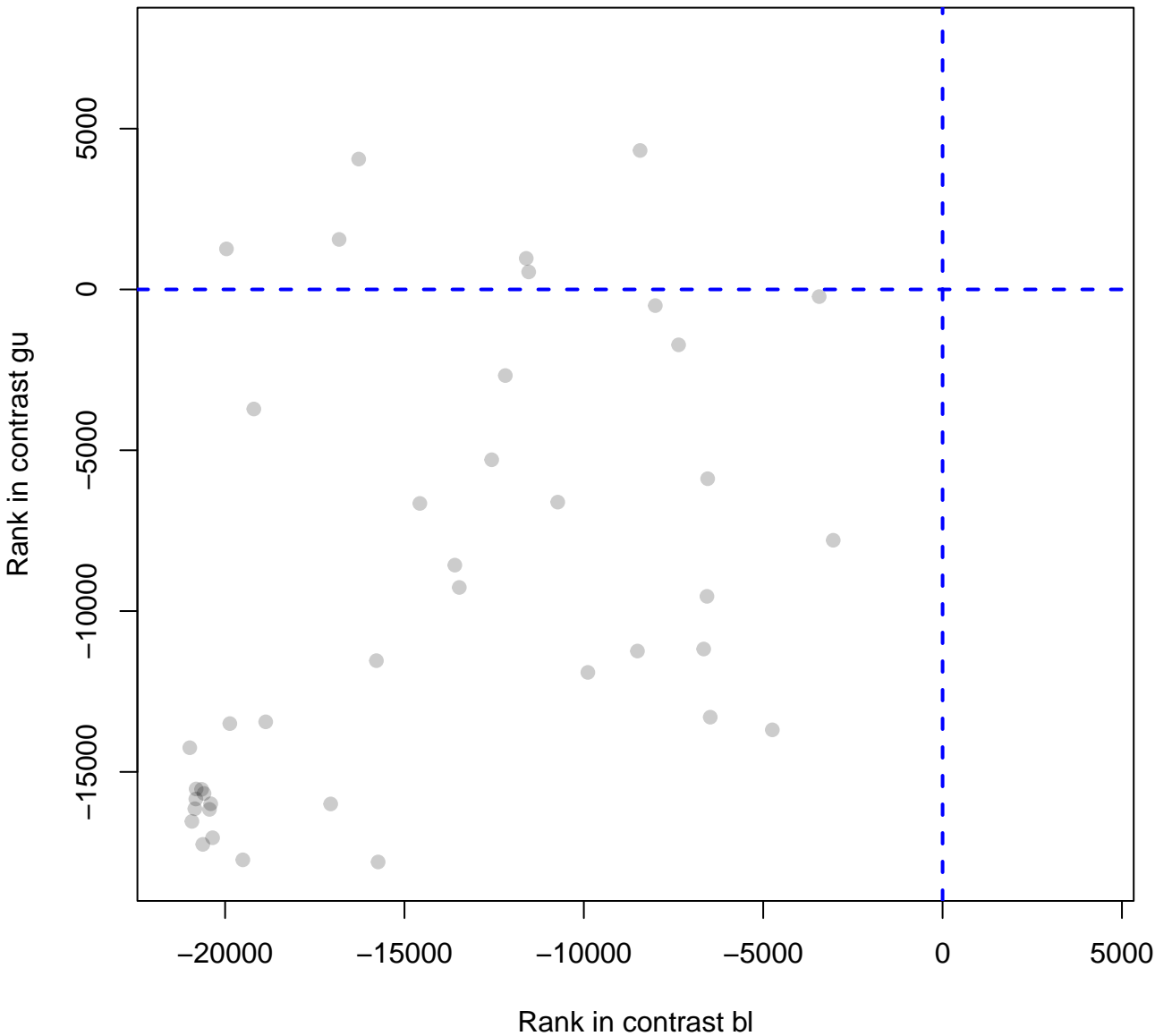
SARS-CoV-1 modulates host translation machinery



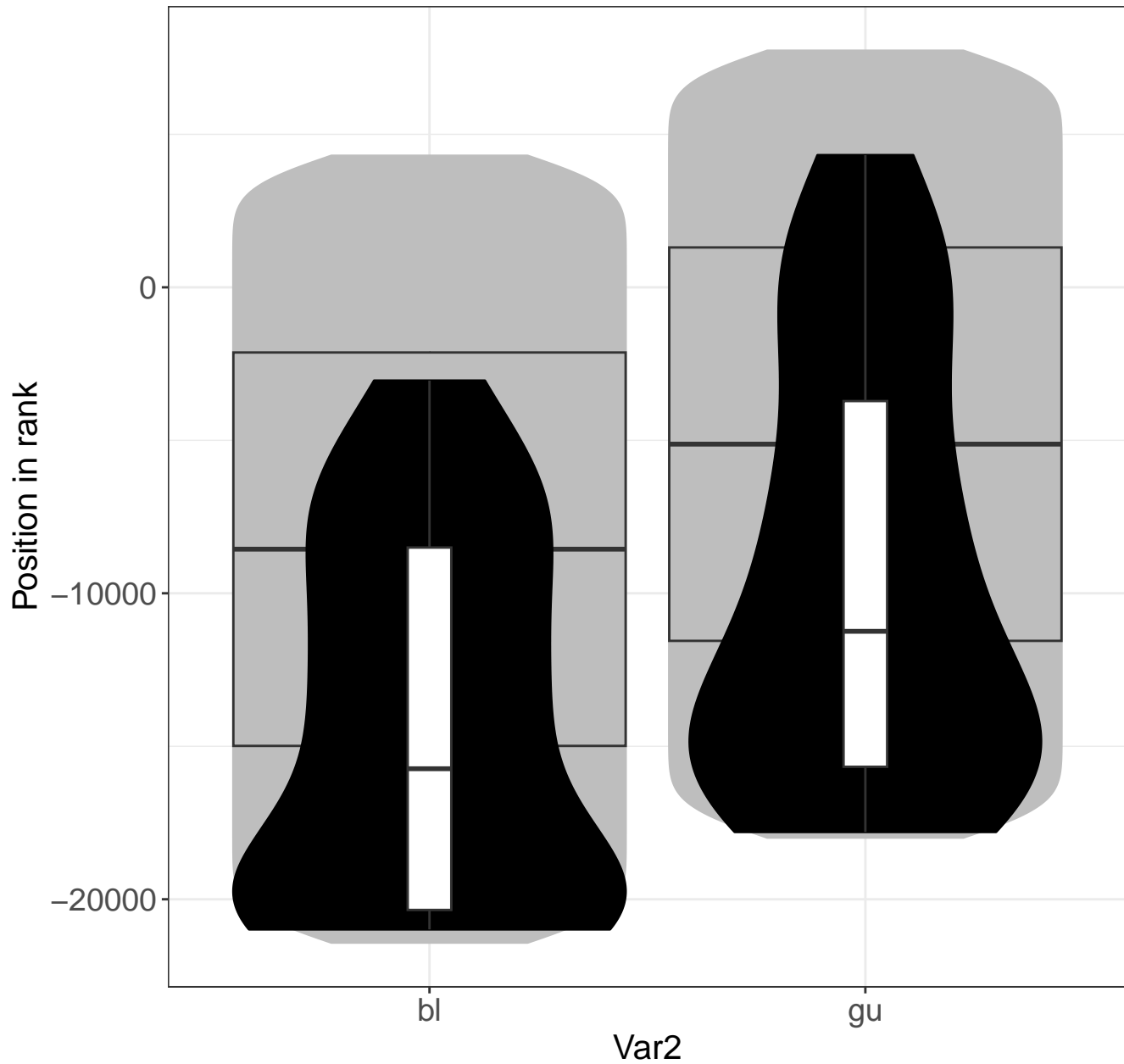
Sensory perception of sweet, bitter, and umami (glutamate)



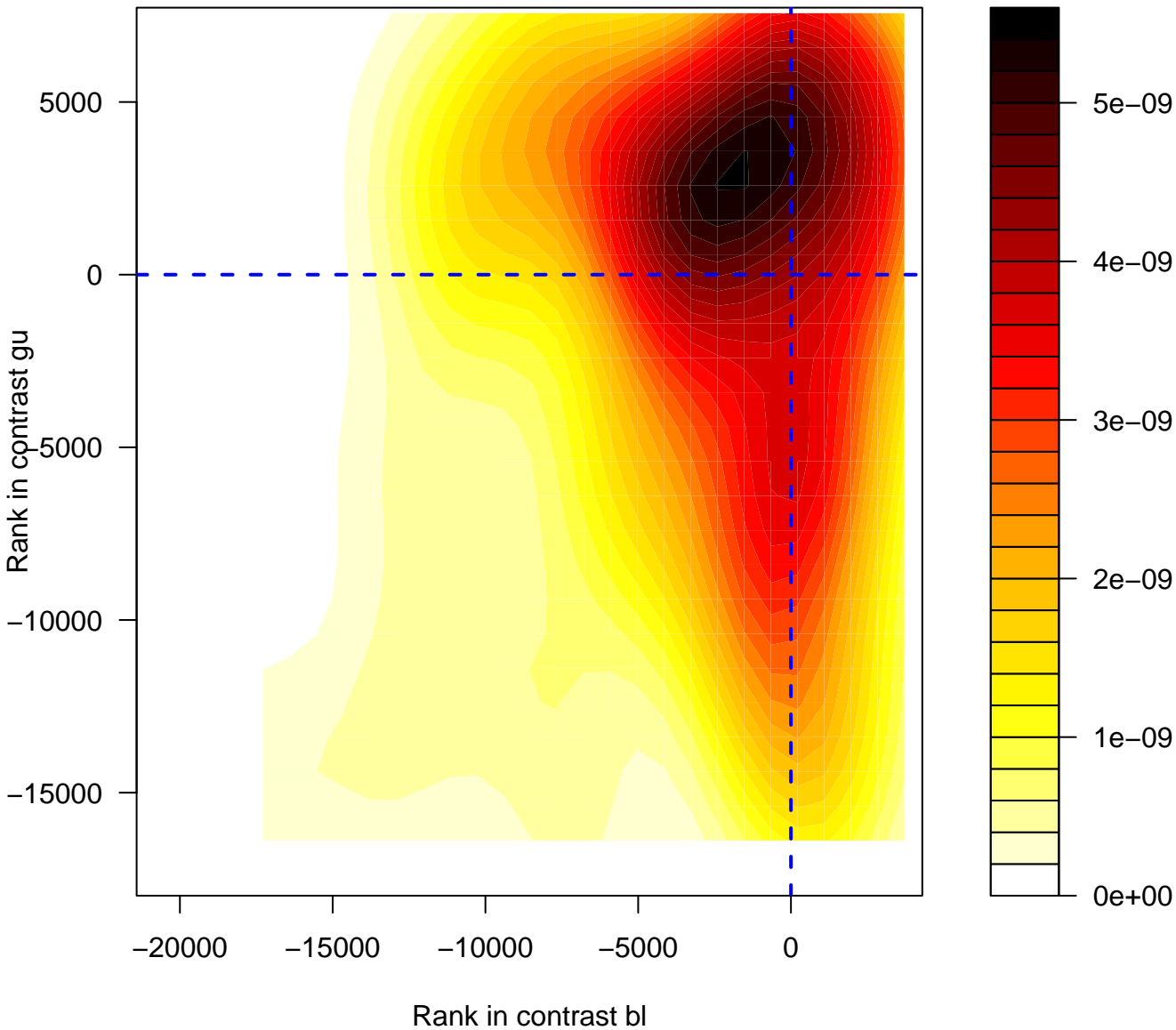
Sensory perception of sweet, bitter, and umami (glutamate) taste



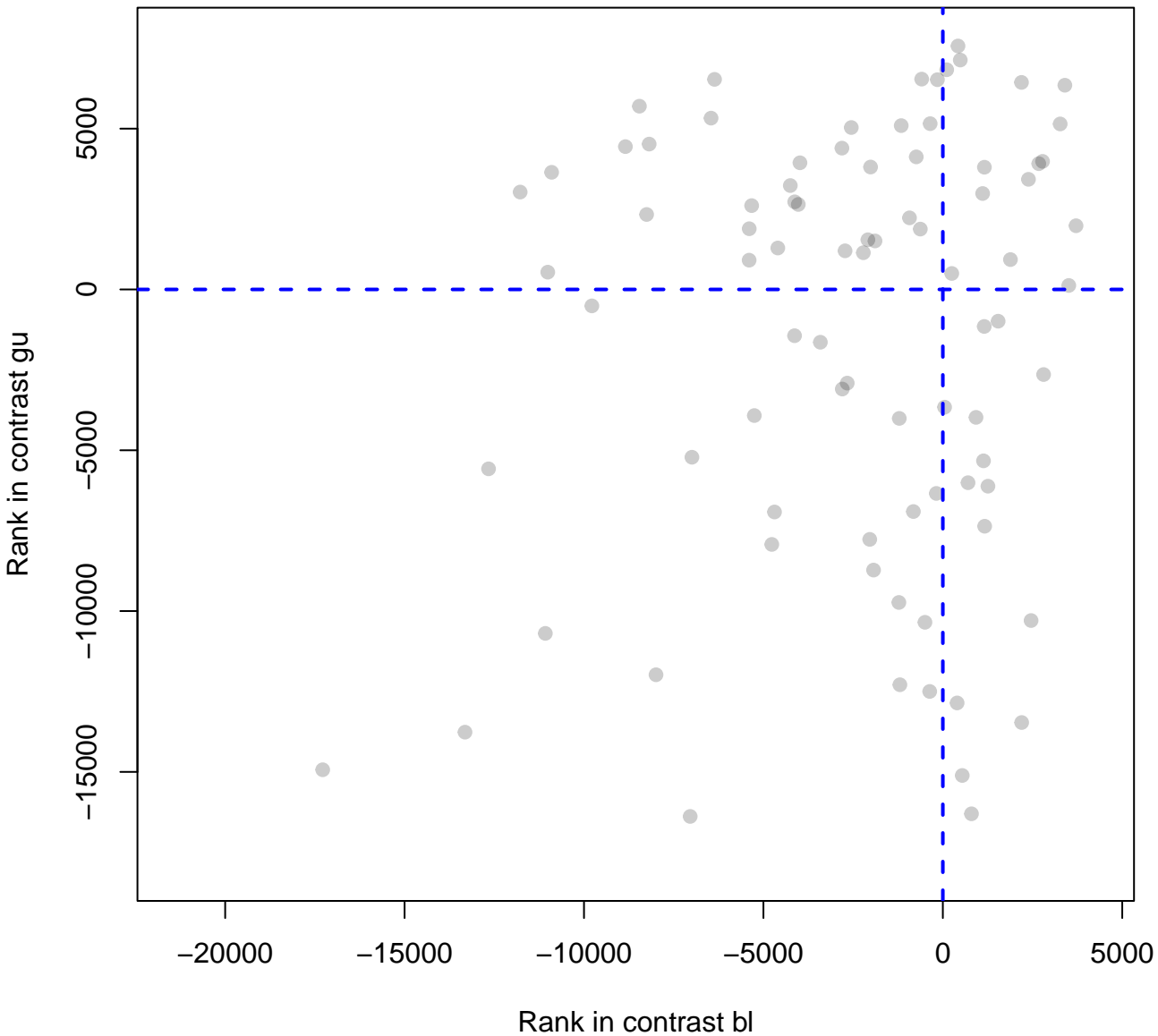
Sensory perception of sweet, bitter, and umami (



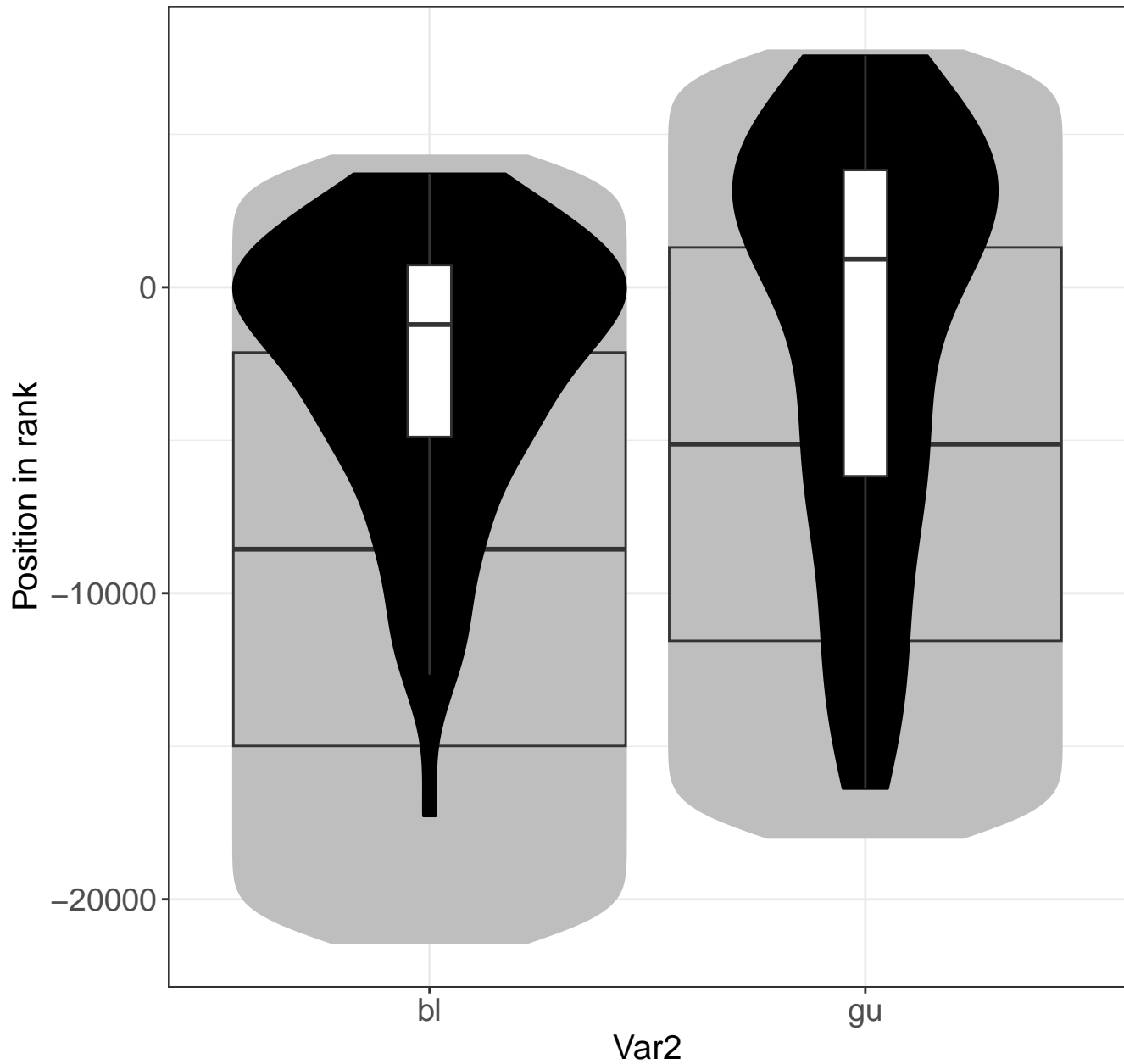
Peptide chain elongation



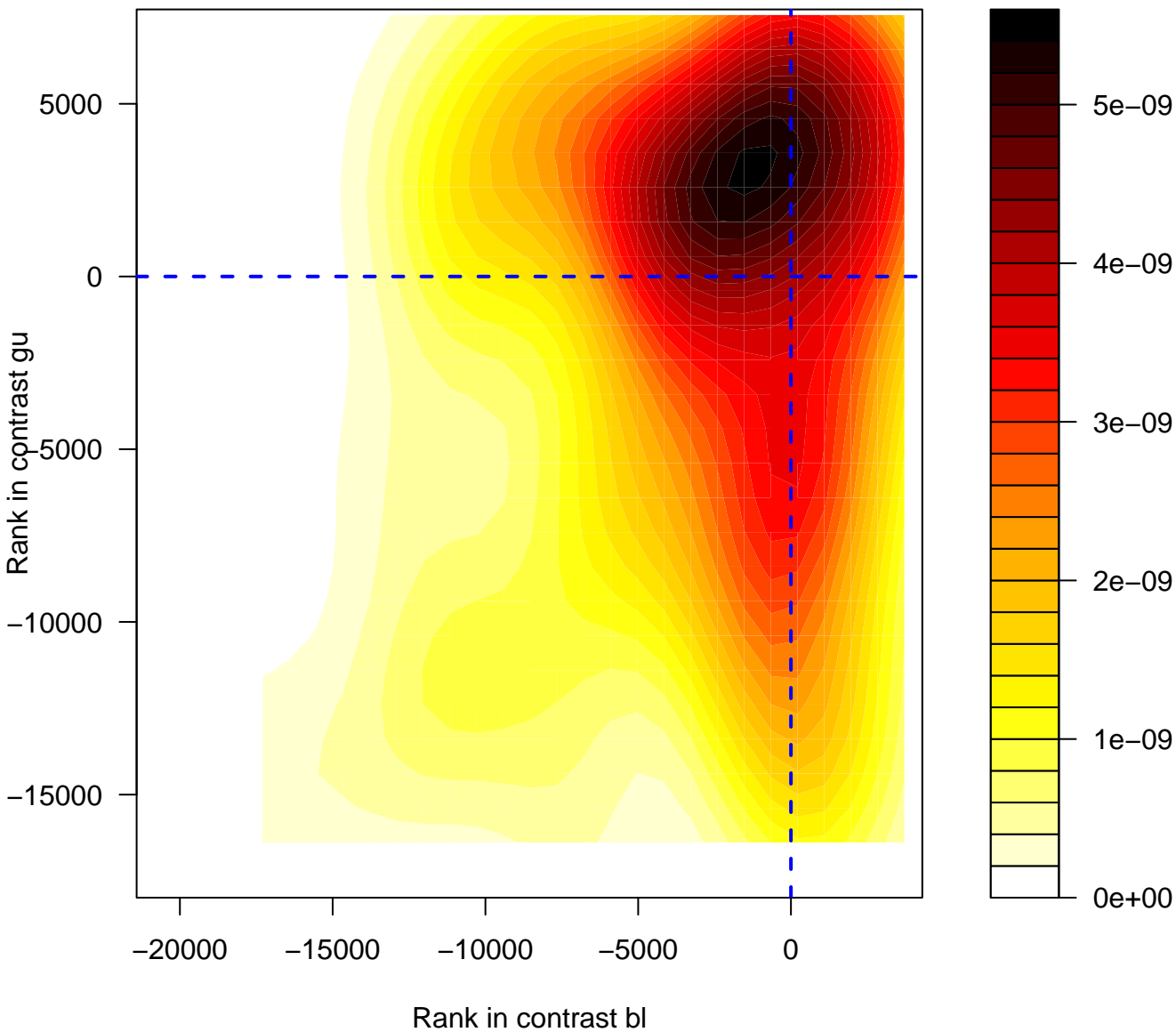
Peptide chain elongation



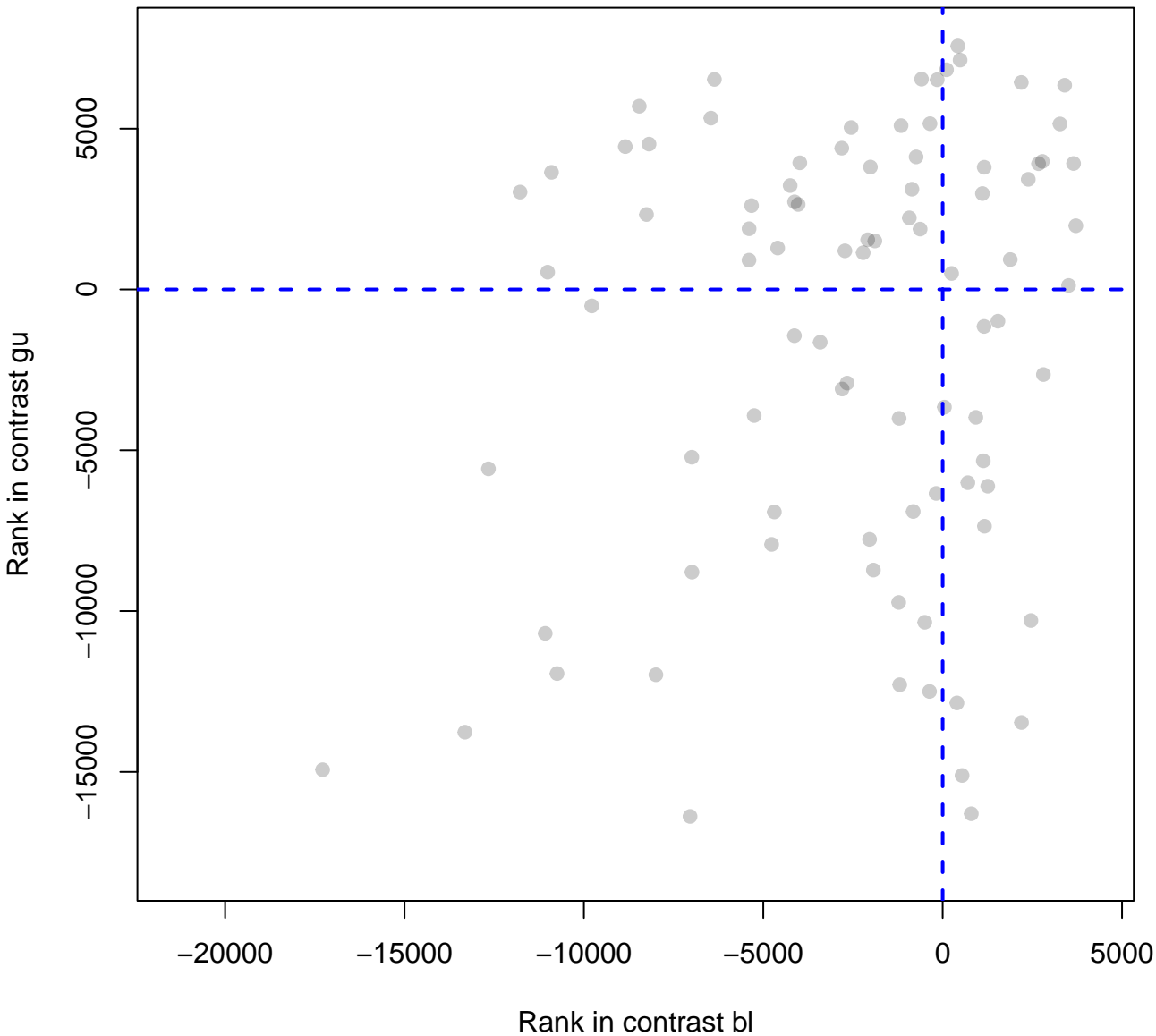
Peptide chain elongation



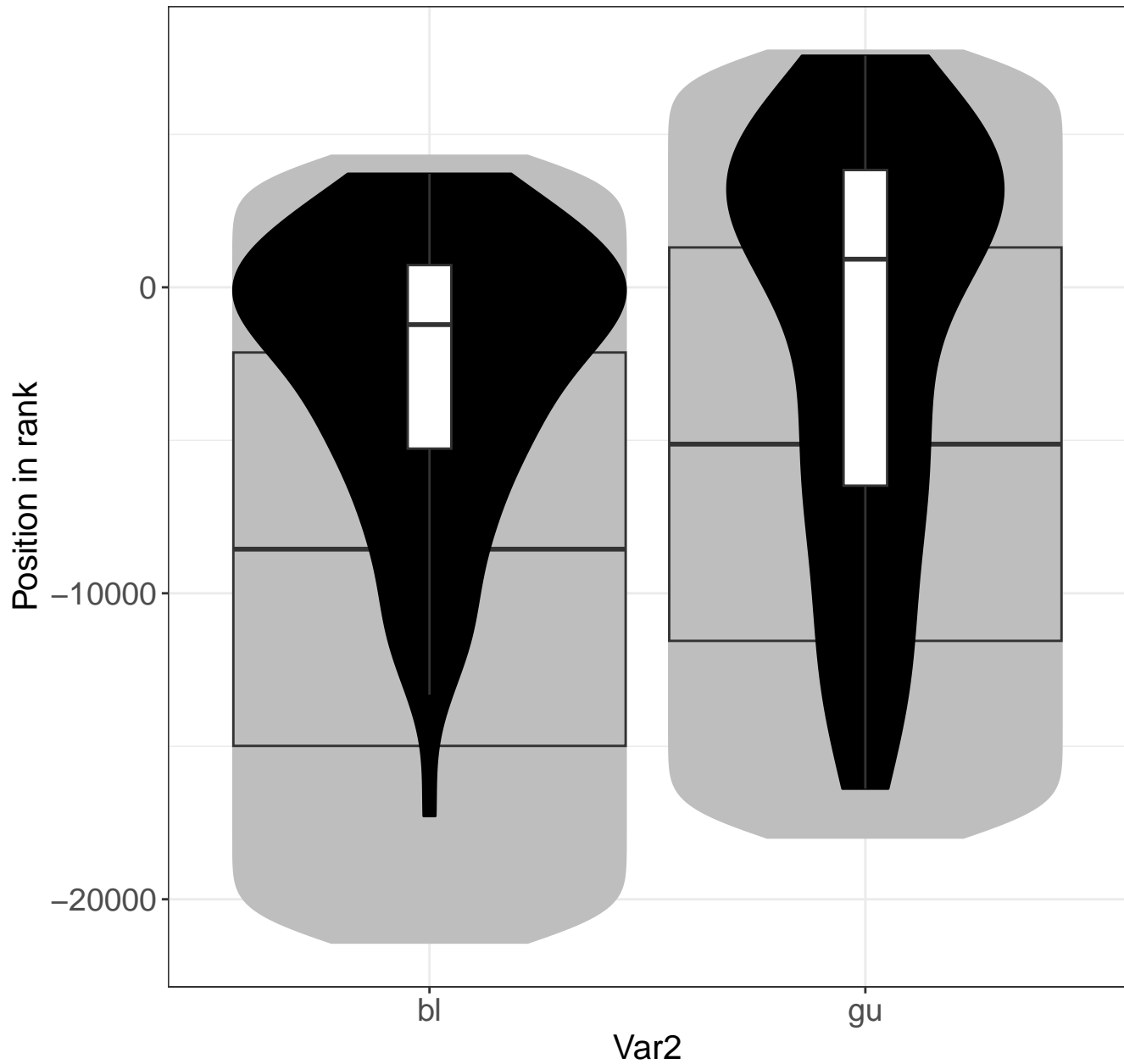
Eukaryotic Translation Elongation



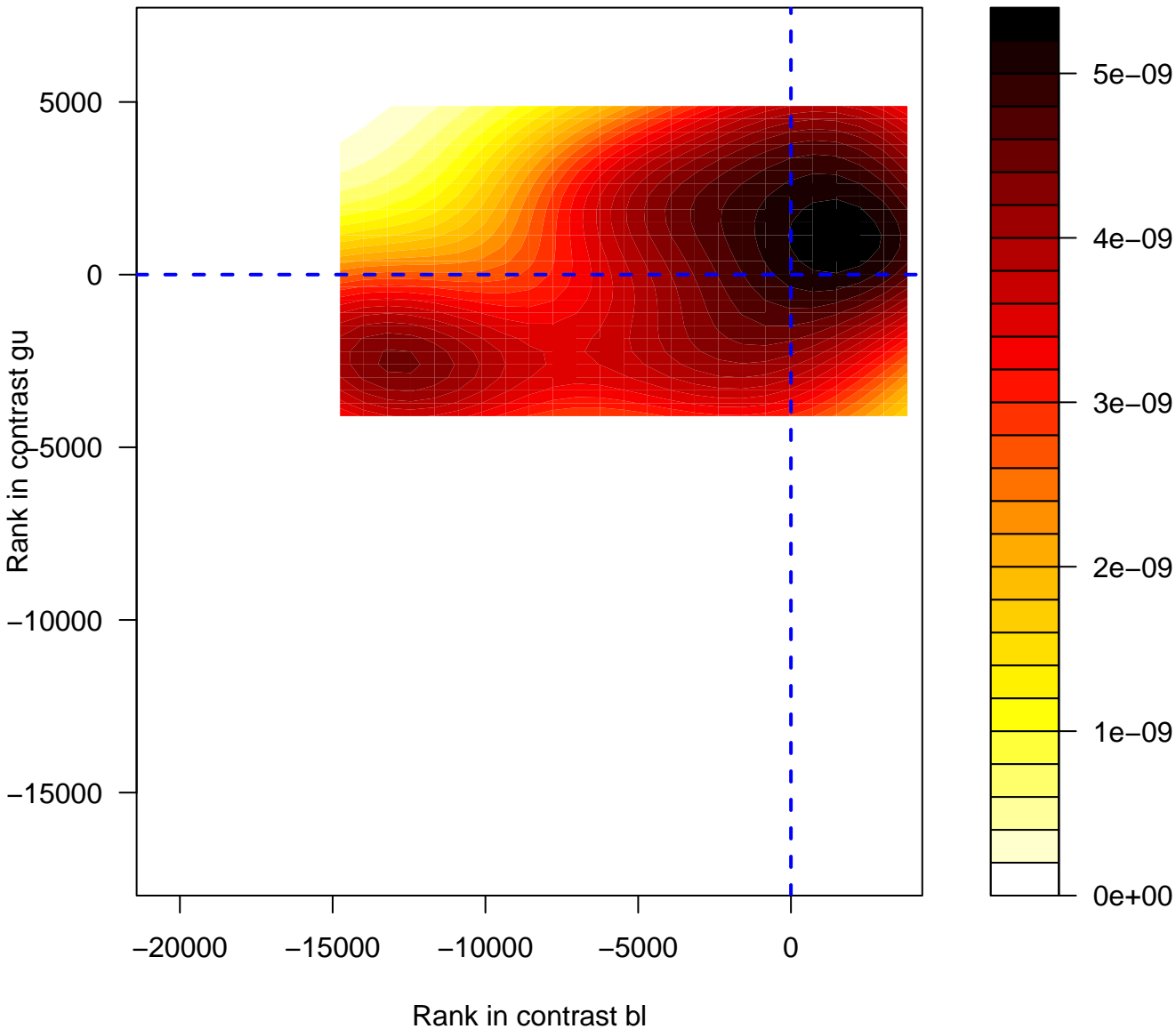
Eukaryotic Translation Elongation



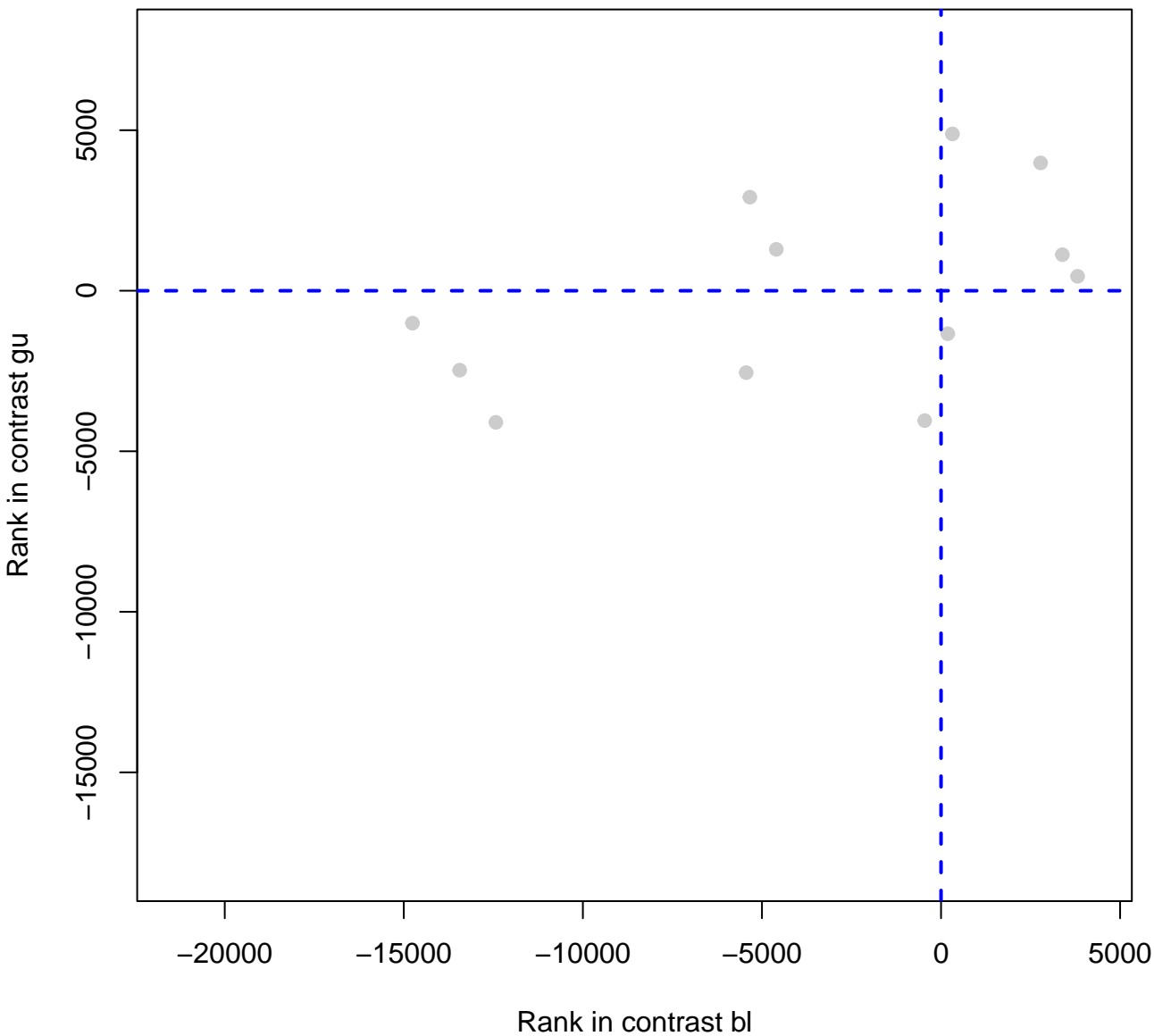
Eukaryotic Translation Elongation



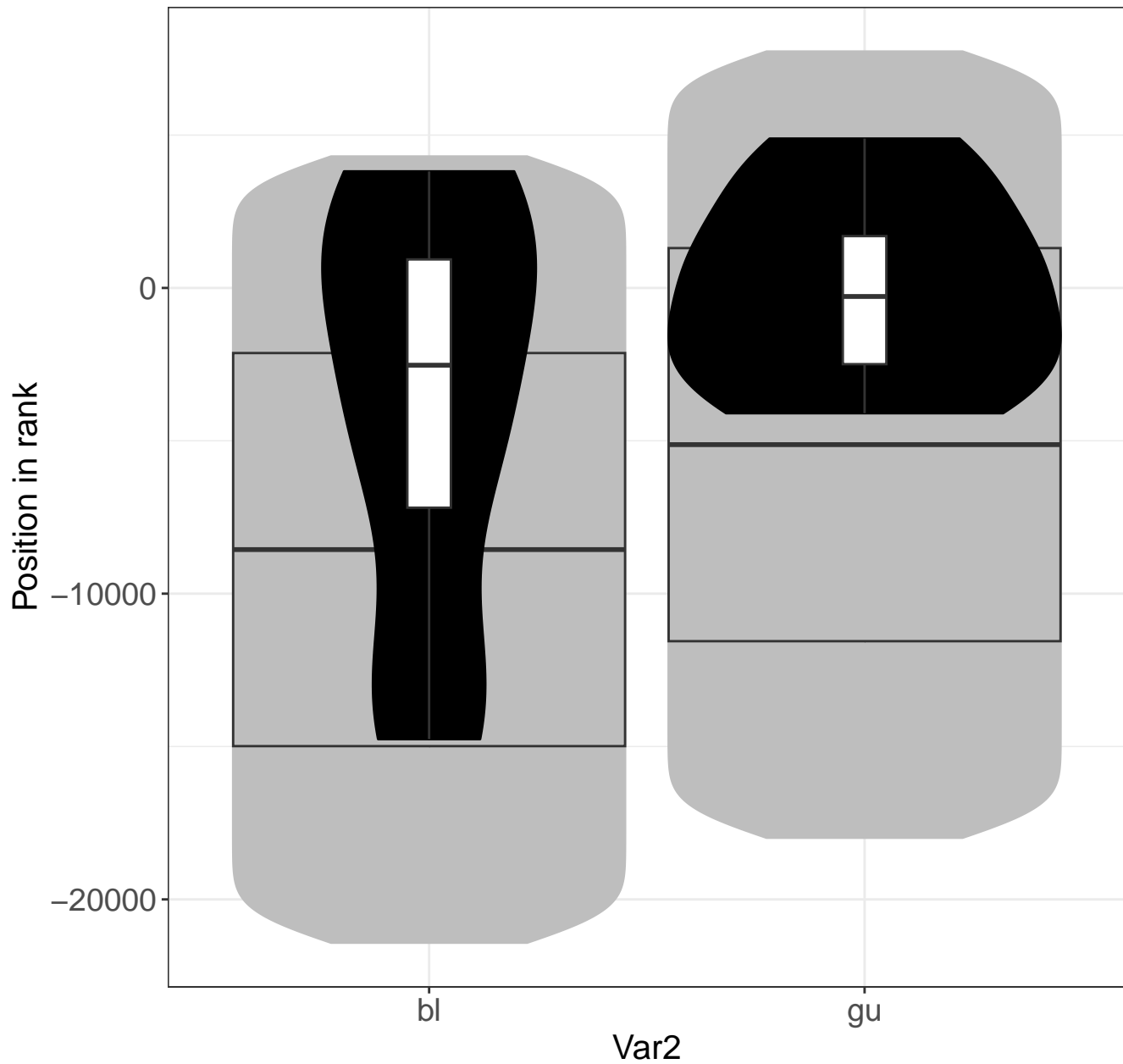
TICAM1-dependent activation of IRF3/IRF7



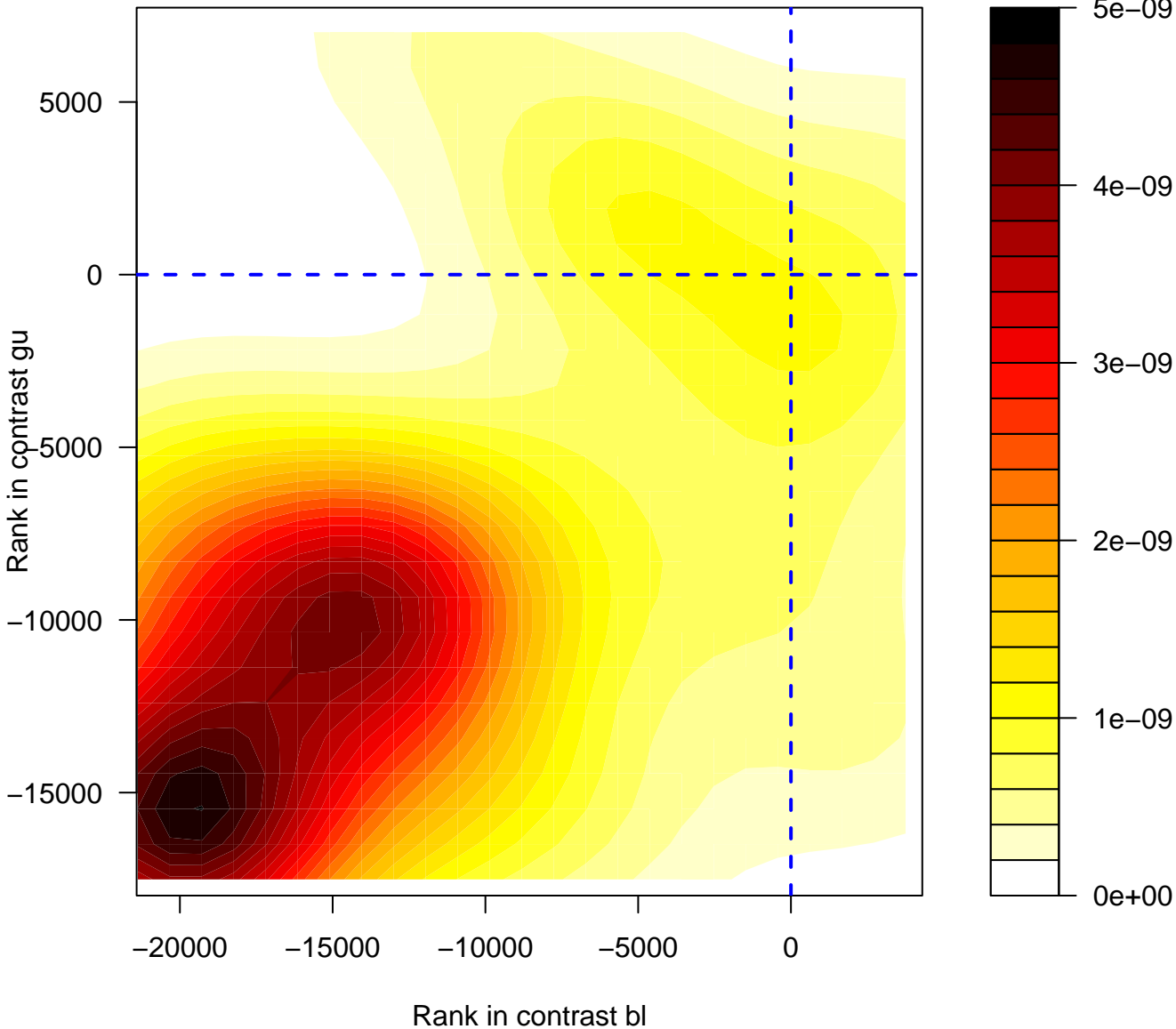
TICAM1-dependent activation of IRF3/IRF7



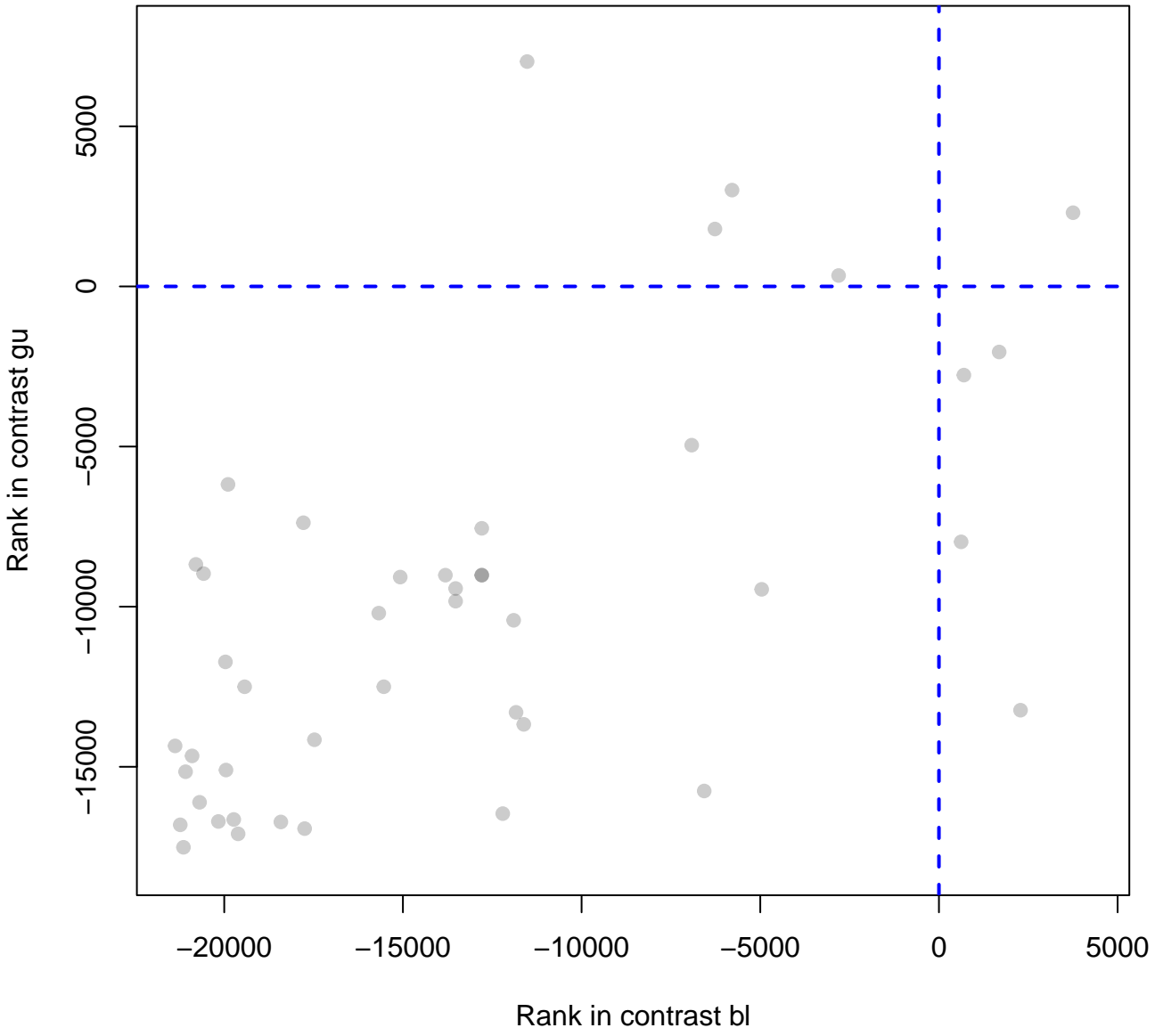
TICAM1-dependent activation of IRF3/IRF7



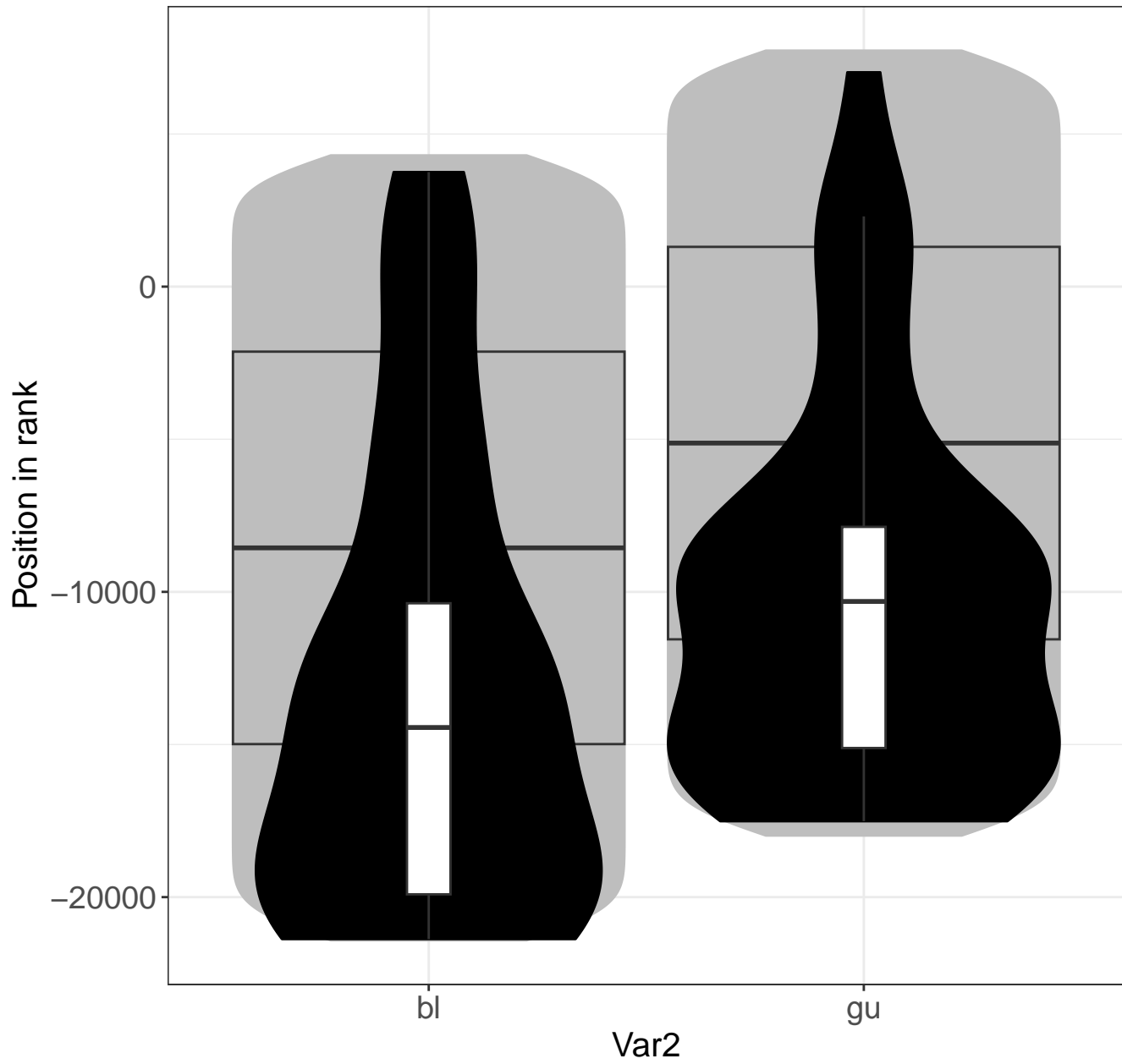
Aspirin ADME



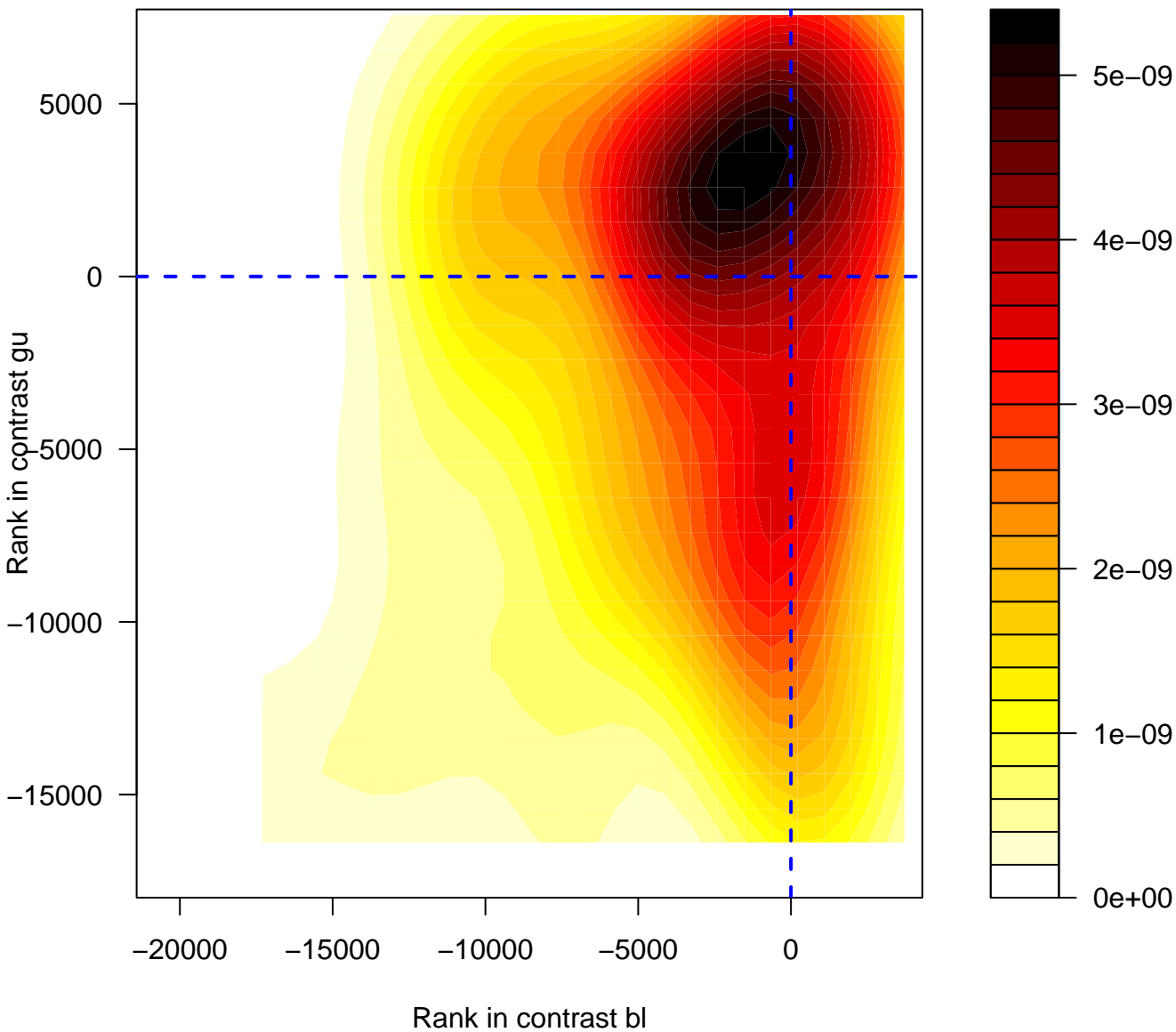
Aspirin ADME



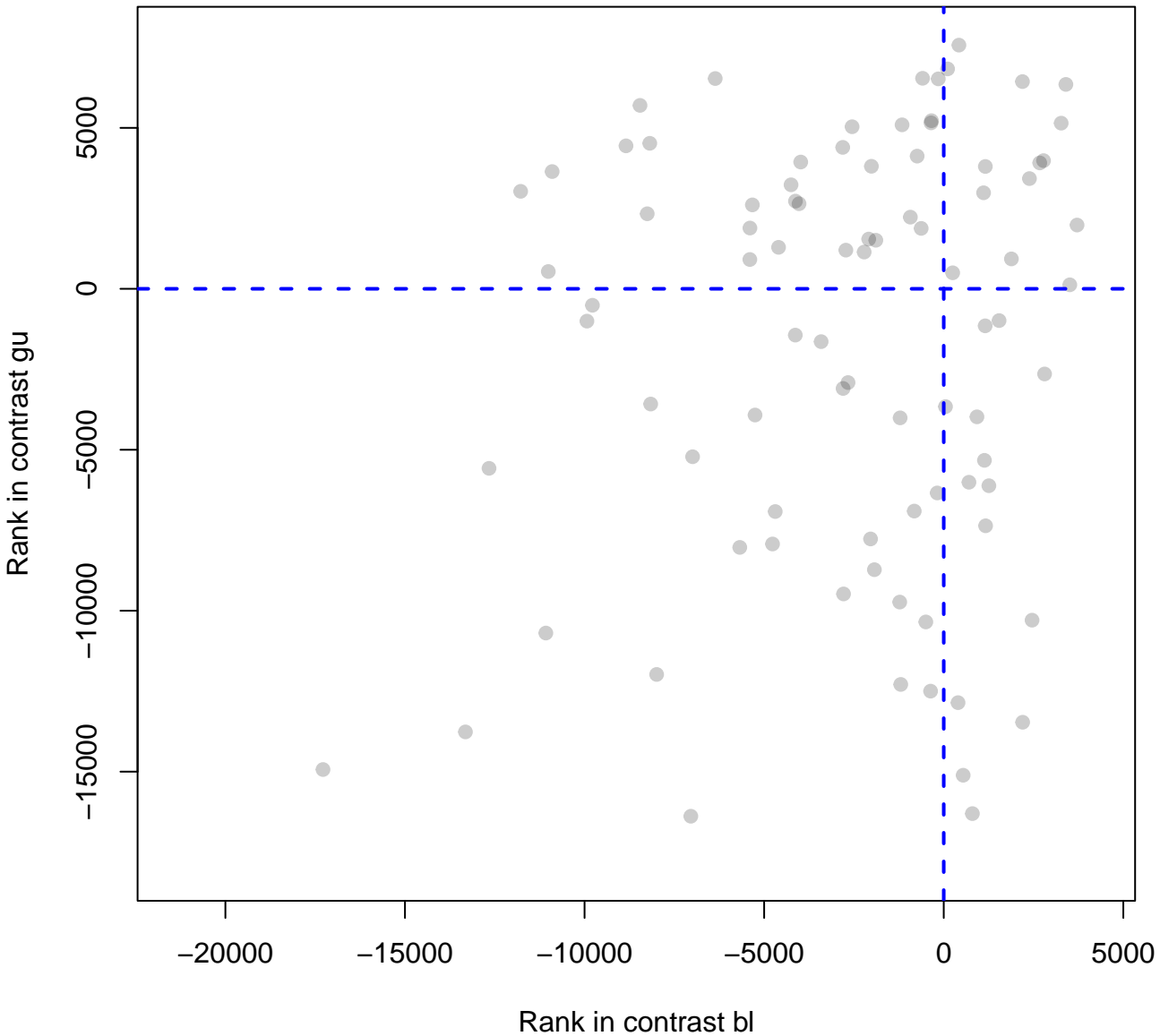
Aspirin ADME



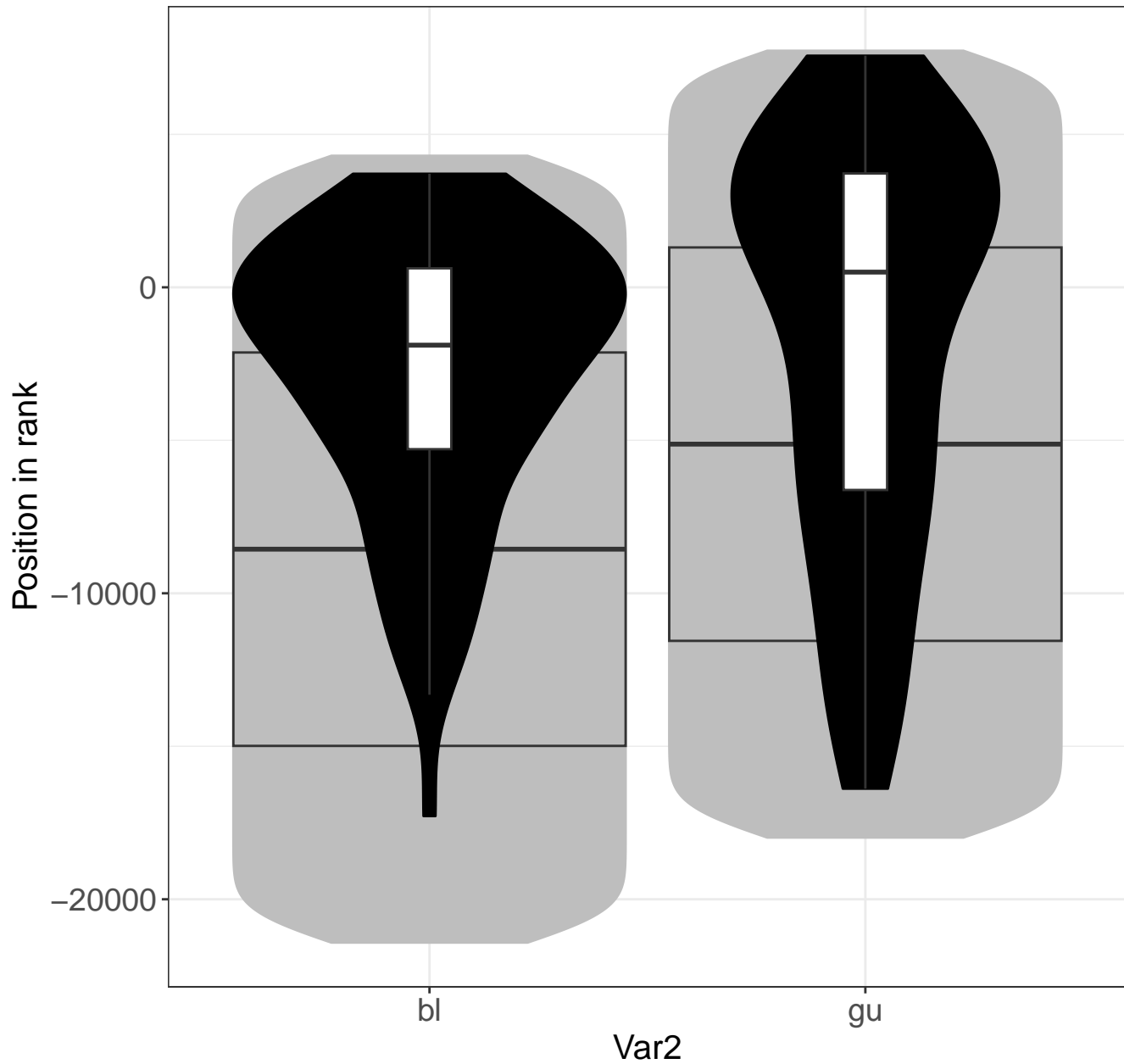
Eukaryotic Translation Termination



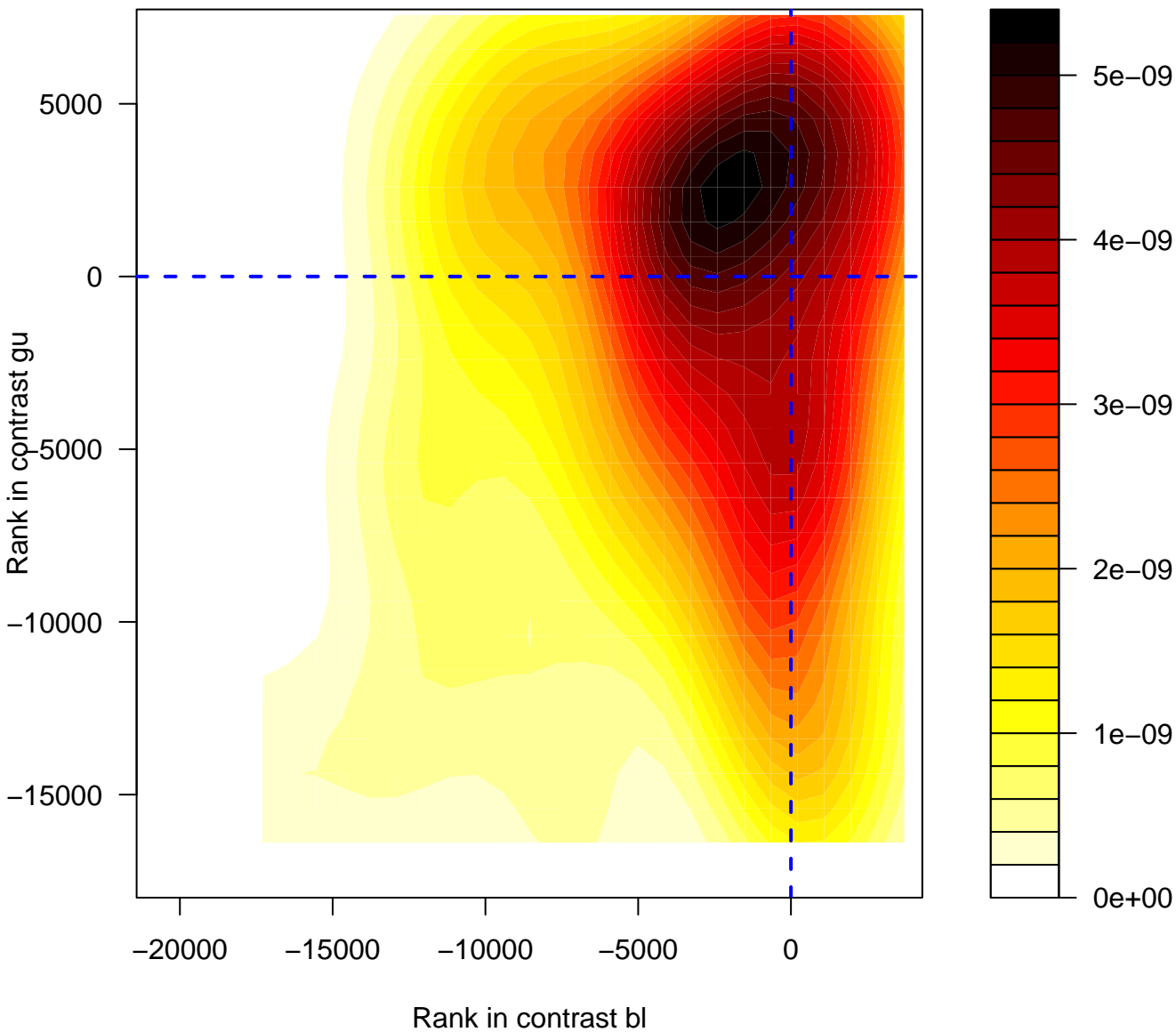
Eukaryotic Translation Termination



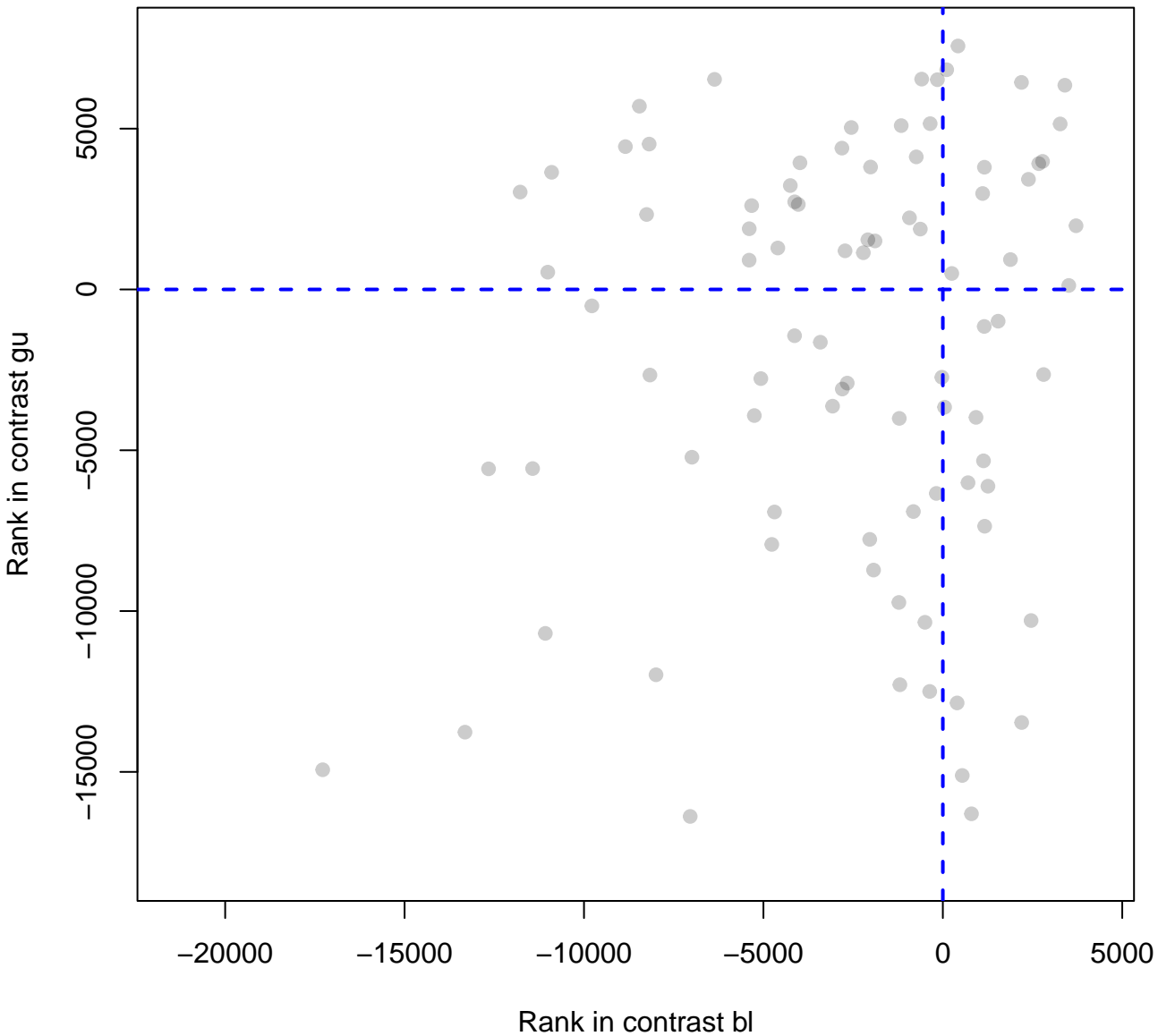
Eukaryotic Translation Termination



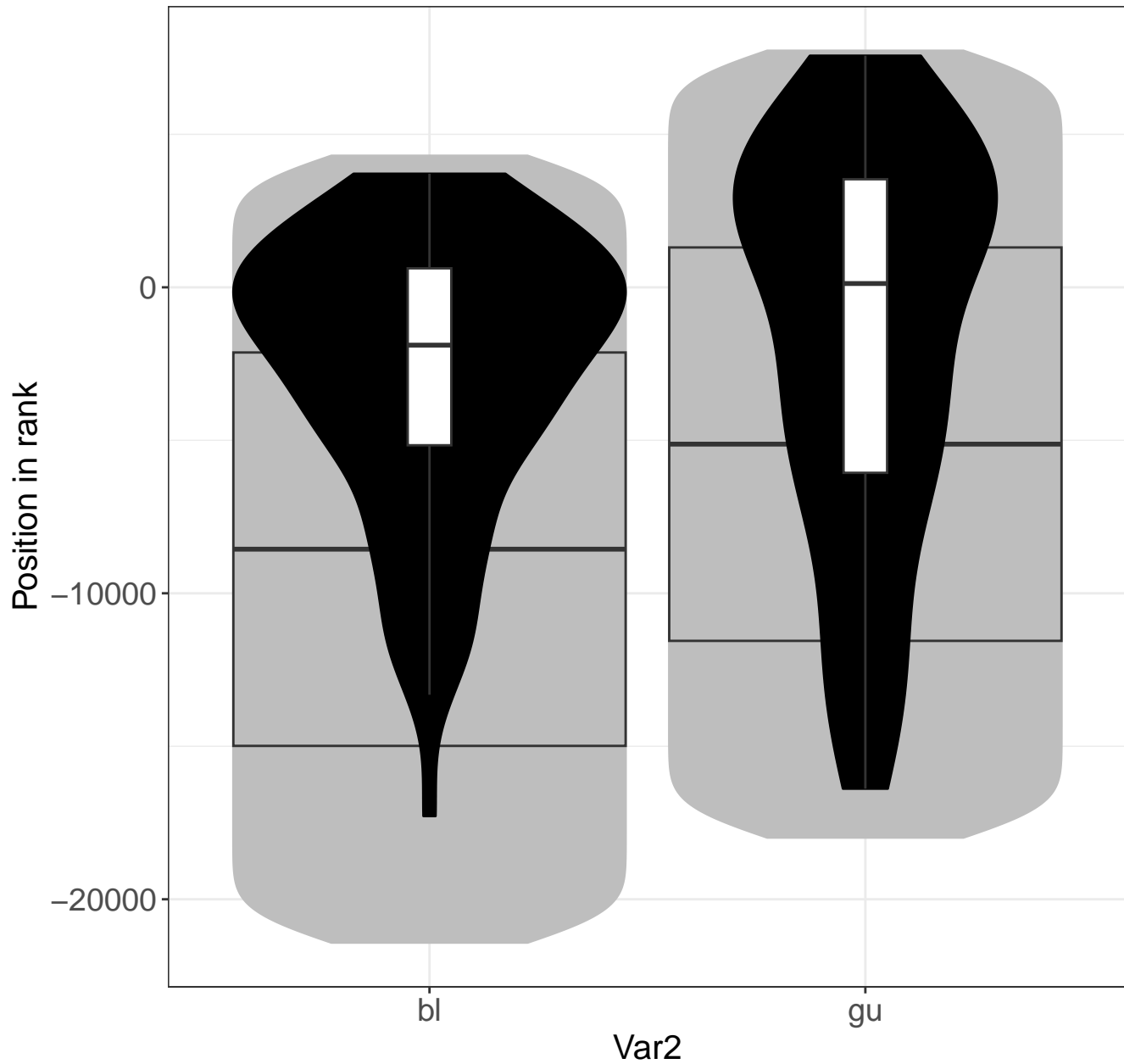
Selenocysteine synthesis



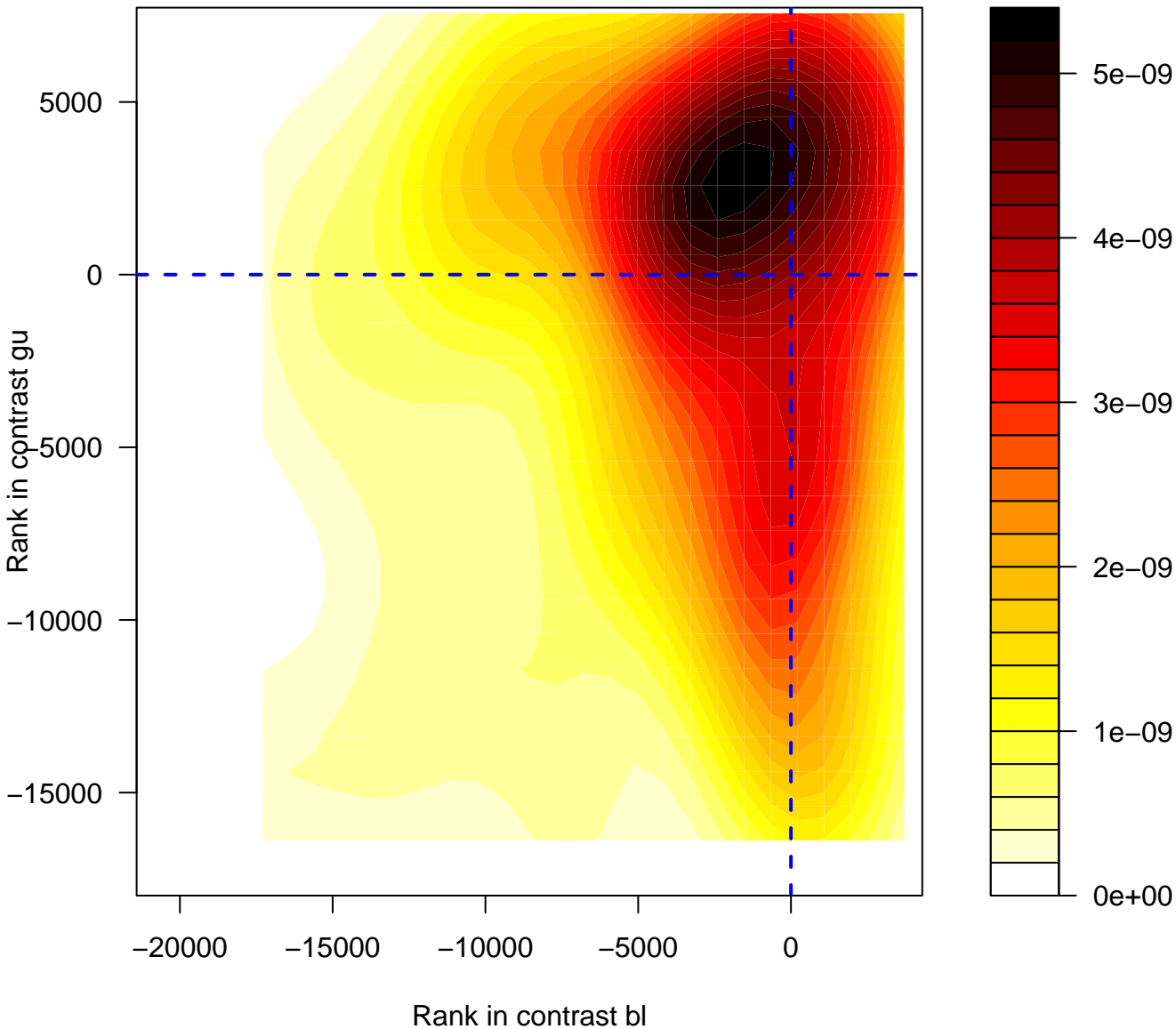
Selenocysteine synthesis



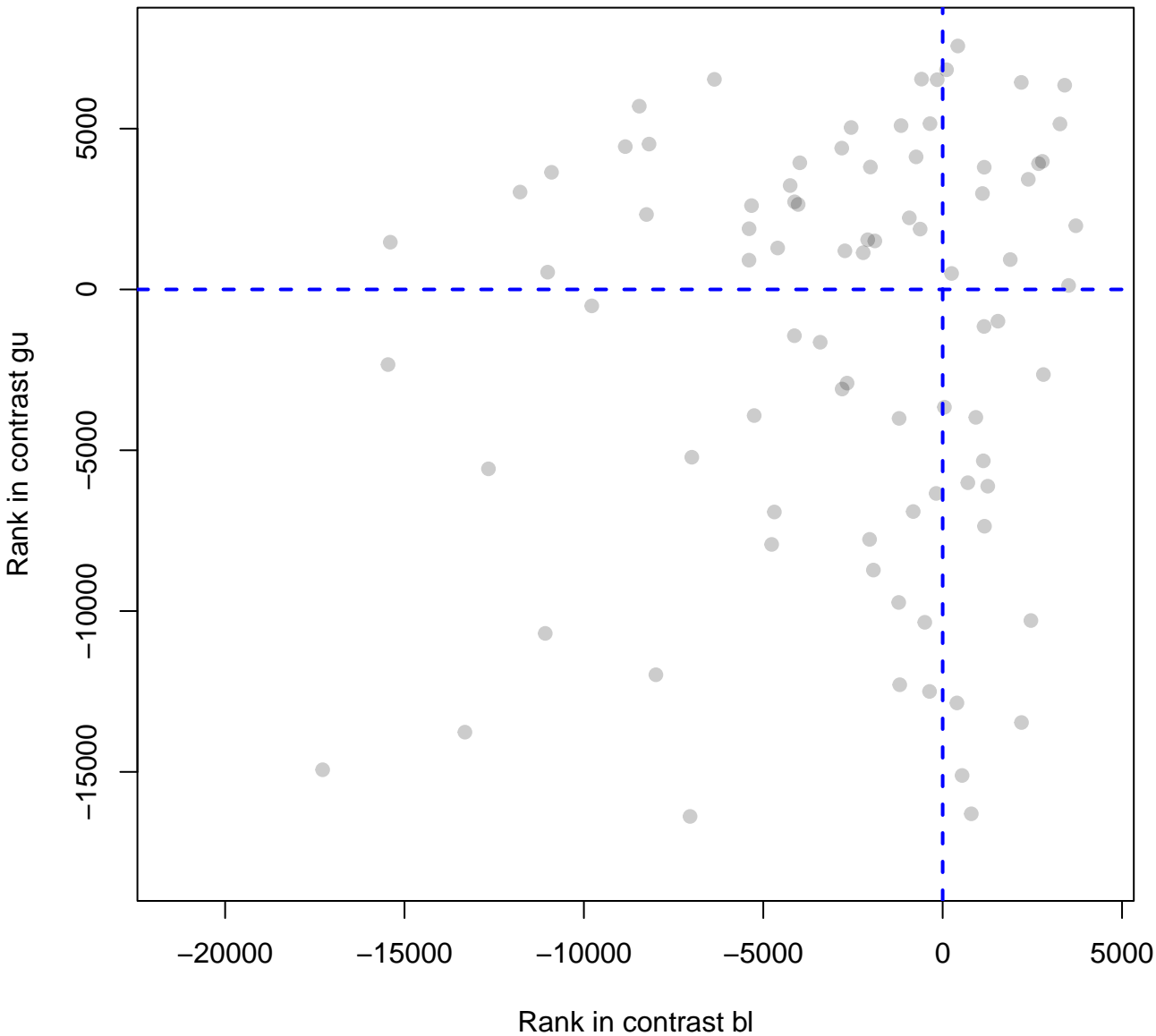
Selenocysteine synthesis



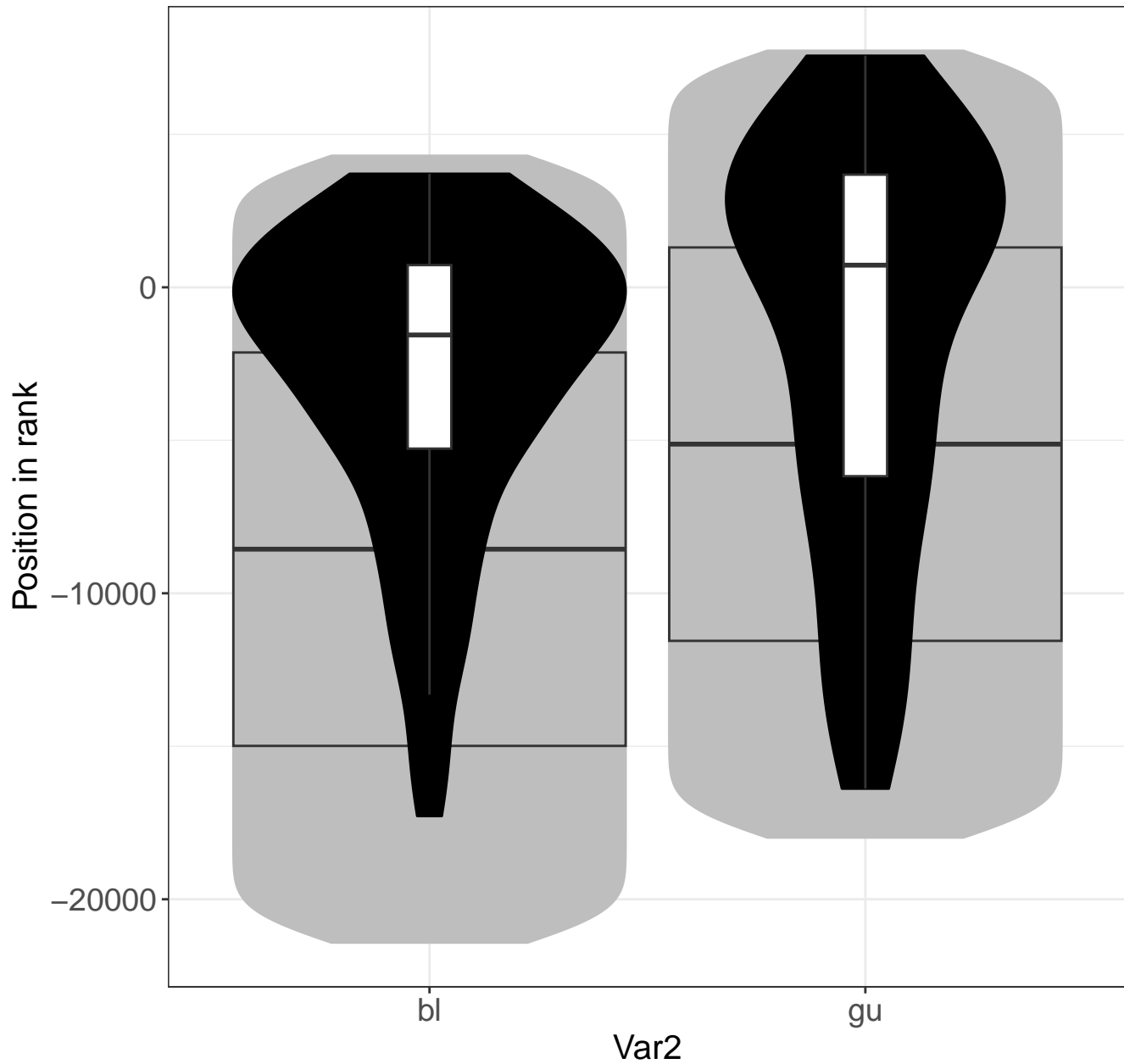
Viral mRNA Translation



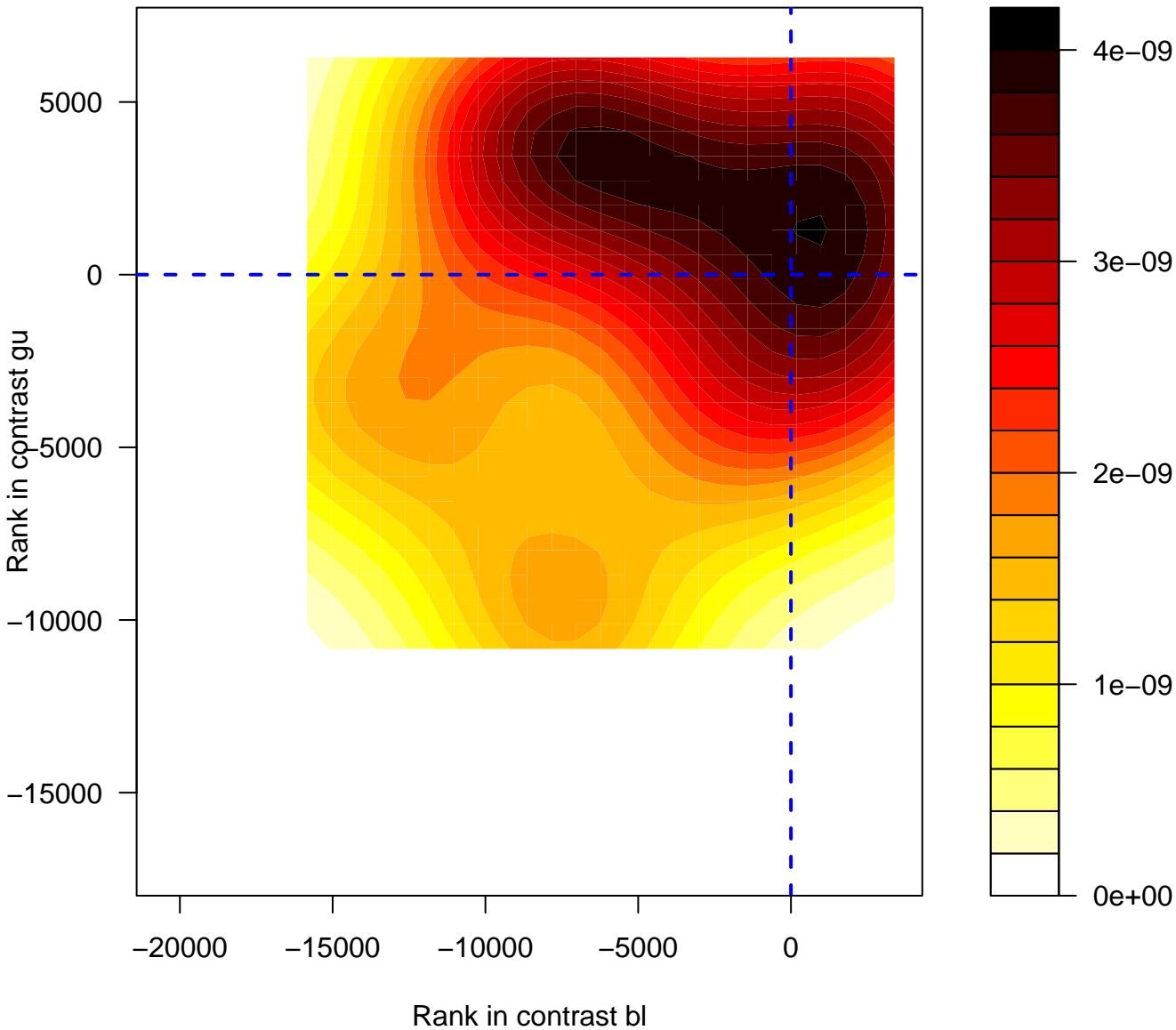
Viral mRNA Translation



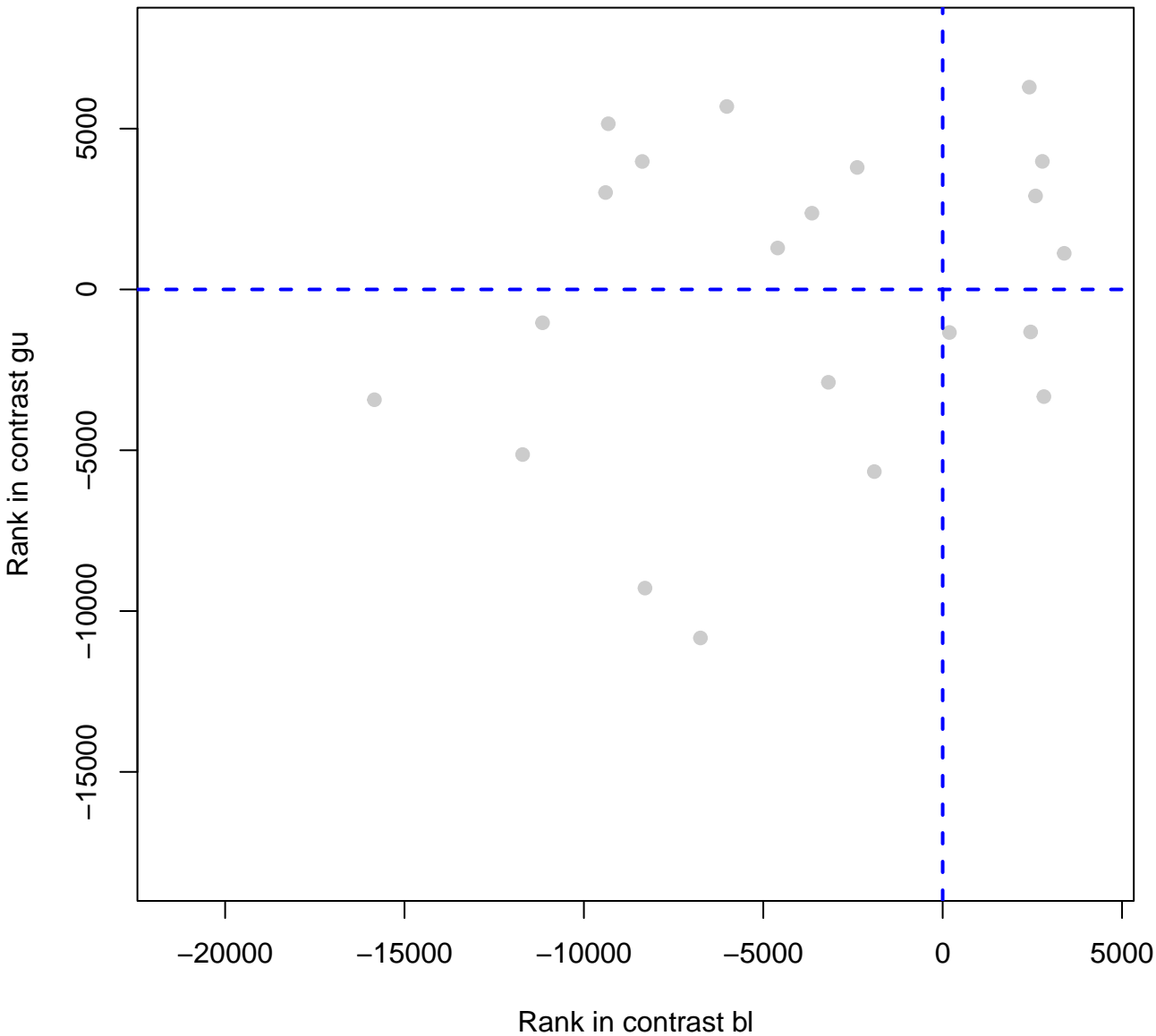
Viral mRNA Translation



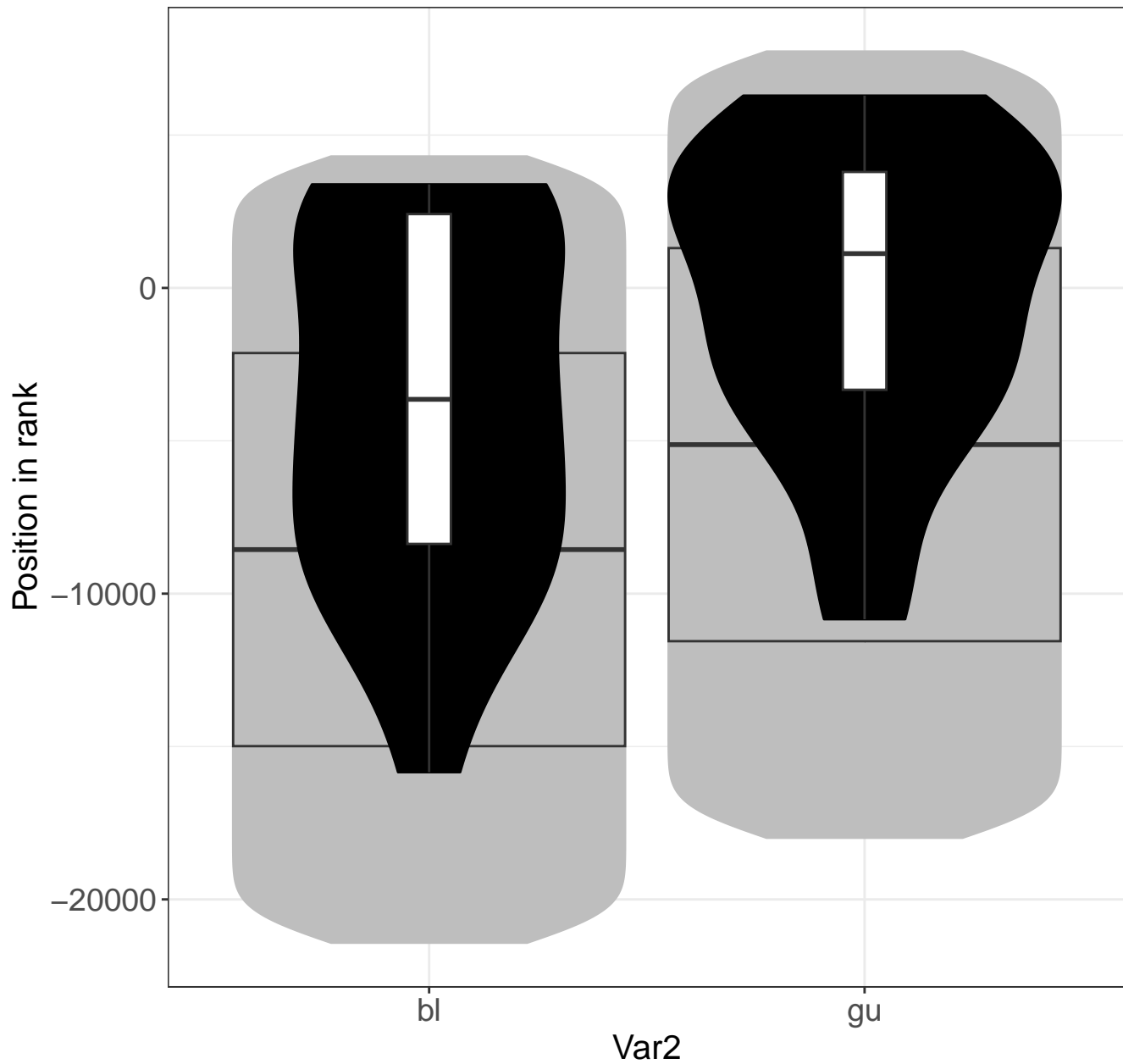
PINK1-PRKN Mediated Mitophagy



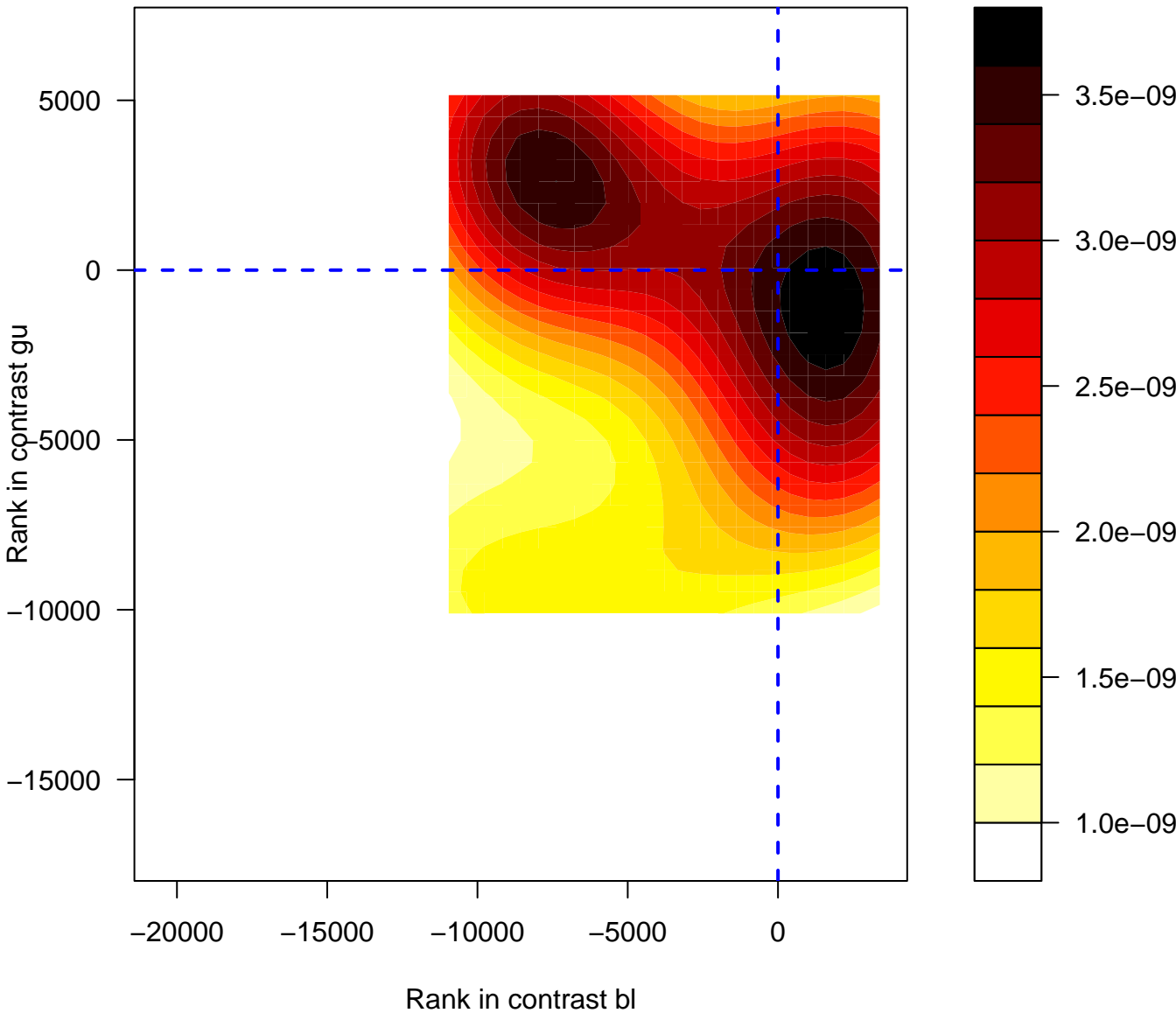
PINK1-PRKN Mediated Mitophagy



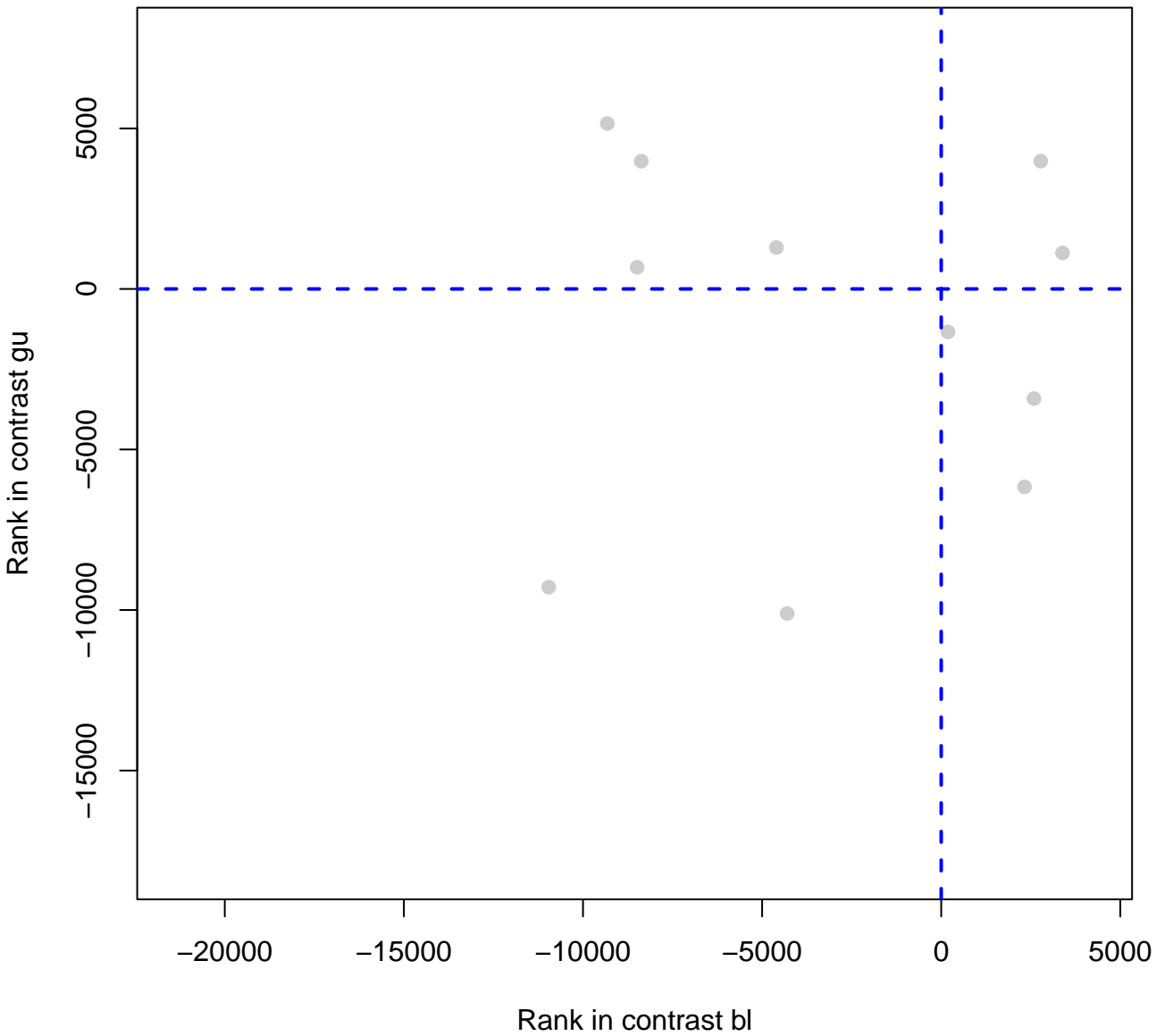
PINK1-PRKN Mediated Mitophagy



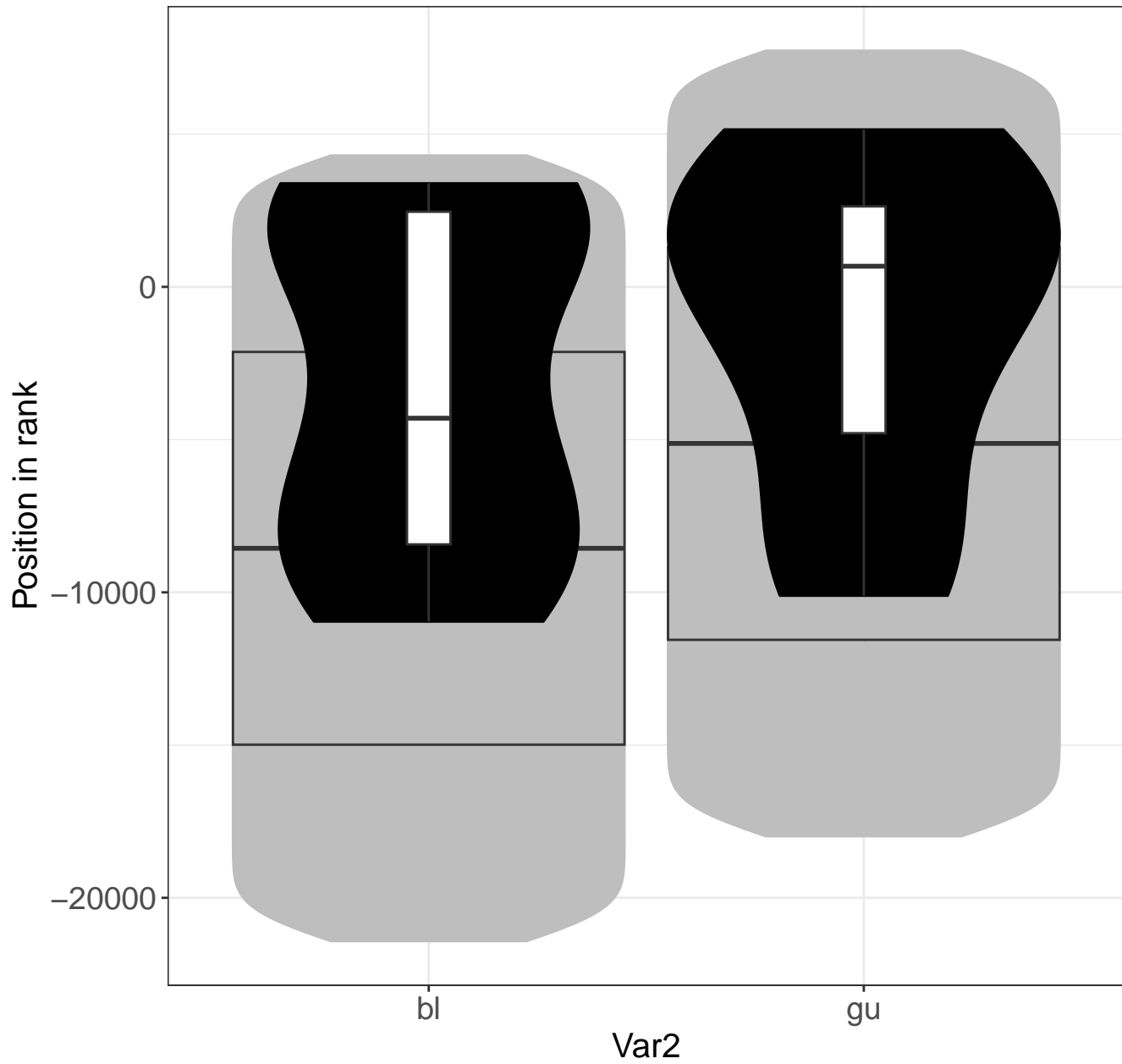
Pexophagy



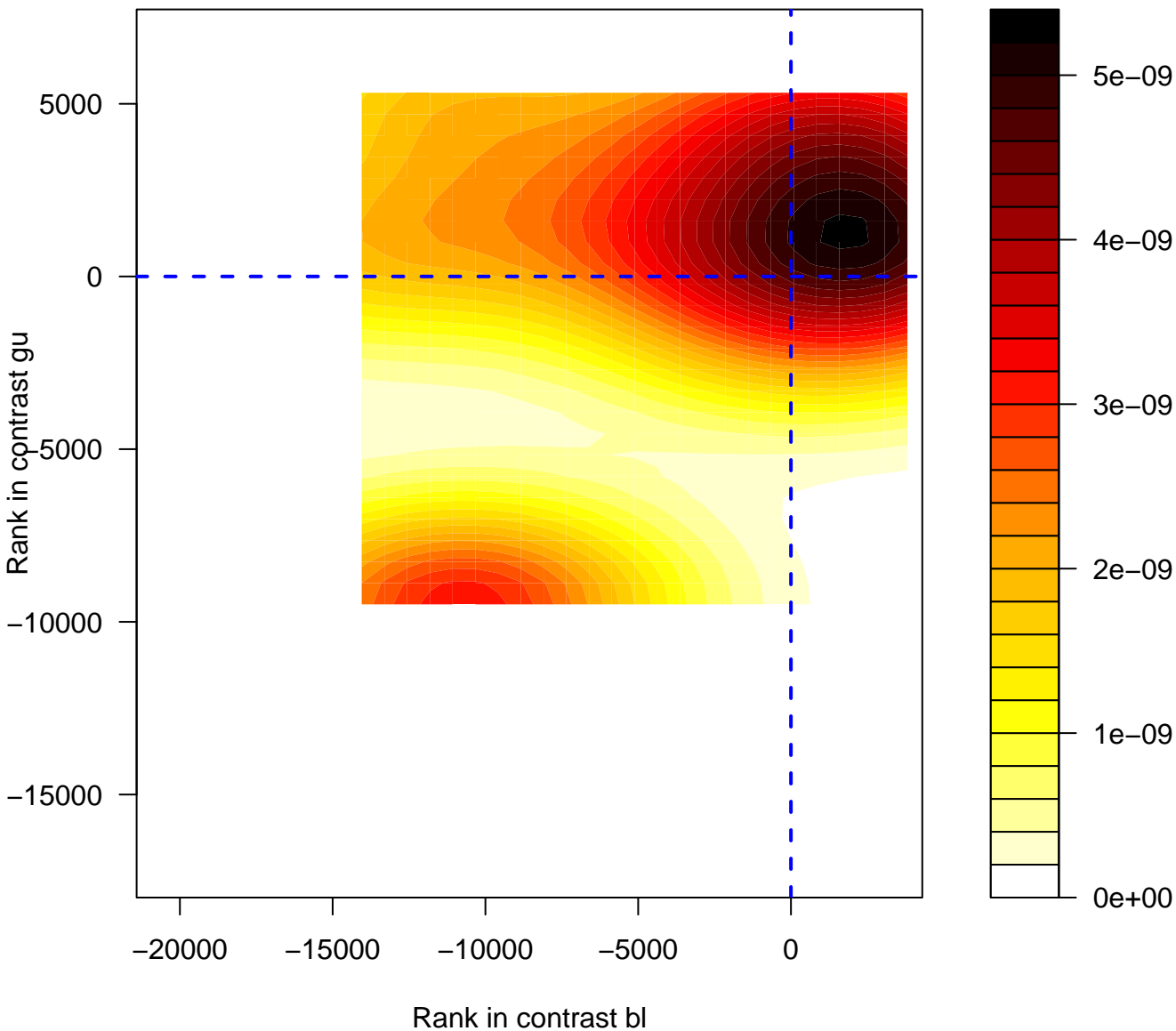
Pexophagy



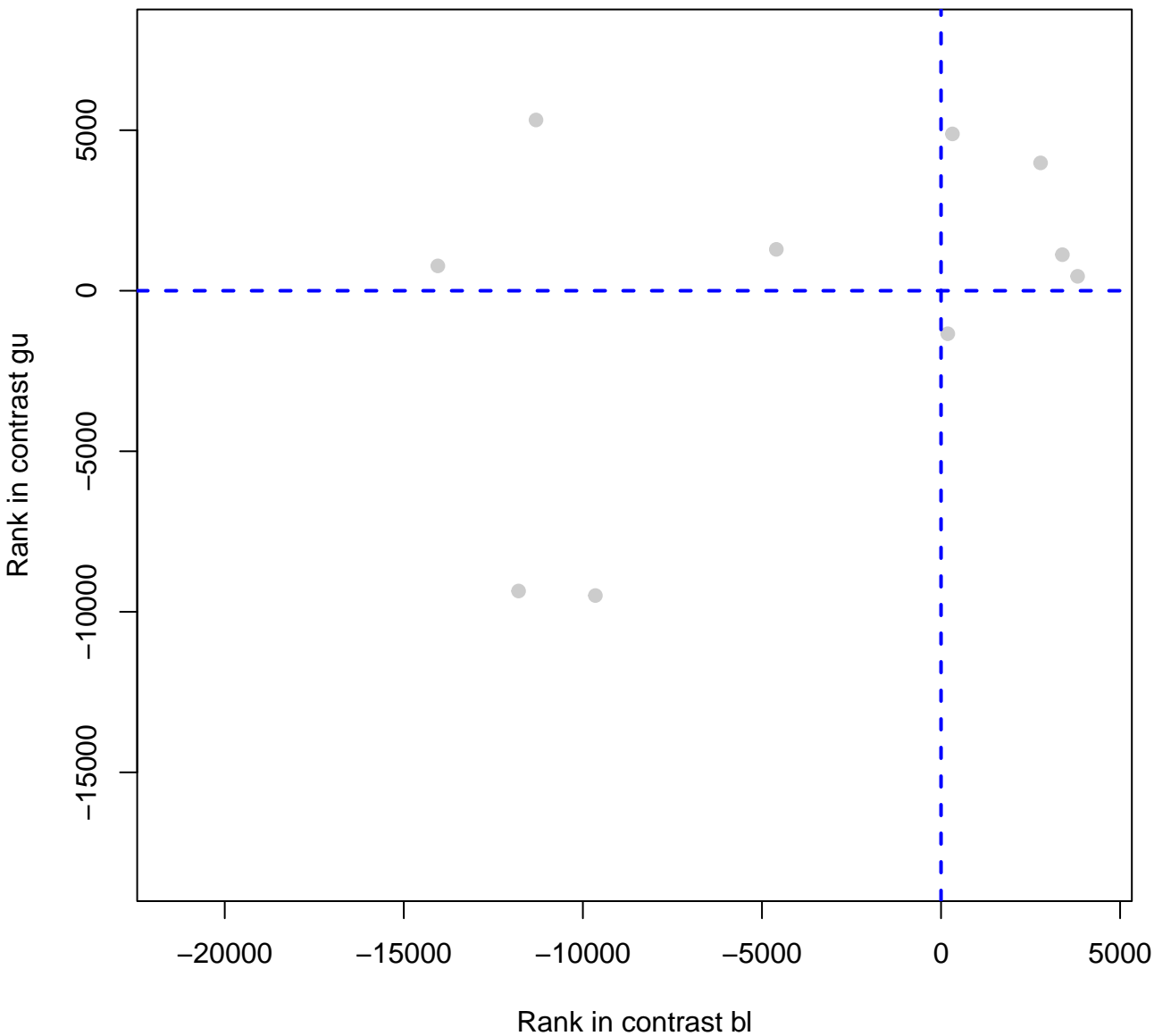
Pexophagy



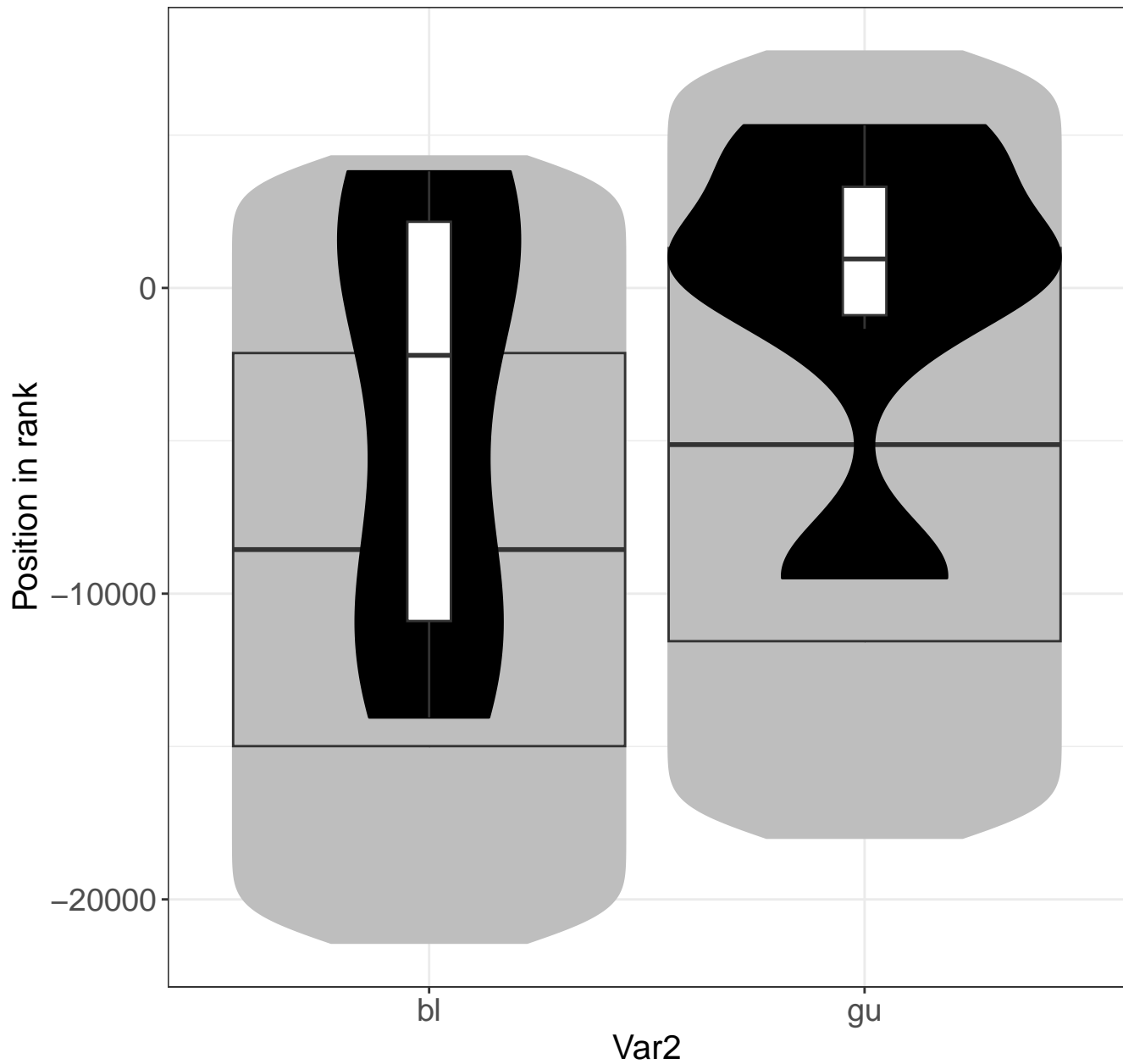
TICAM1,TRAF6-dependent induction of TAK1 complex



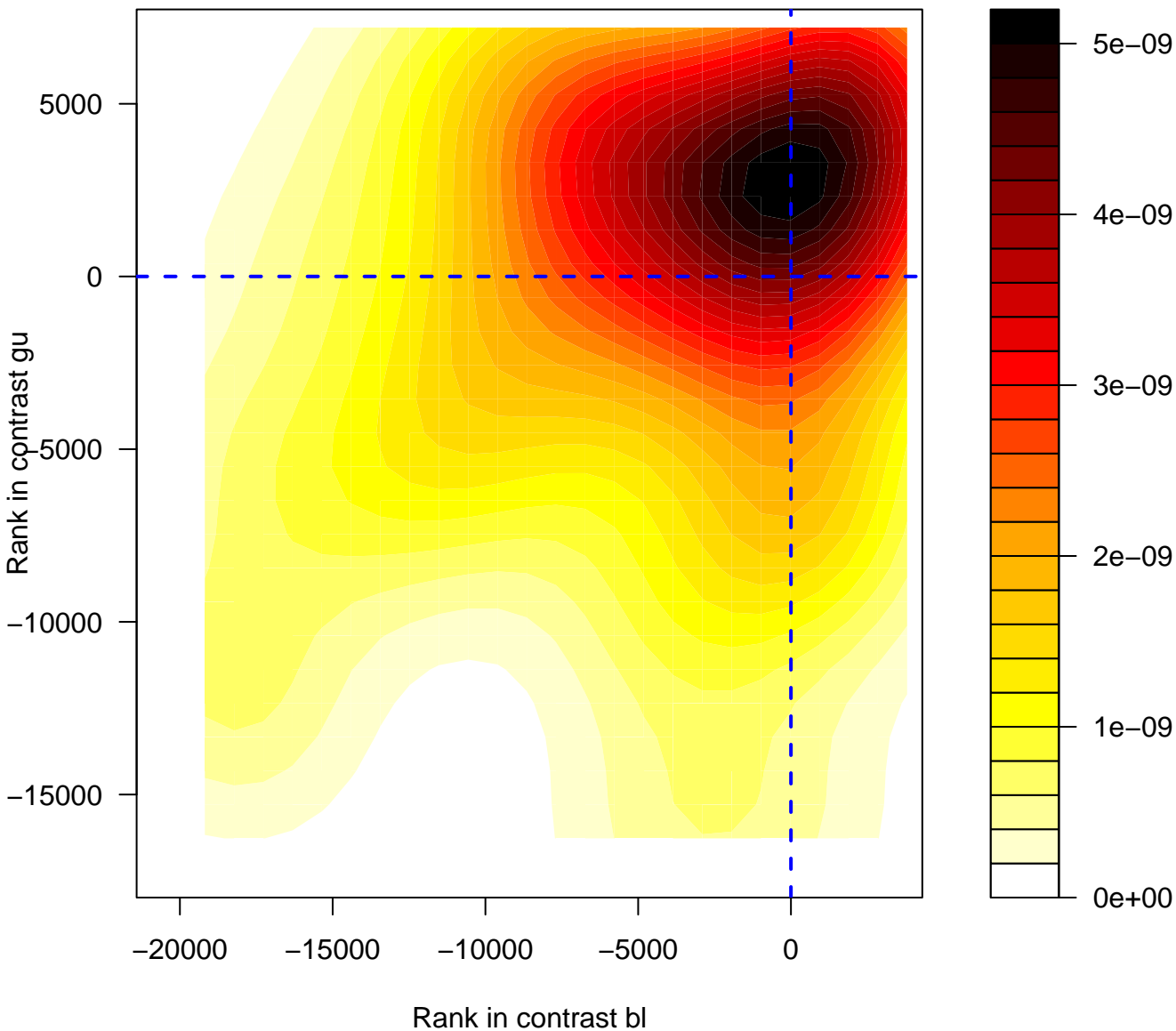
TICAM1, TRAF6-dependent induction of TAK1 complex



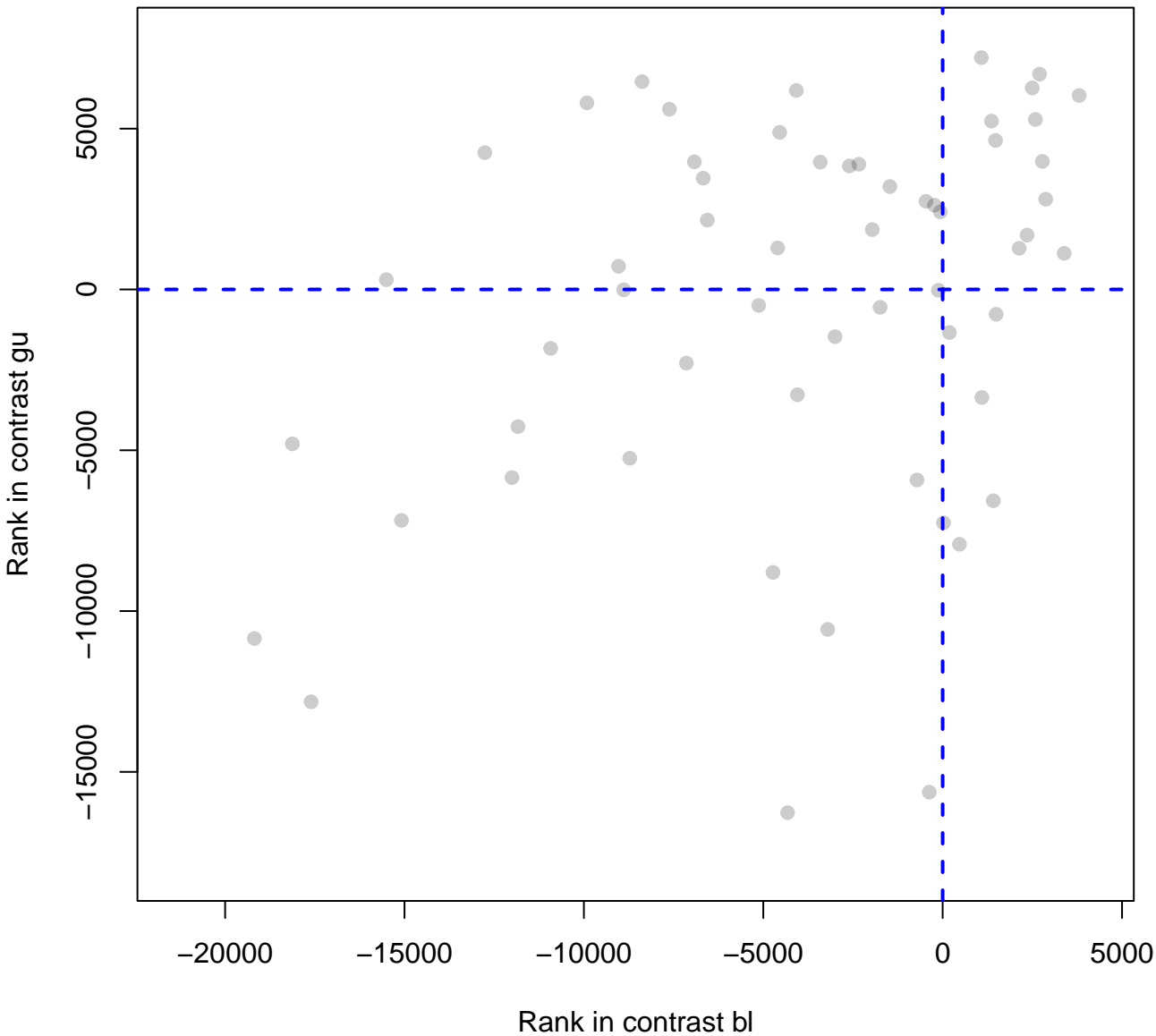
TICAM1, TRAF6-dependent induction of TAK1 co



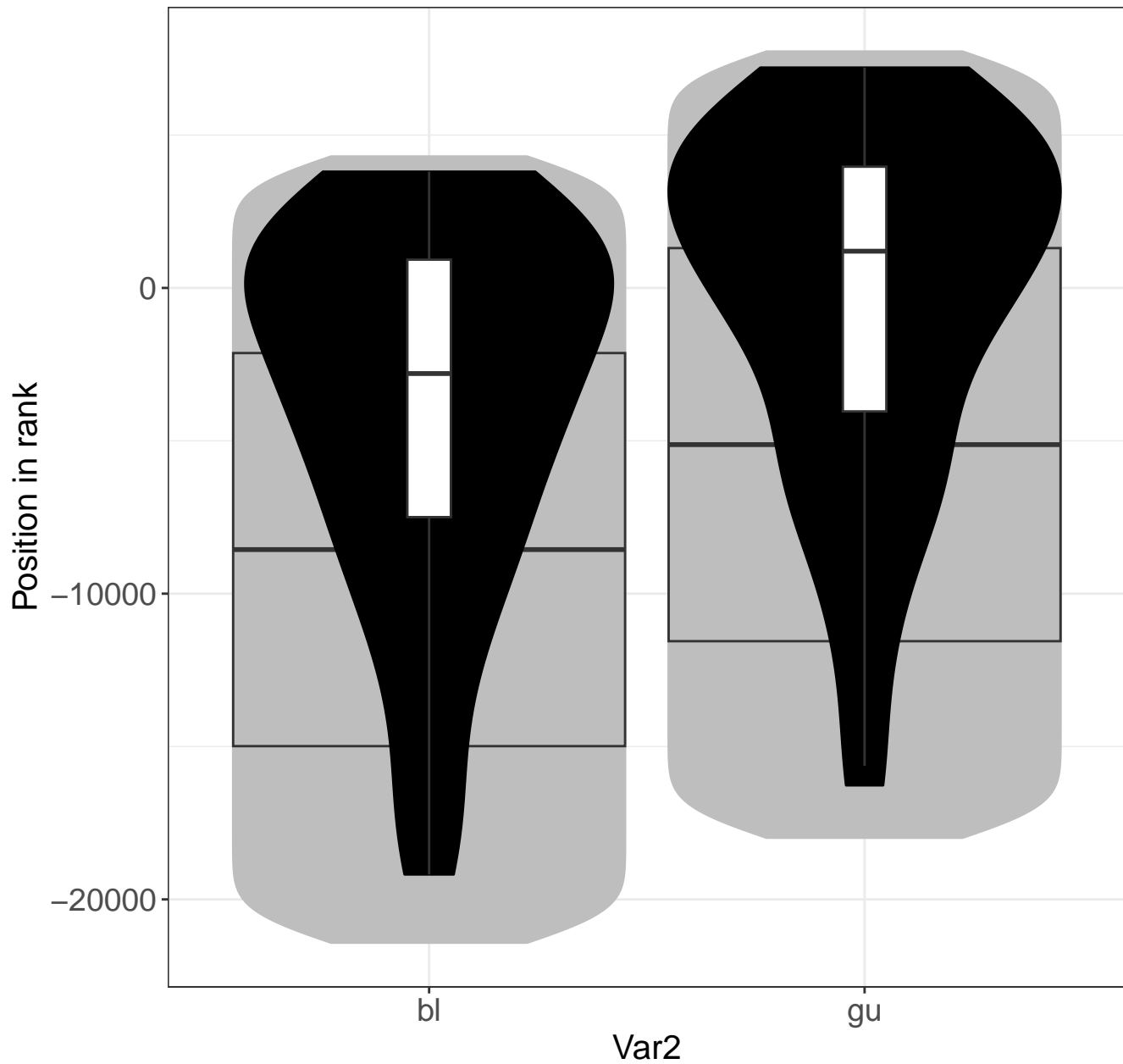
SCF(Skp2)-mediated degradation of p27/p21



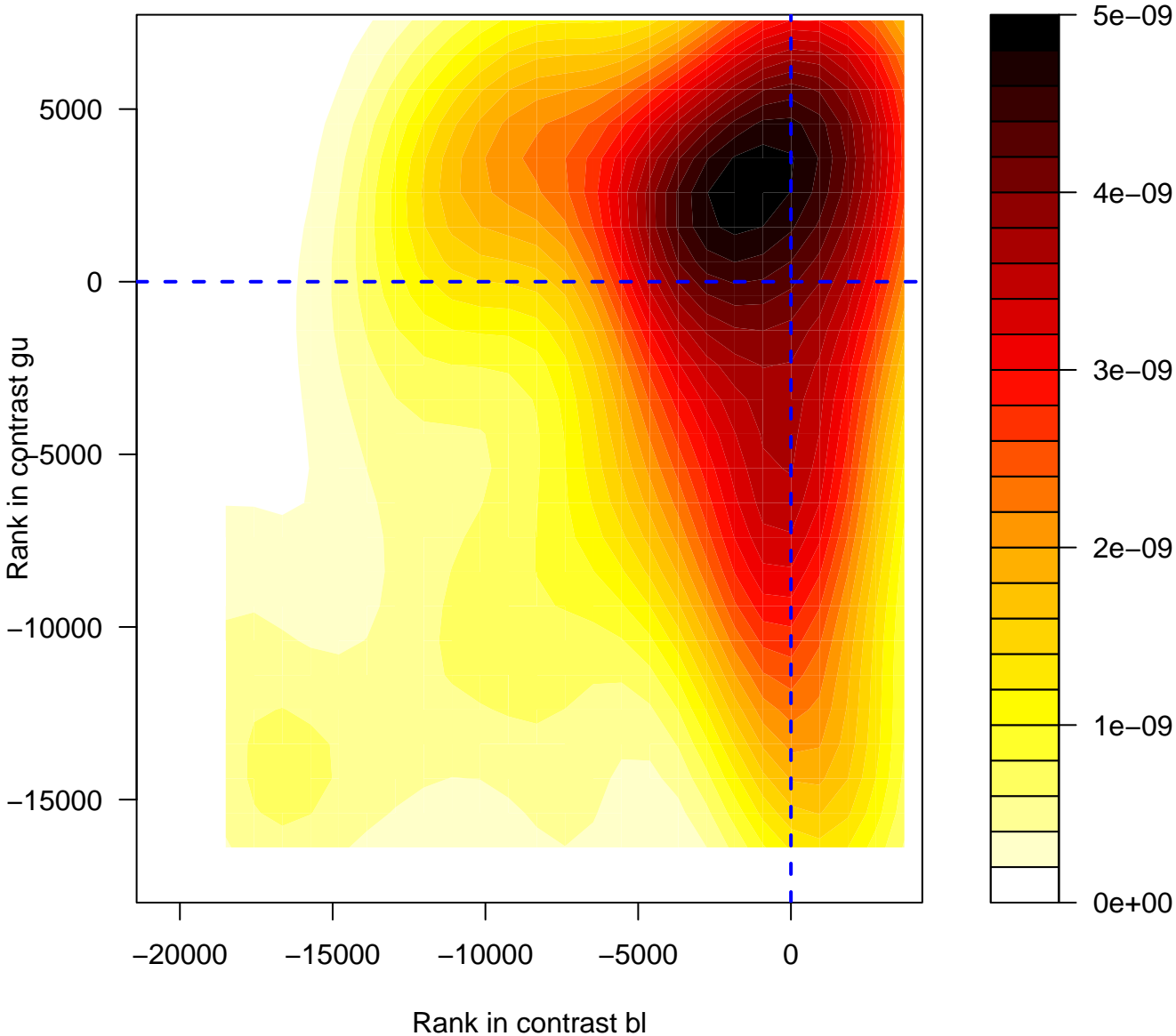
SCF(Skp2)-mediated degradation of p27/p21



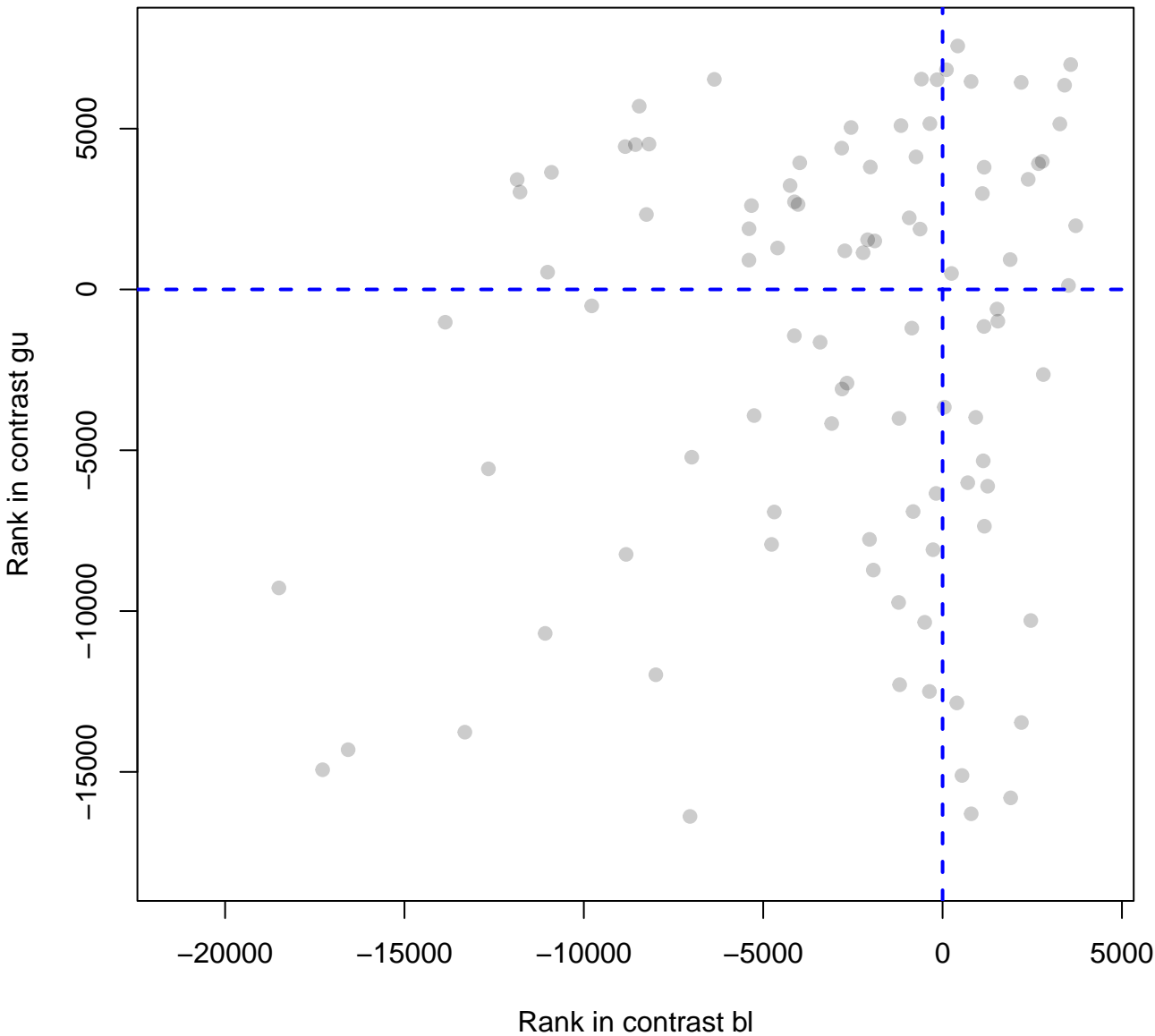
SCF(Skp2)-mediated degradation of p27/p21



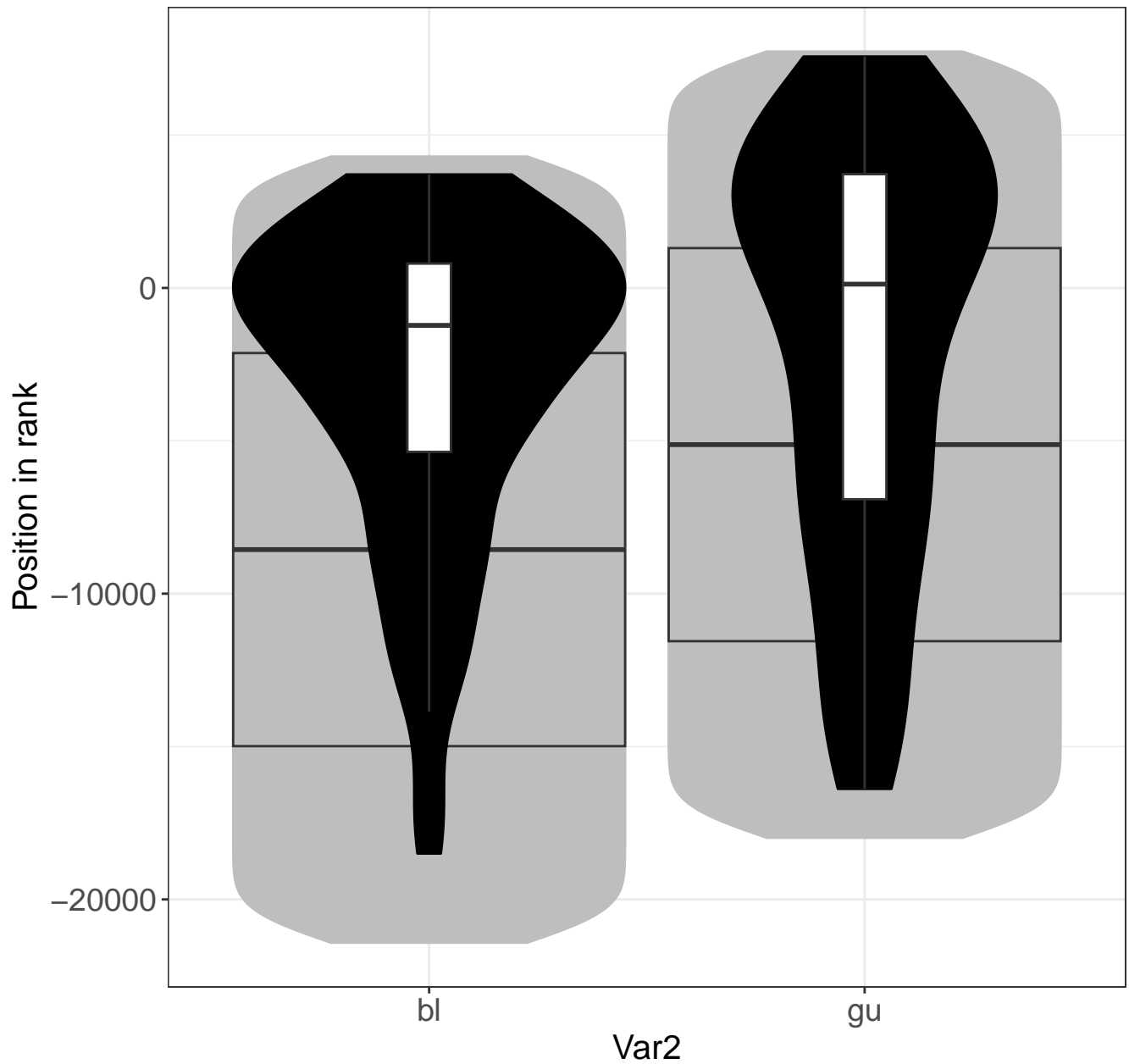
Response of EIF2AK4 (GCN2) to amino acid deficiency



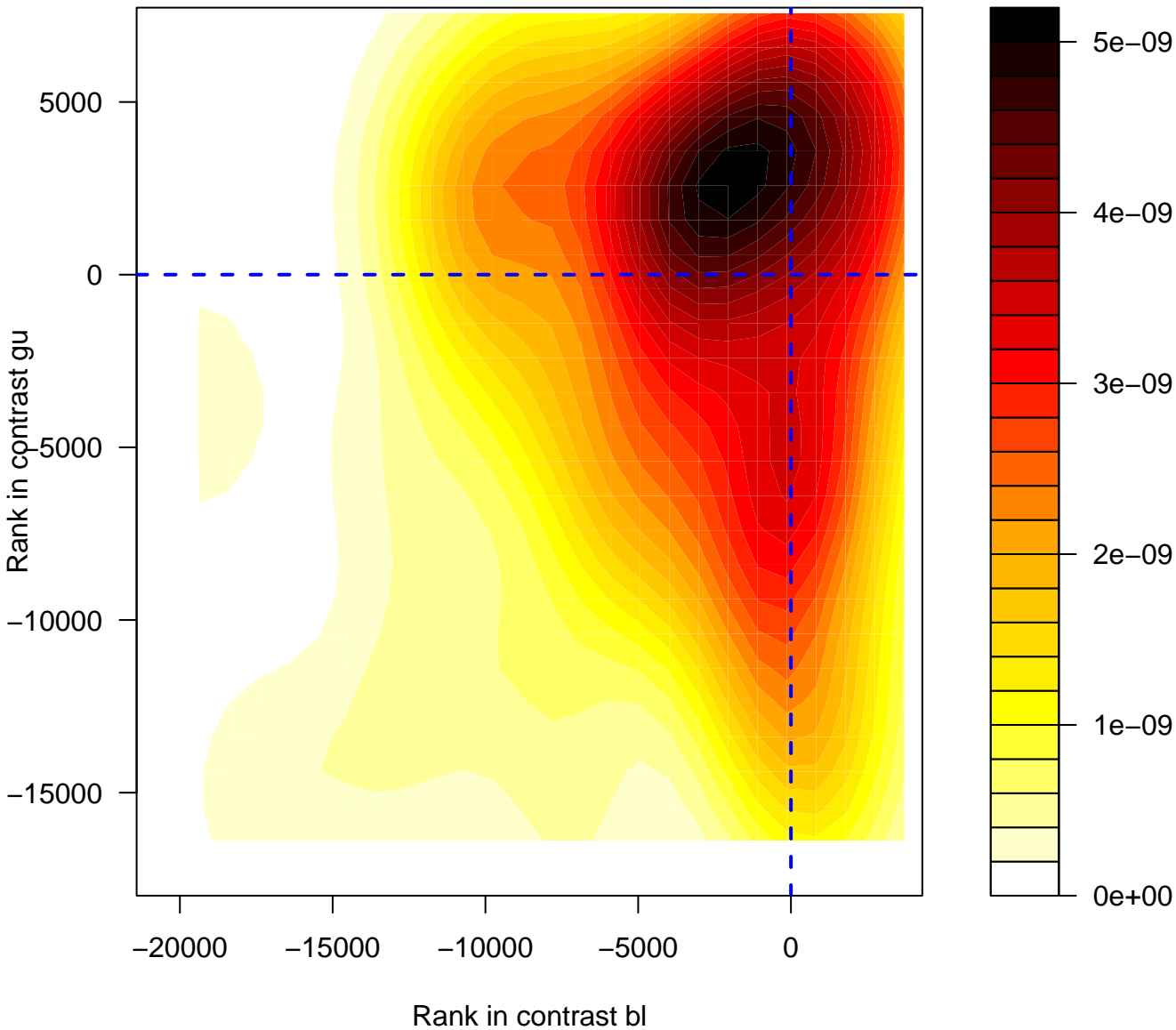
Response of EIF2AK4 (GCN2) to amino acid deficiency



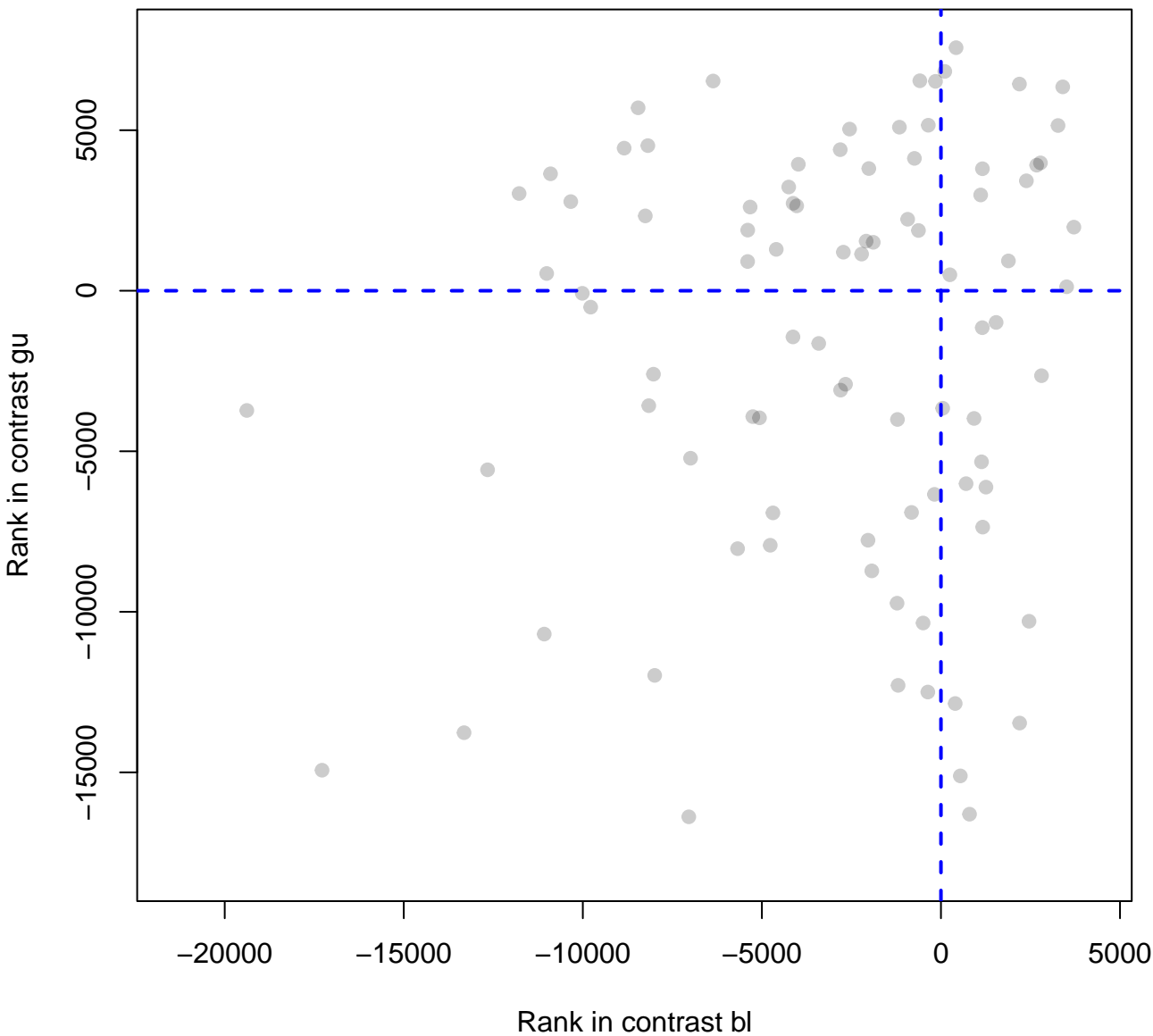
Response of EIF2AK4 (GCN2) to amino acid def



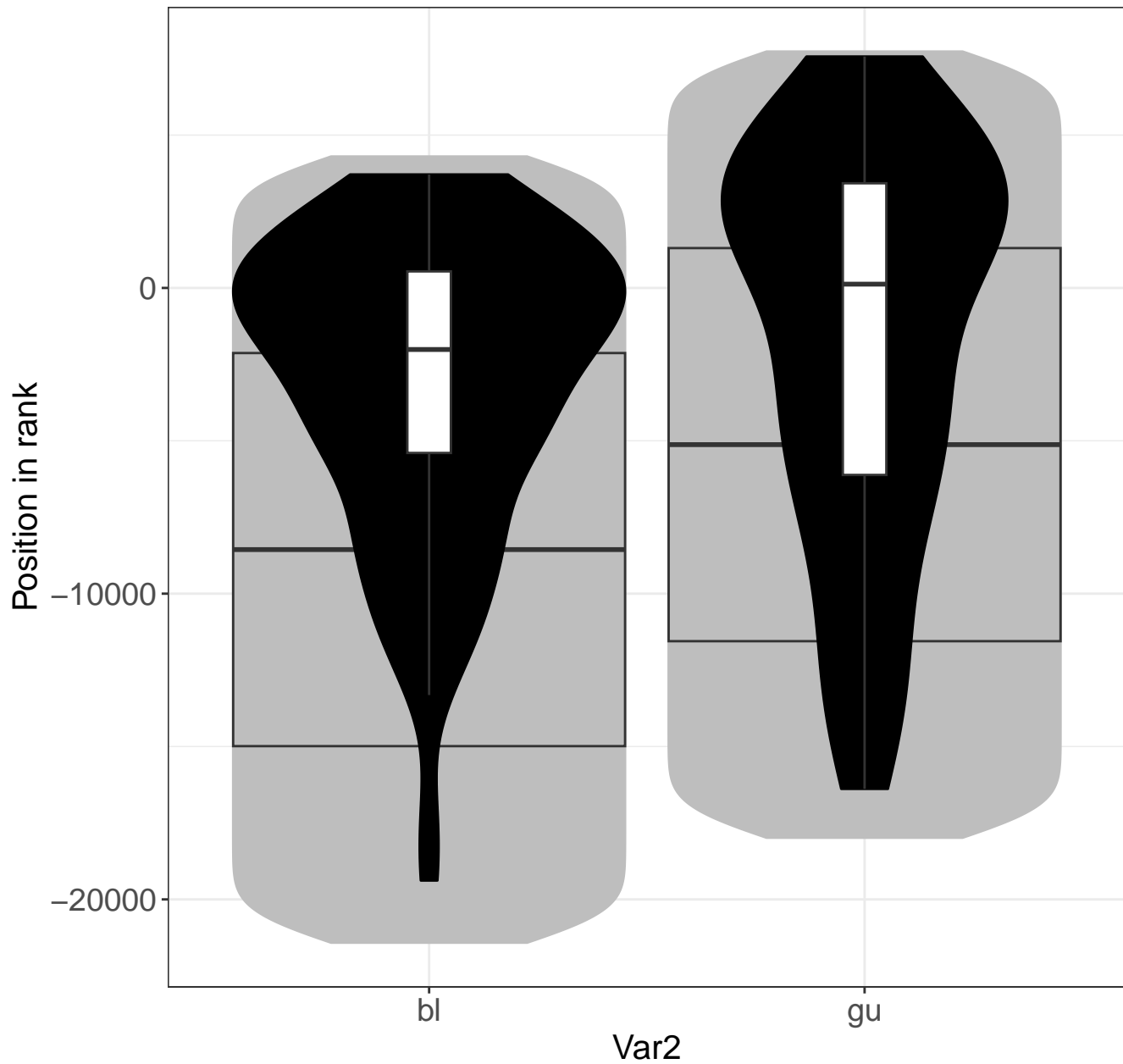
Non-Mediated Decay (NMD) independent of the Exon Junction (EJ)



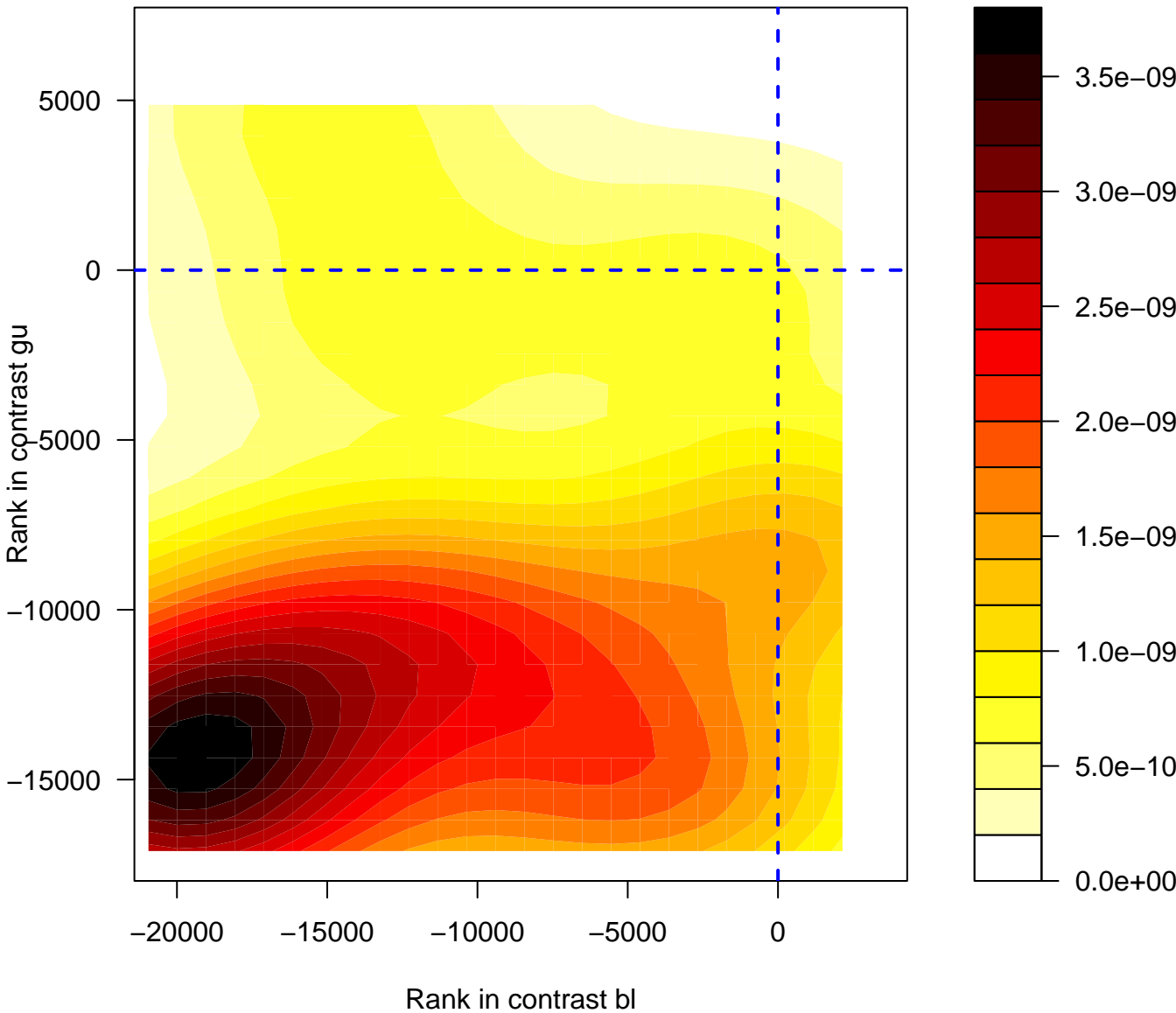
Sense Mediated Decay (NMD) independent of the Exon Junction Complex



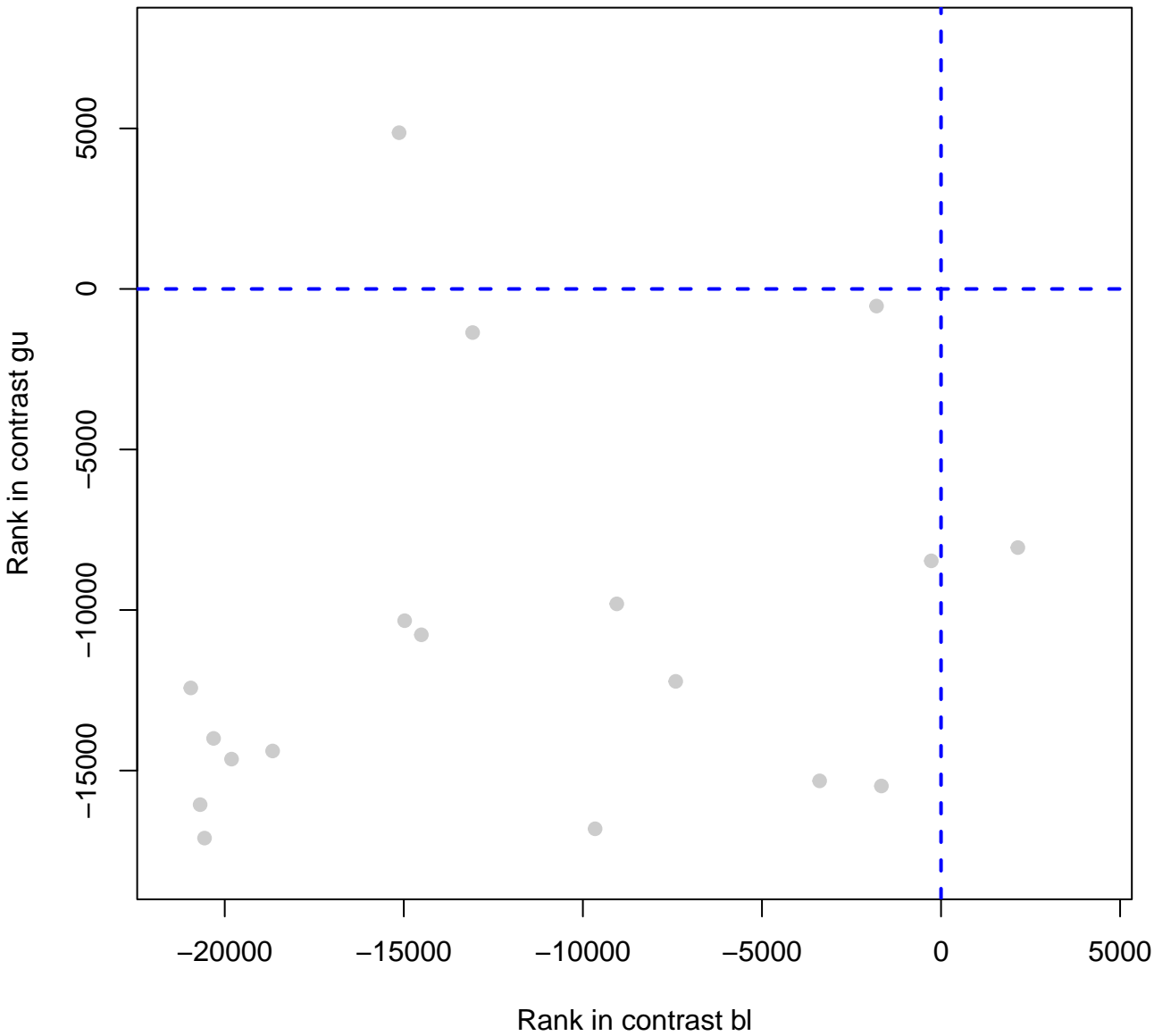
Nonsense Mediated Decay (NMD) independent of



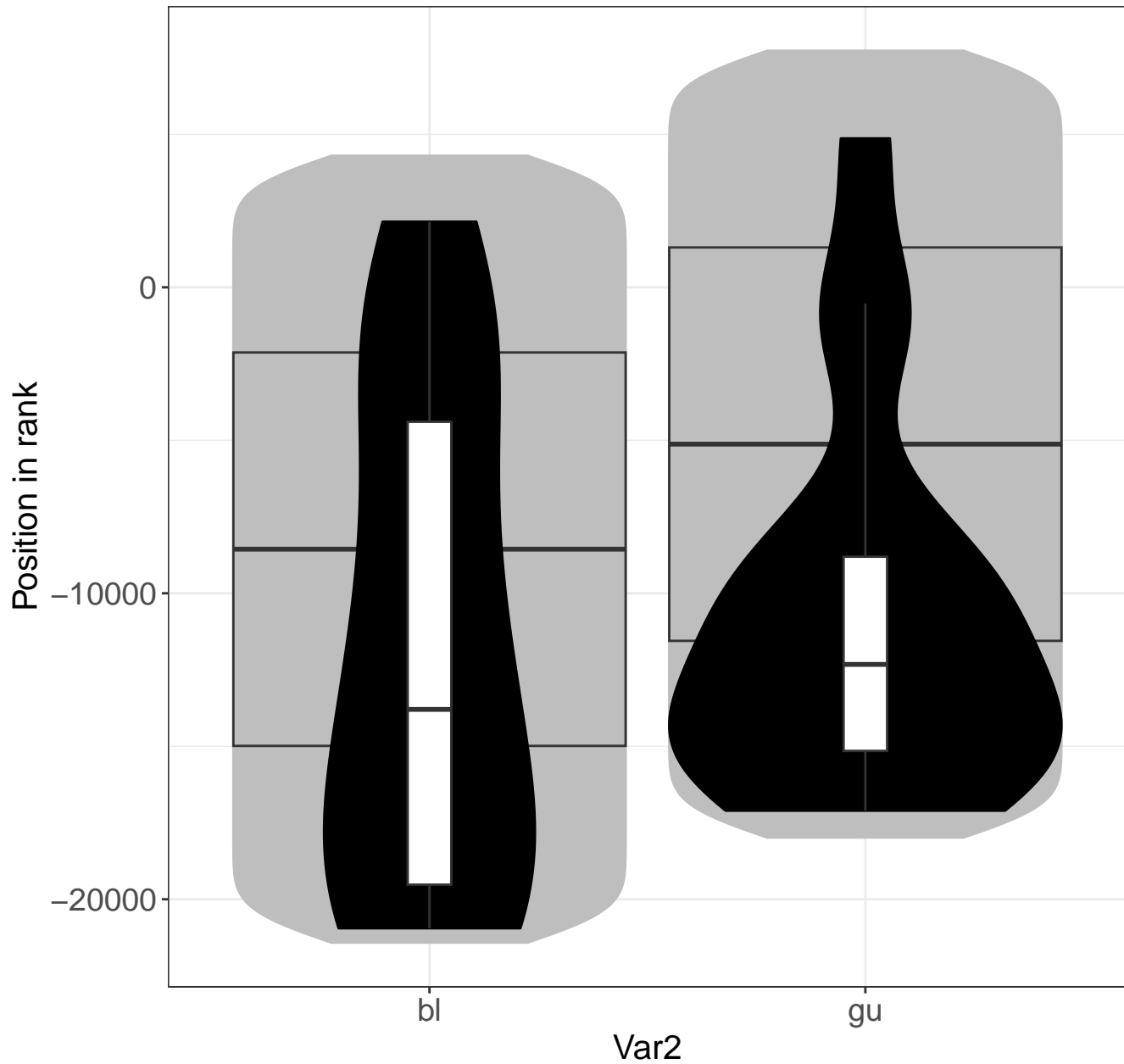
Digestion



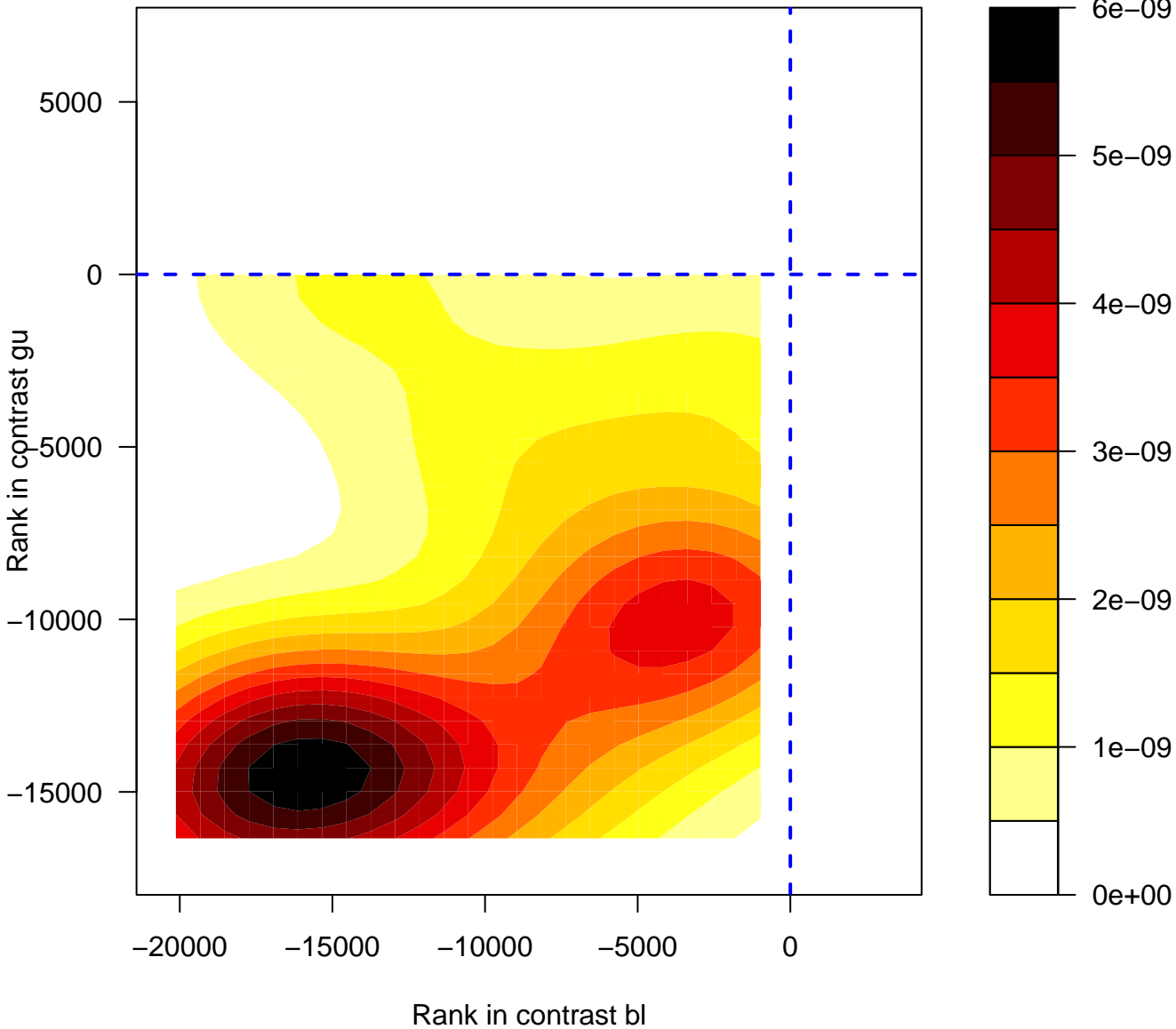
Digestion



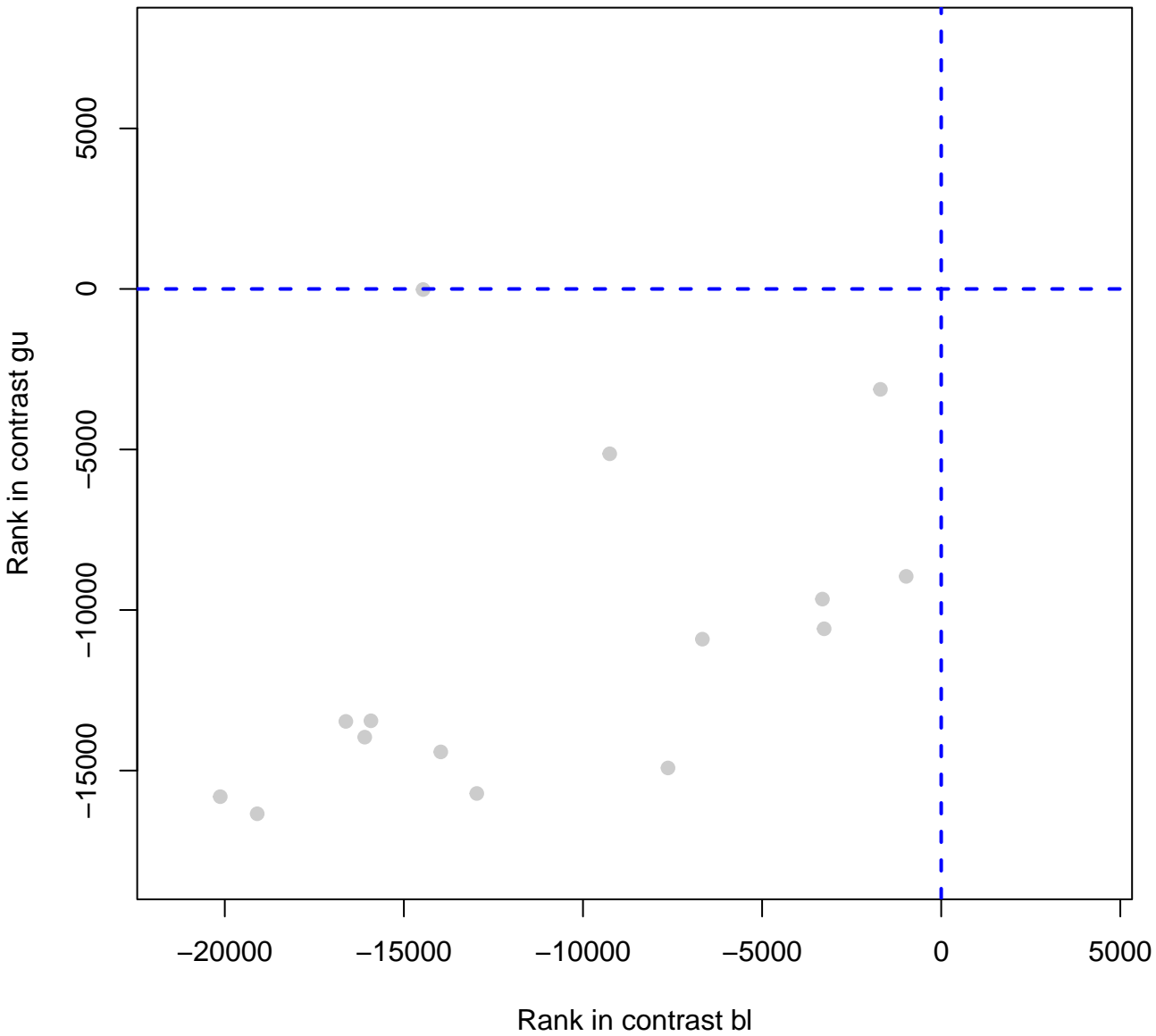
Digestion



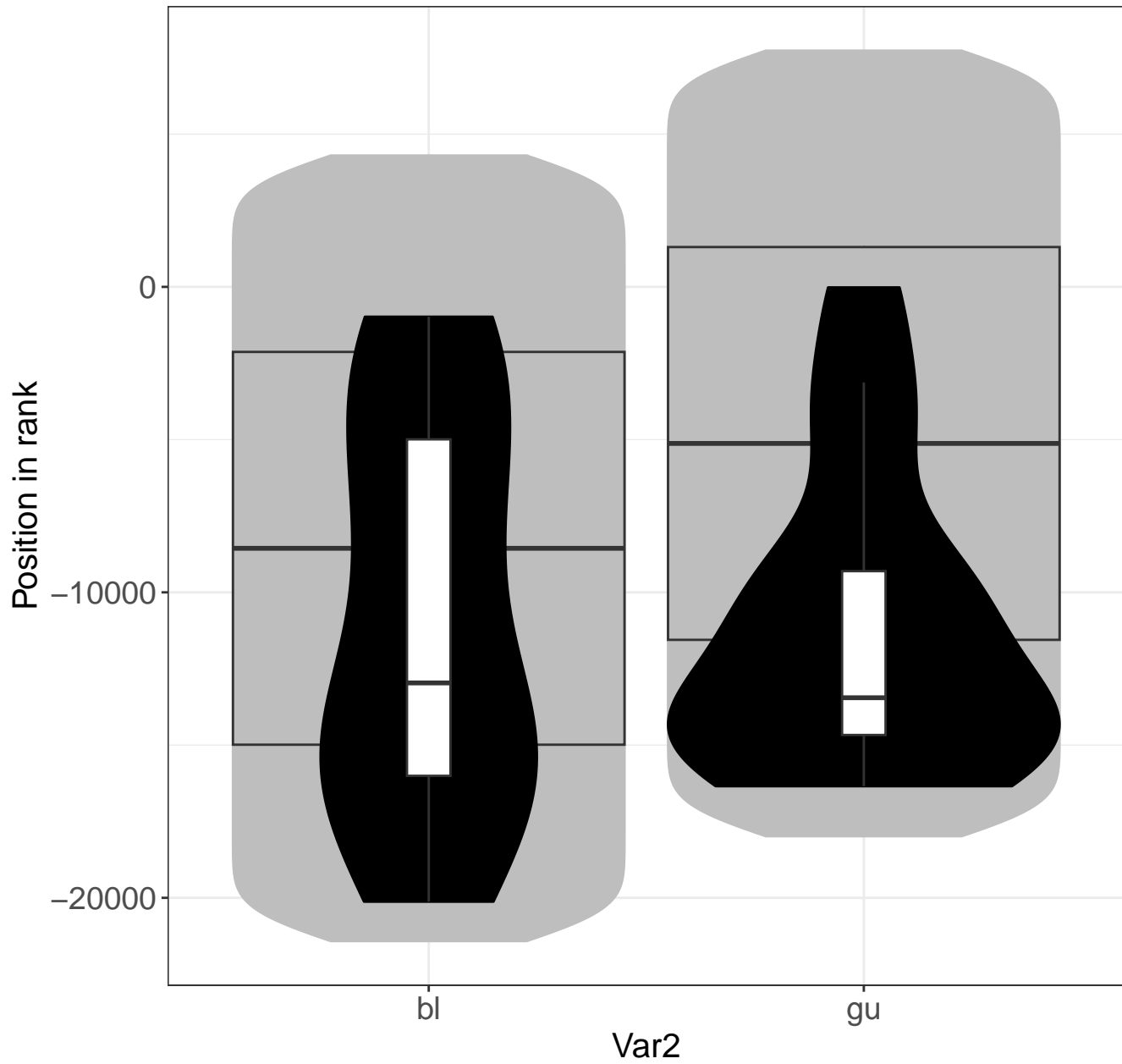
cGMP effects



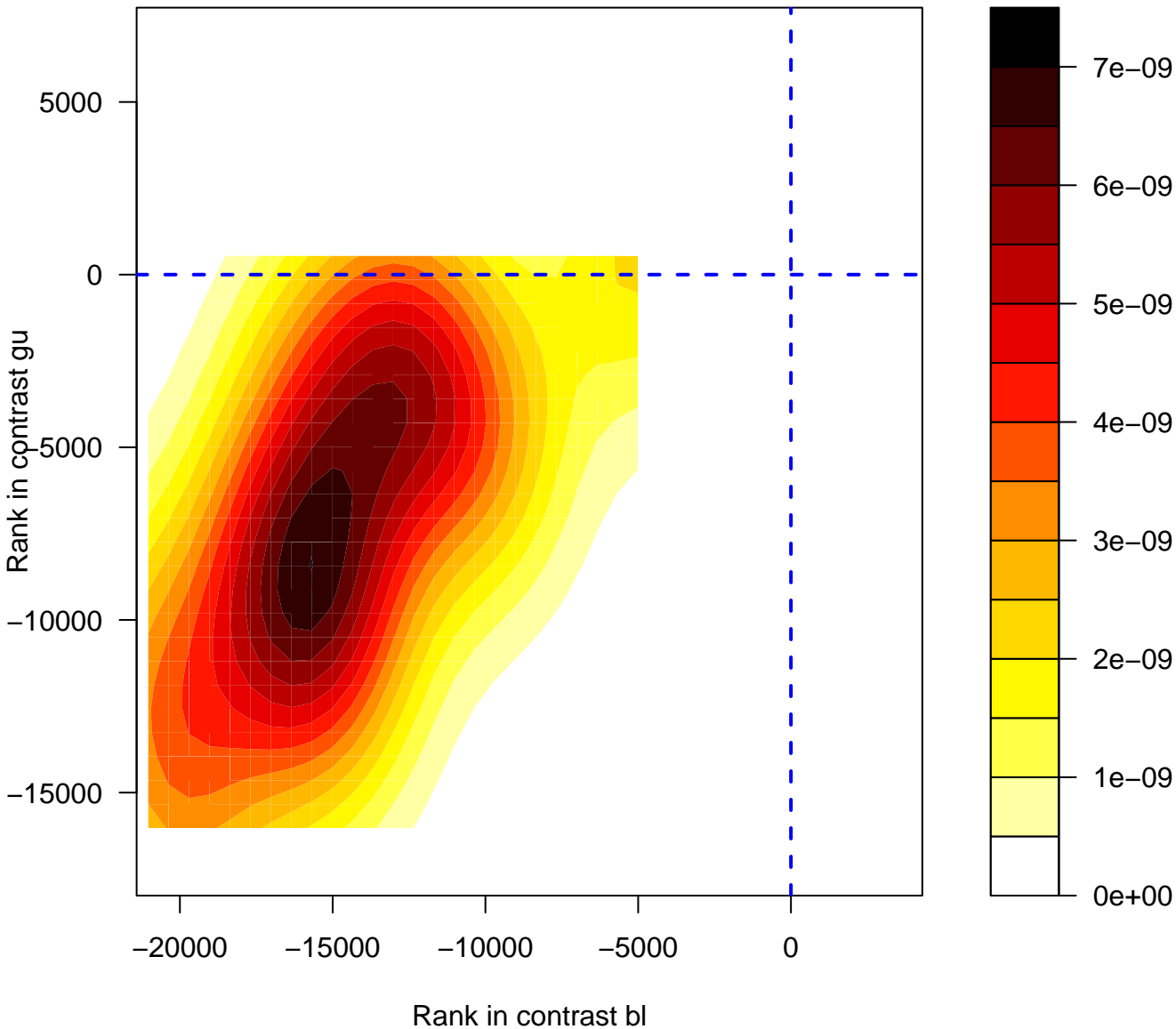
cGMP effects



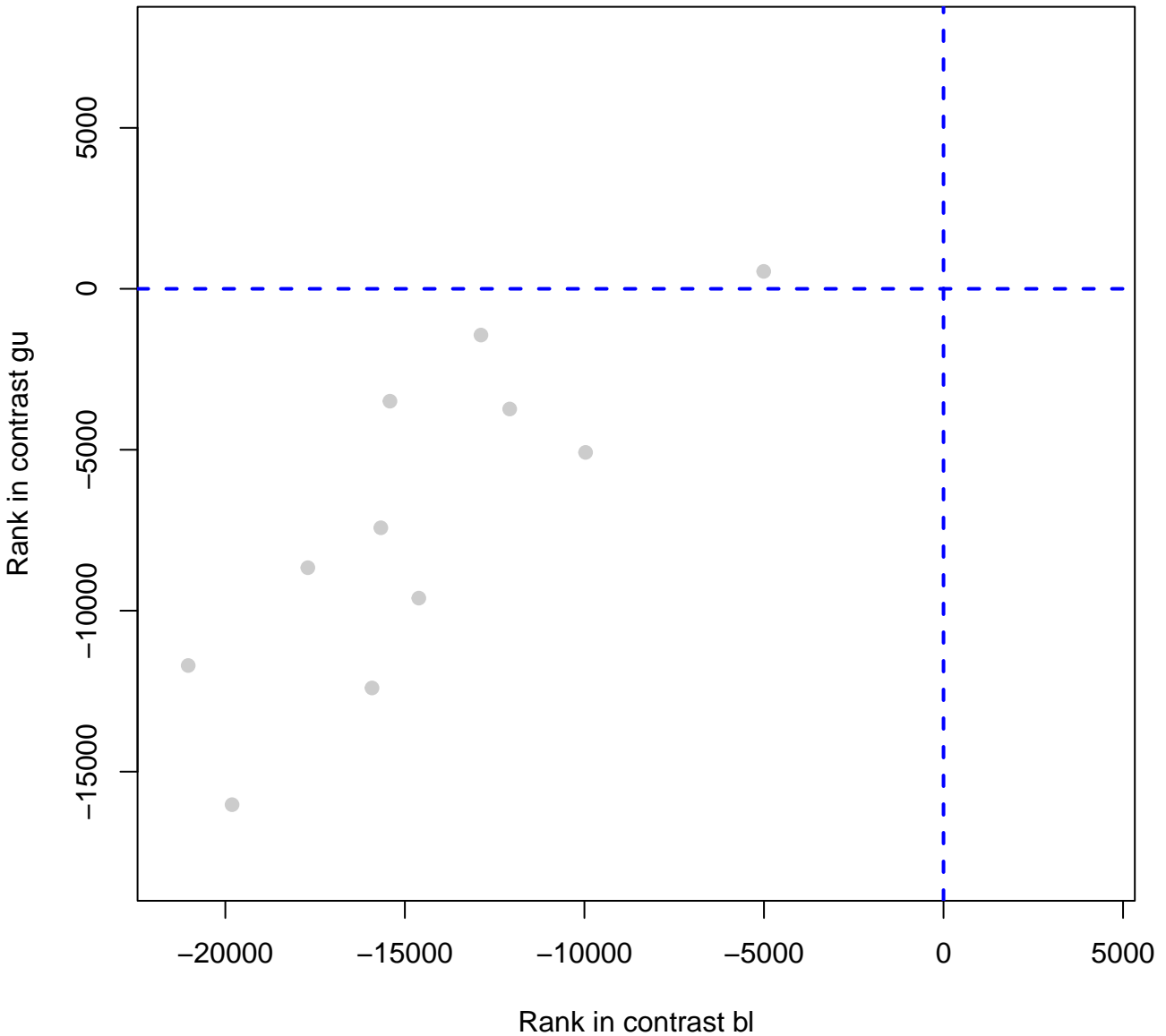
cGMP effects



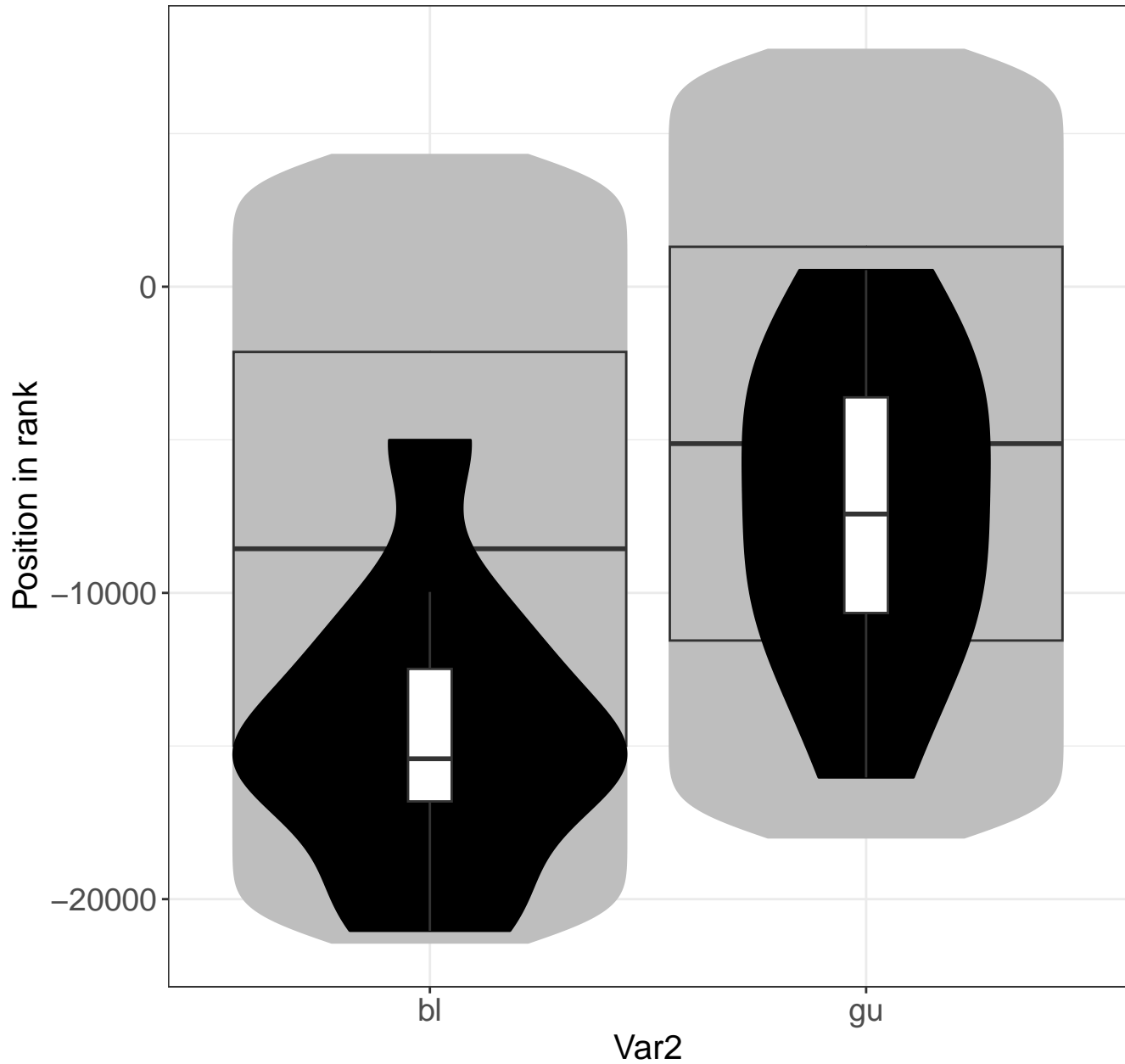
Apoptotic cleavage of cell adhesion proteins



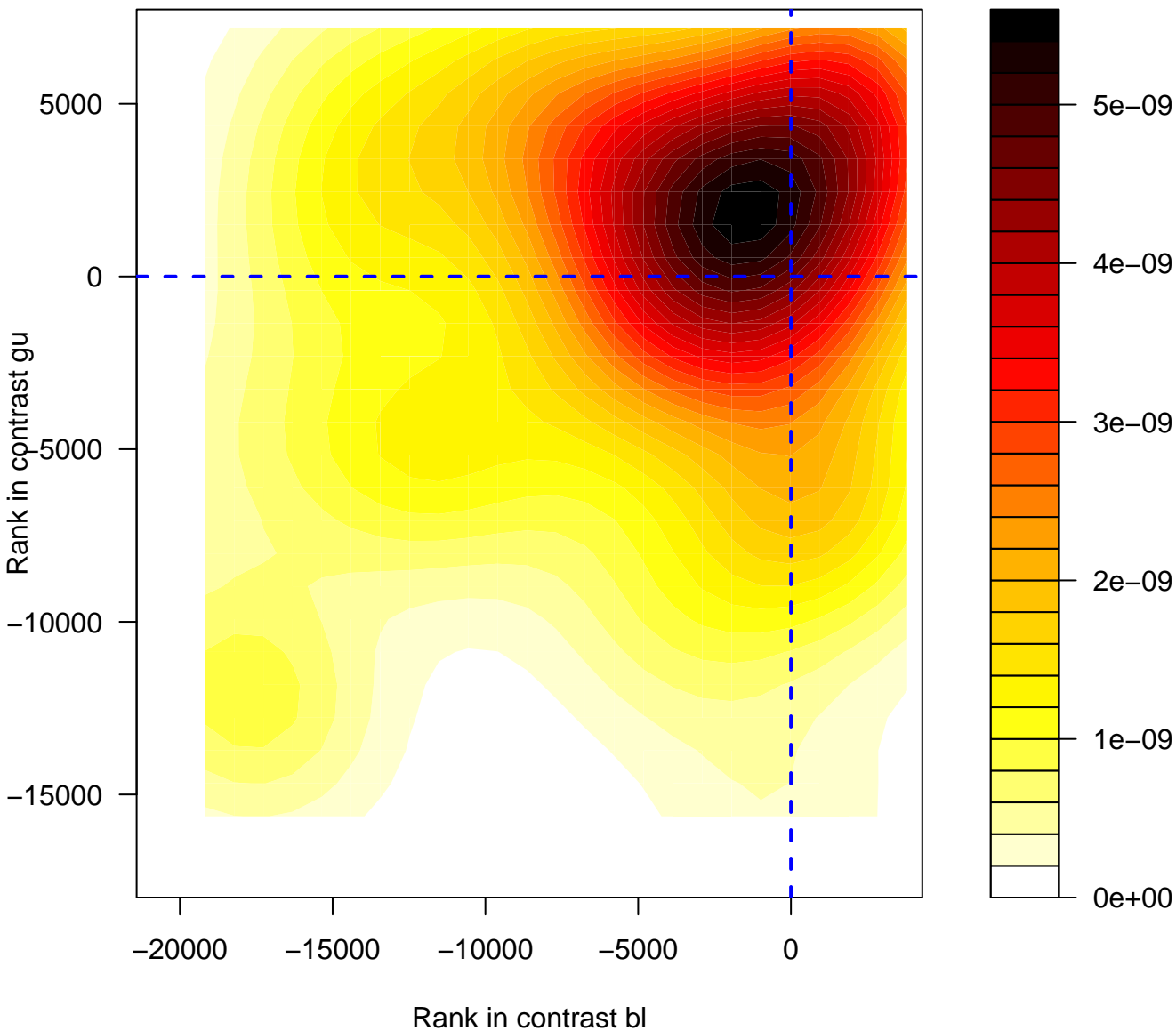
Apoptotic cleavage of cell adhesion proteins



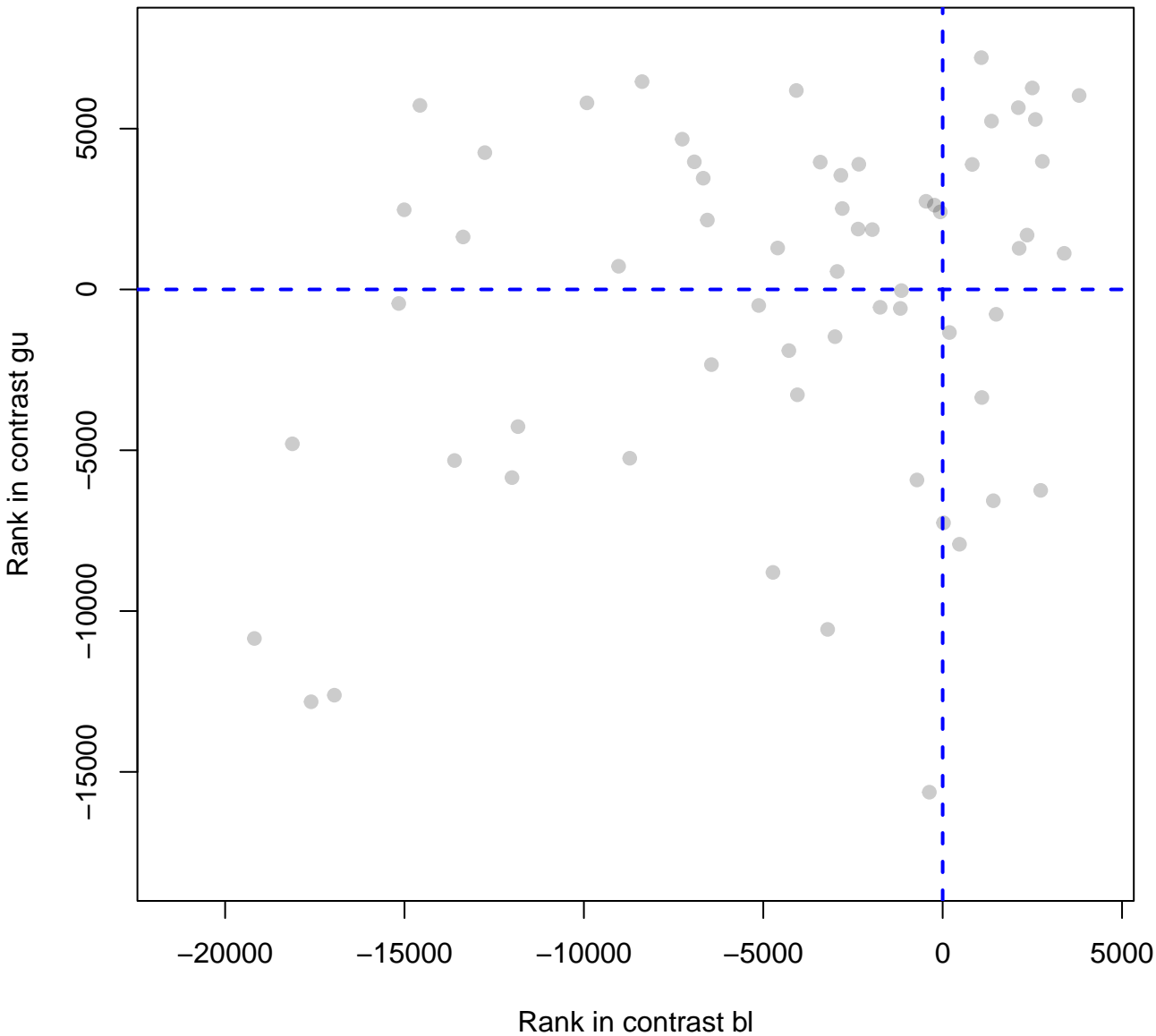
Apoptotic cleavage of cell adhesion proteins



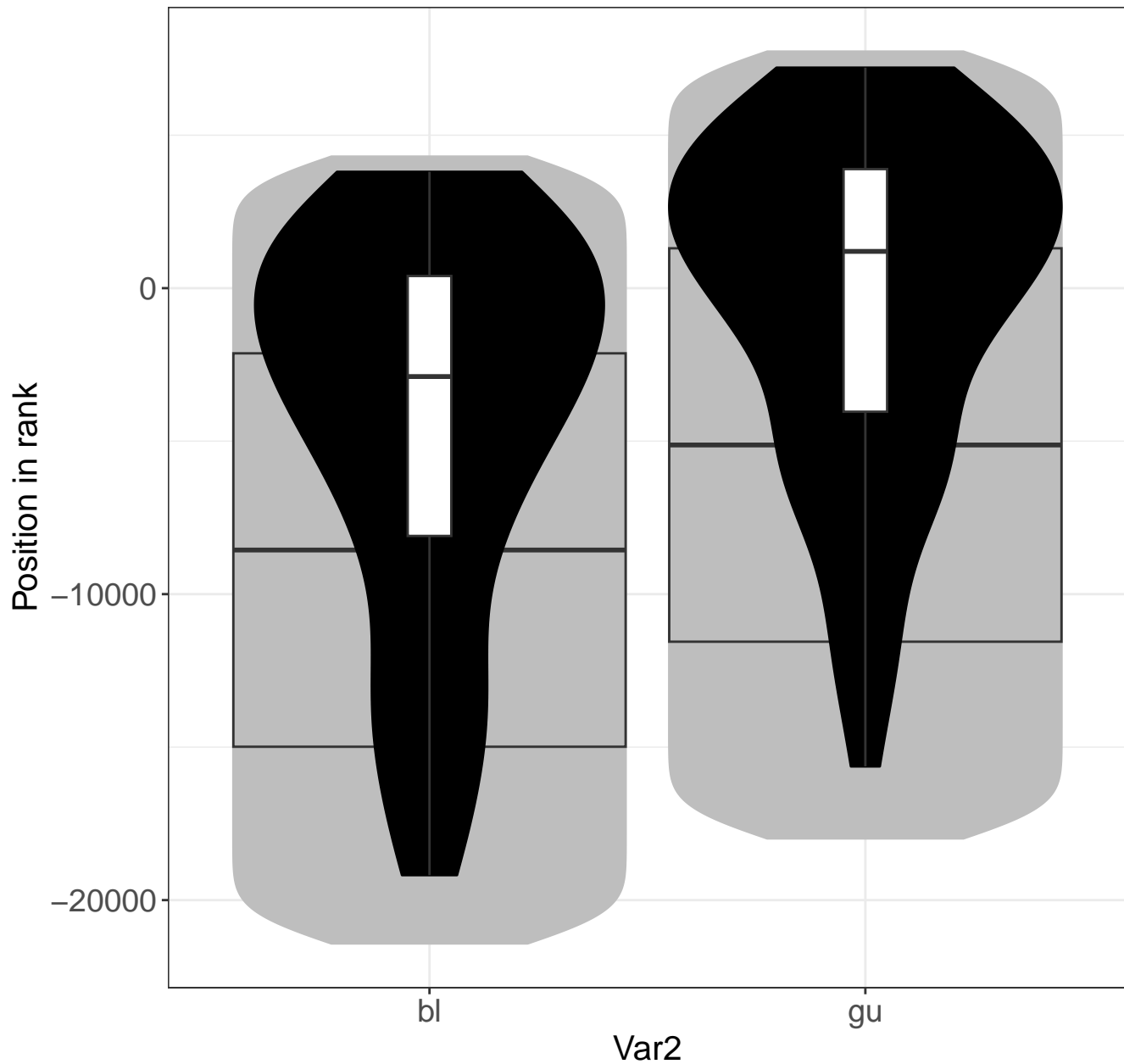
Autodegradation of Cdh1 by Cdh1:APC/C



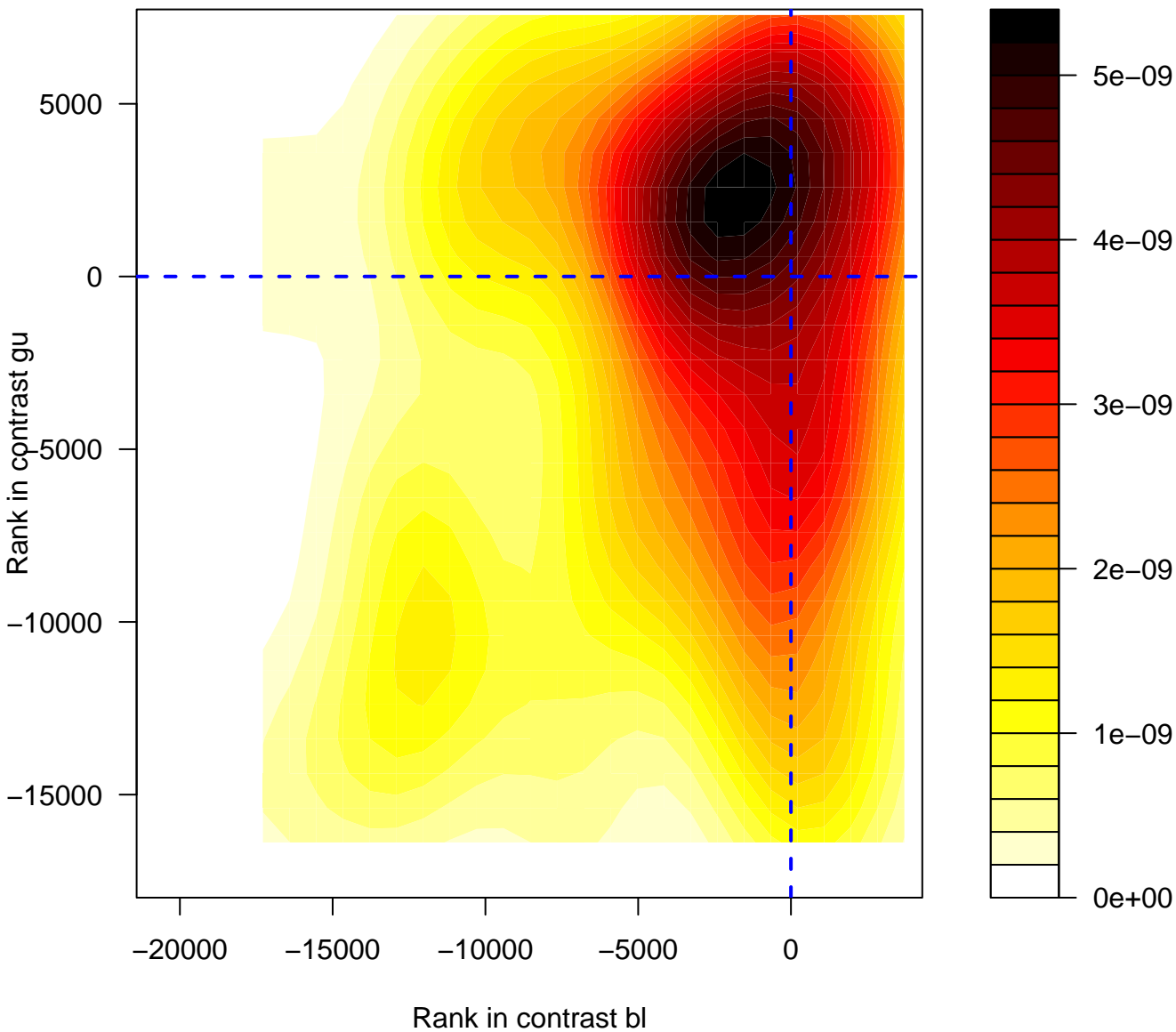
Autodegradation of Cdh1 by Cdh1:APC/C



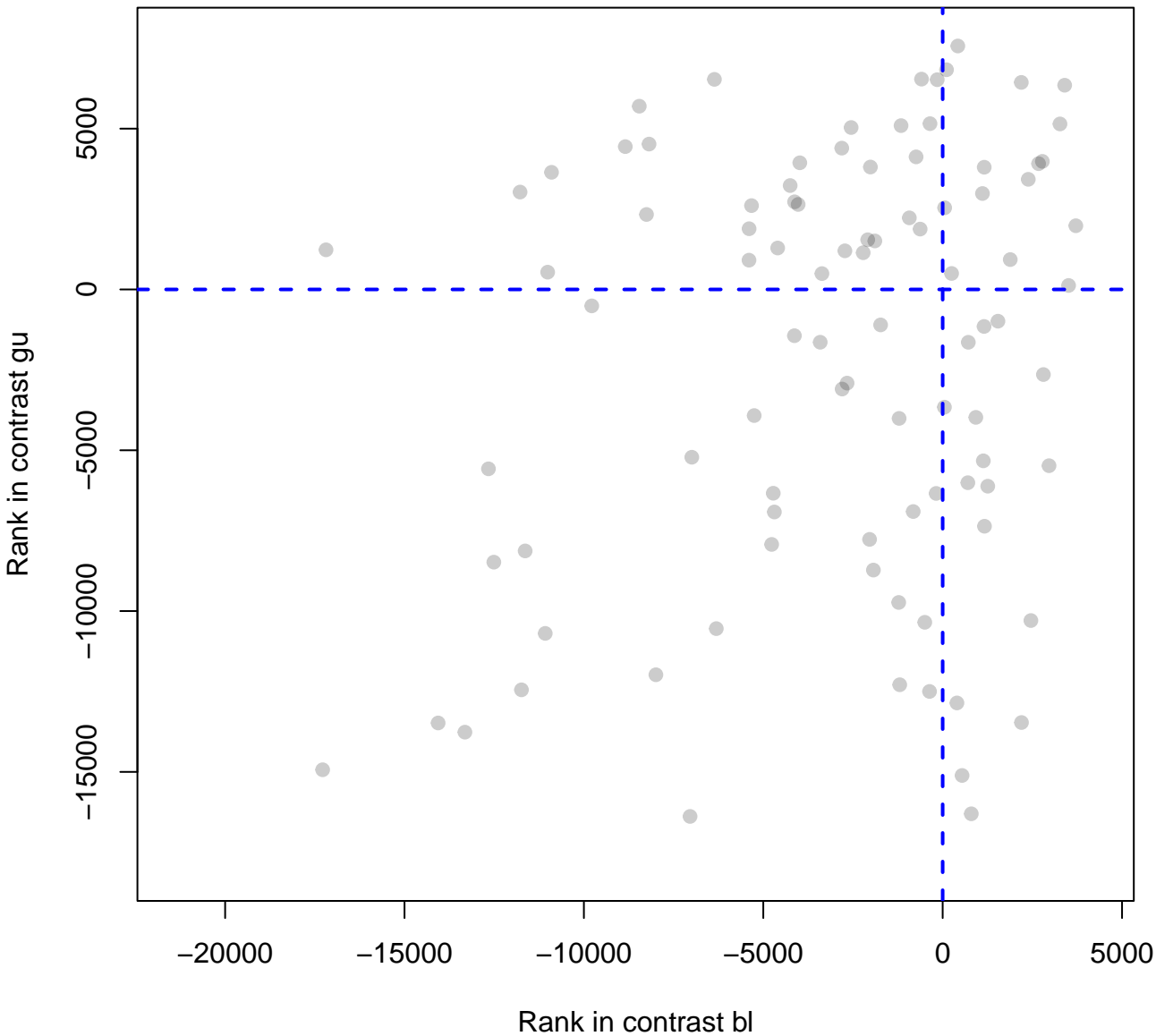
Autodegradation of Cdh1 by Cdh1:APC/C



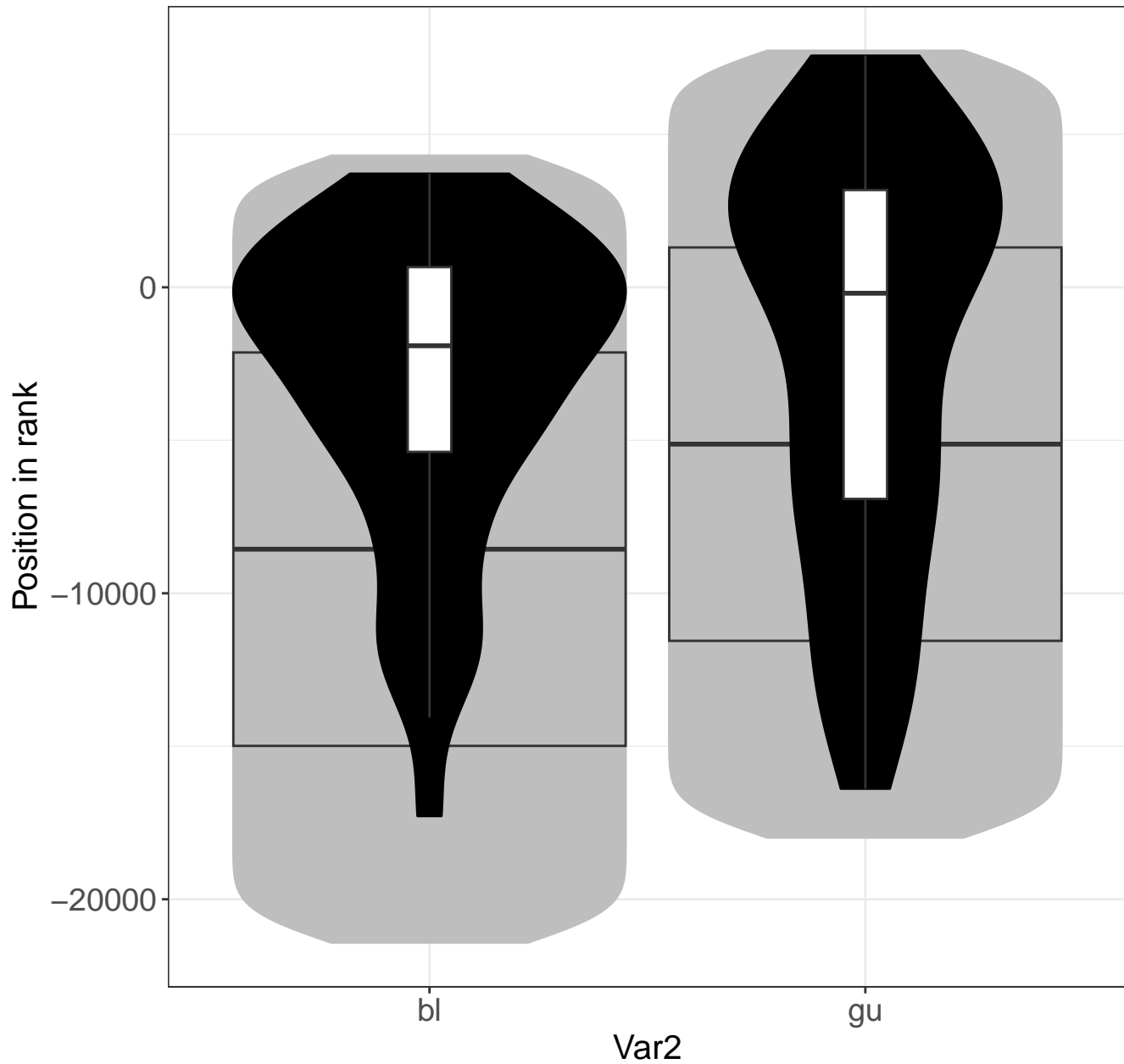
Formation of a pool of free 40S subunits



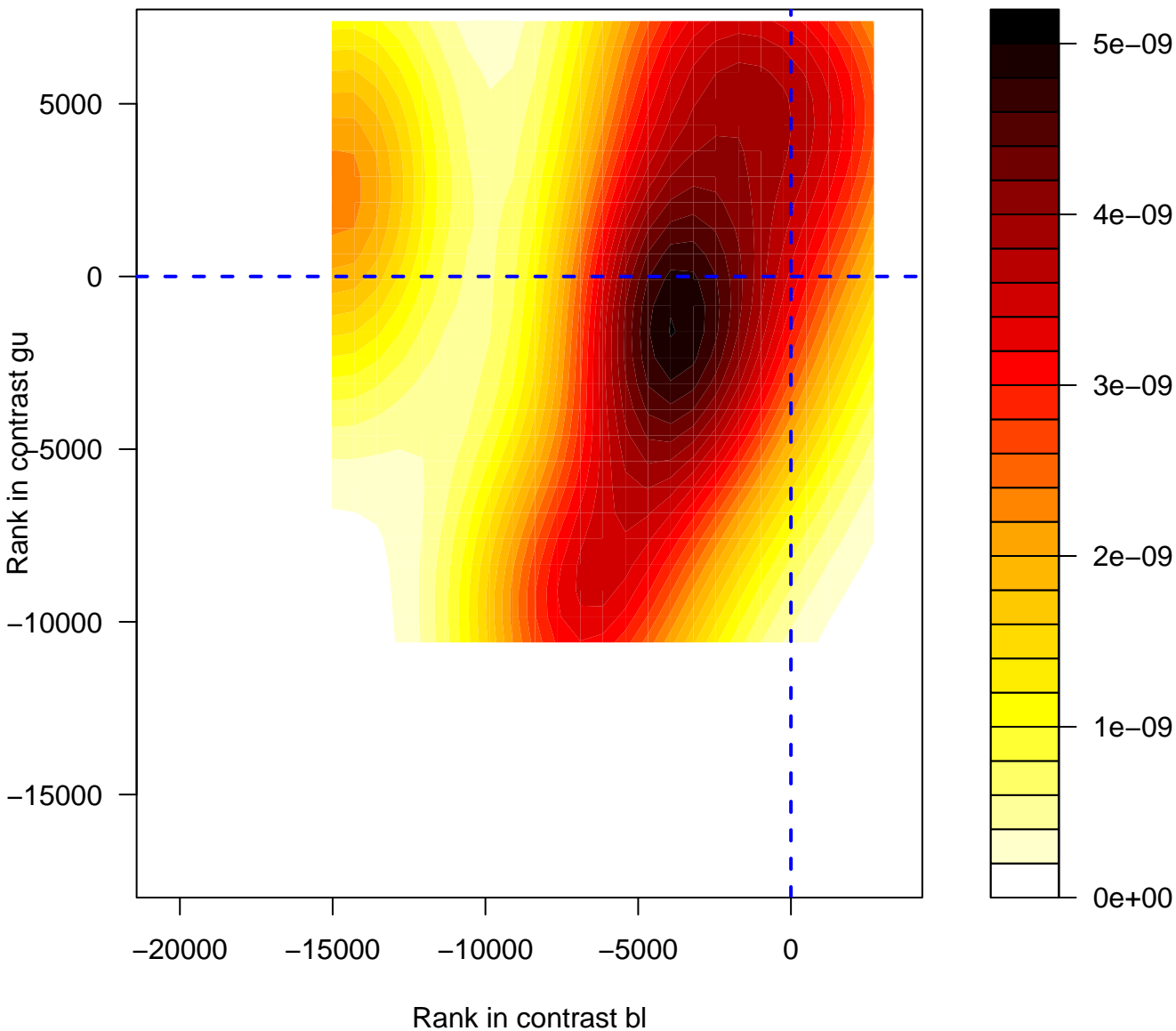
Formation of a pool of free 40S subunits



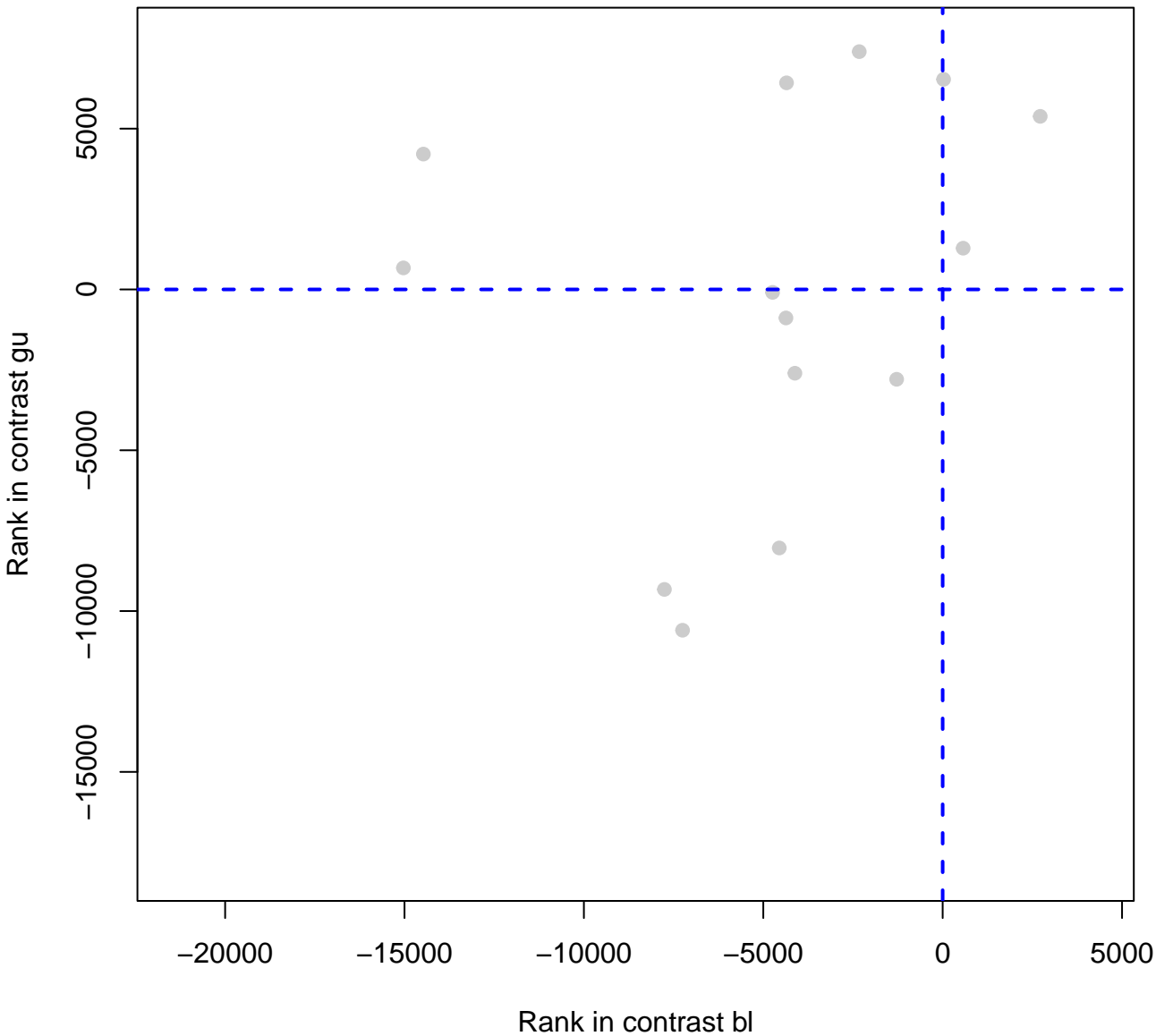
Formation of a pool of free 40S subunits



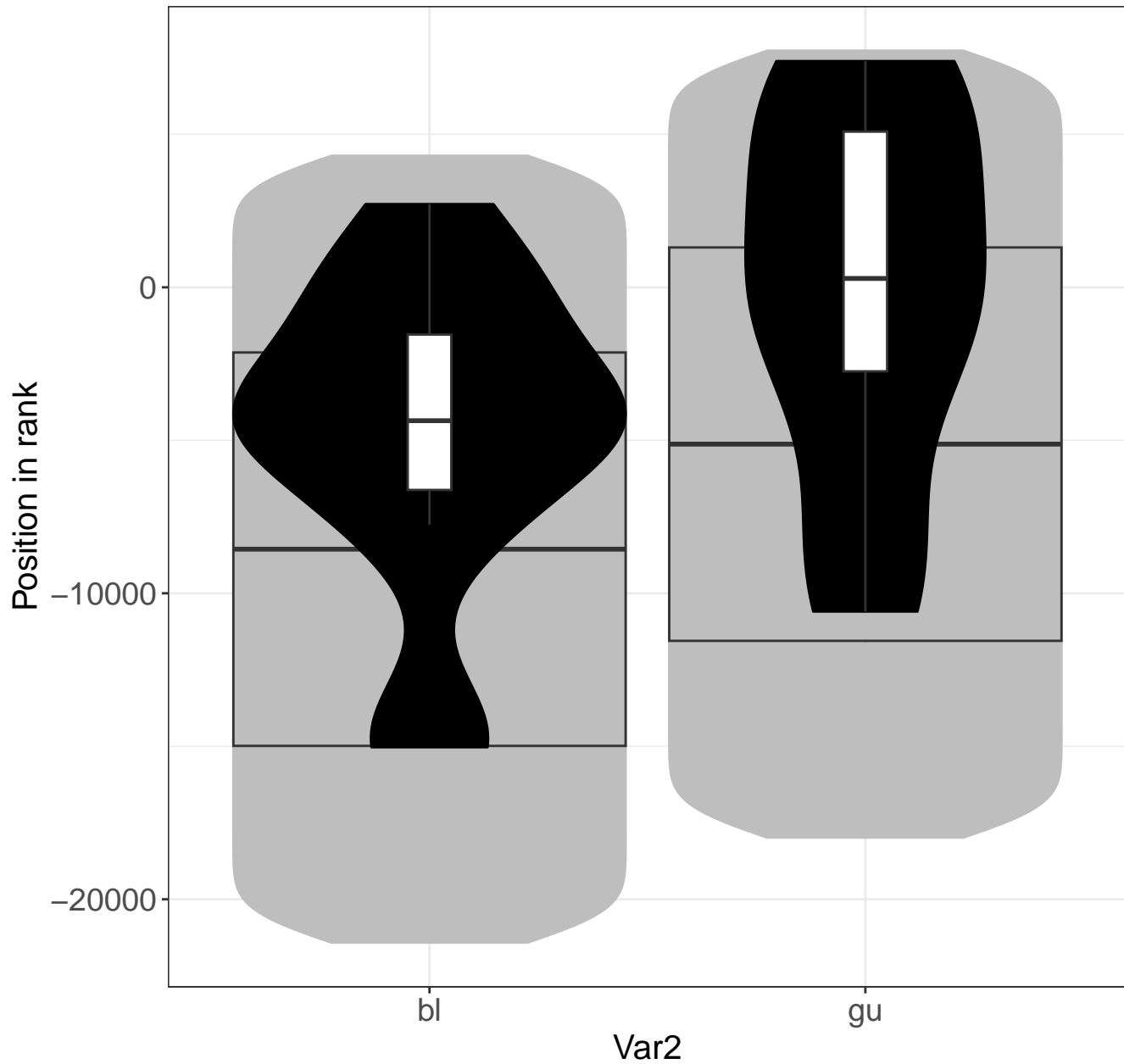
Golgi Cisternae Pericentriolar Stack Reorganization



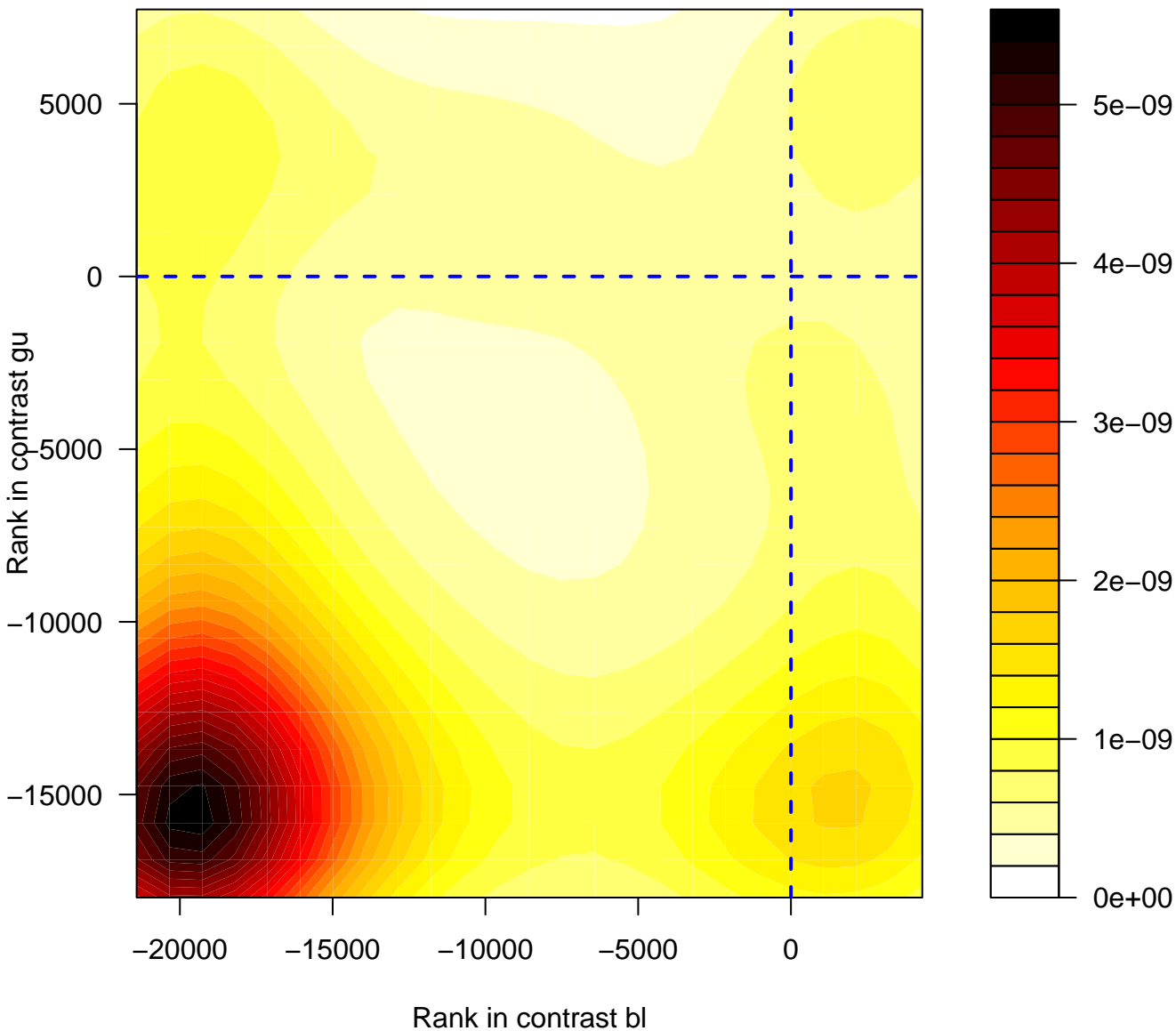
Golgi Cisternae Pericentriolar Stack Reorganization



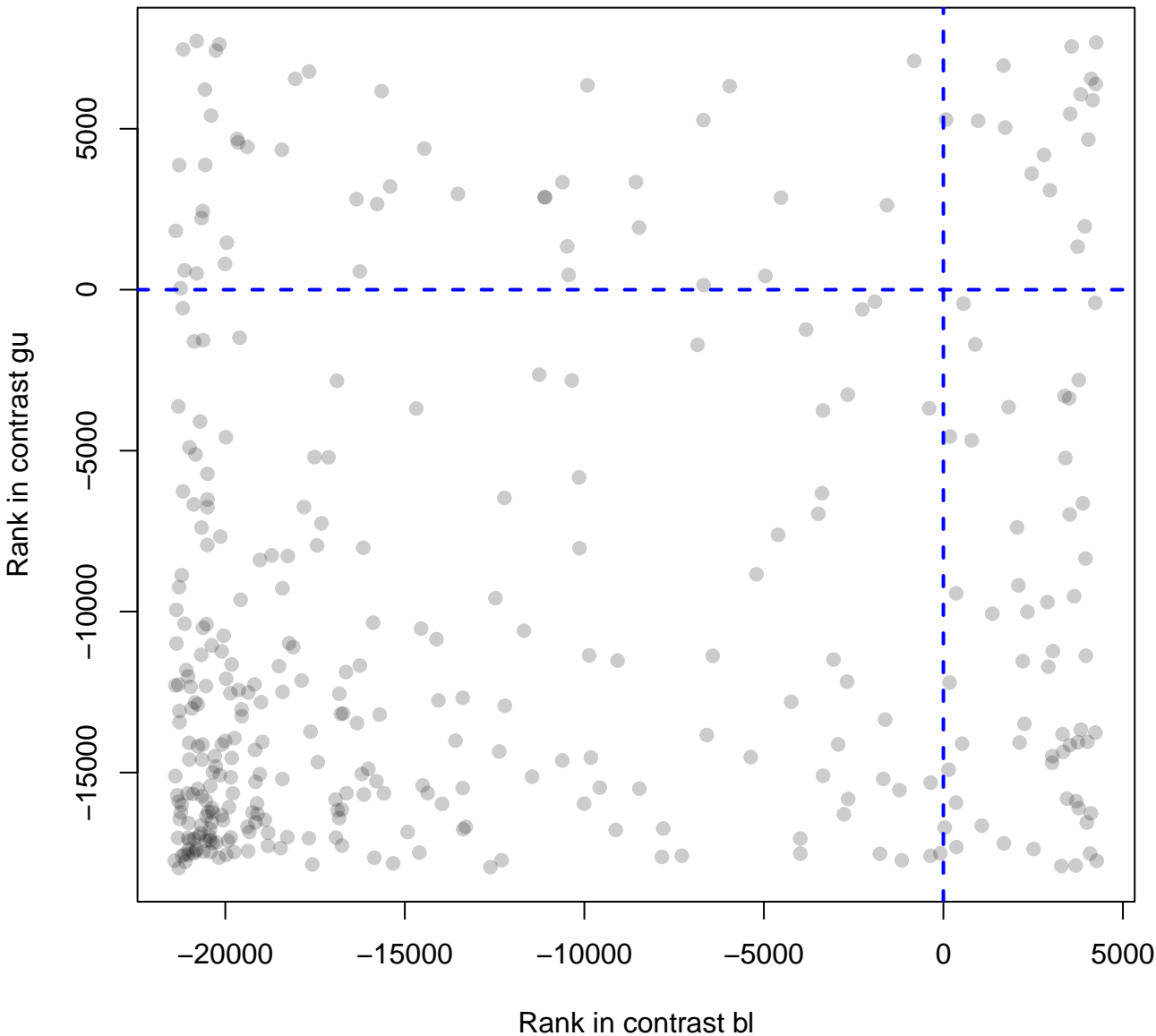
Golgi Cisternae Pericentriolar Stack Reorganizat



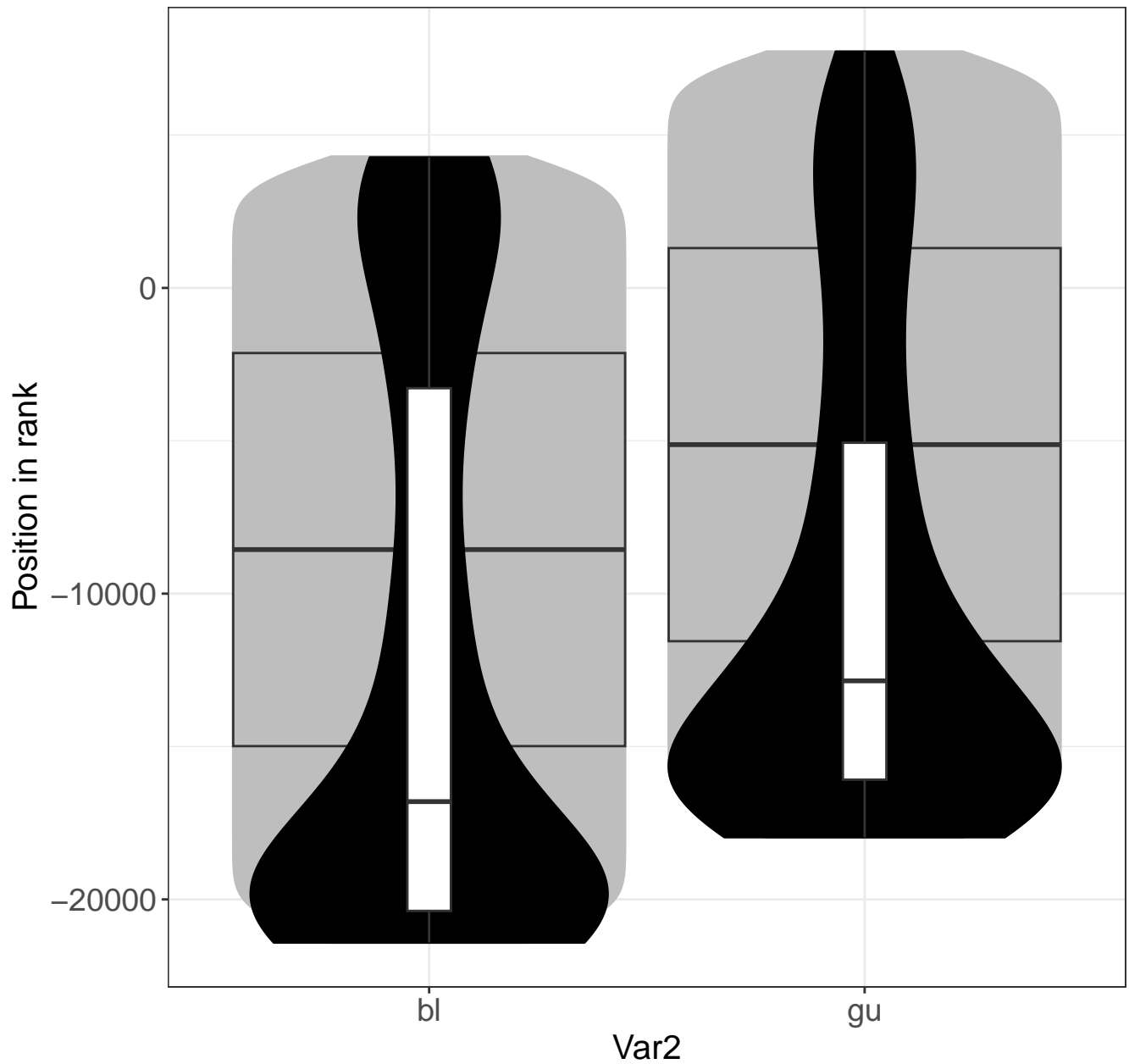
Expression and translocation of olfactory receptors



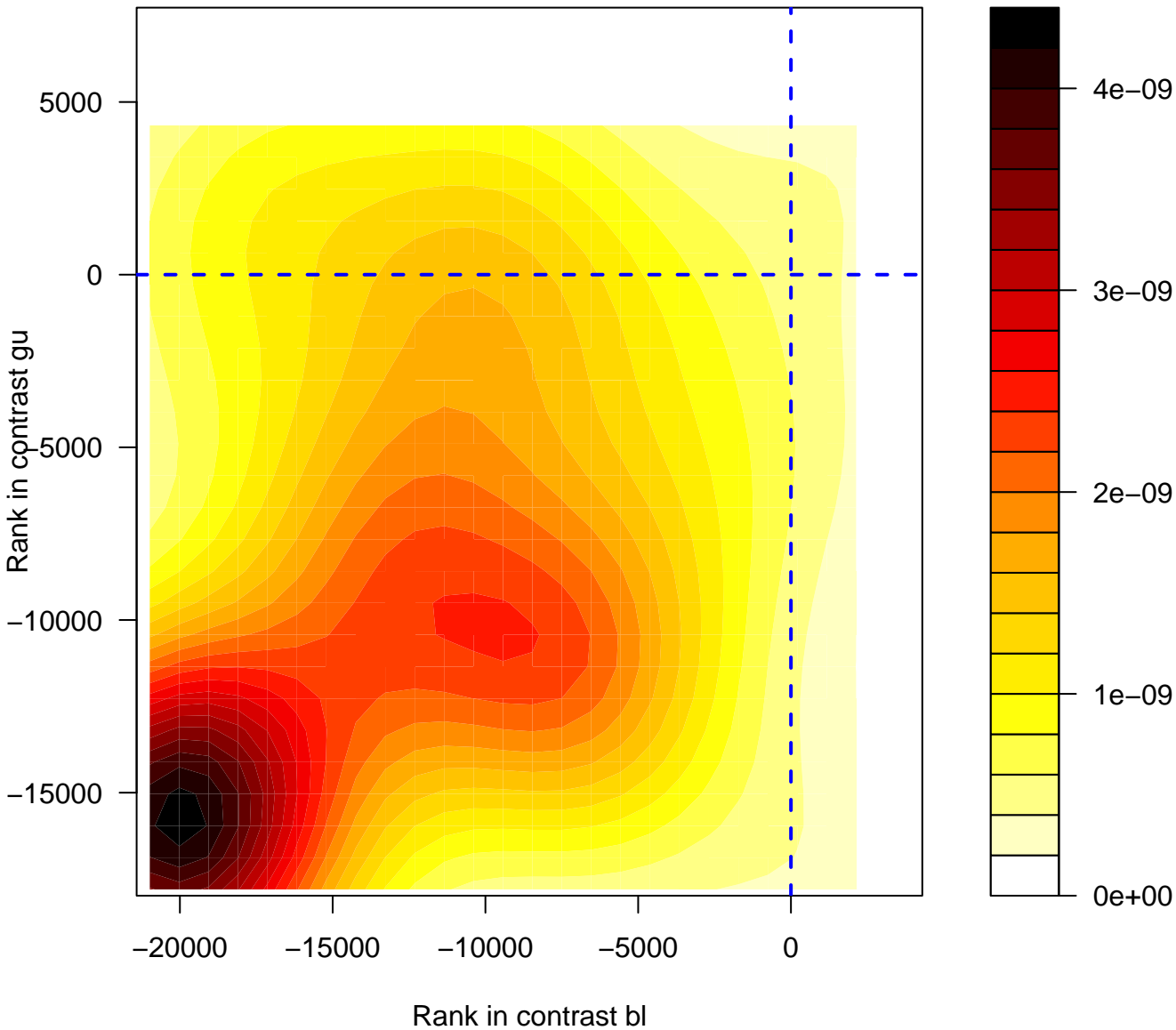
Expression and translocation of olfactory receptors



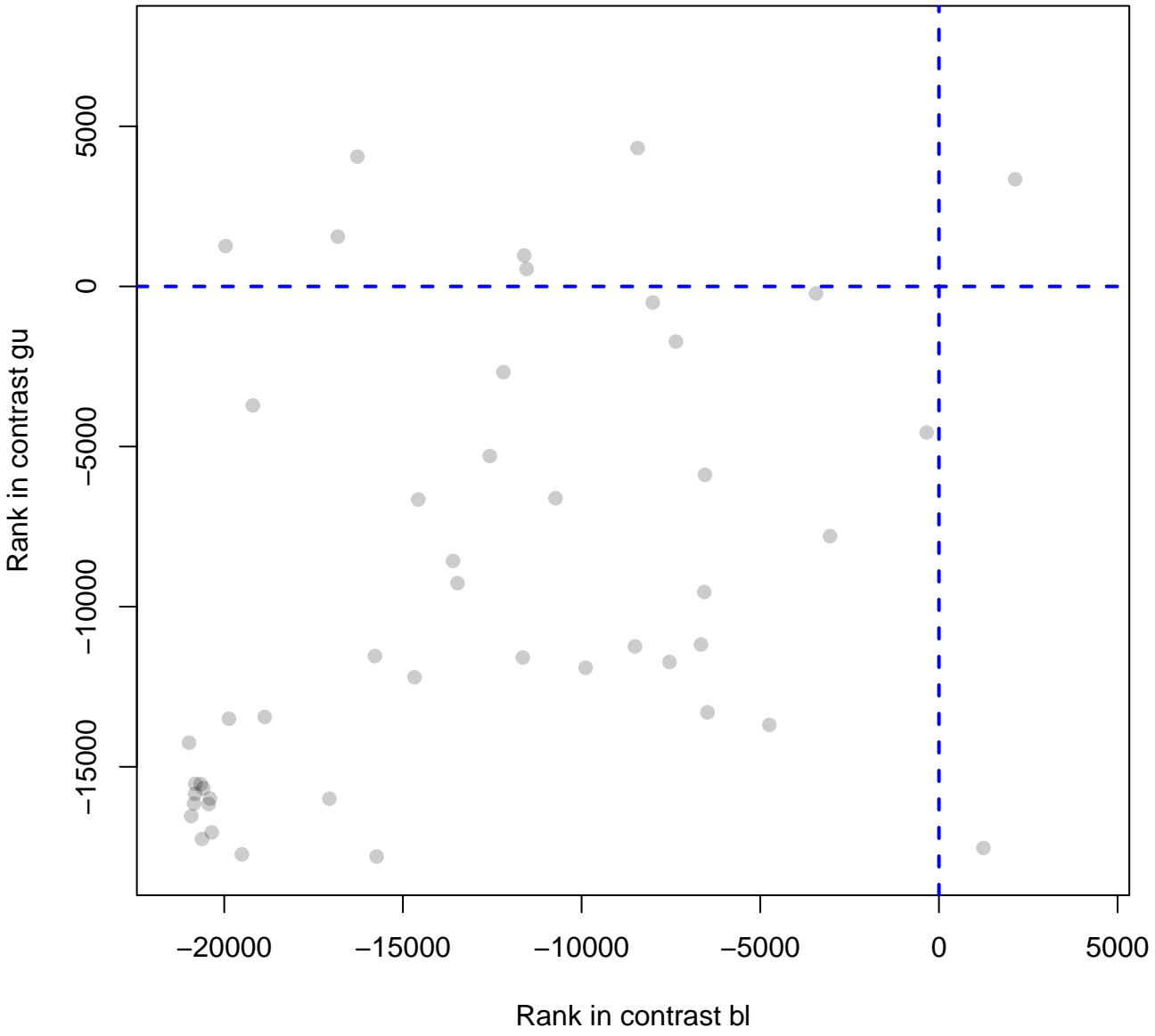
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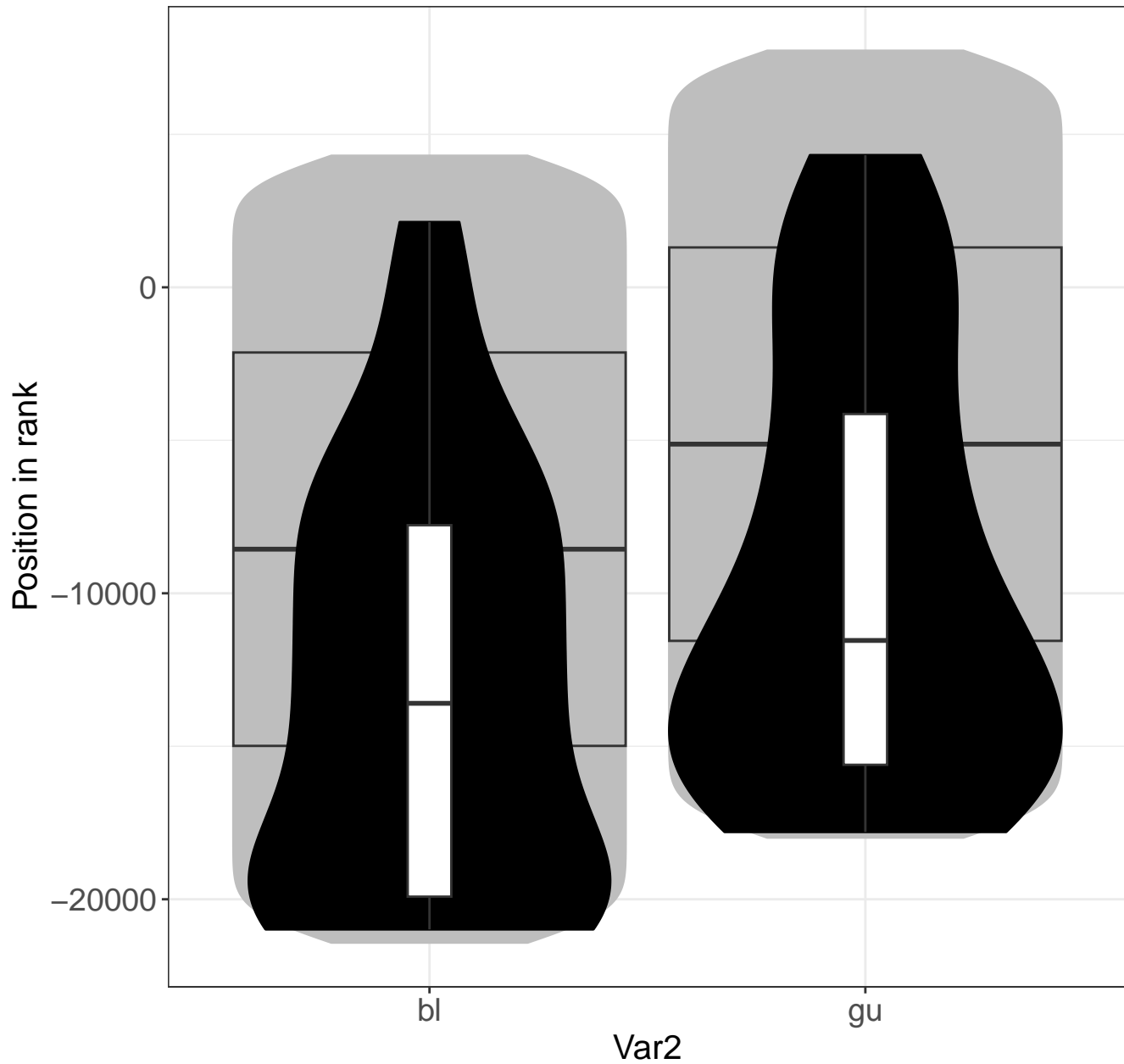
Sensory perception of taste



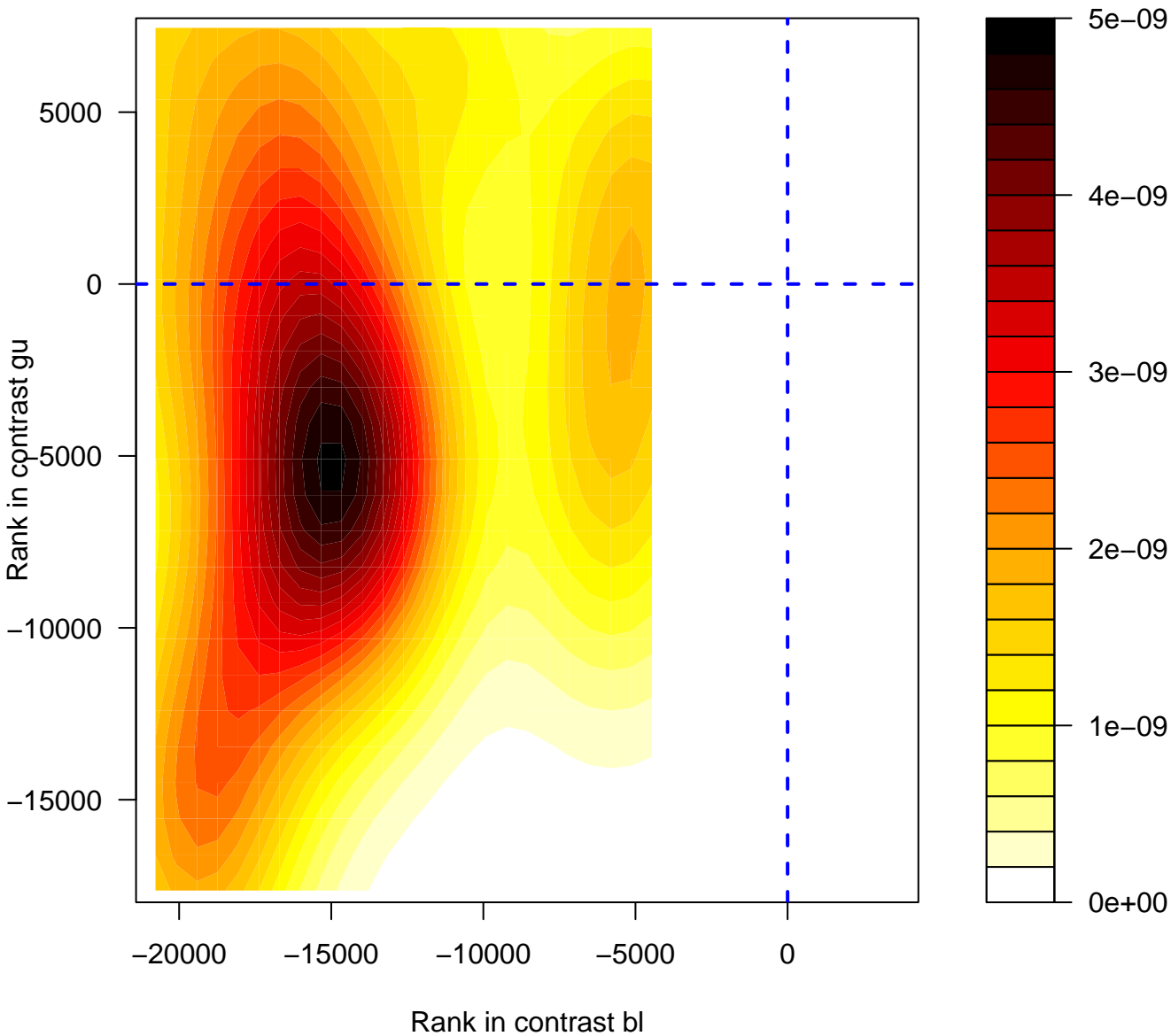
Sensory perception of taste



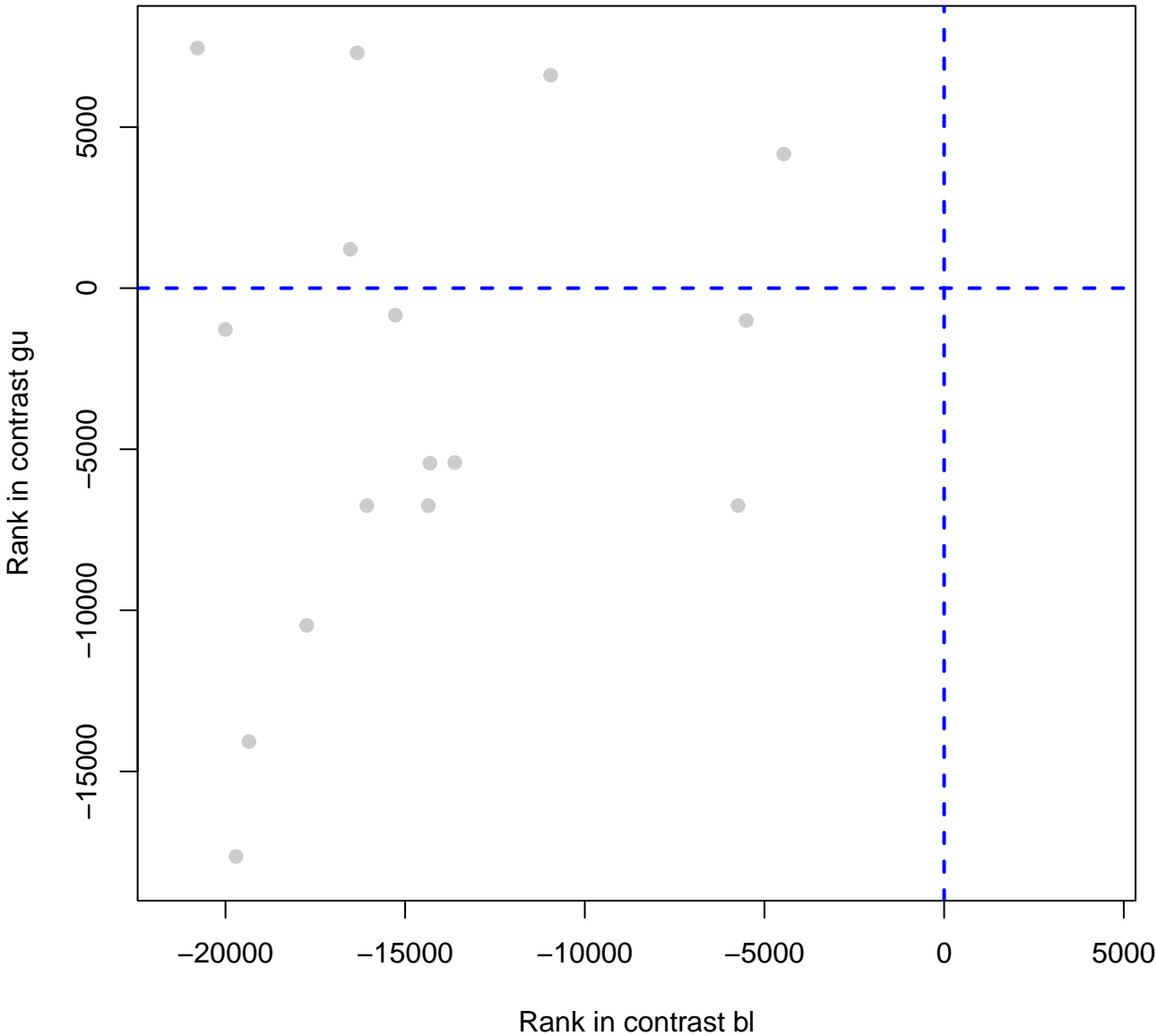
Sensory perception of taste



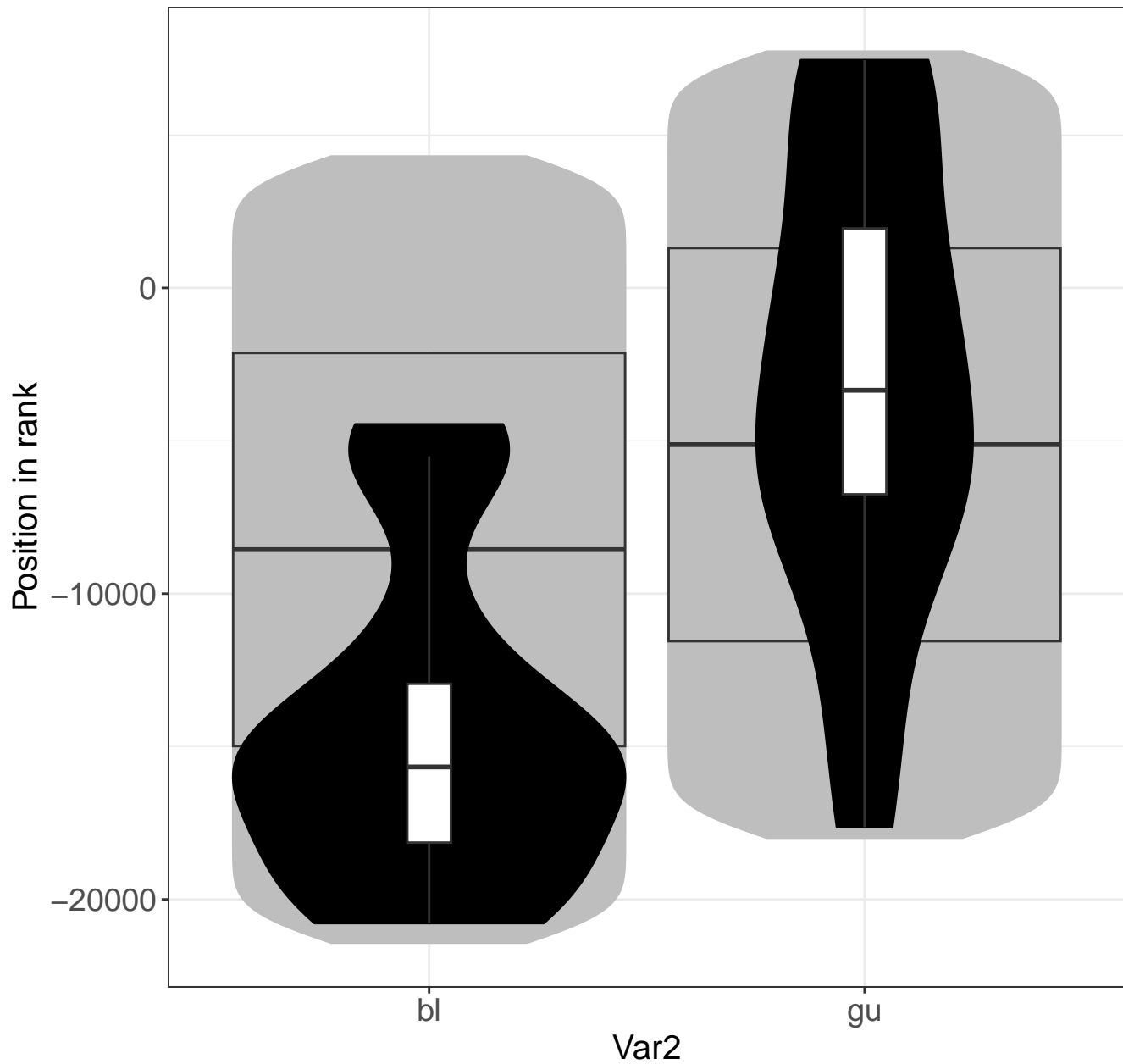
Competing endogenous RNAs (ceRNAs) regulate PTEN trans



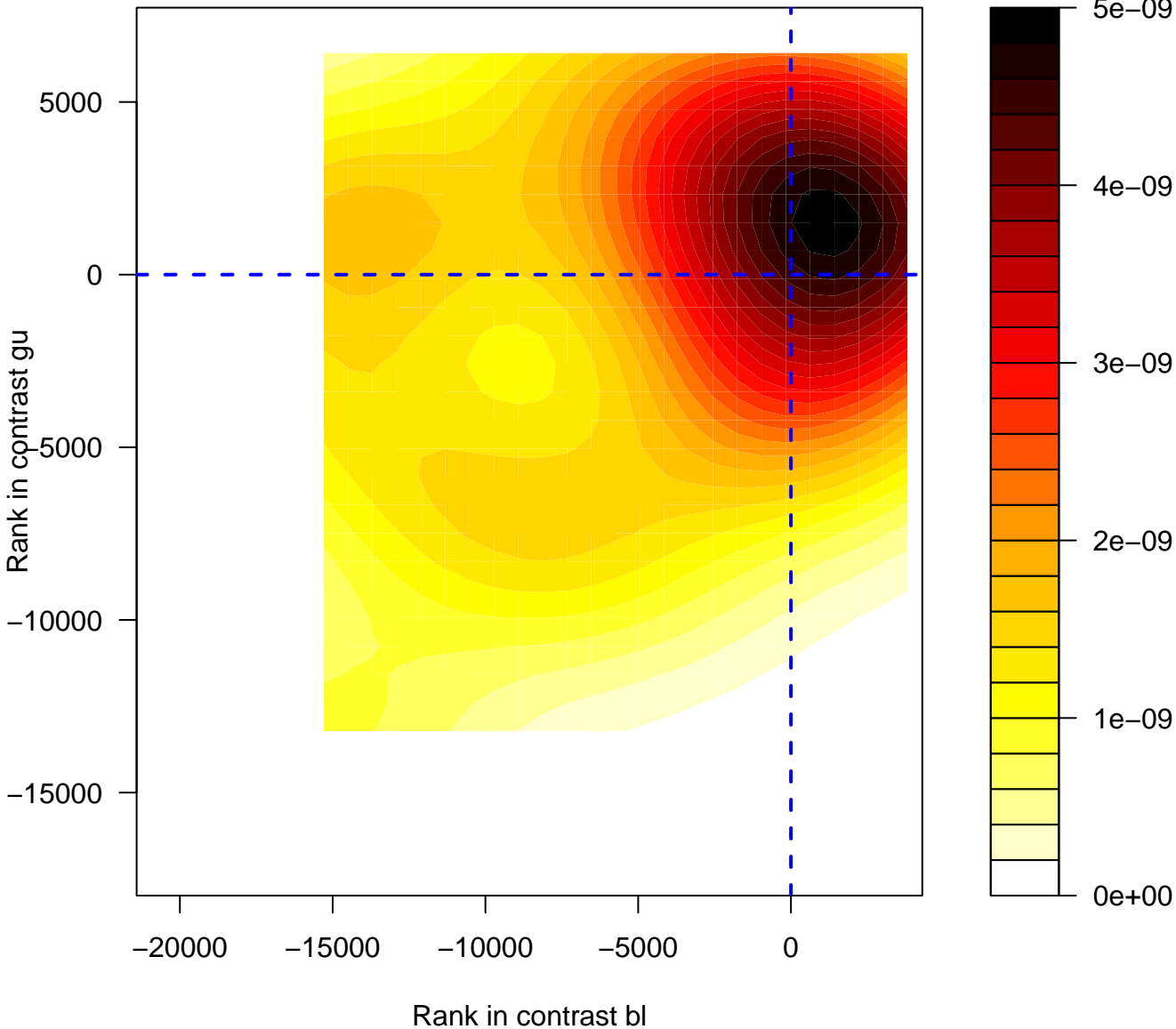
Competing endogenous RNAs (ceRNAs) regulate PTEN translation



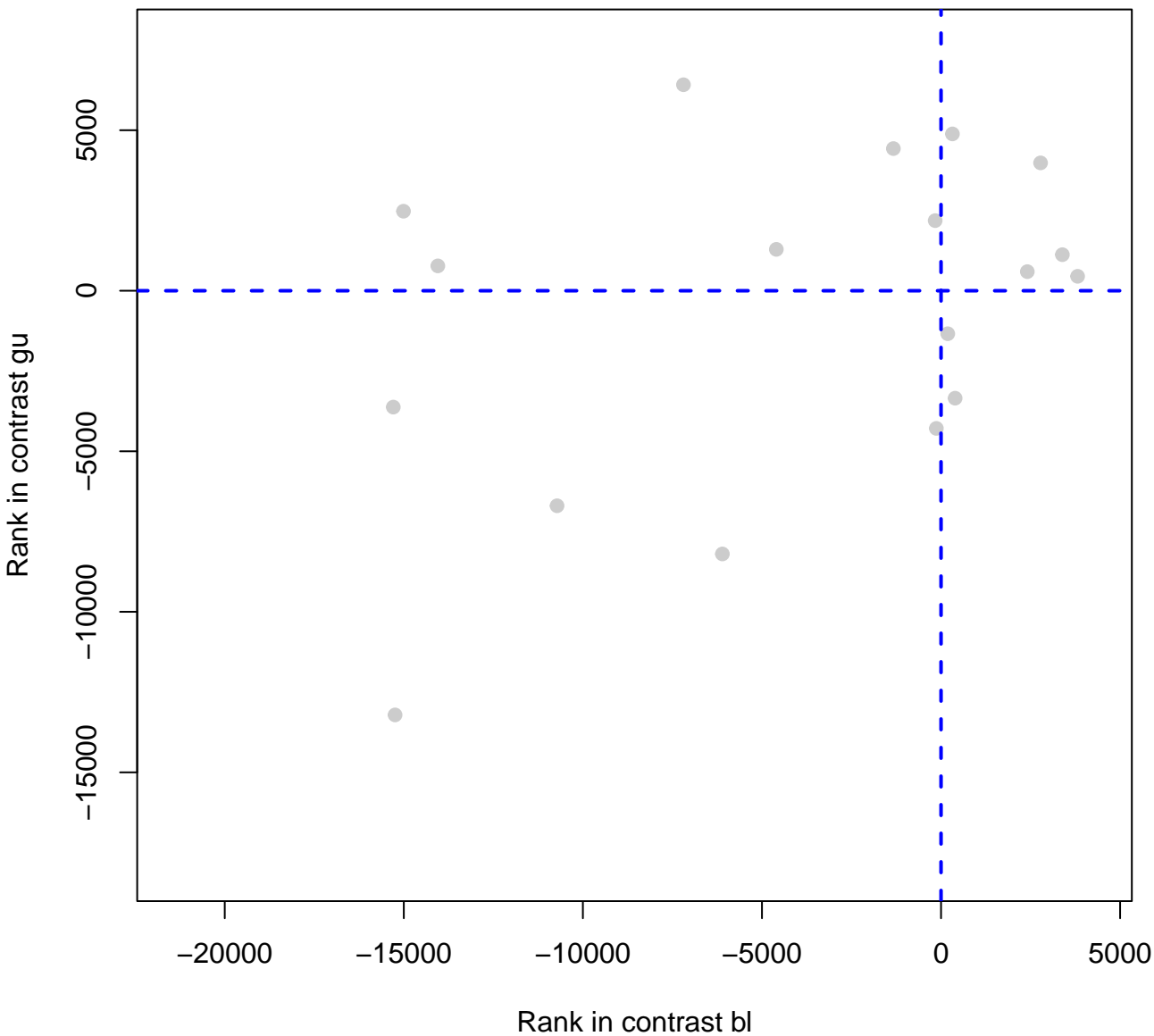
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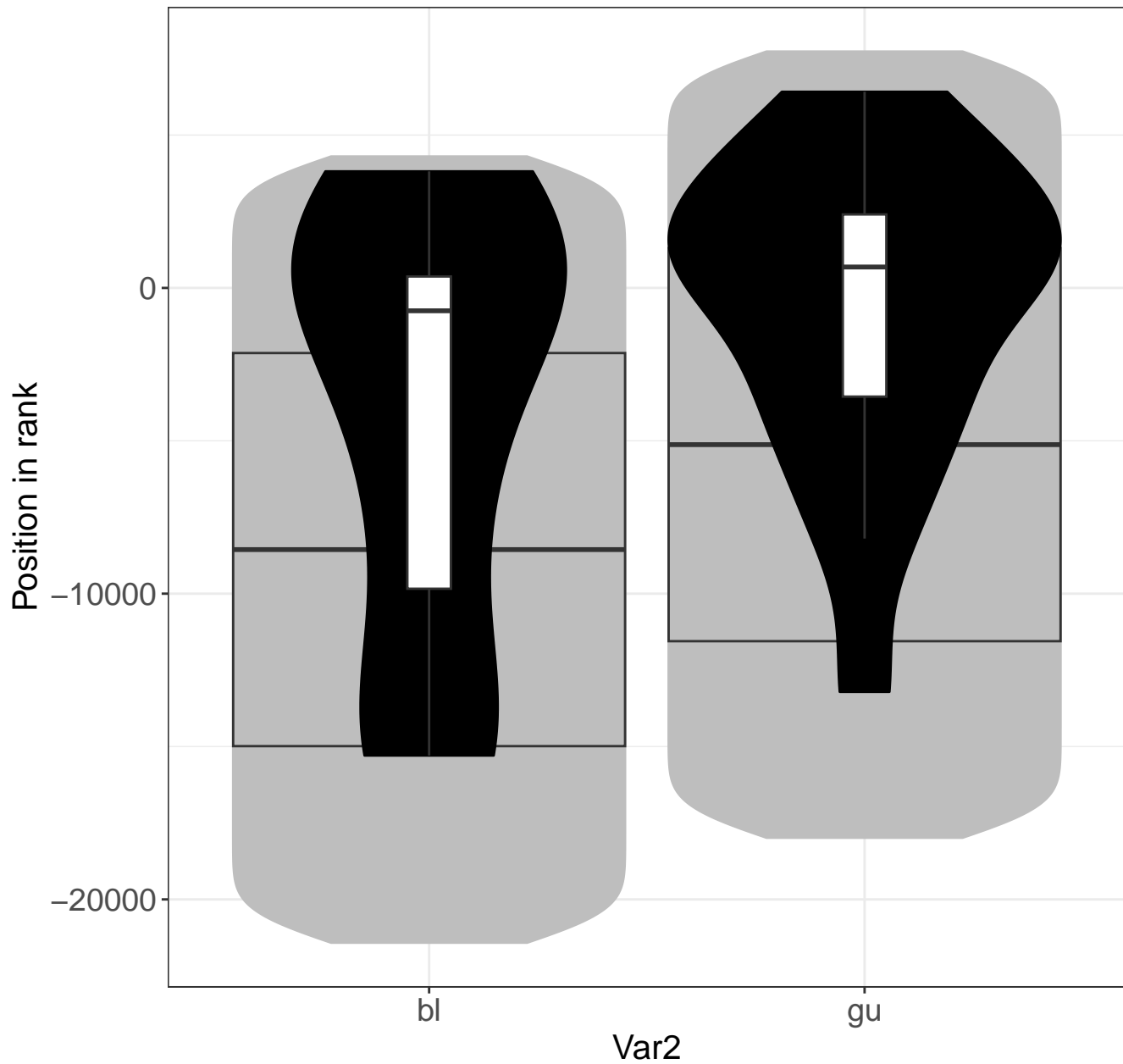
TICAM1, RIP1-mediated IKK complex recruitment



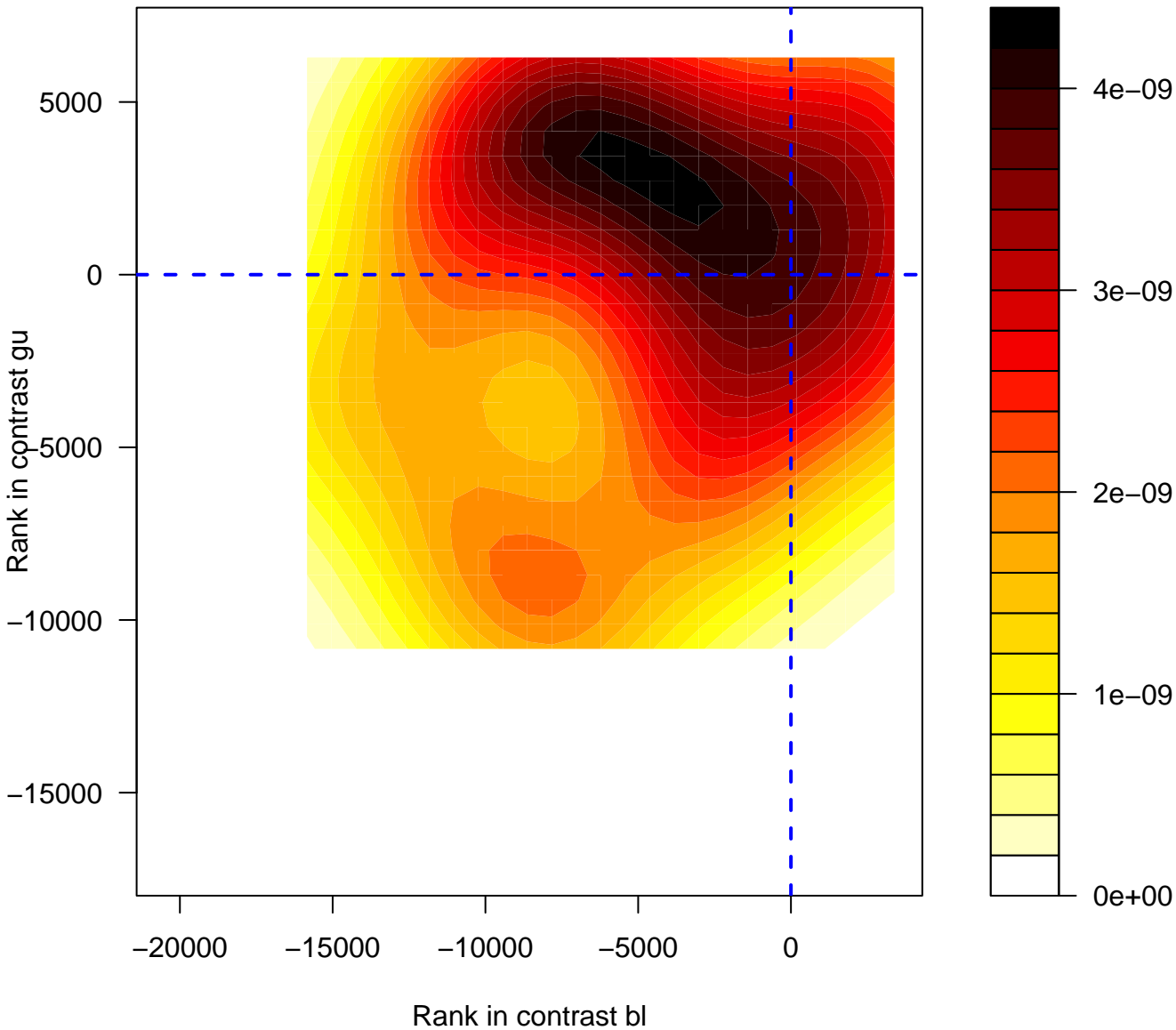
TICAM1, RIP1-mediated IKK complex recruitment



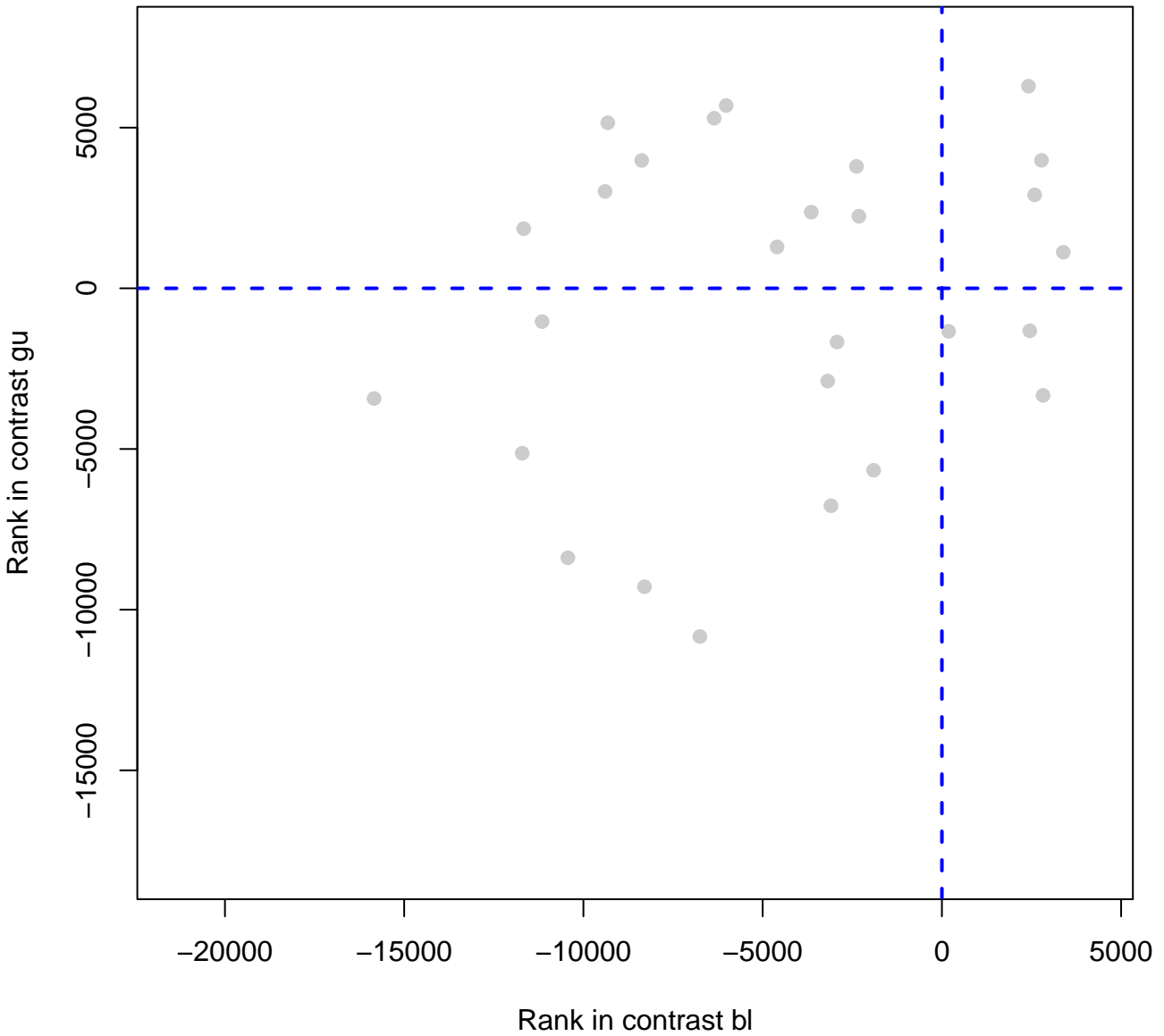
TICAM1, RIP1-mediated IKK complex recruitme



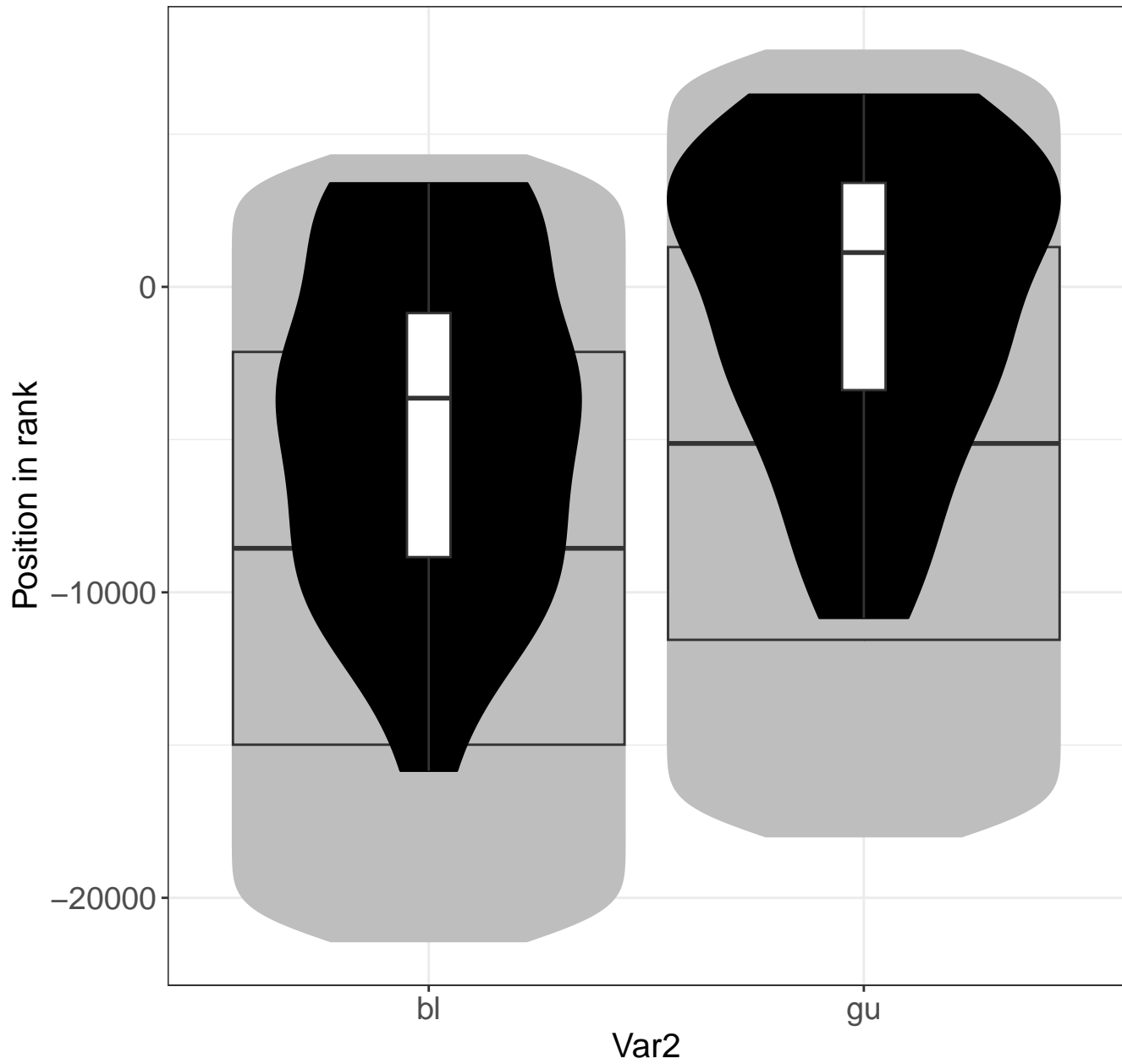
Mitophagy



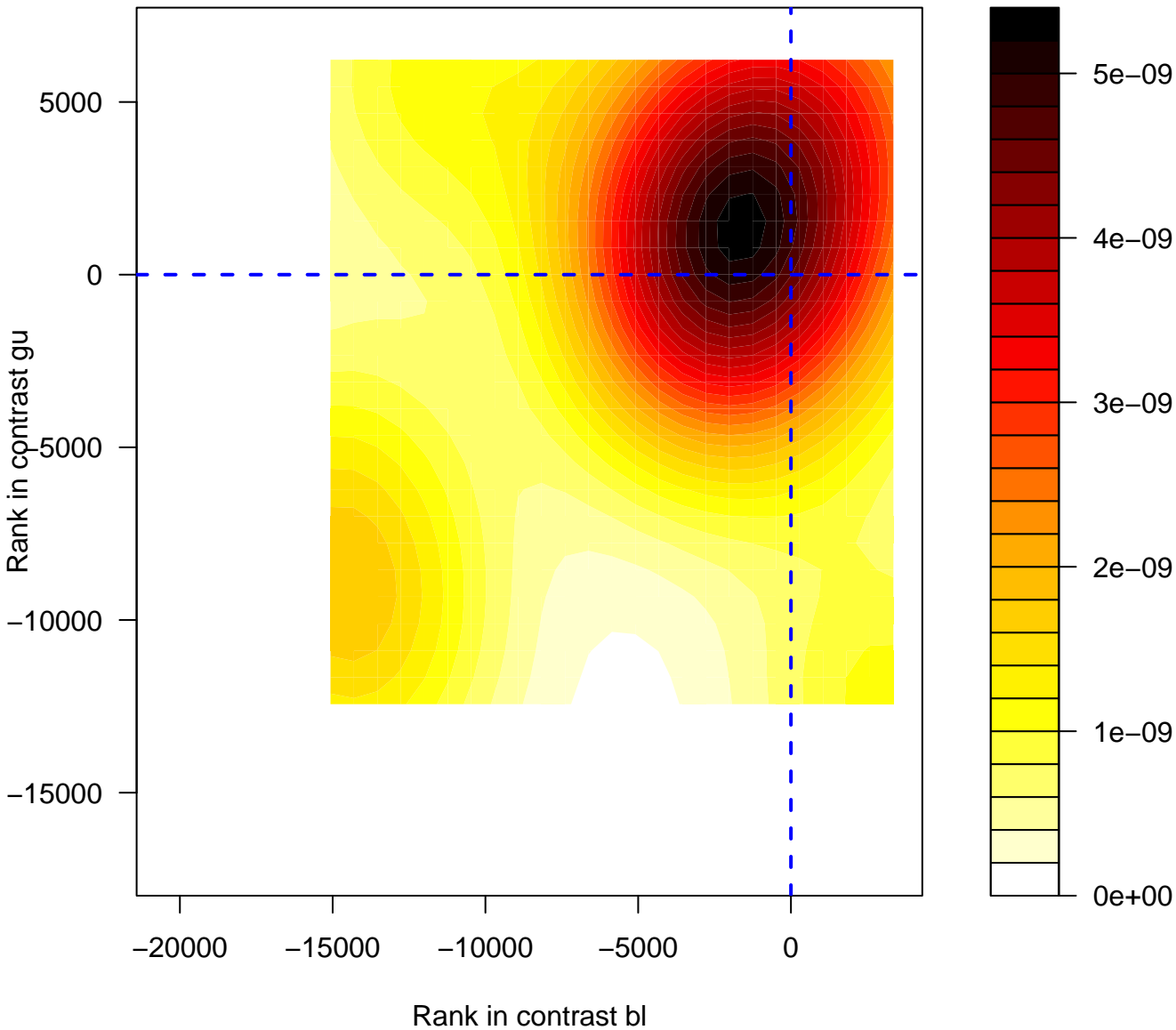
Mitophagy



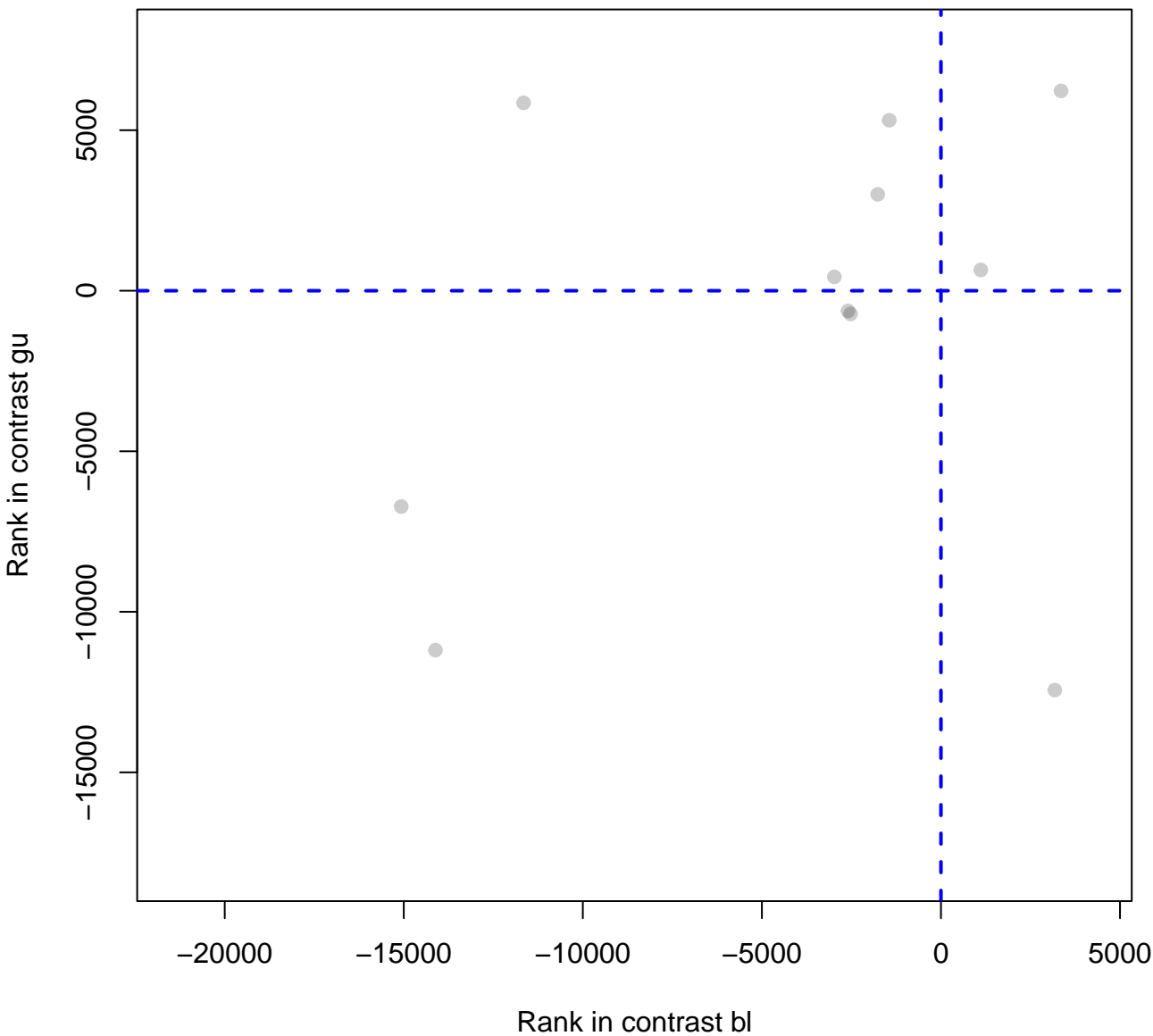
Mitophagy



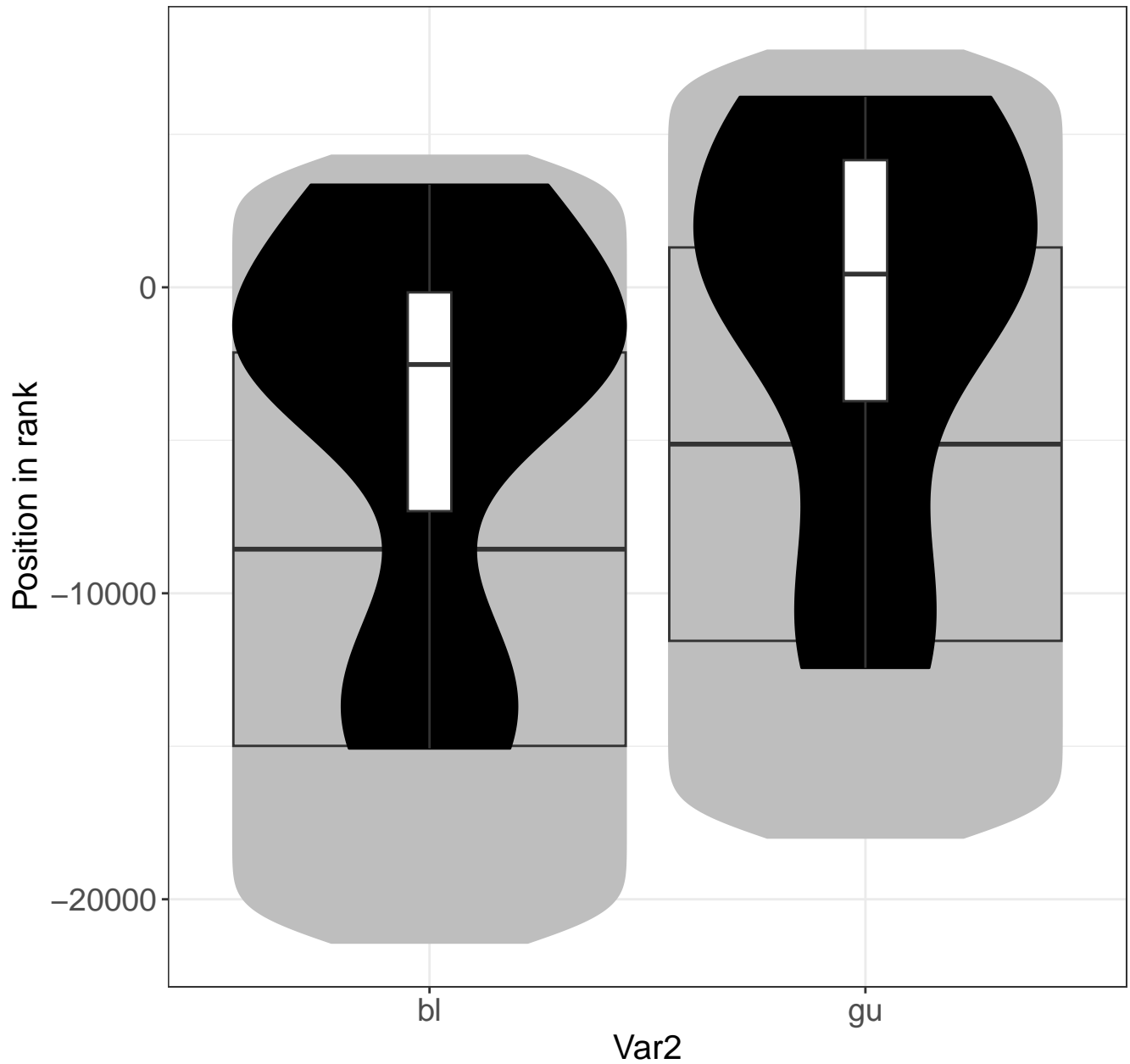
Activation of the TFAP2 (AP-2) family of transcription fact



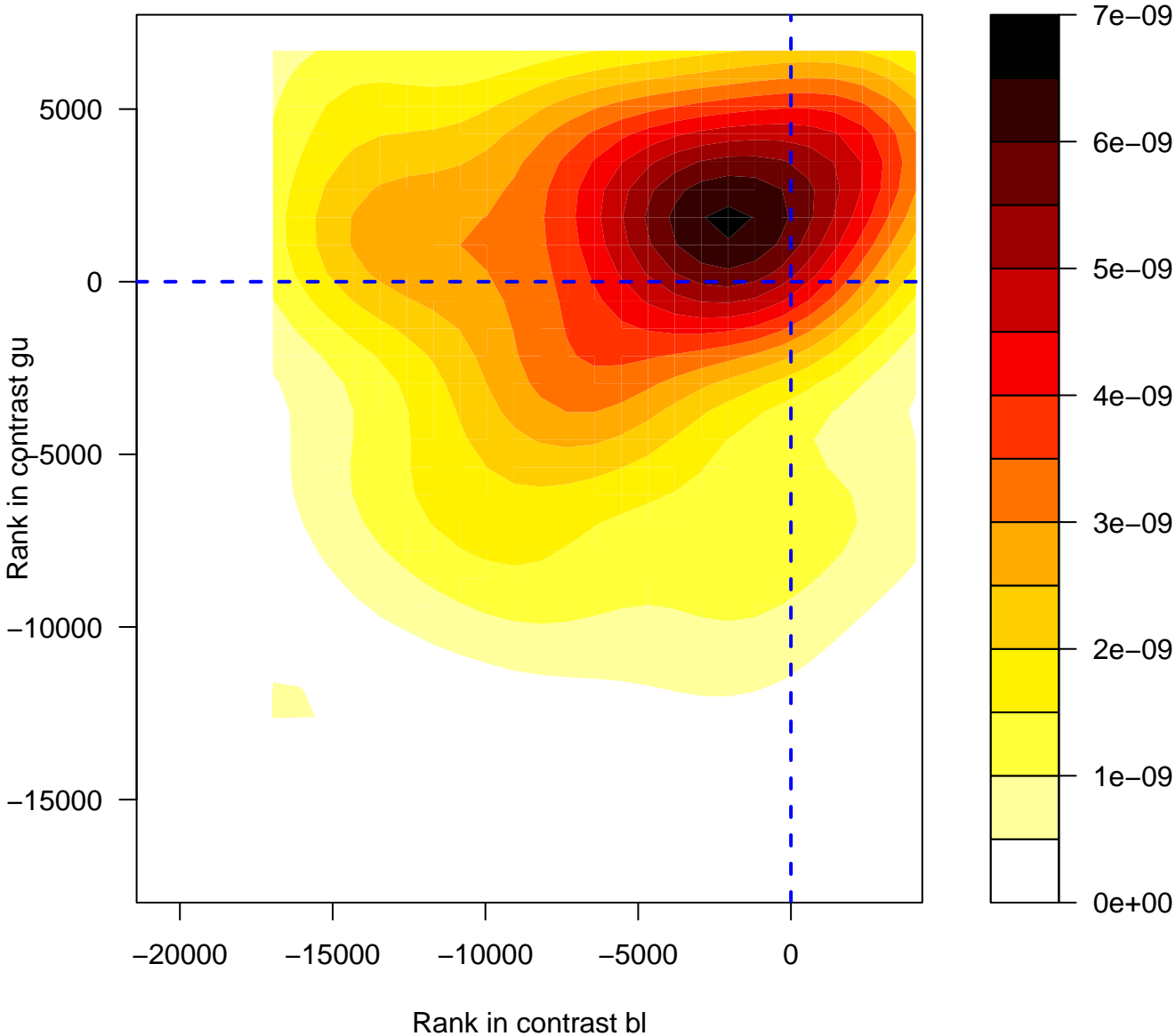
Activation of the TFAP2 (AP-2) family of transcription factors



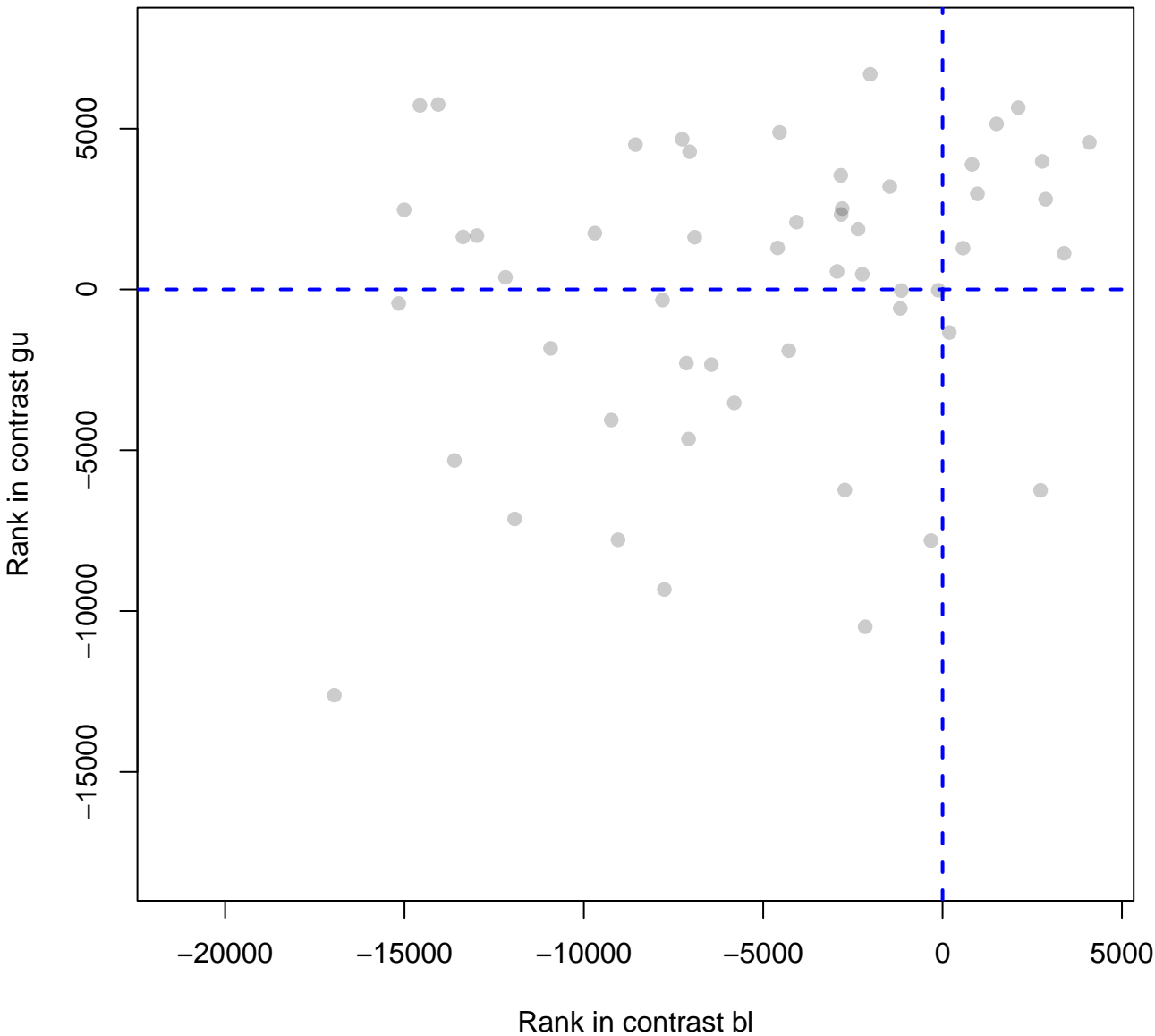
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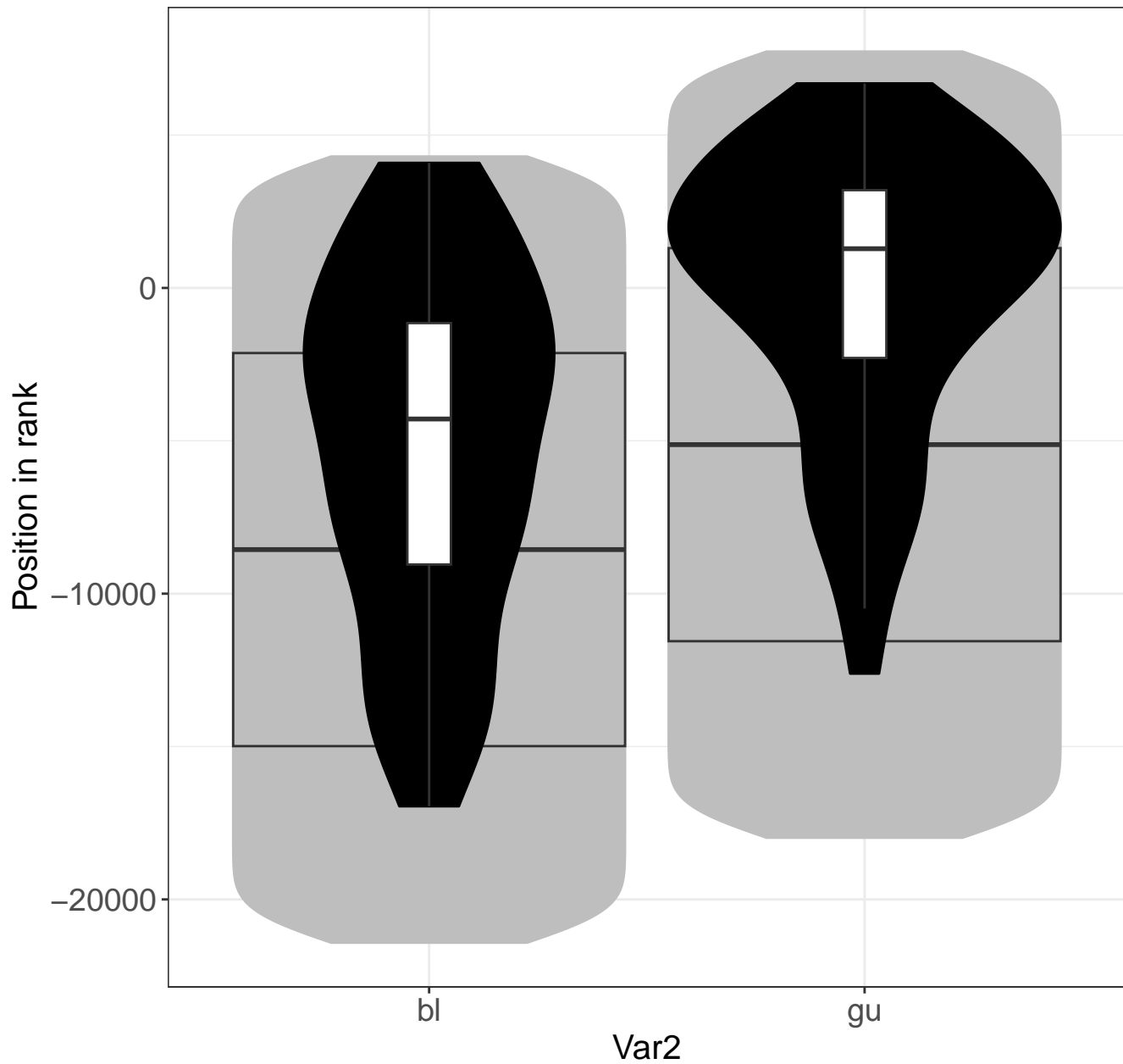
Senescence-Associated Secretory Phenotype (SASP)



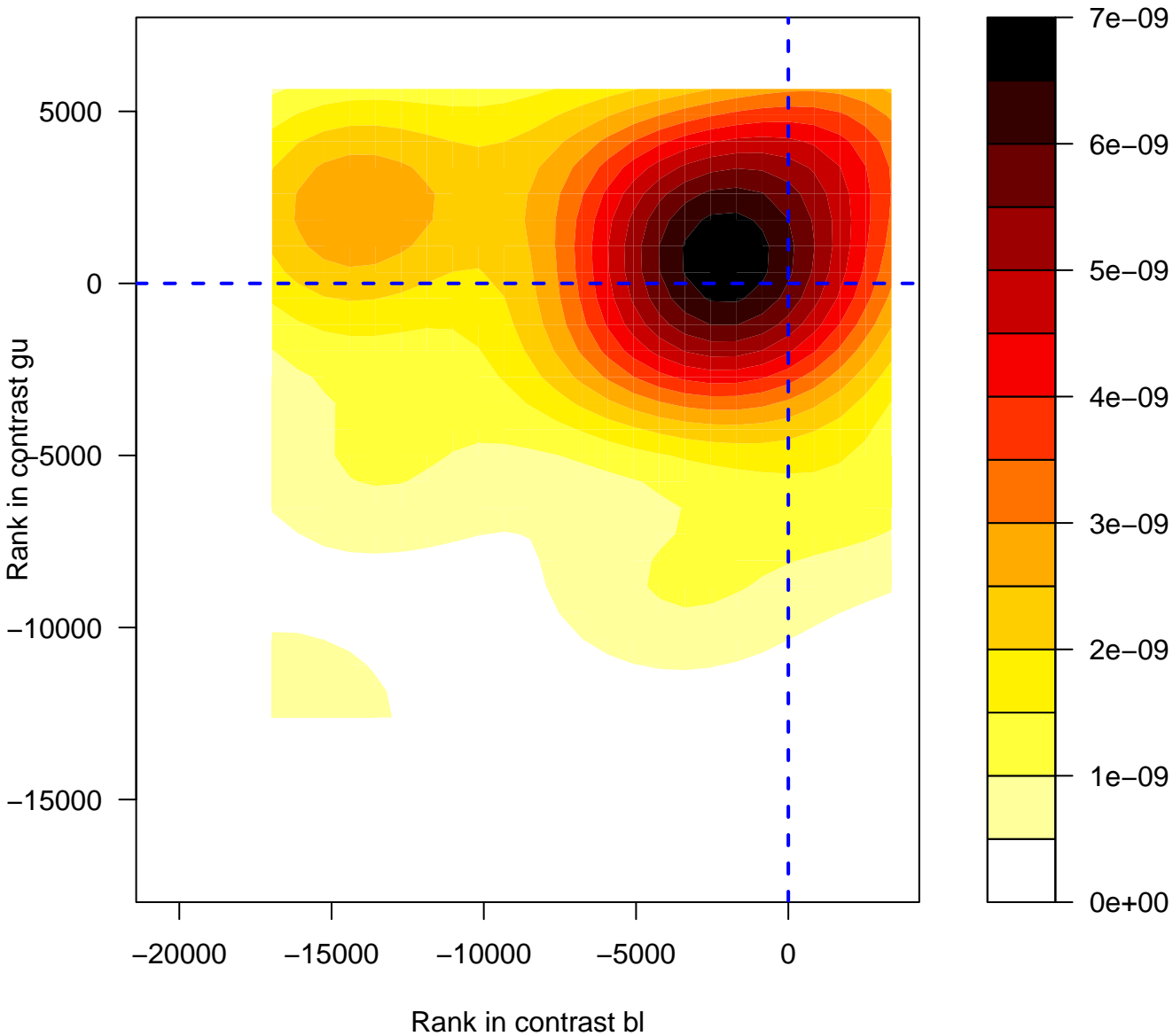
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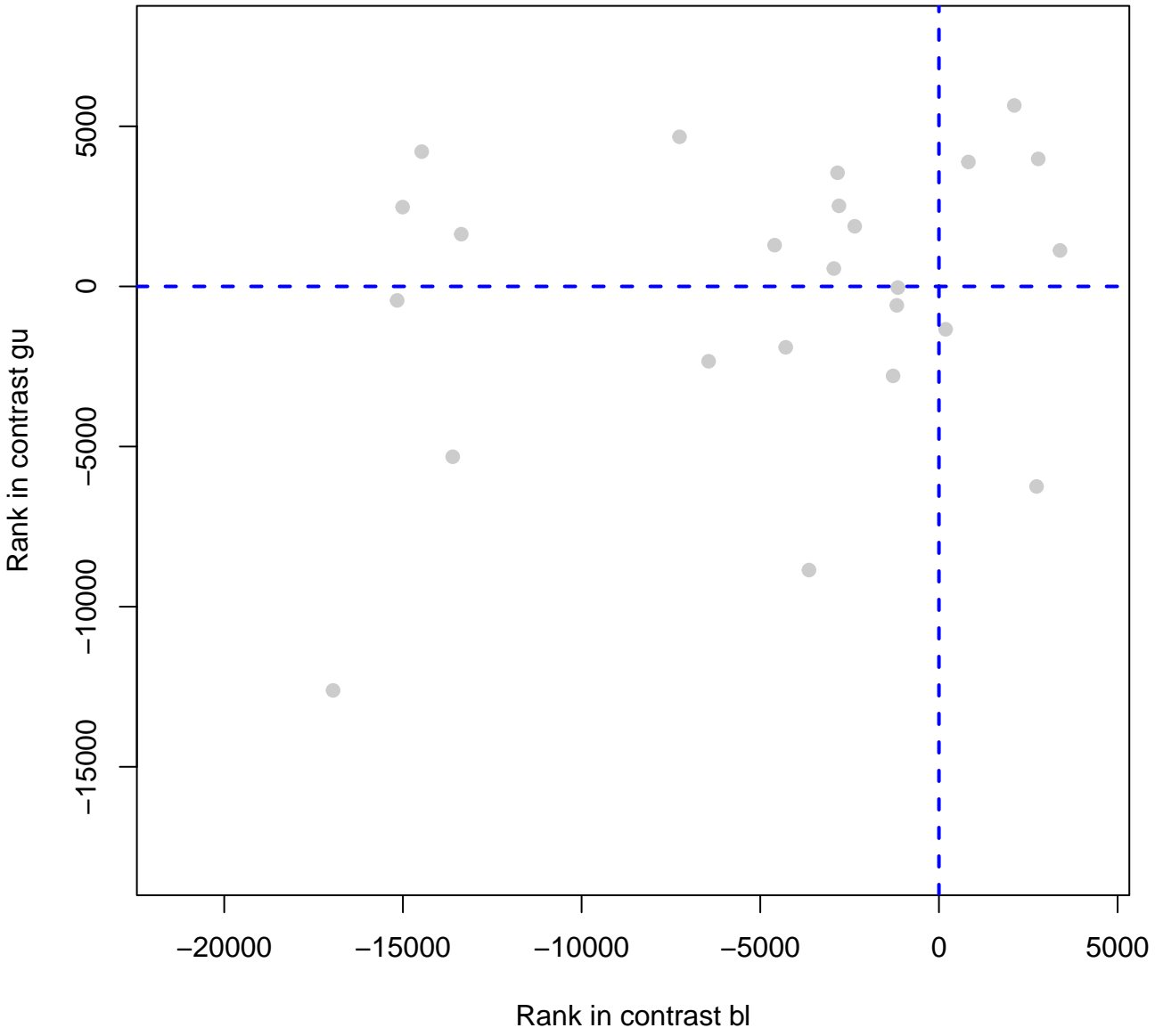
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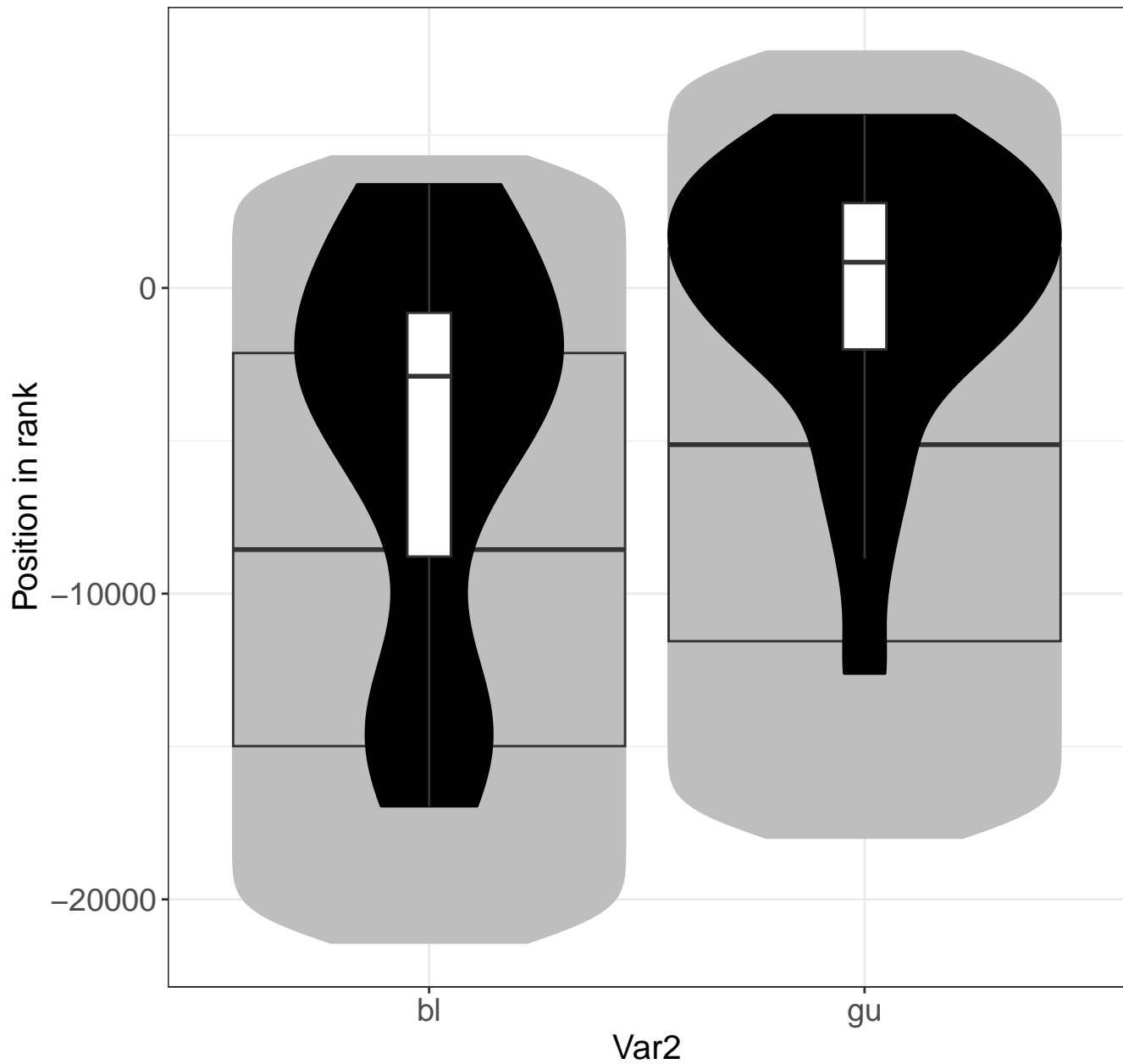
APC/C:Cdc20 mediated degradation of Cyclin B



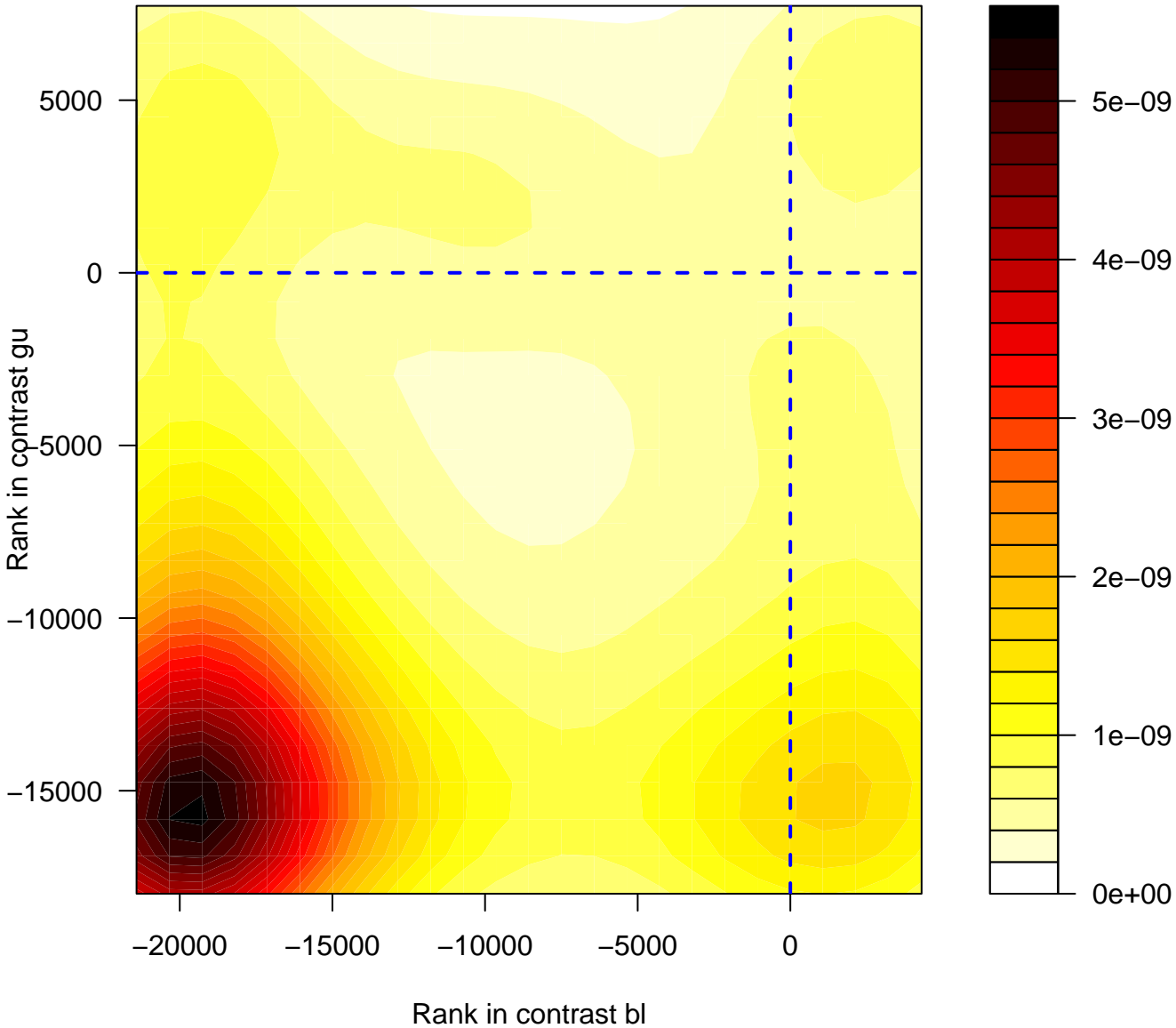
APC/C:Cdc20 mediated degradation of Cyclin B



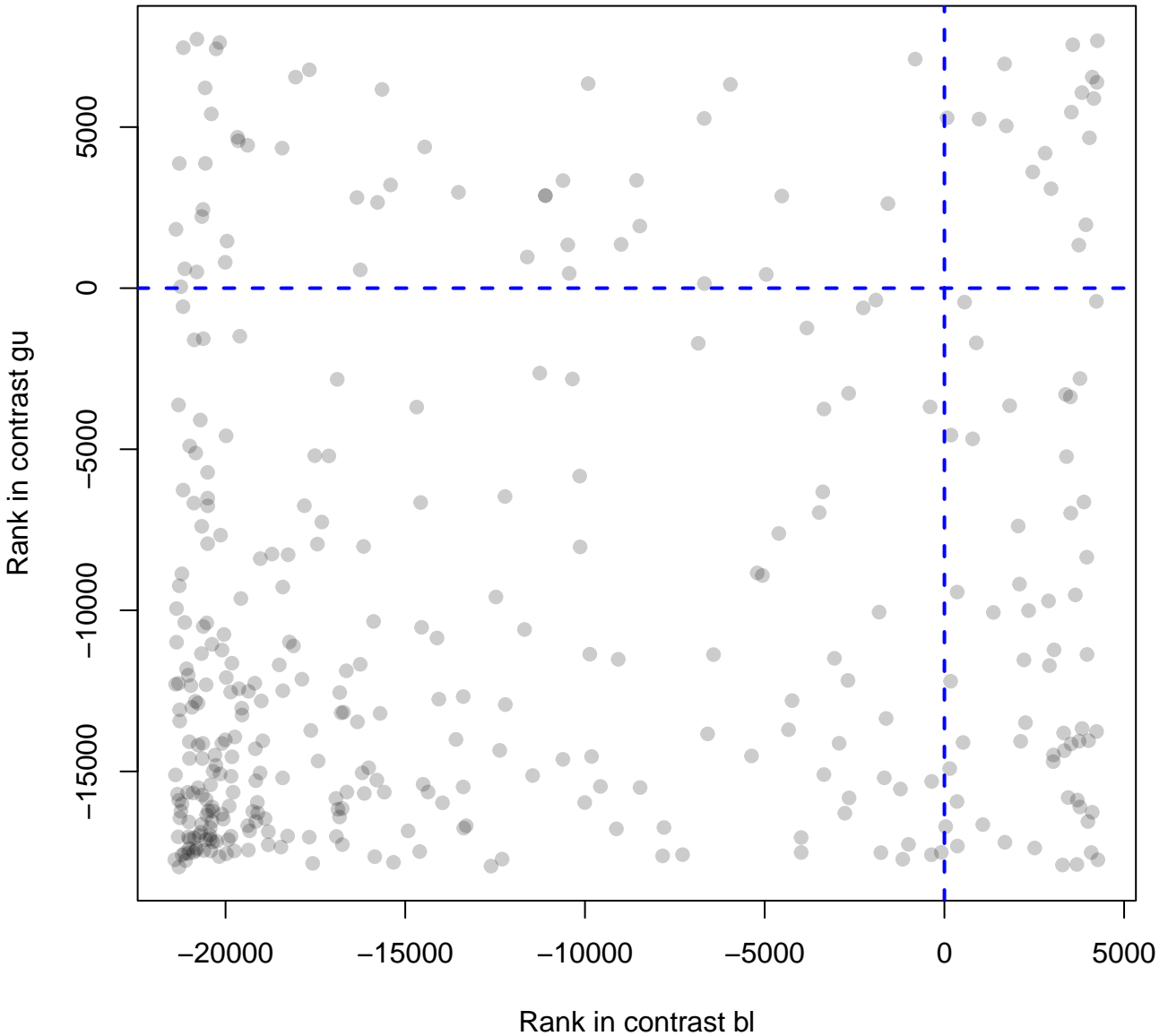
APC/C:Cdc20 mediated degradation of Cyclin B



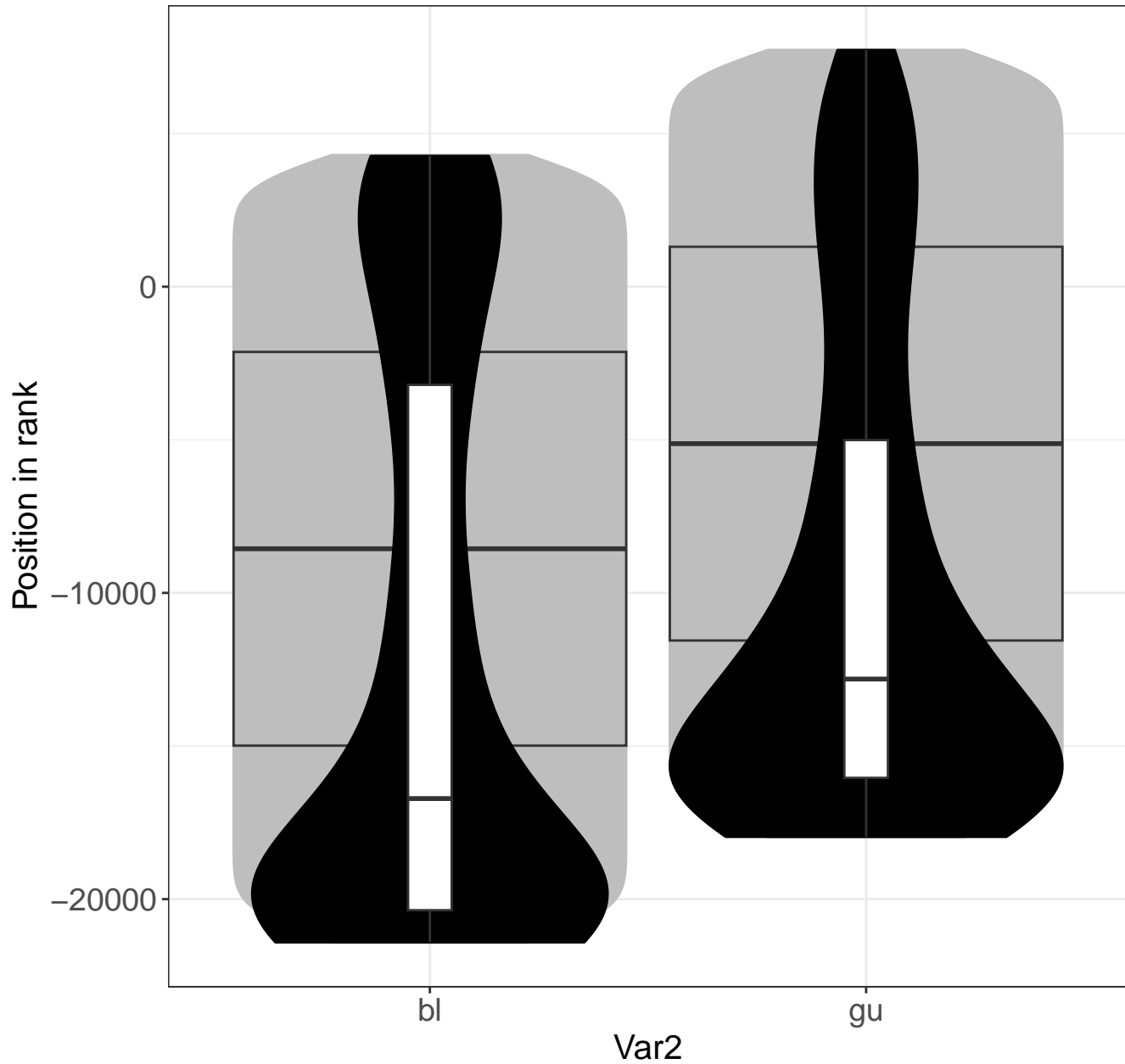
Olfactory Signaling Pathway



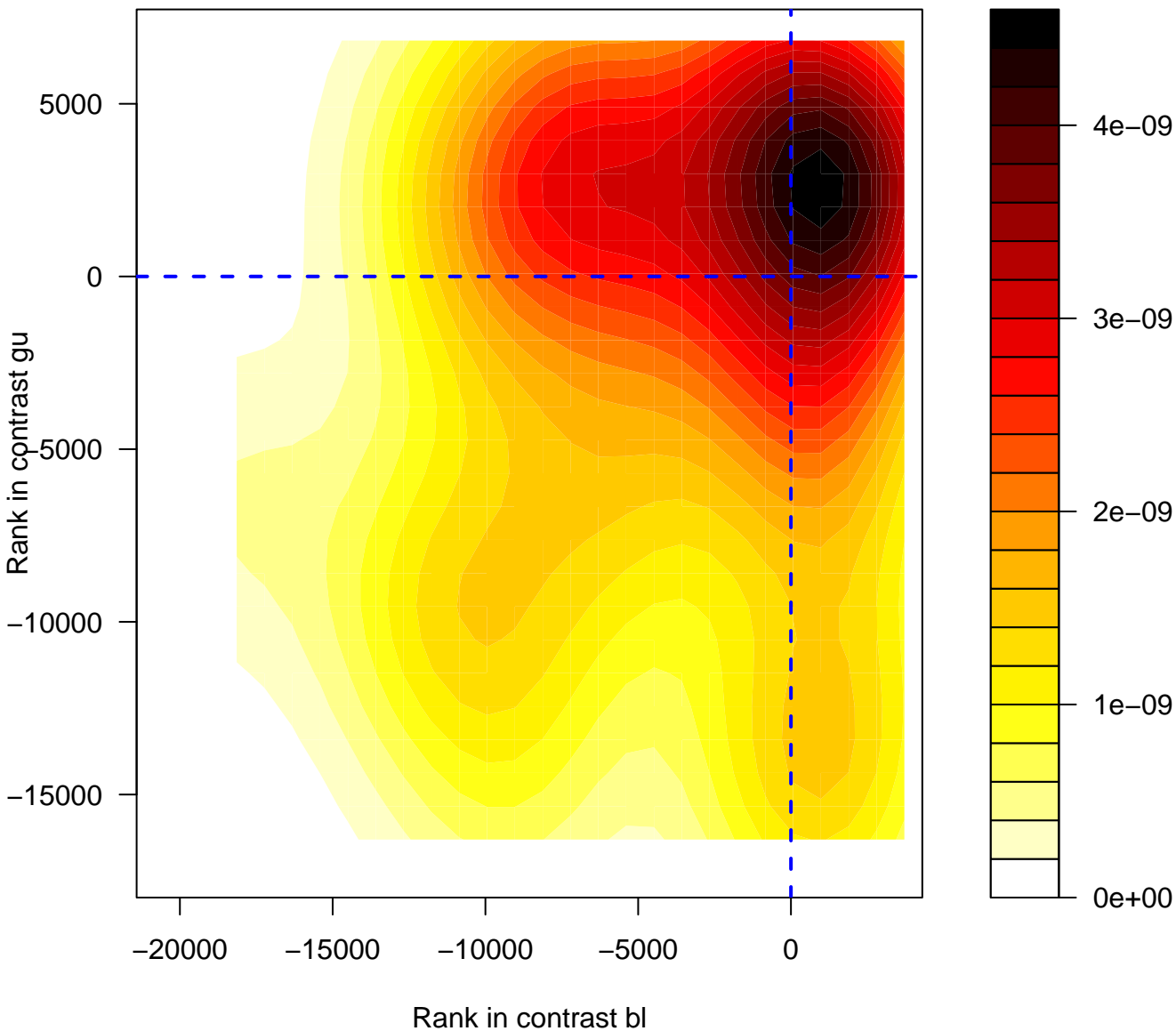
Olfactory Signaling Pathway



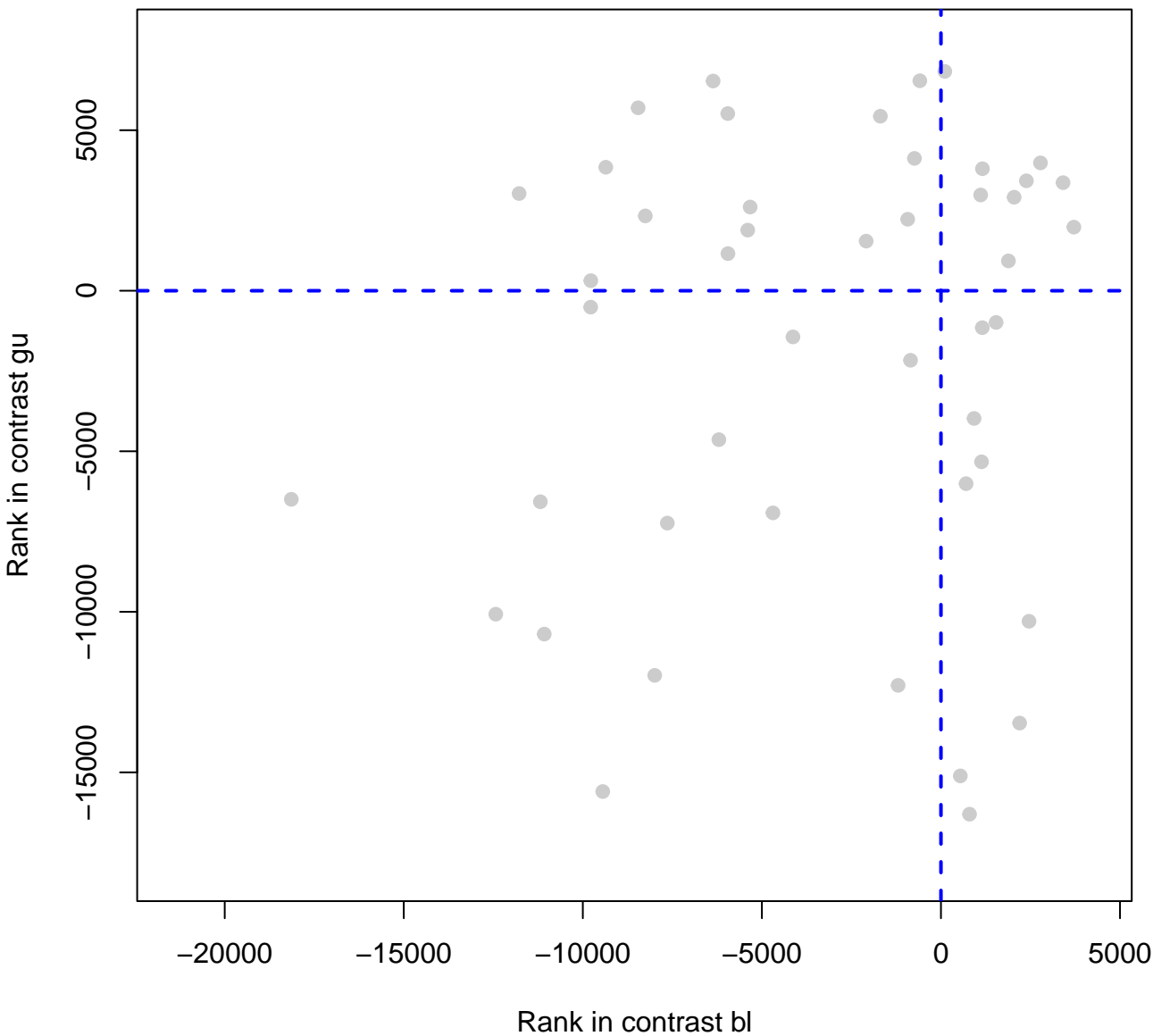
Olfactory Signaling Pathway



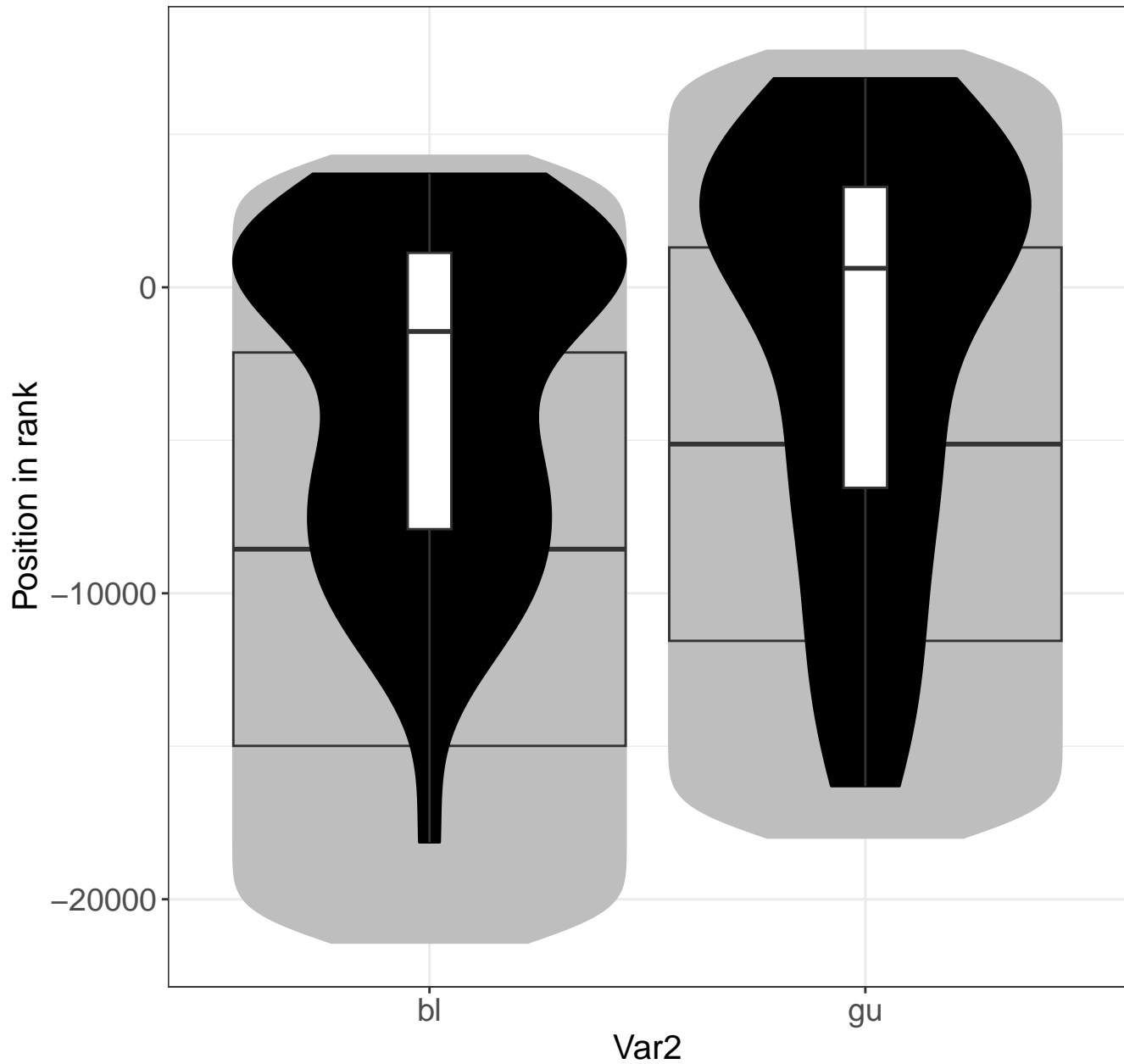
SARS-CoV-2 modulates host translation machinery



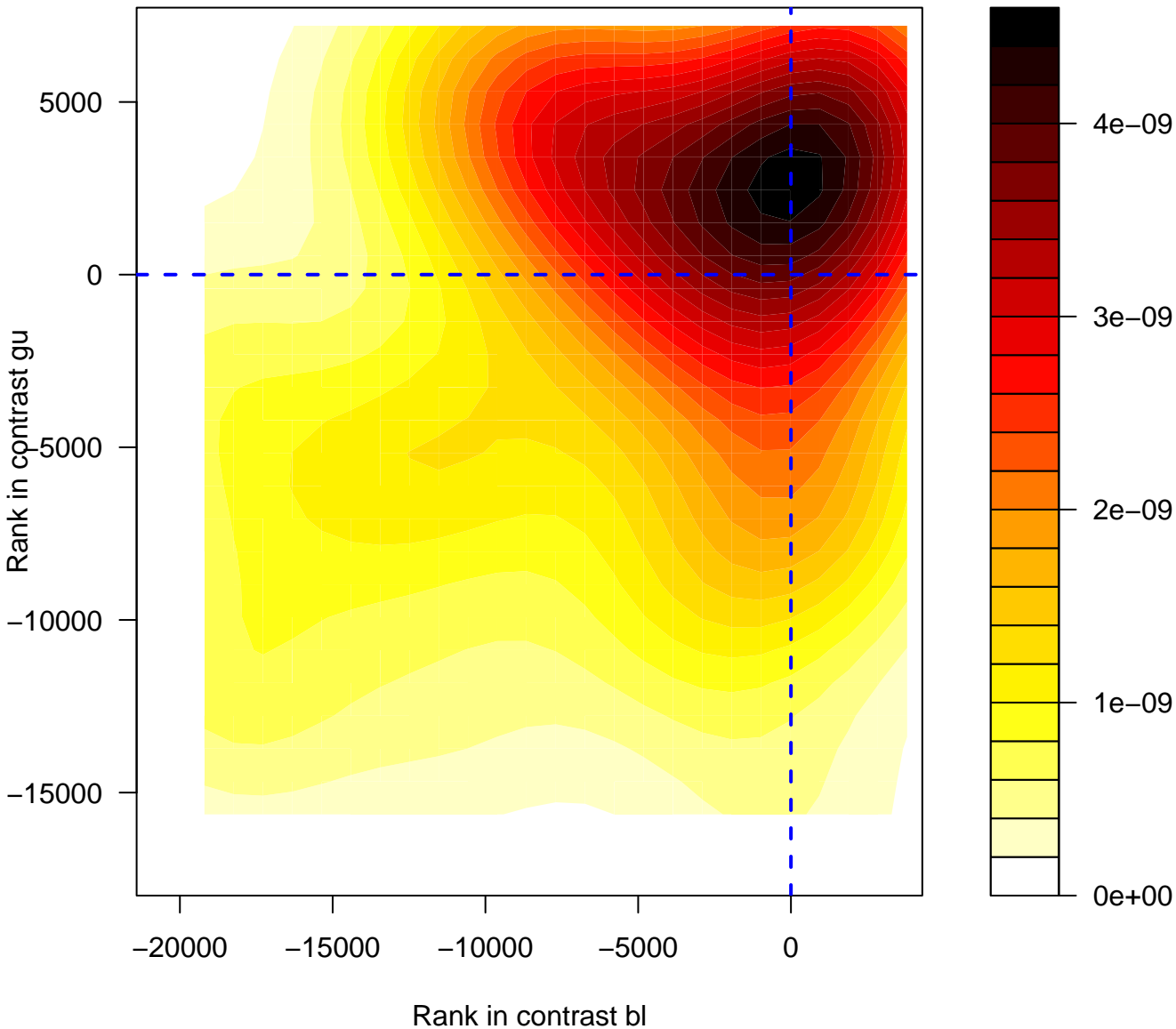
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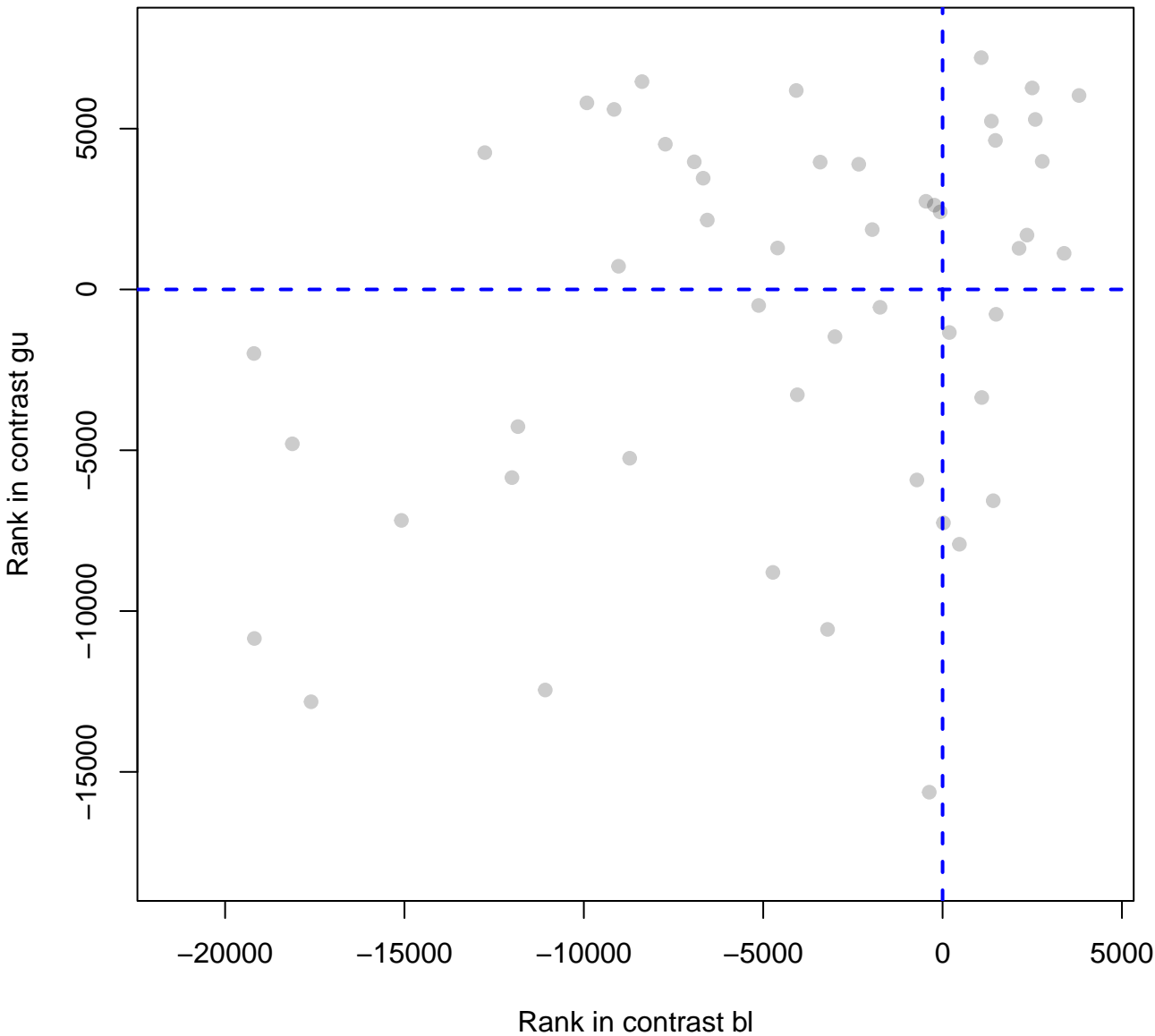
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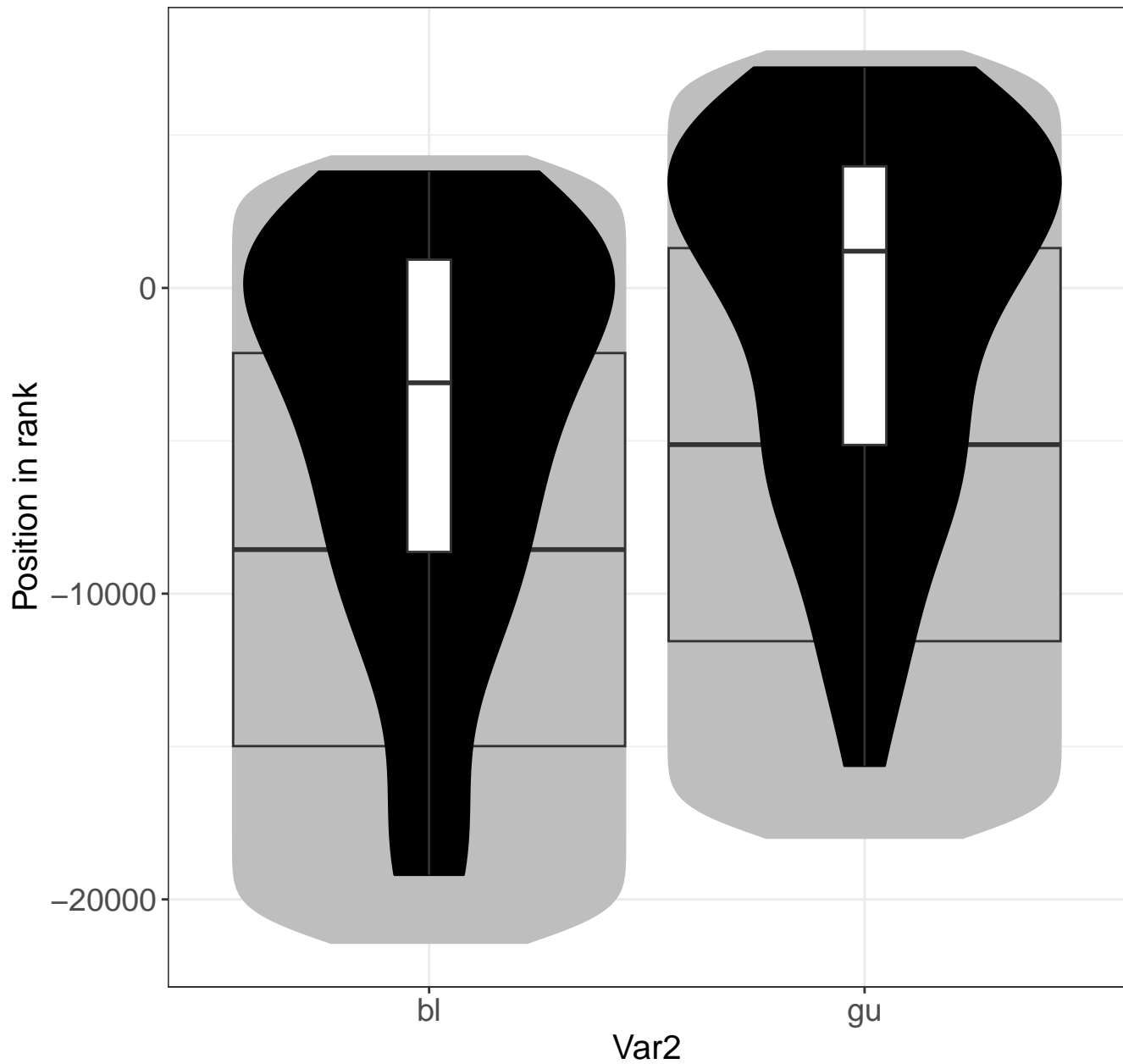
GSK3B and BTRC:CUL1-mediated-degradation of NFE2



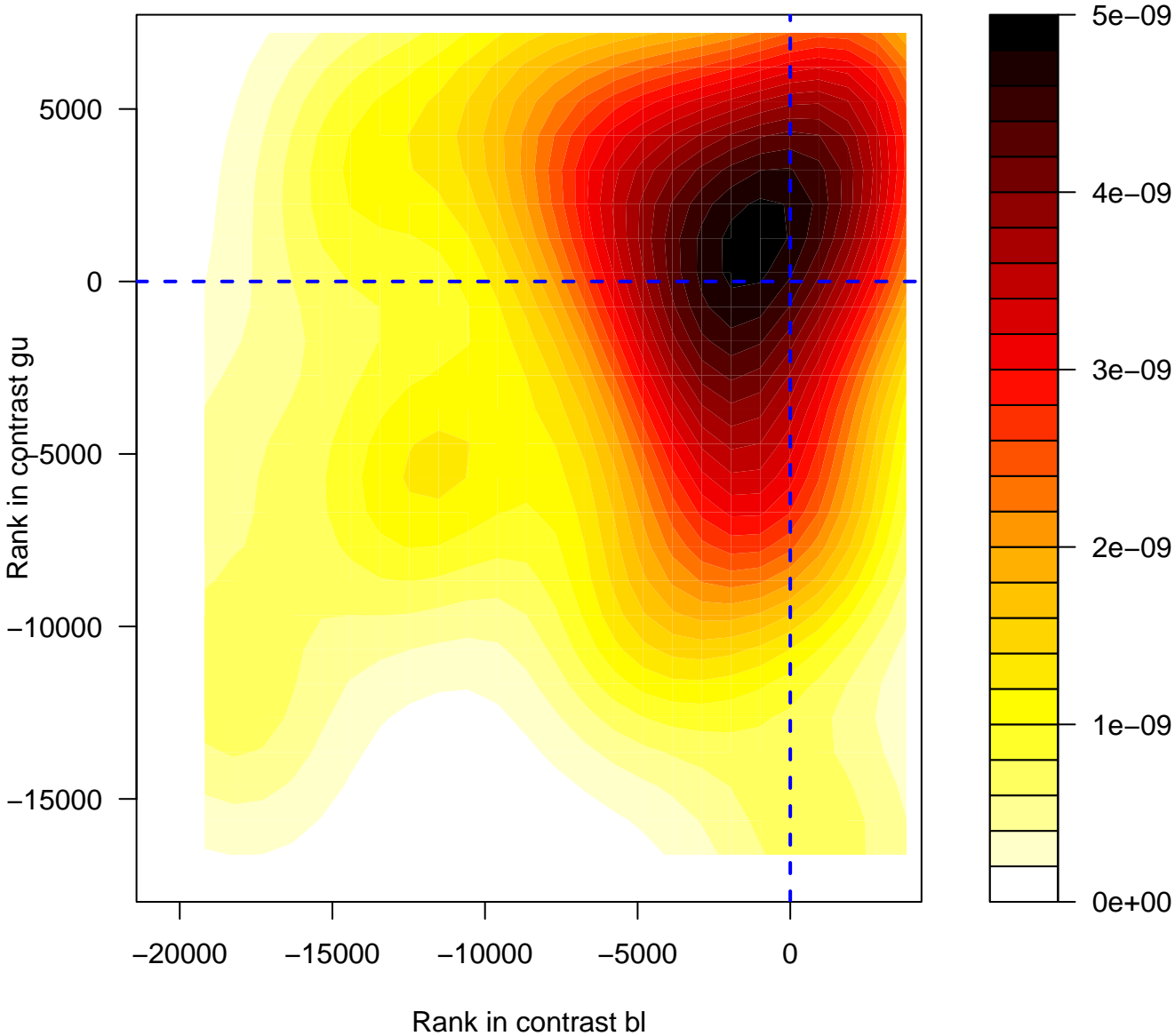
GSK3B and BTRC:CUL1-mediated-degradation of NFE2L2



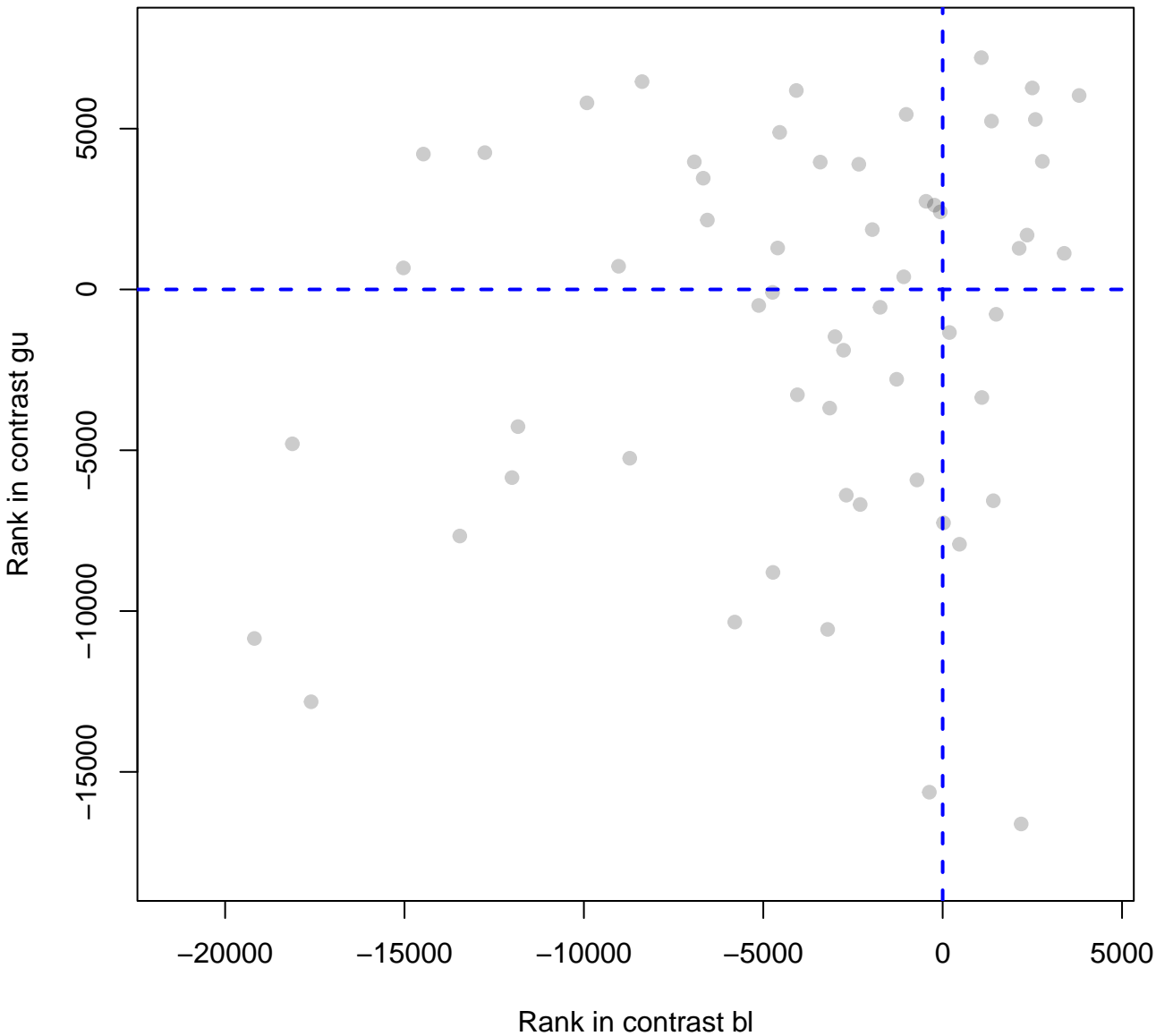
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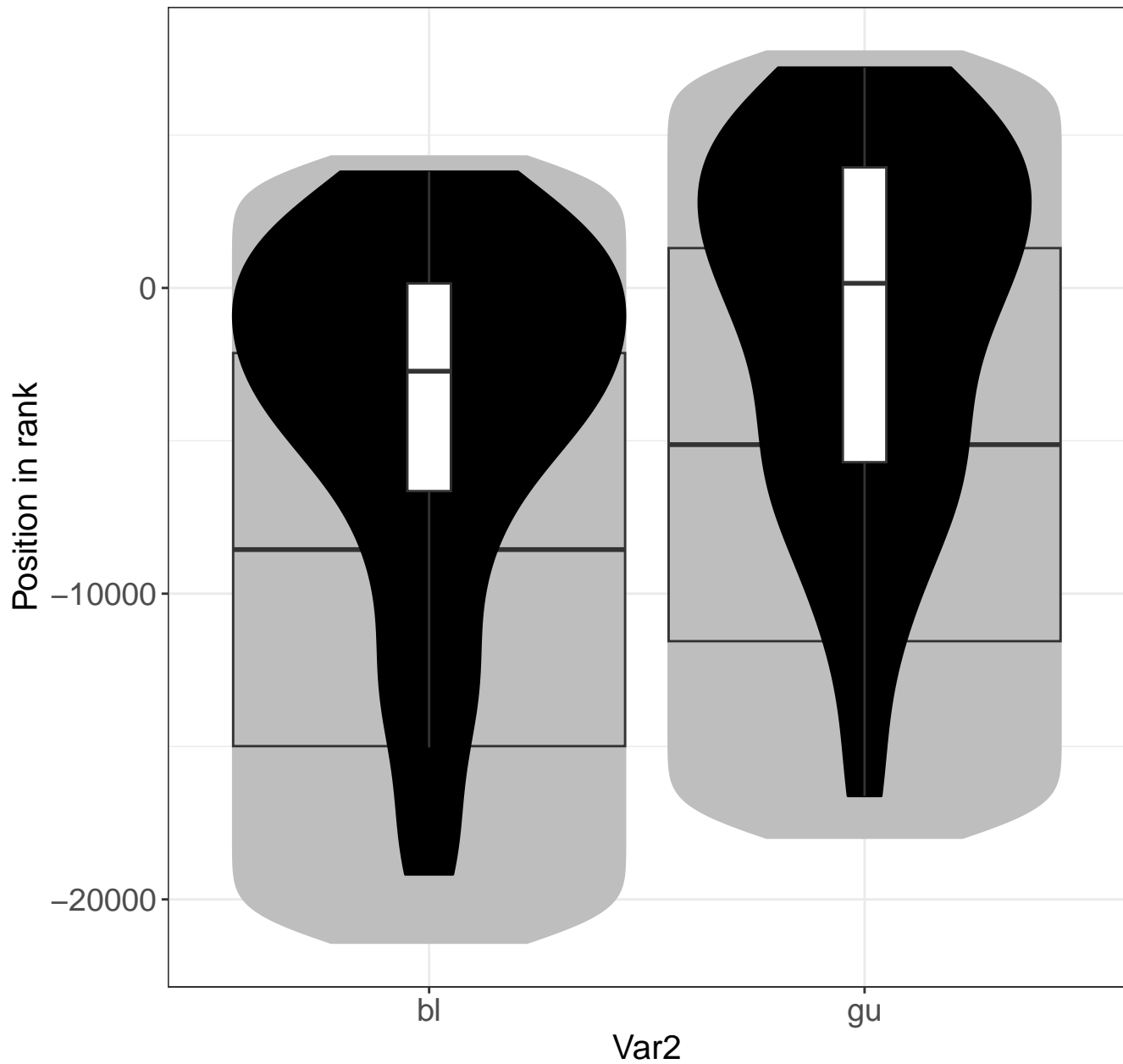
The role of GTSE1 in G2/M progression after G2 checkpo



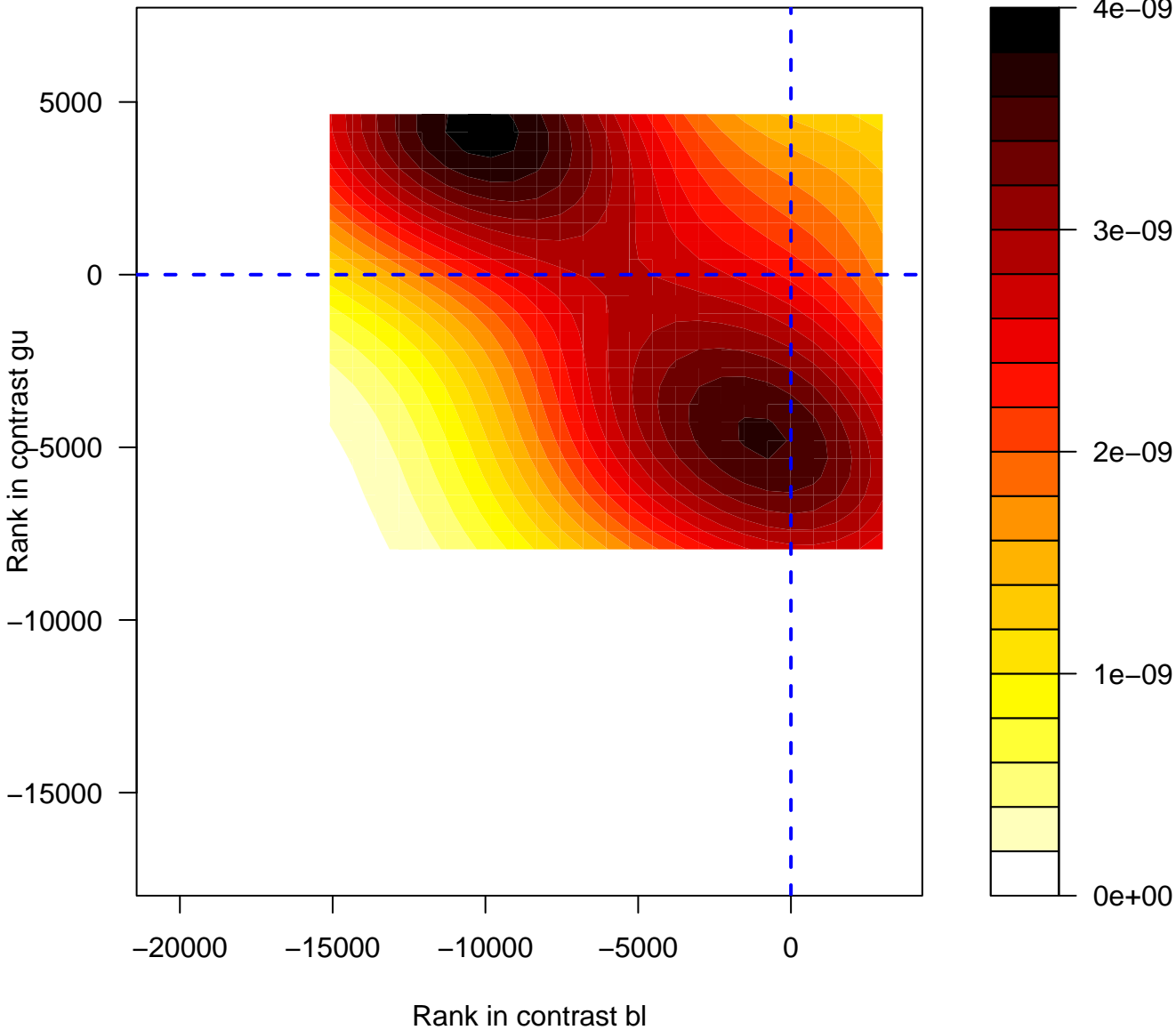
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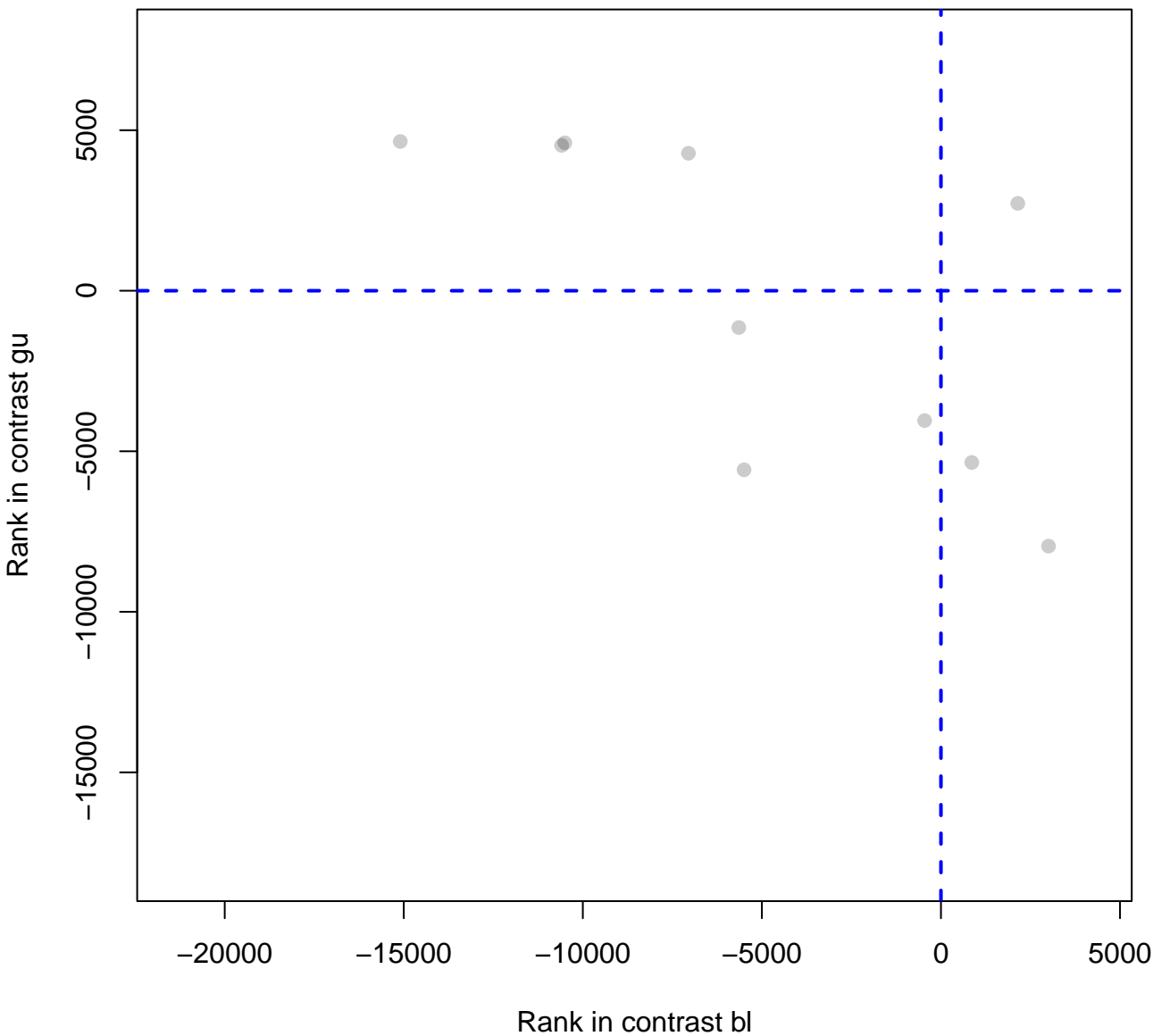
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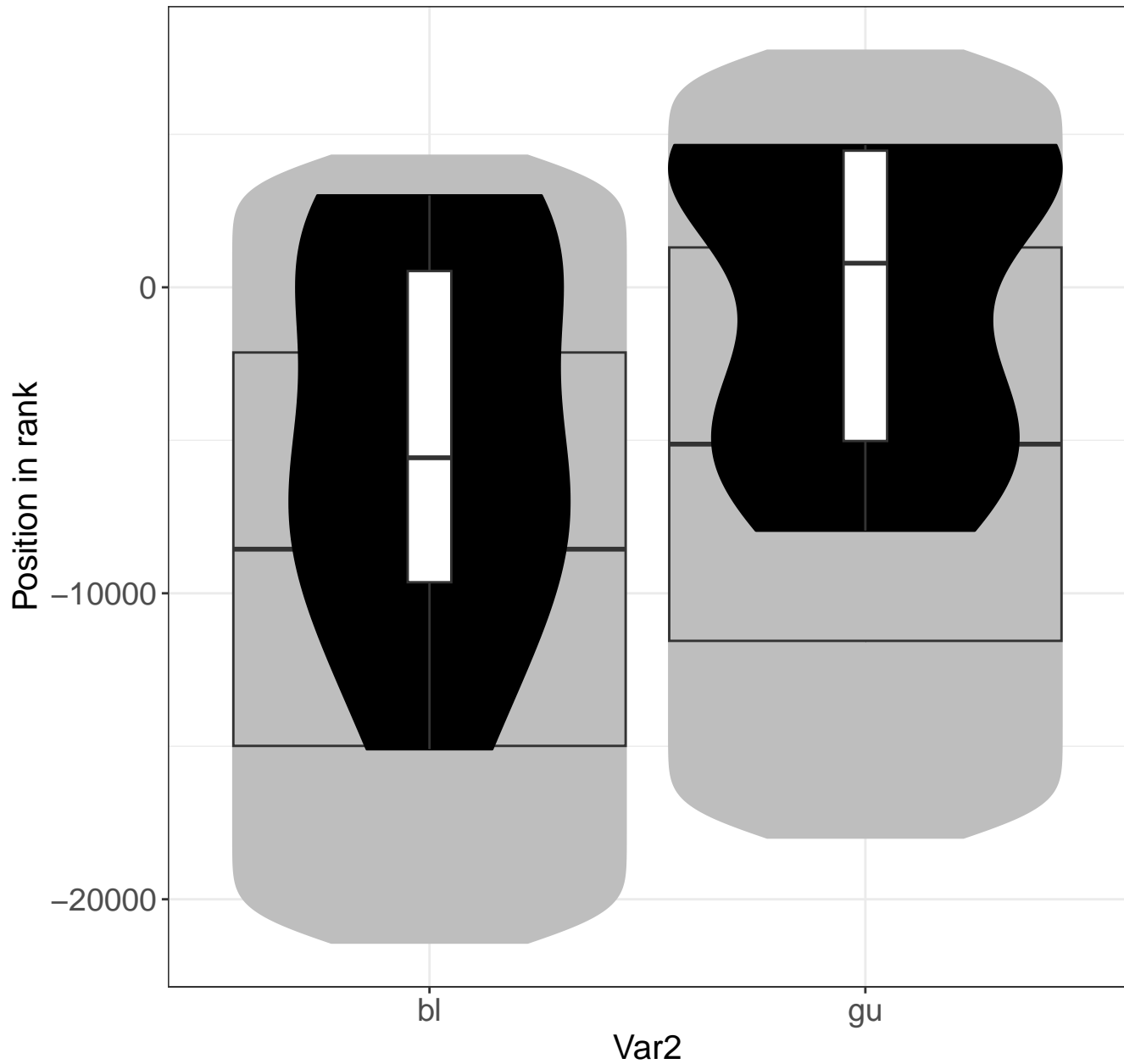
SUMOylation of immune response proteins



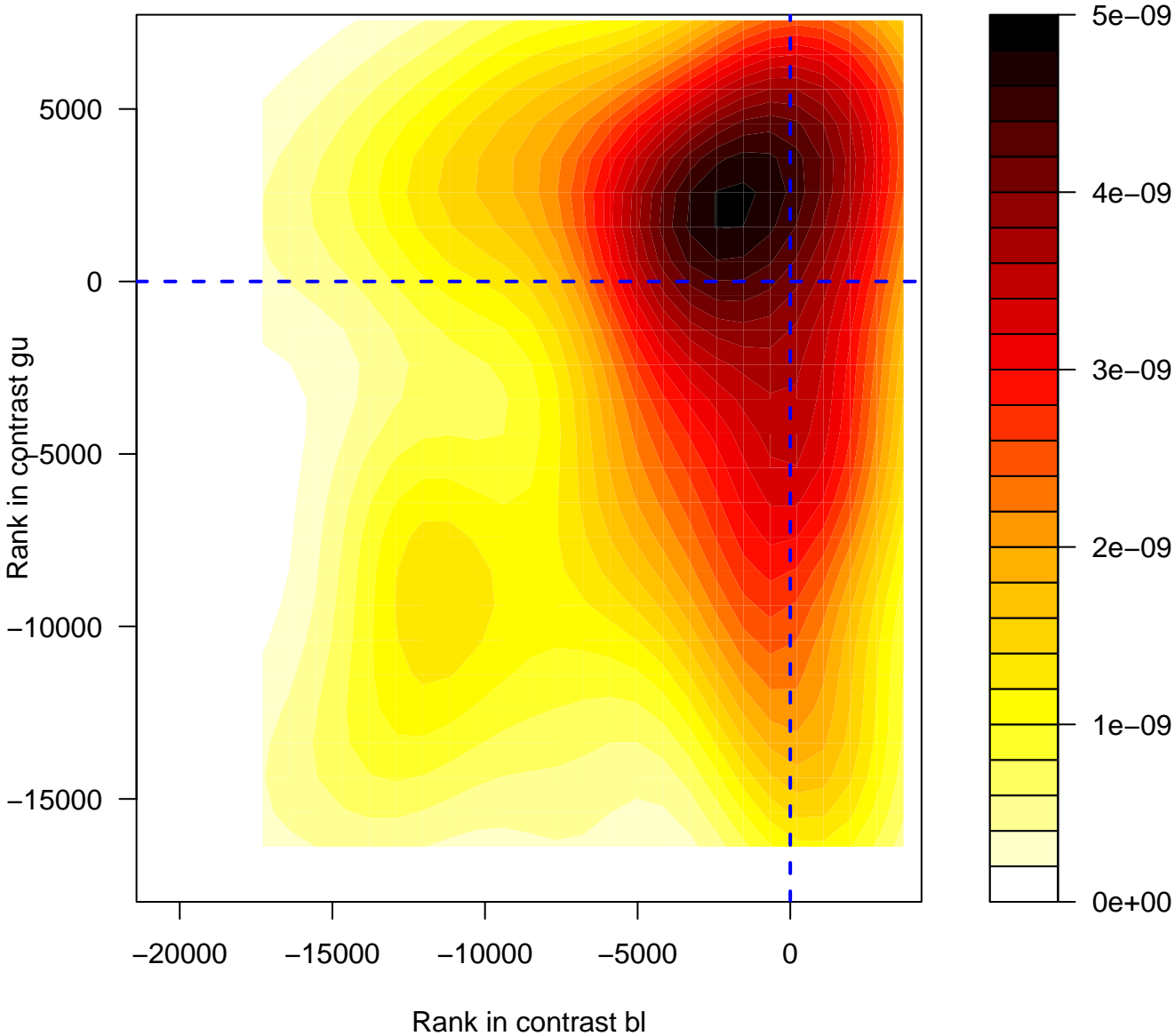
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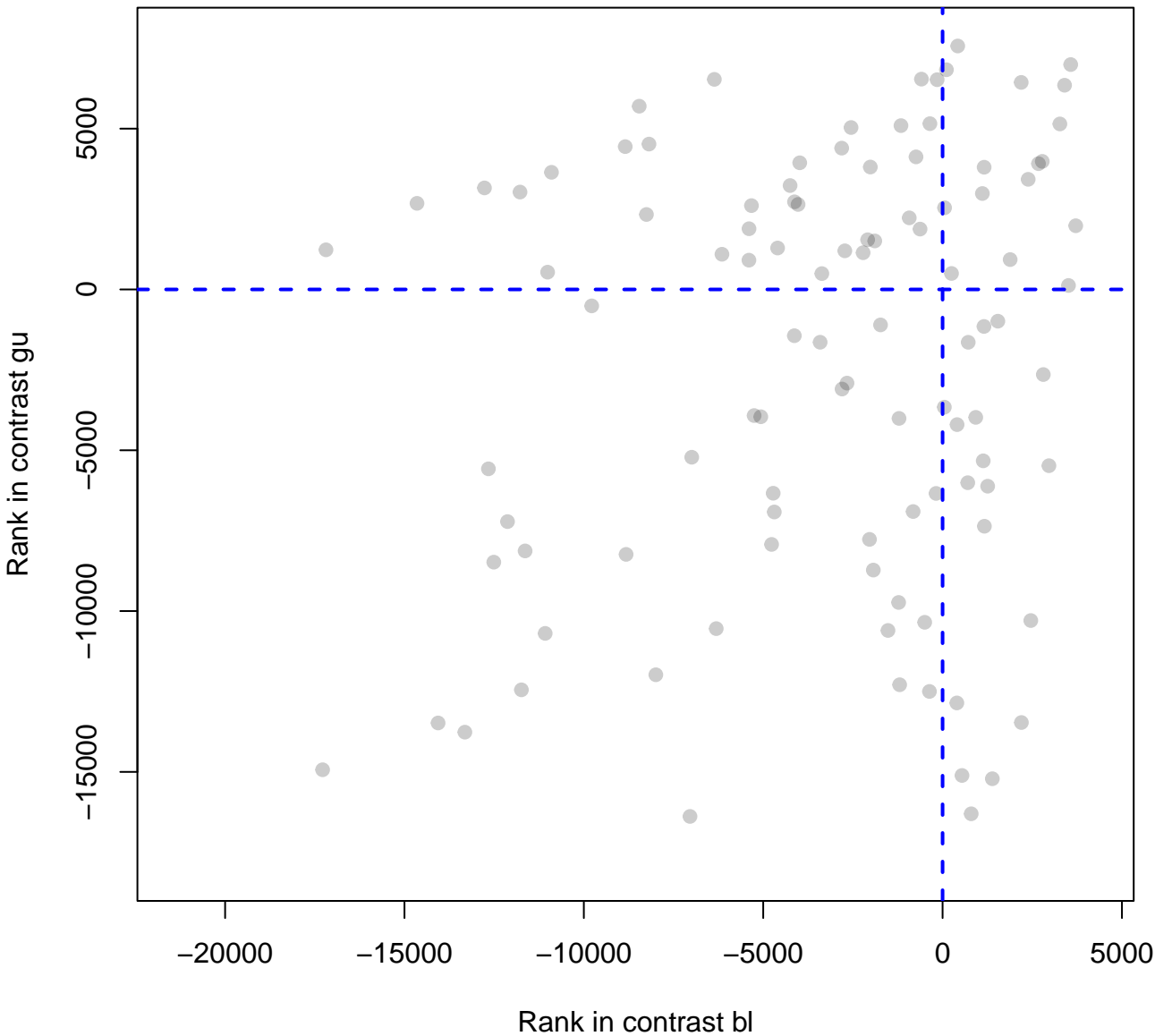
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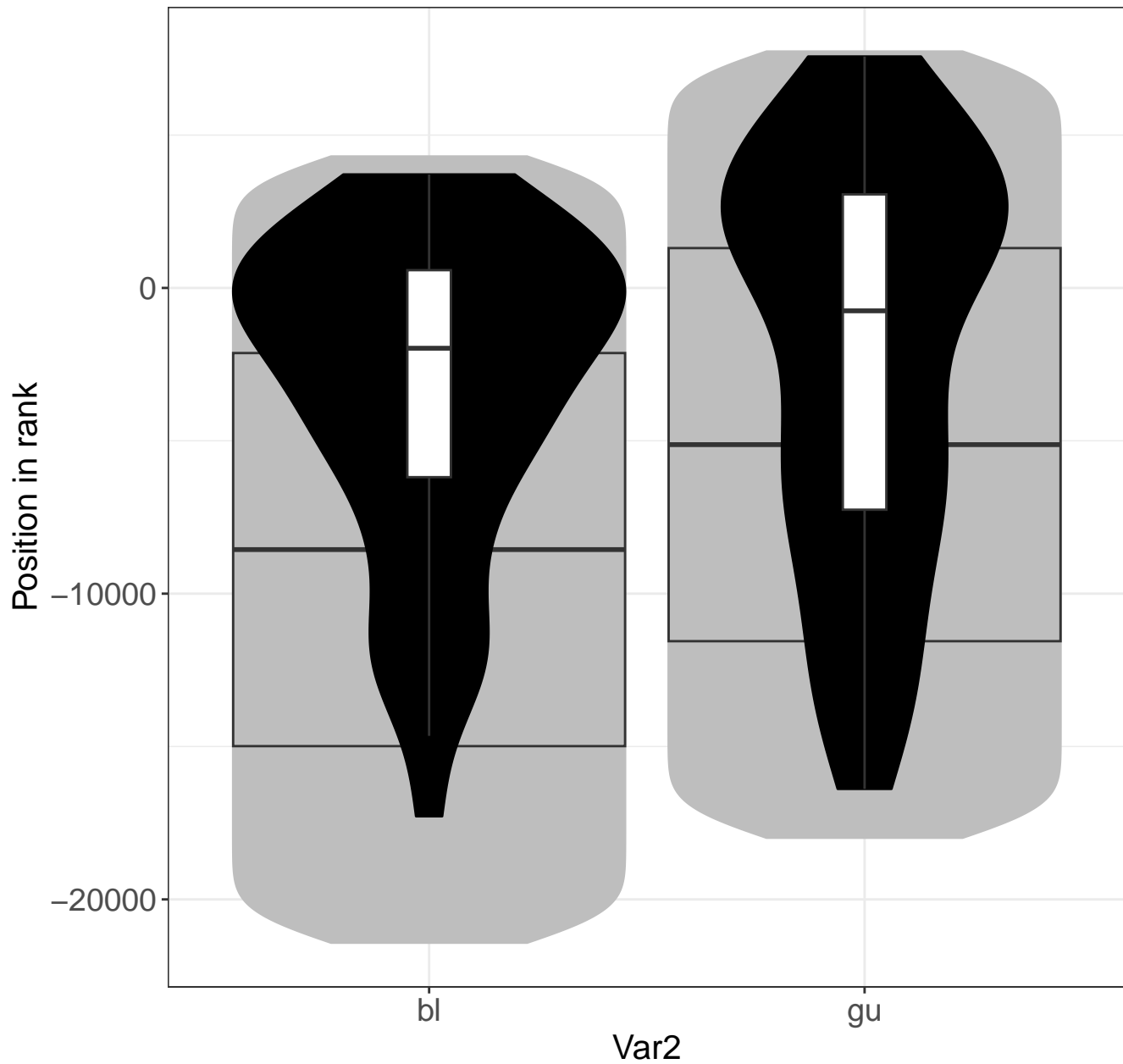
GTP hydrolysis and joining of the 60S ribosomal subun



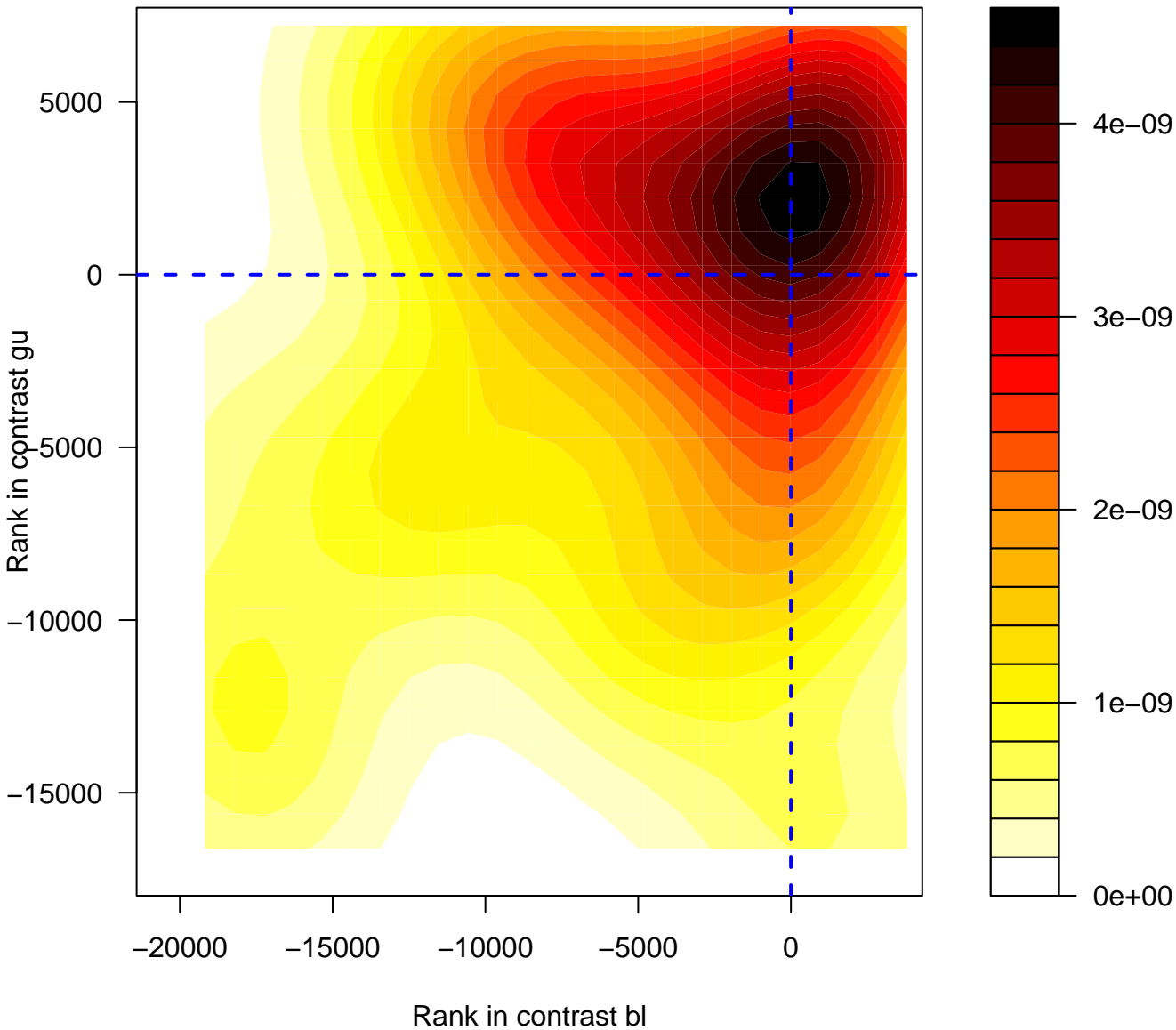
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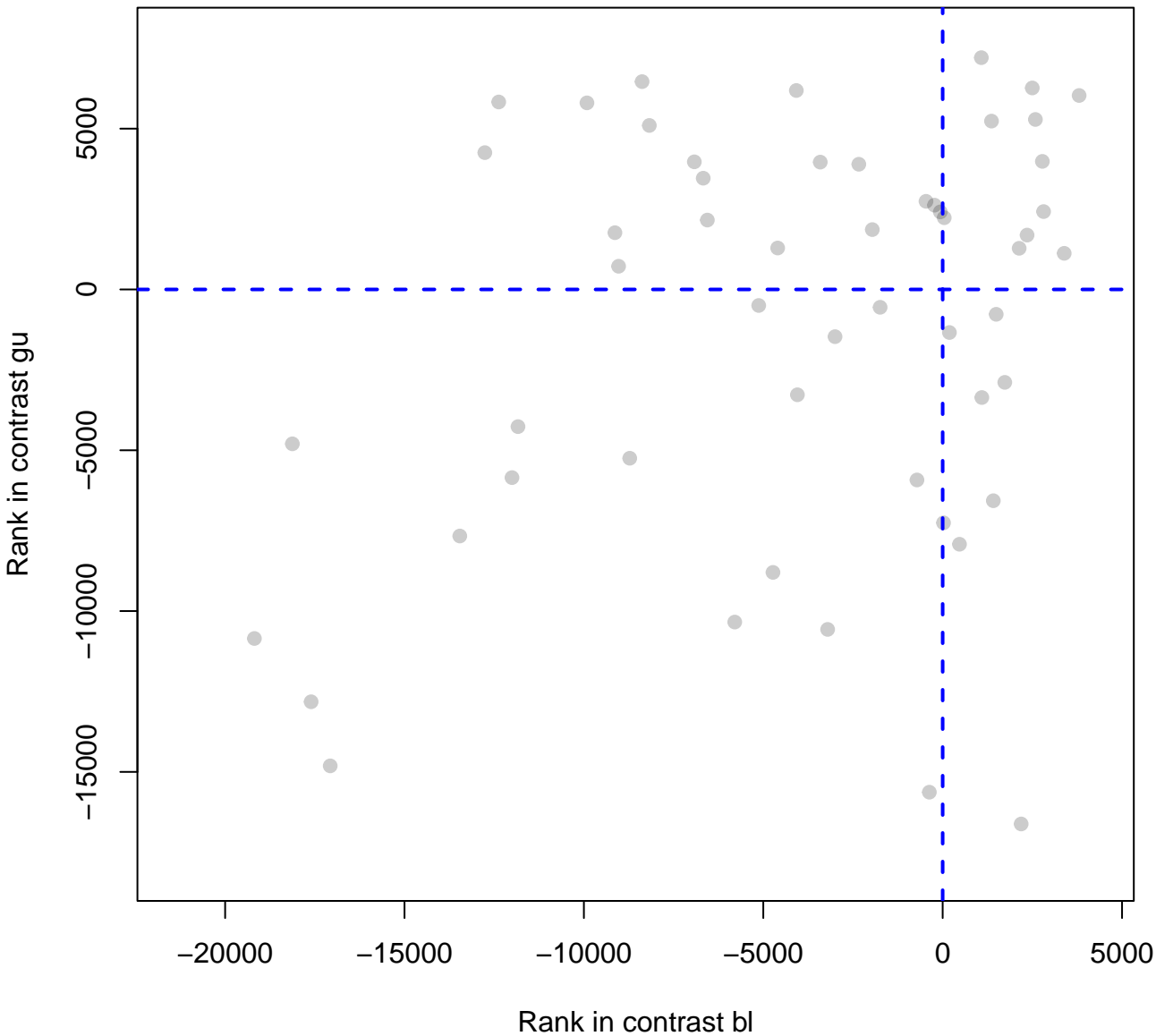
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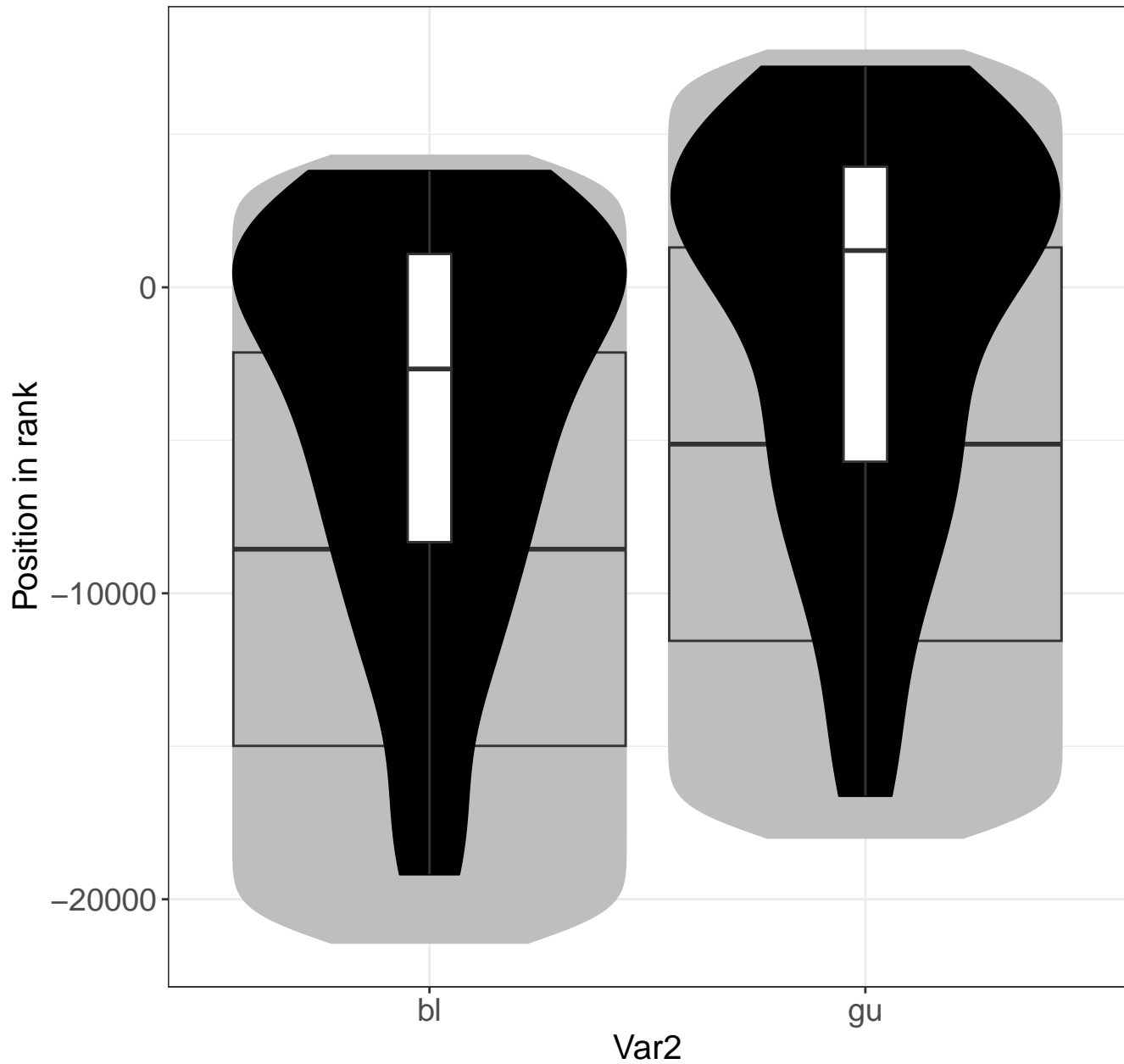
Hh mutants are degraded by ERAD



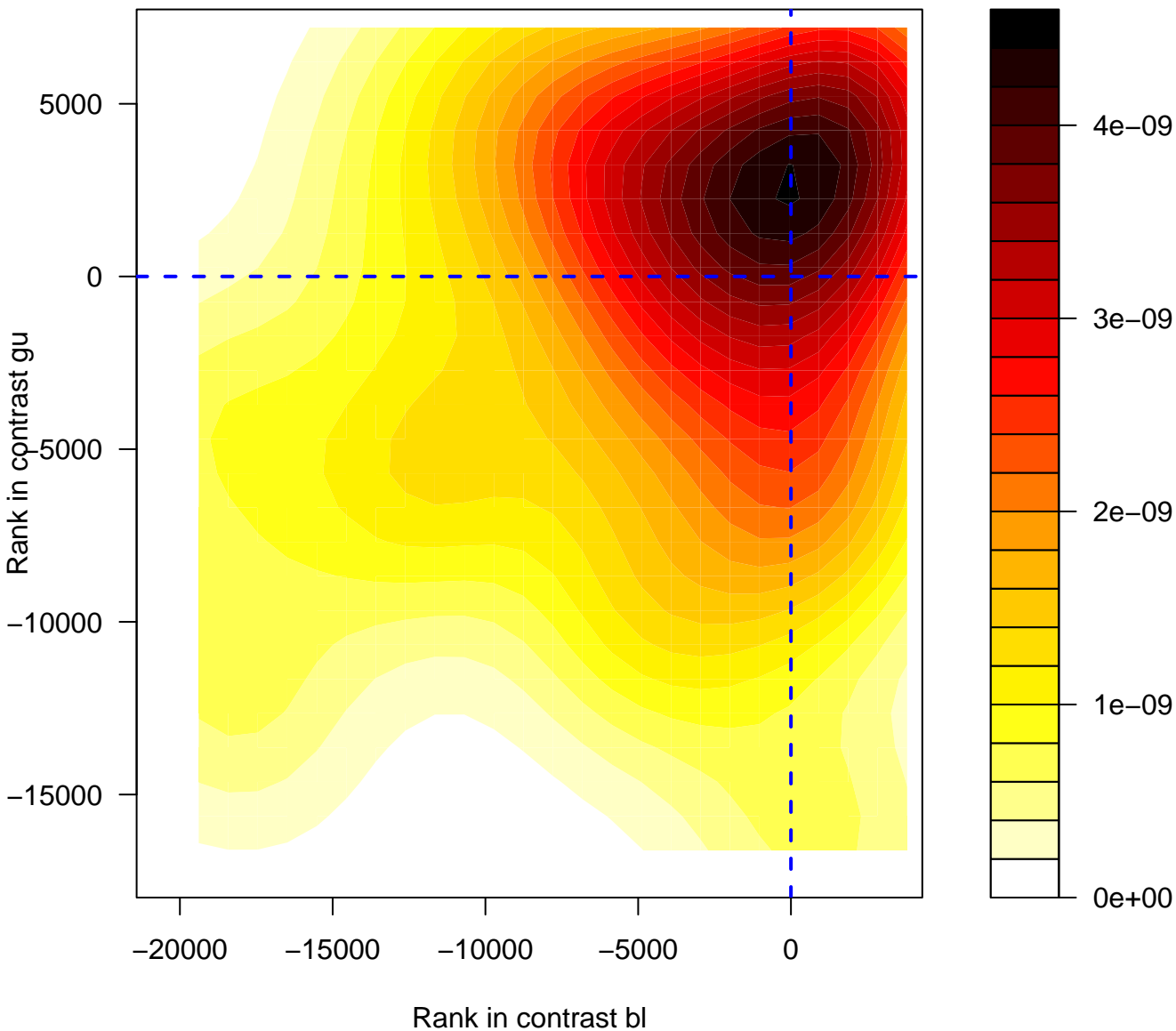
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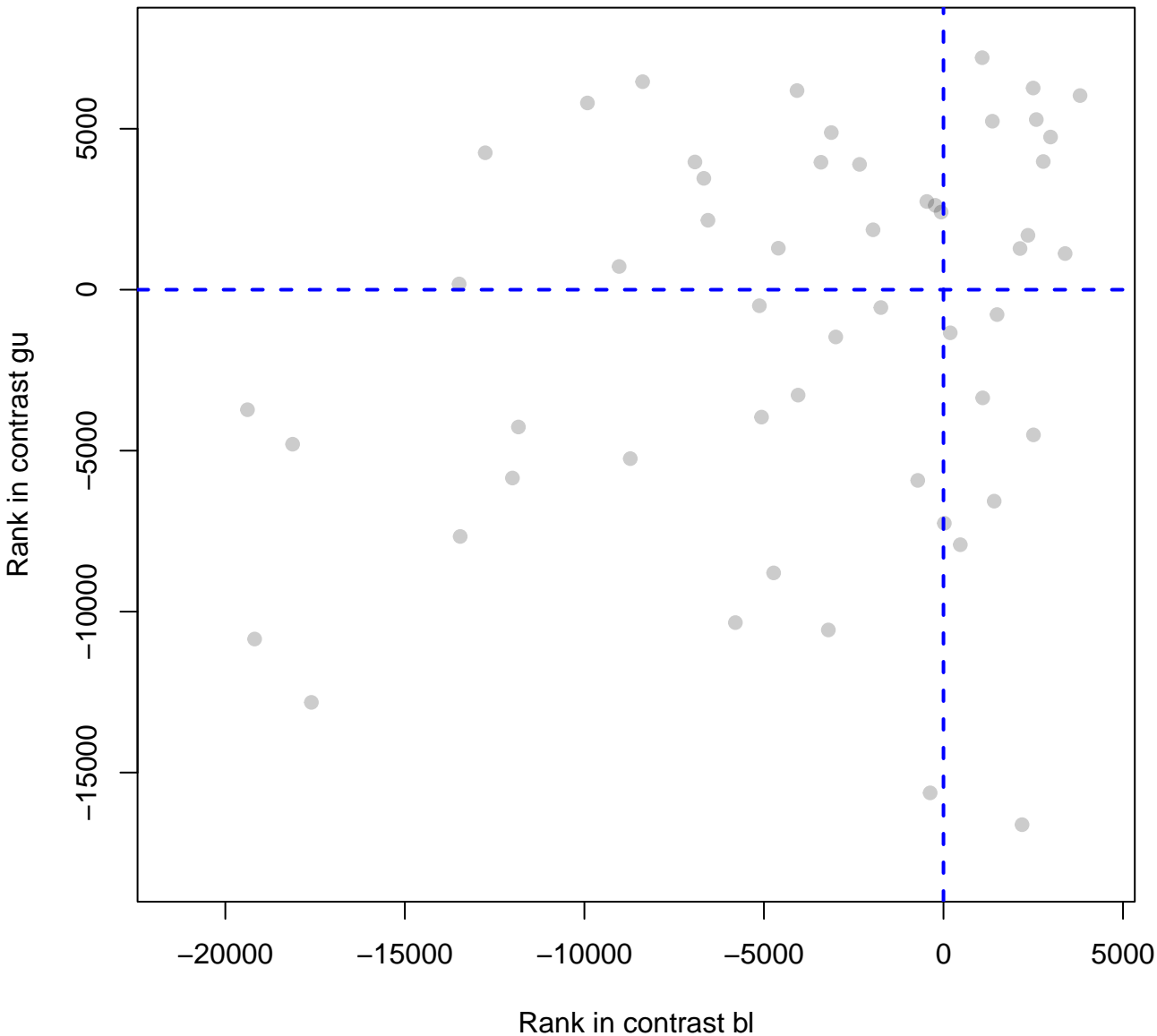
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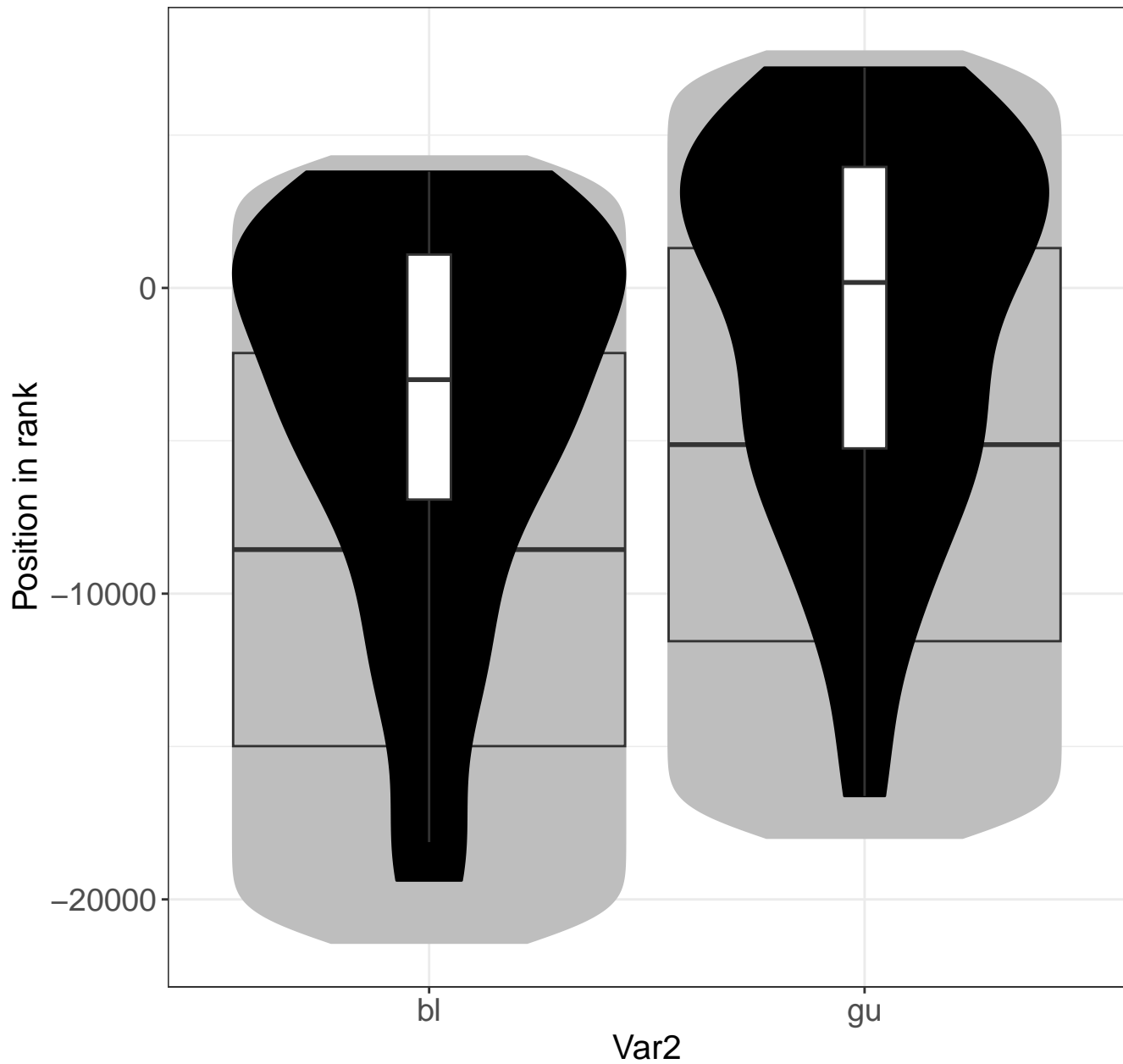
AUF1 (hnRNP D0) binds and destabilizes mRNA



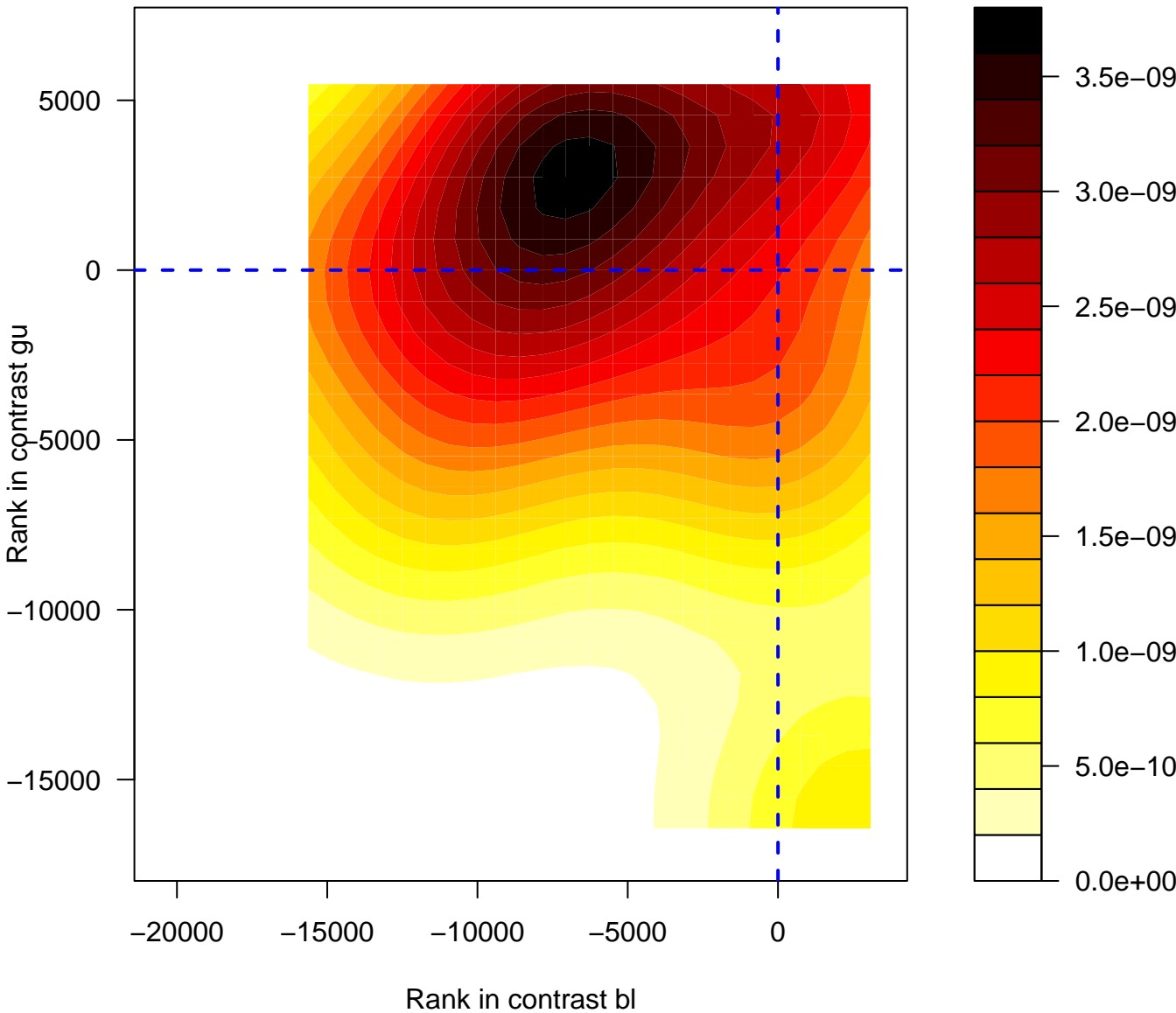
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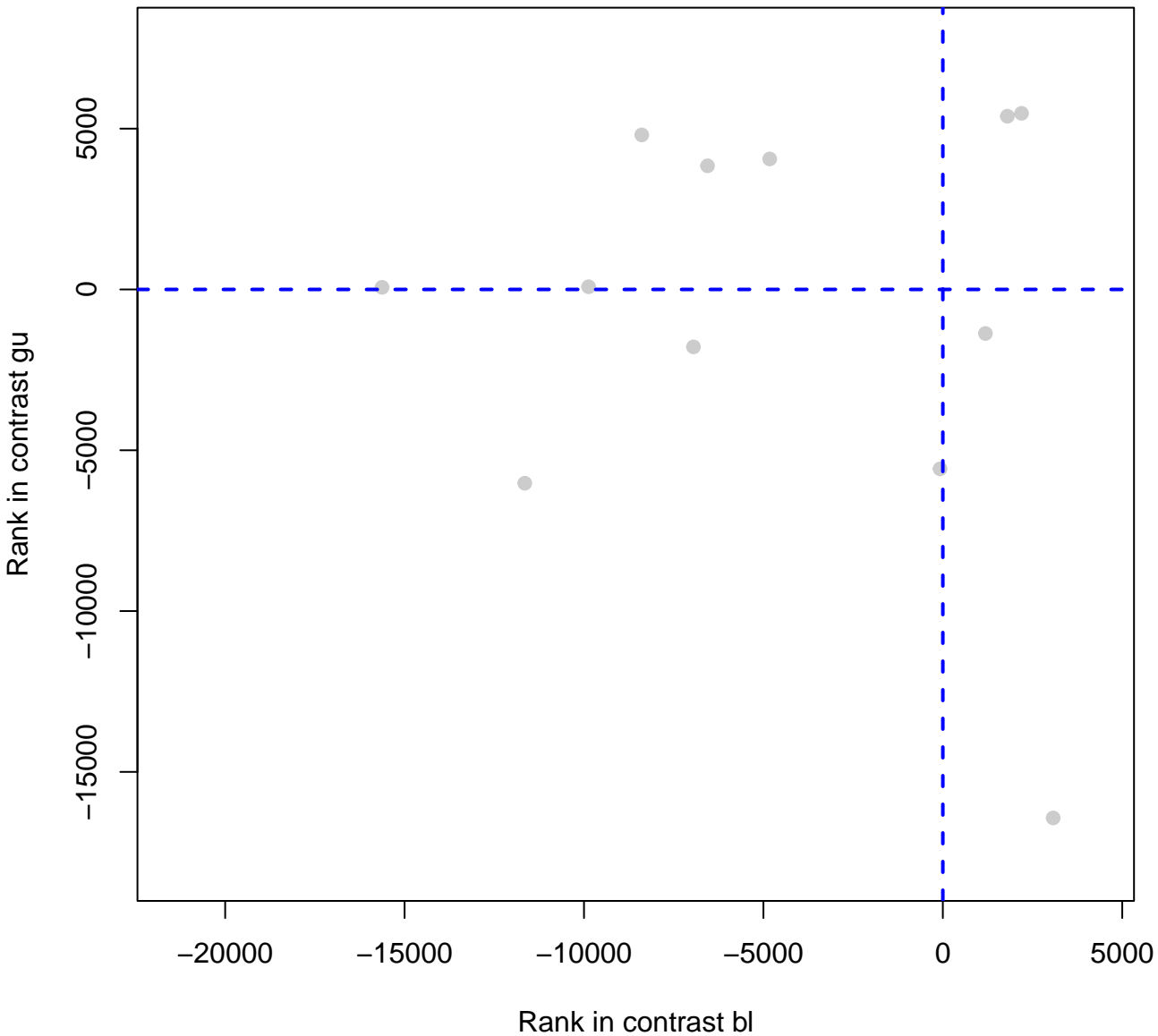
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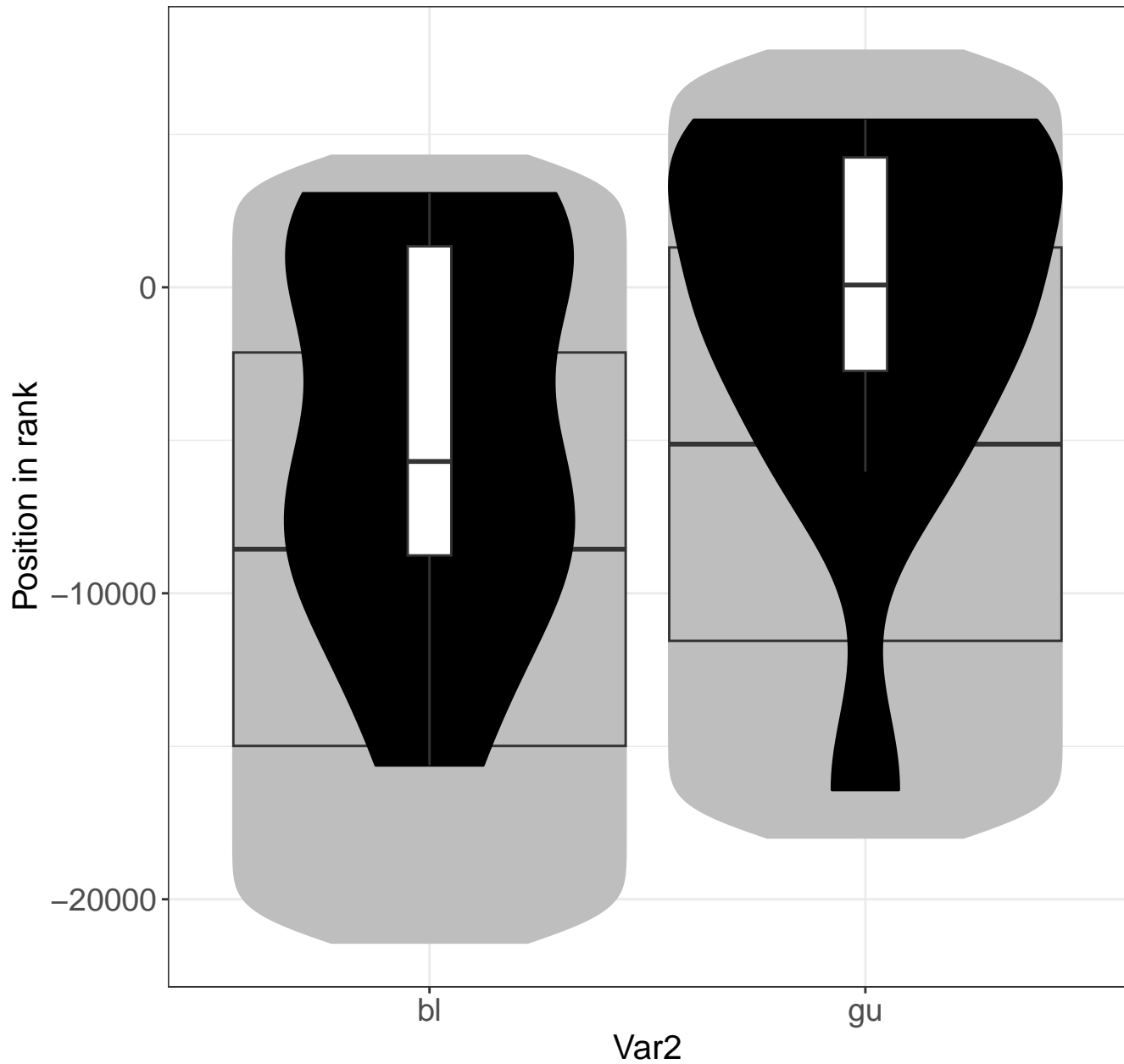
Mitochondrial iron-sulfur cluster biogenesis



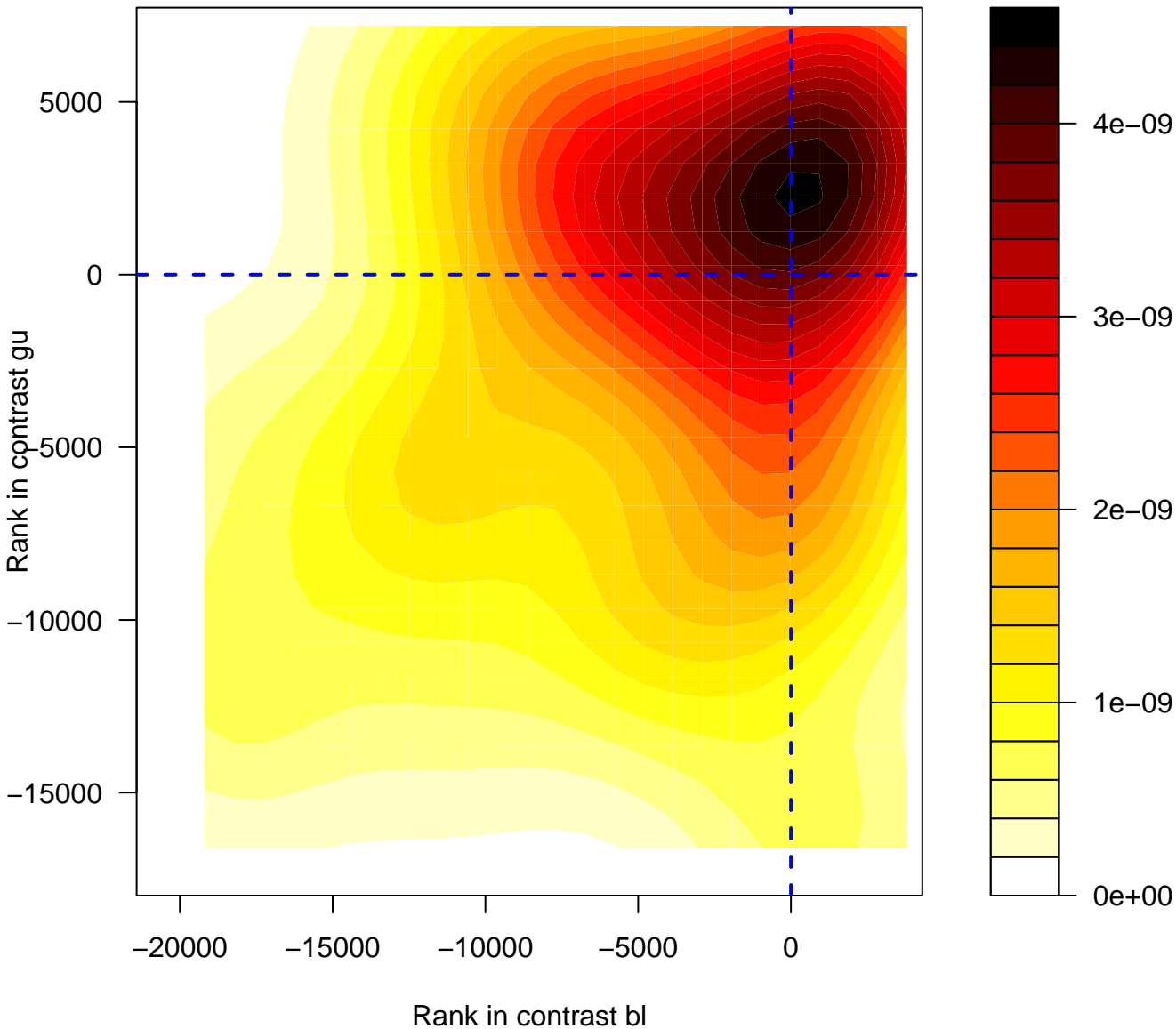
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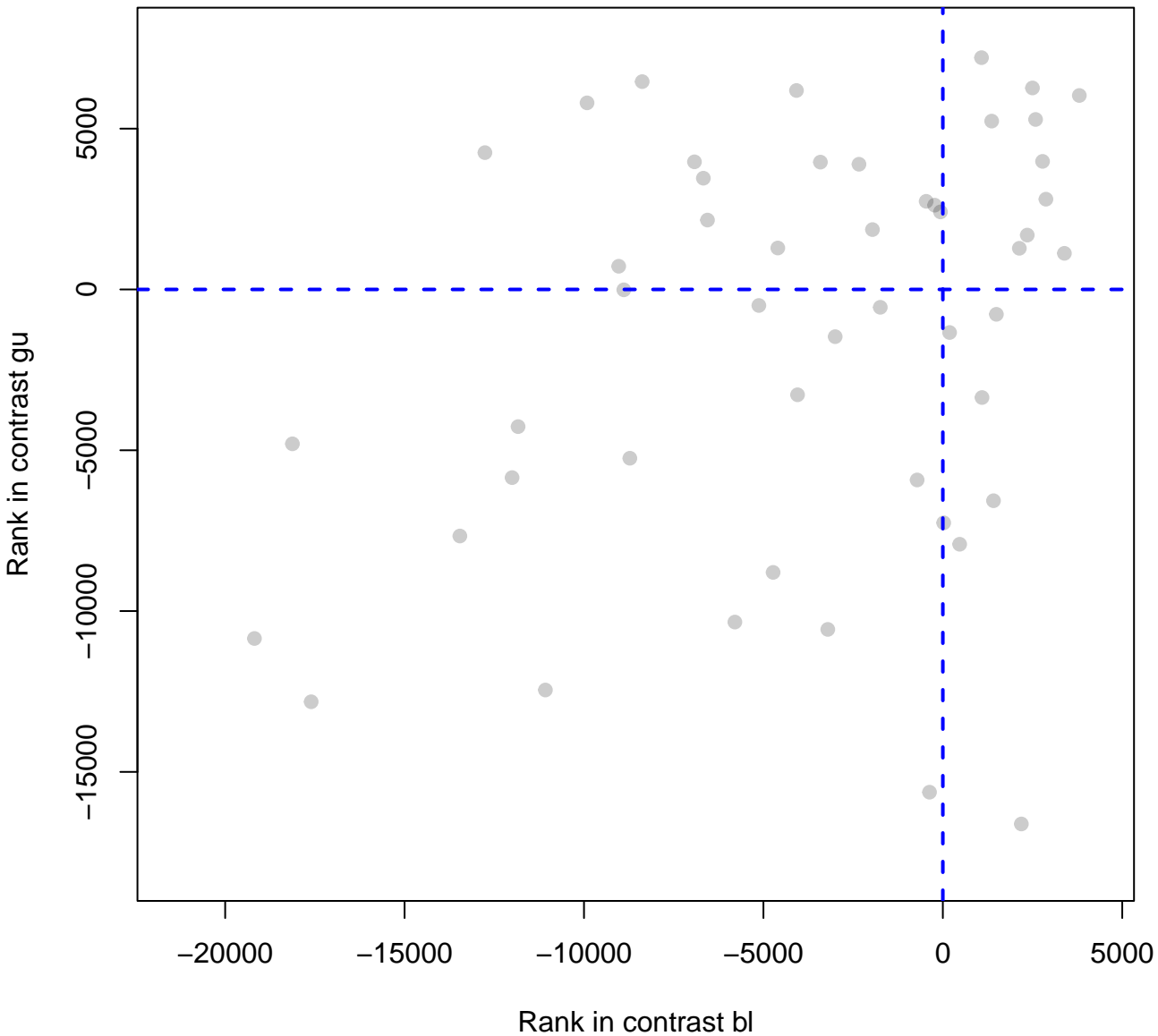
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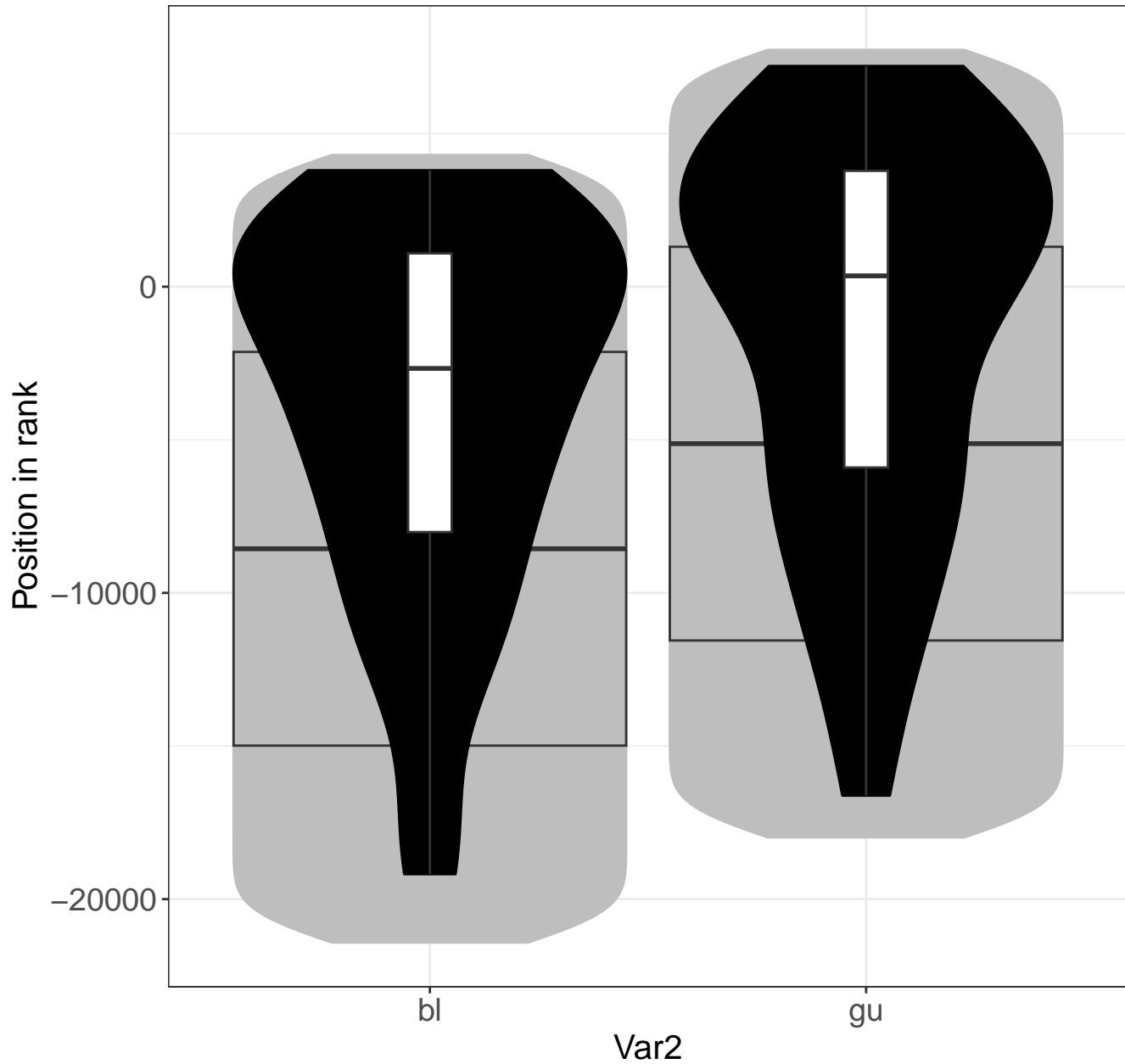
Ubiquitin-dependent degradation of Cyclin D



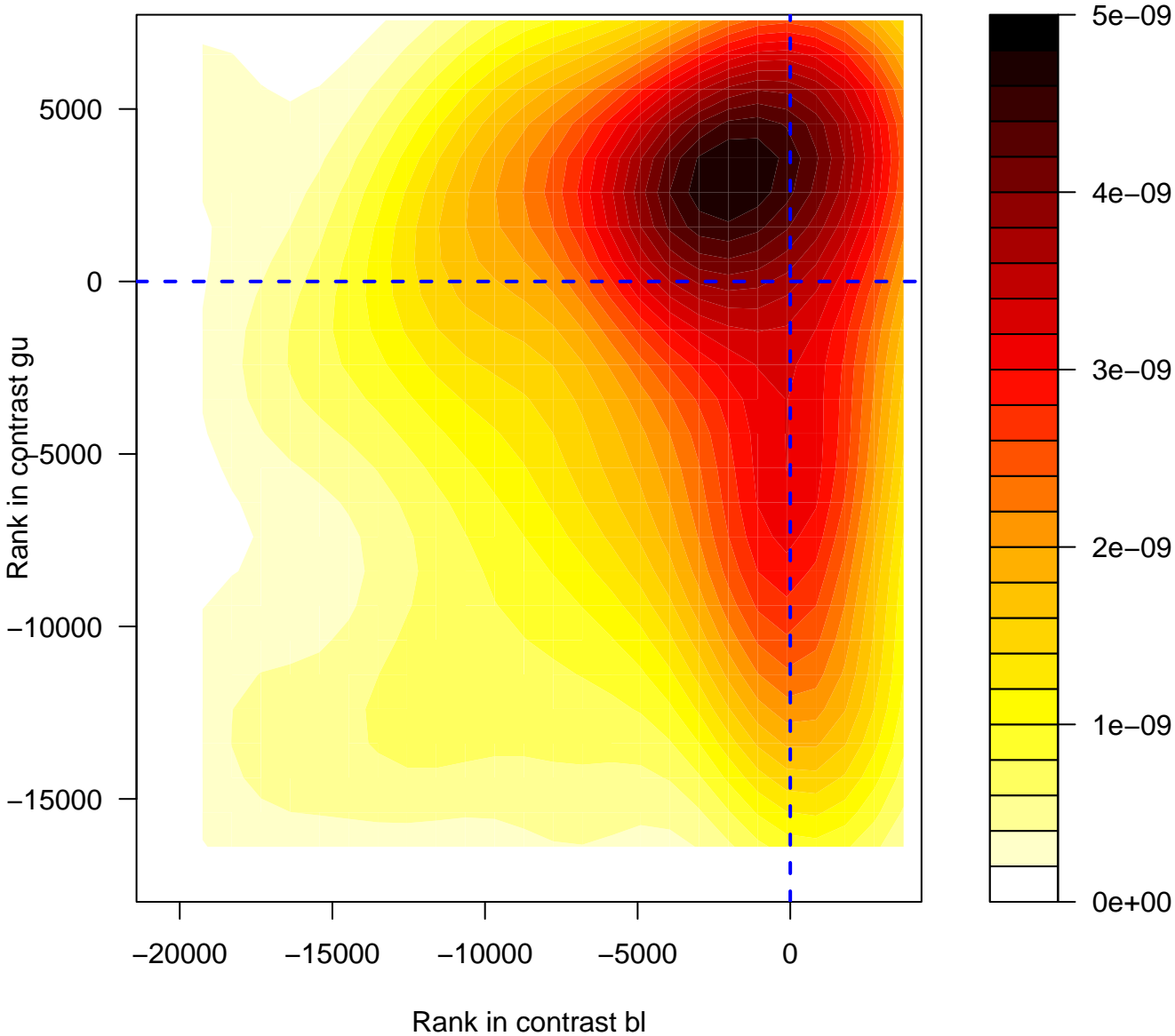
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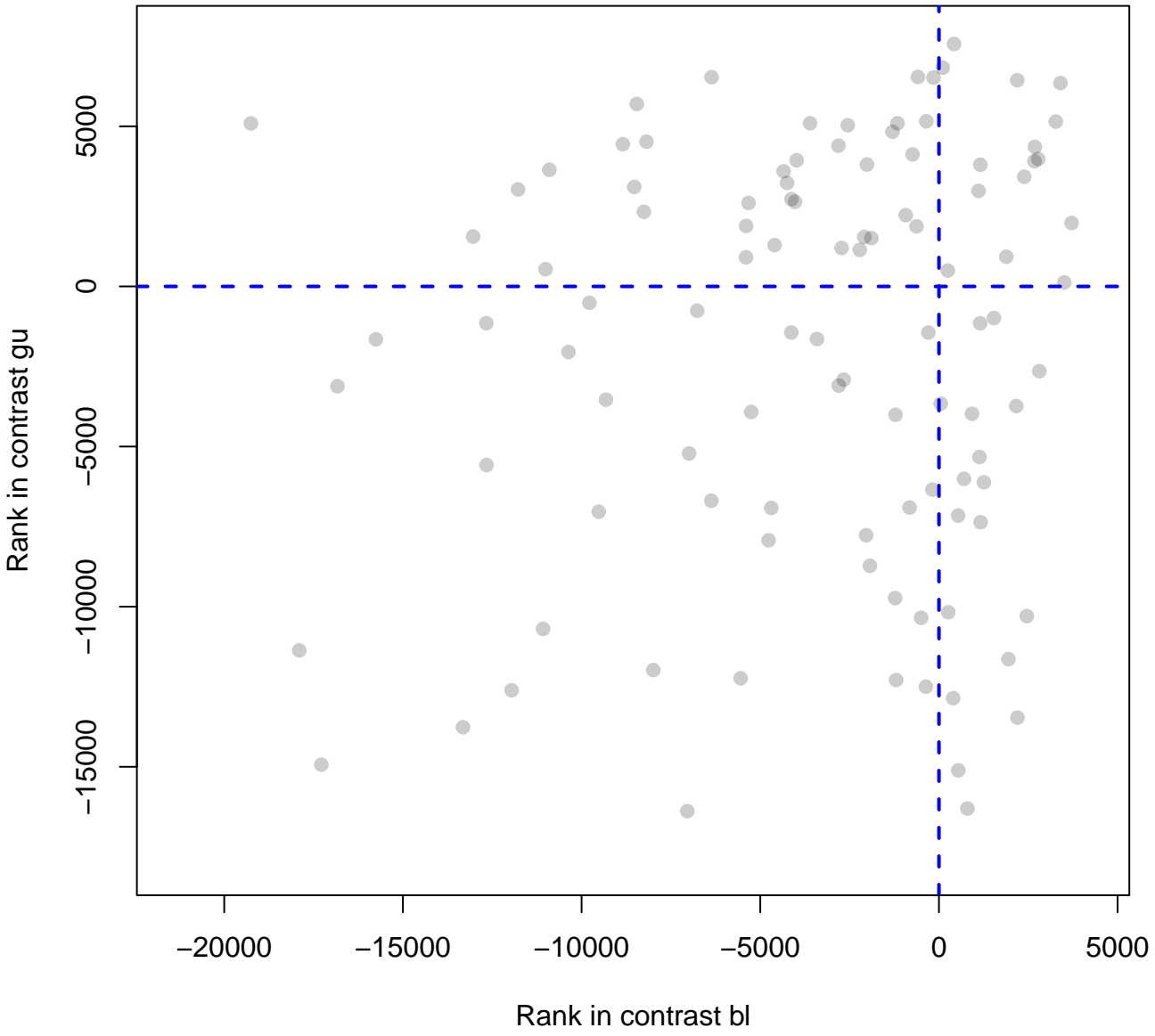
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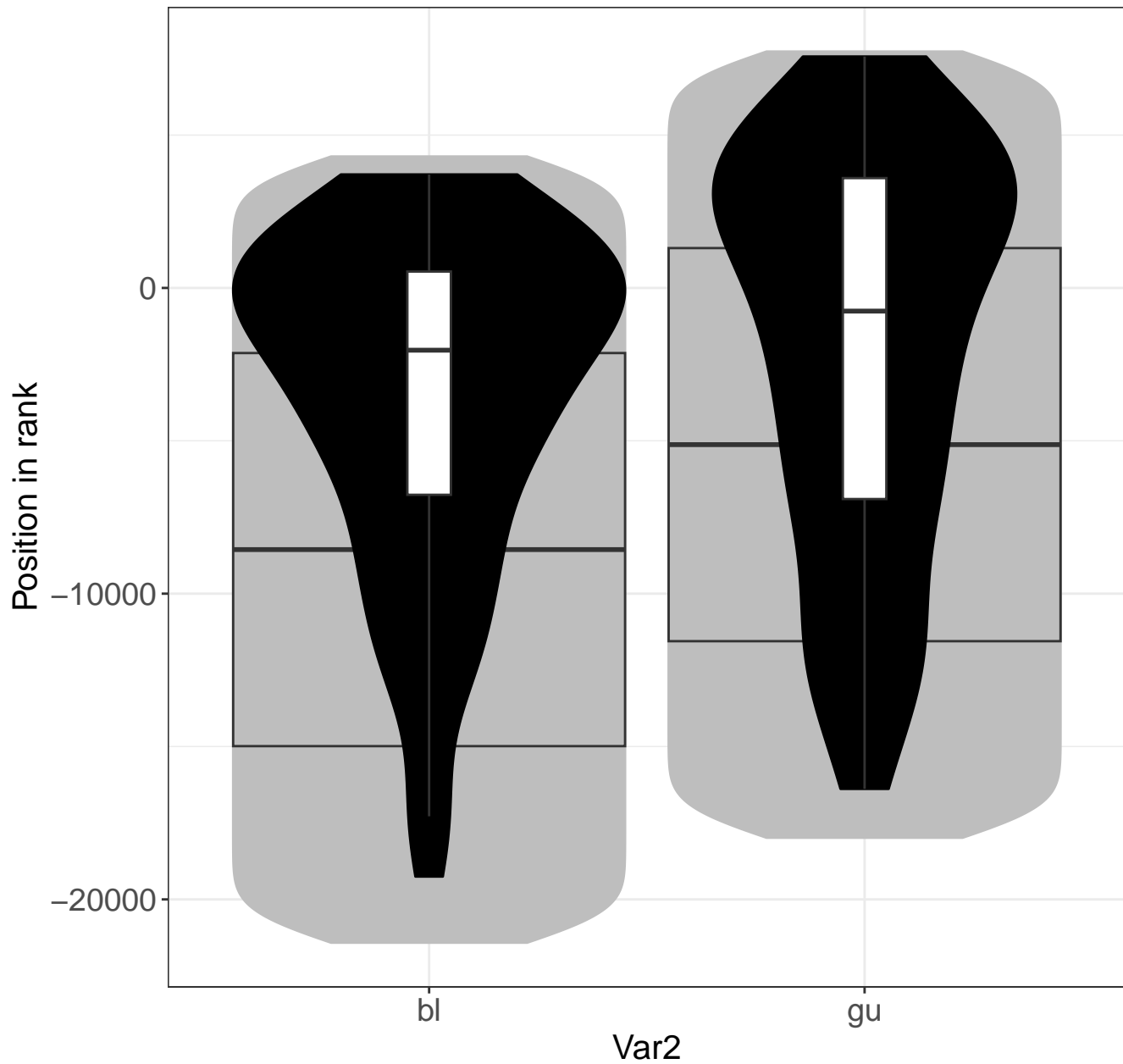
SRP-dependent cotranslational protein targeting to memb



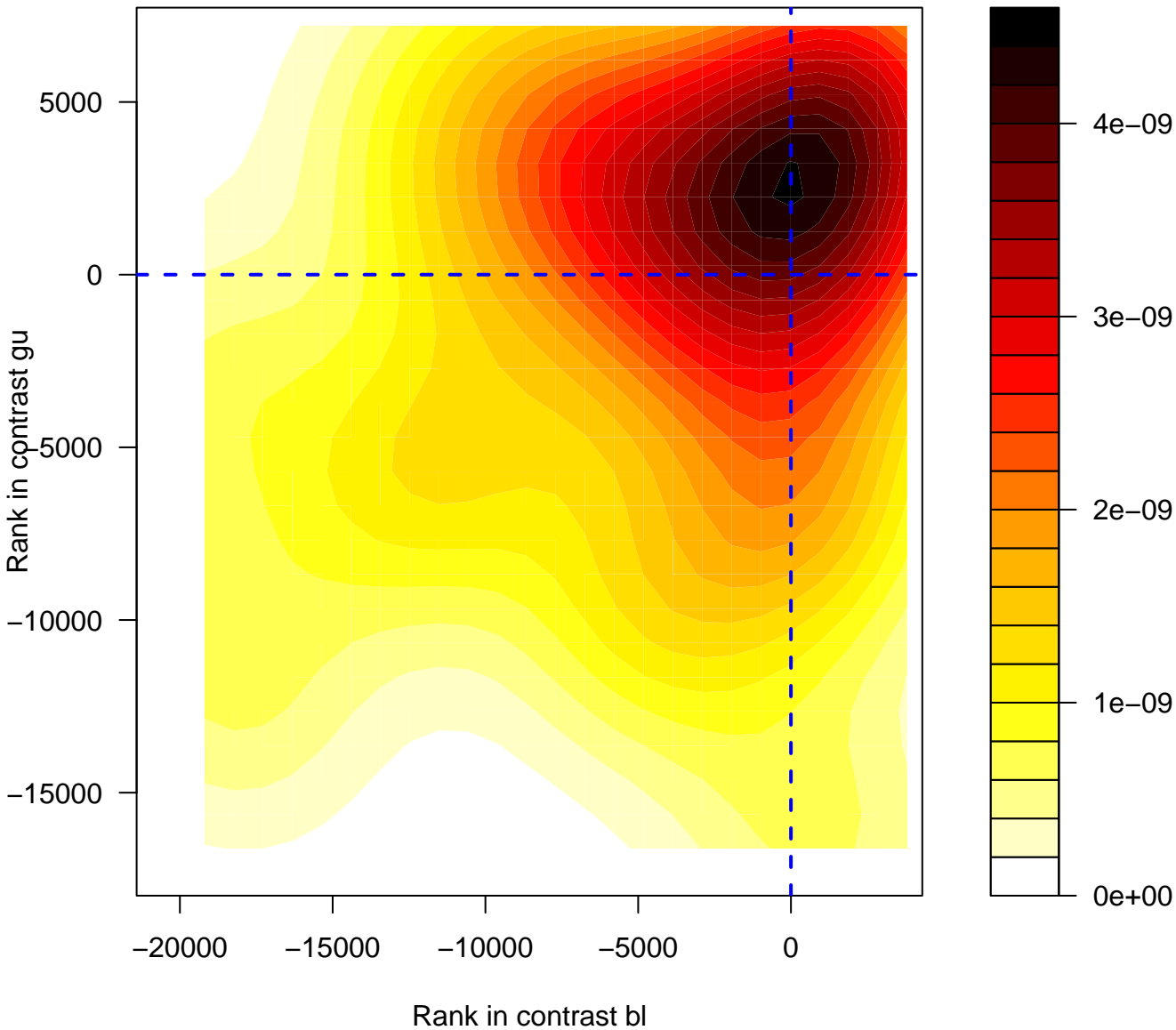
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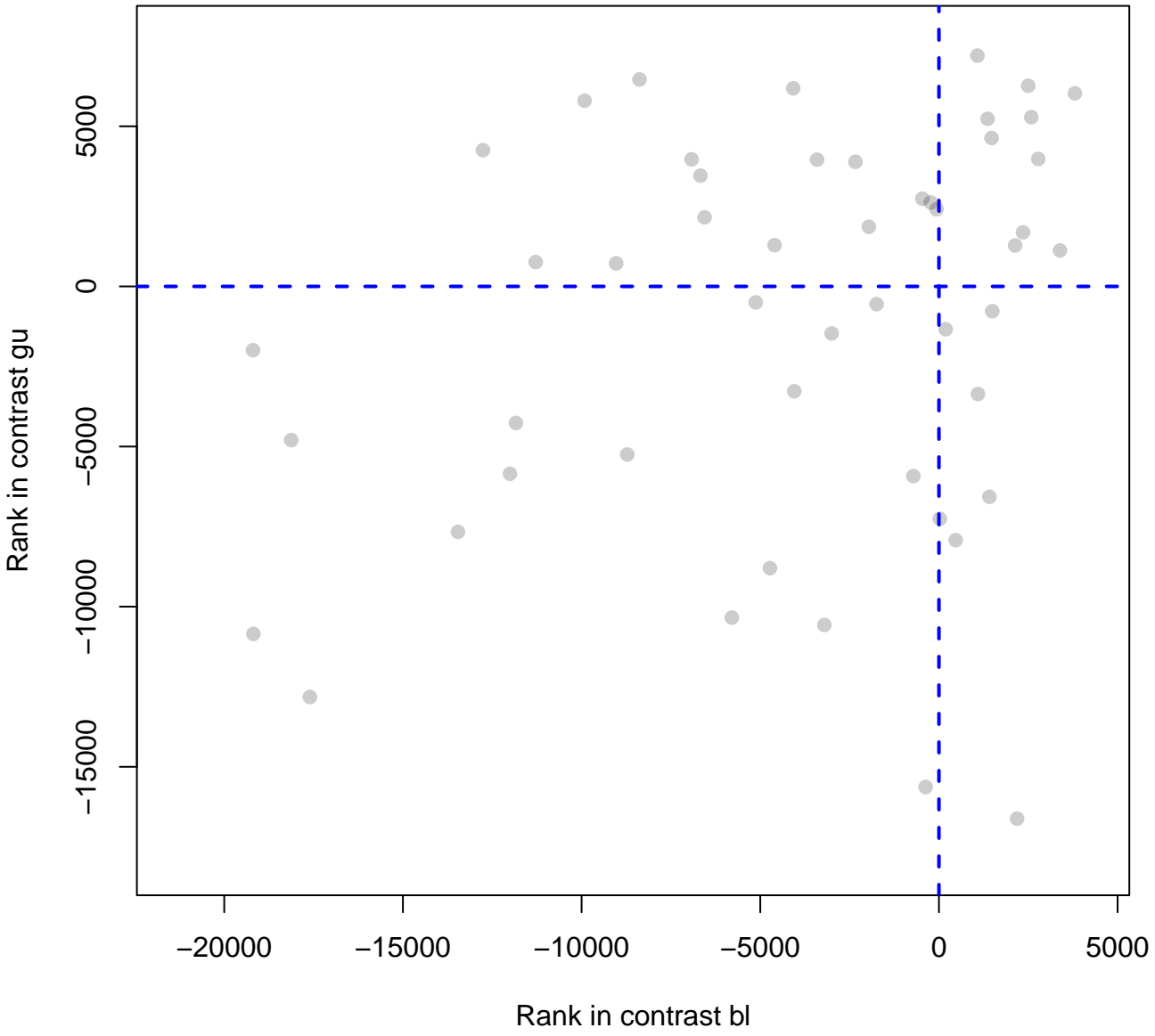
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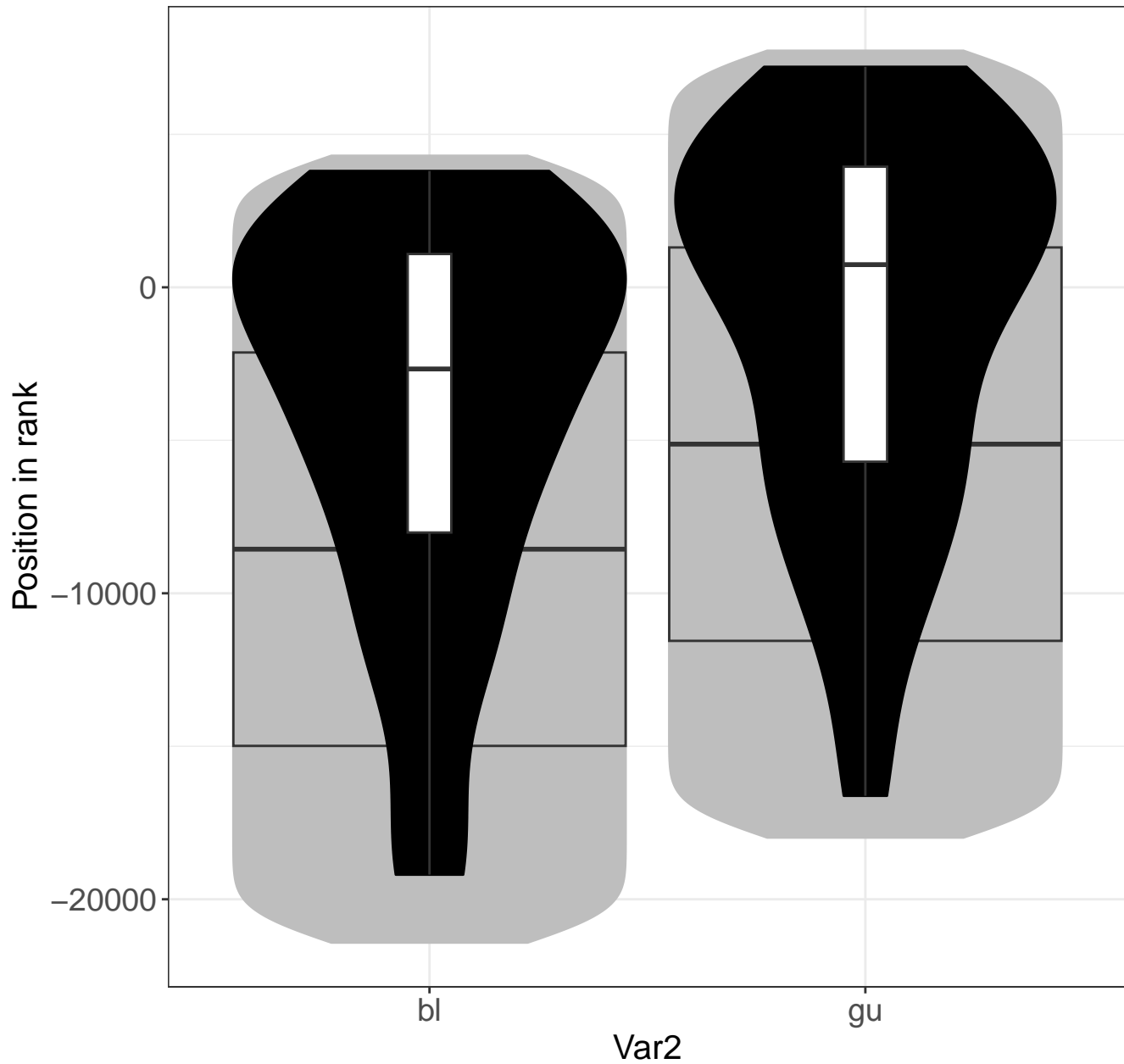
Vpu mediated degradation of CD4



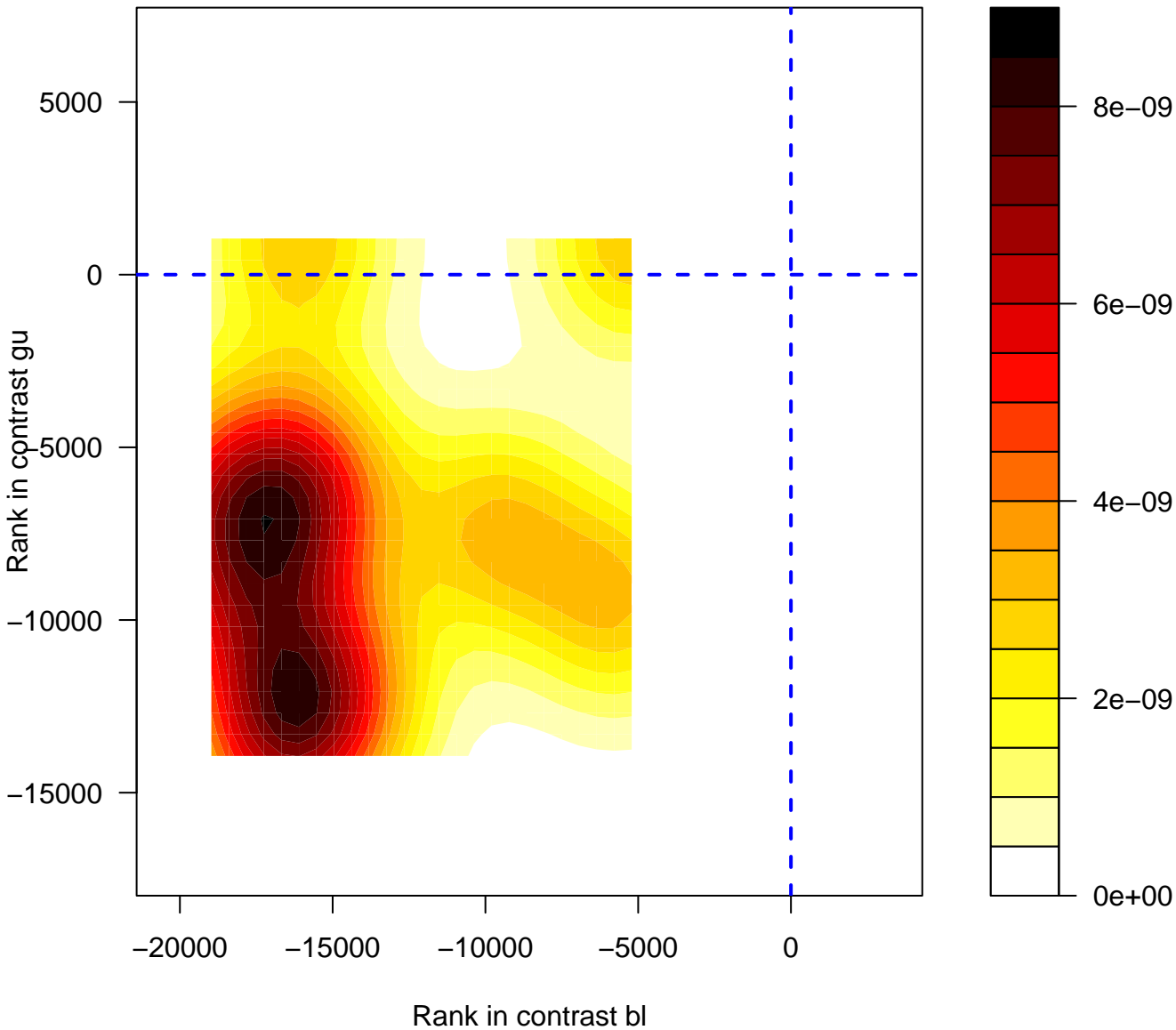
Vpu mediated degradation of CD4



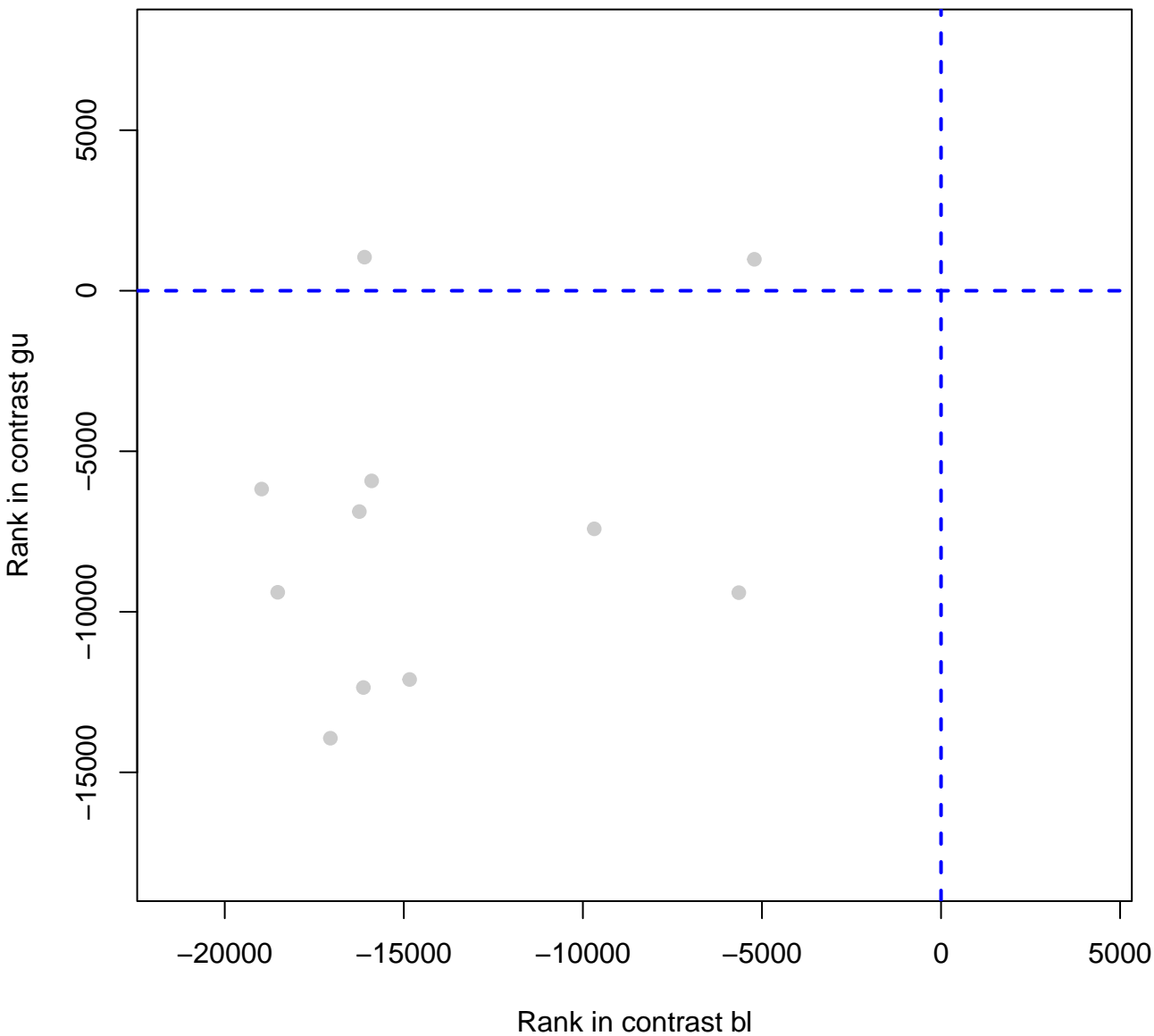
Vpu mediated degradation of CD4



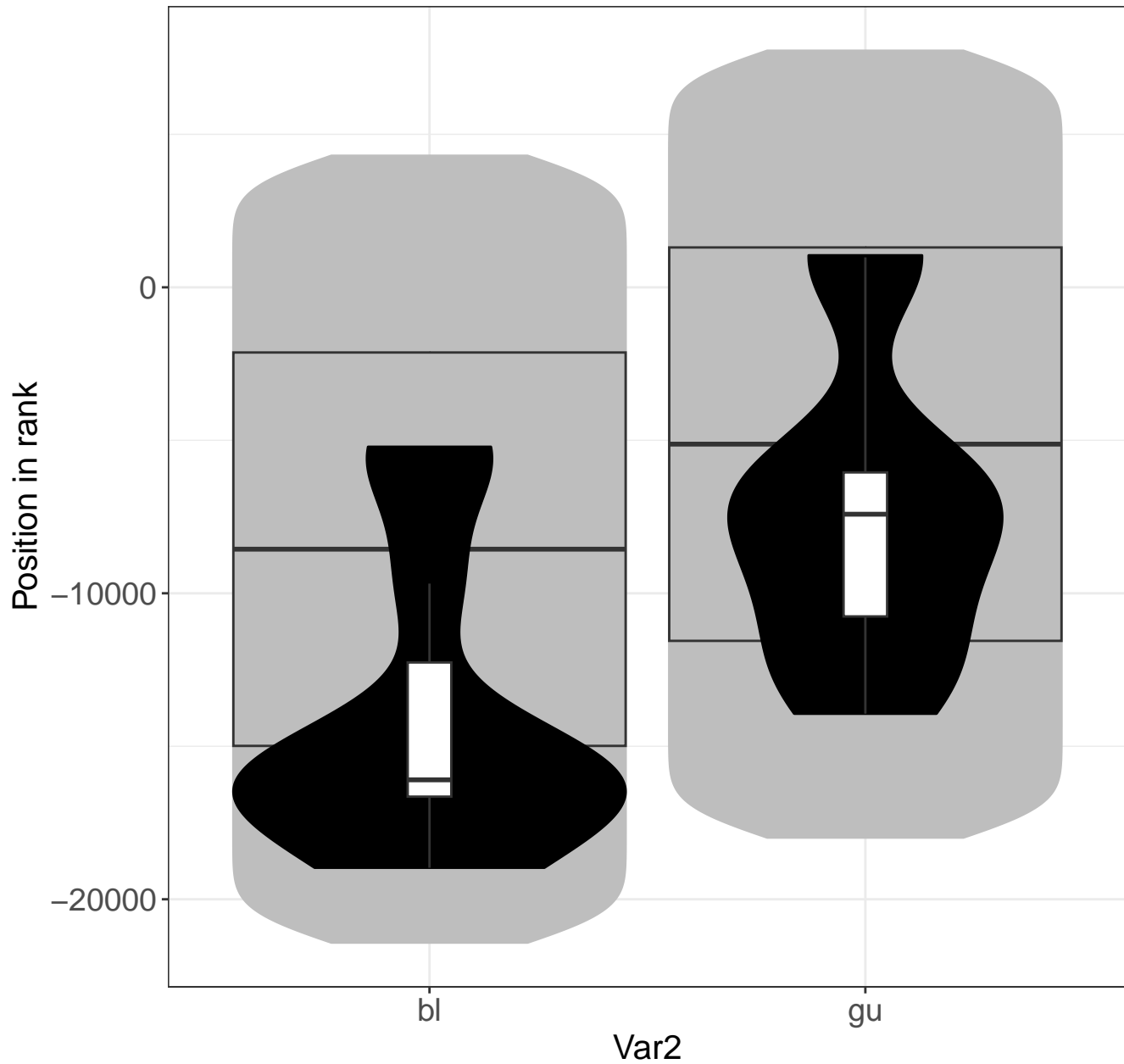
Activation of RAC1



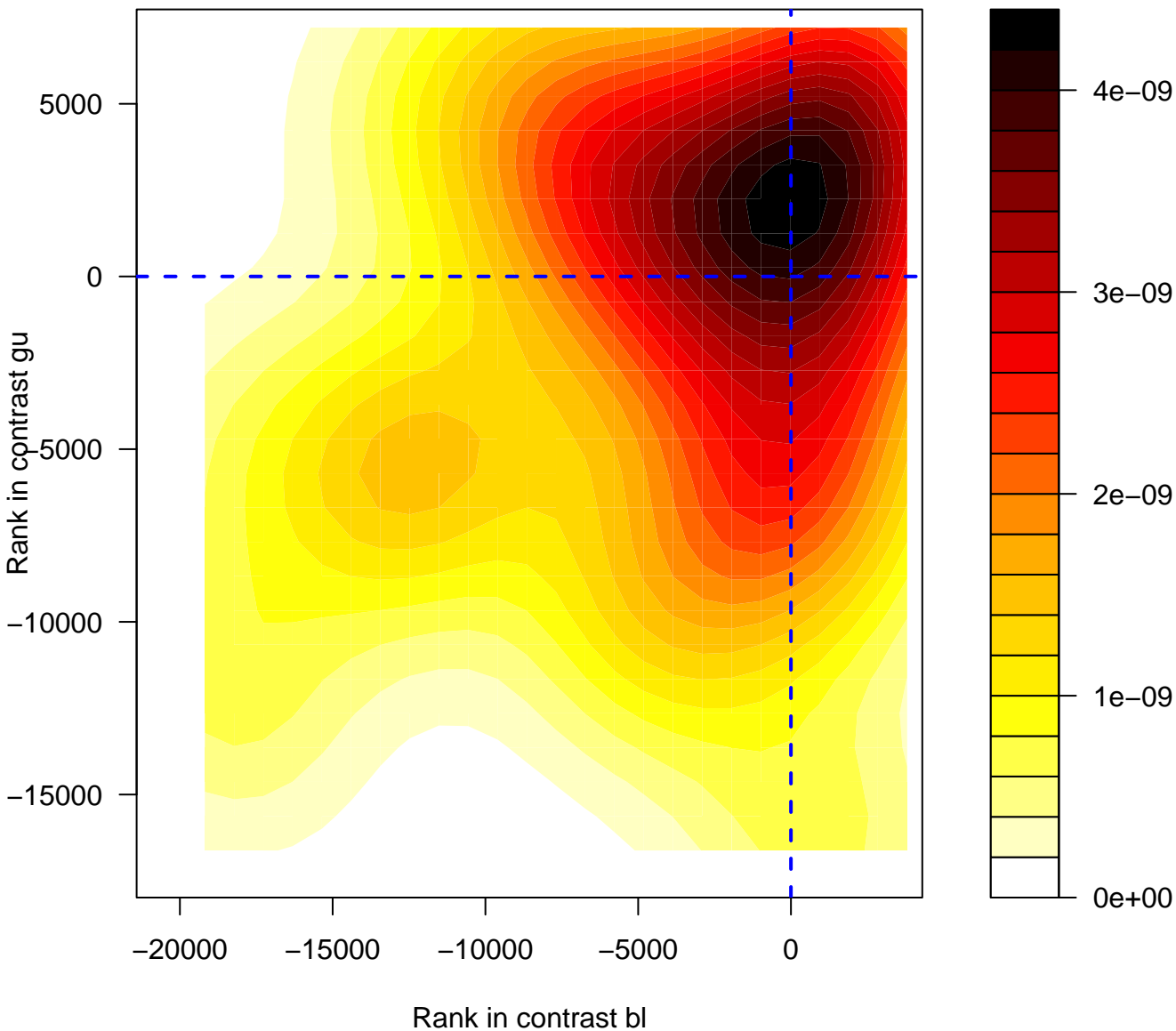
Activation of RAC1



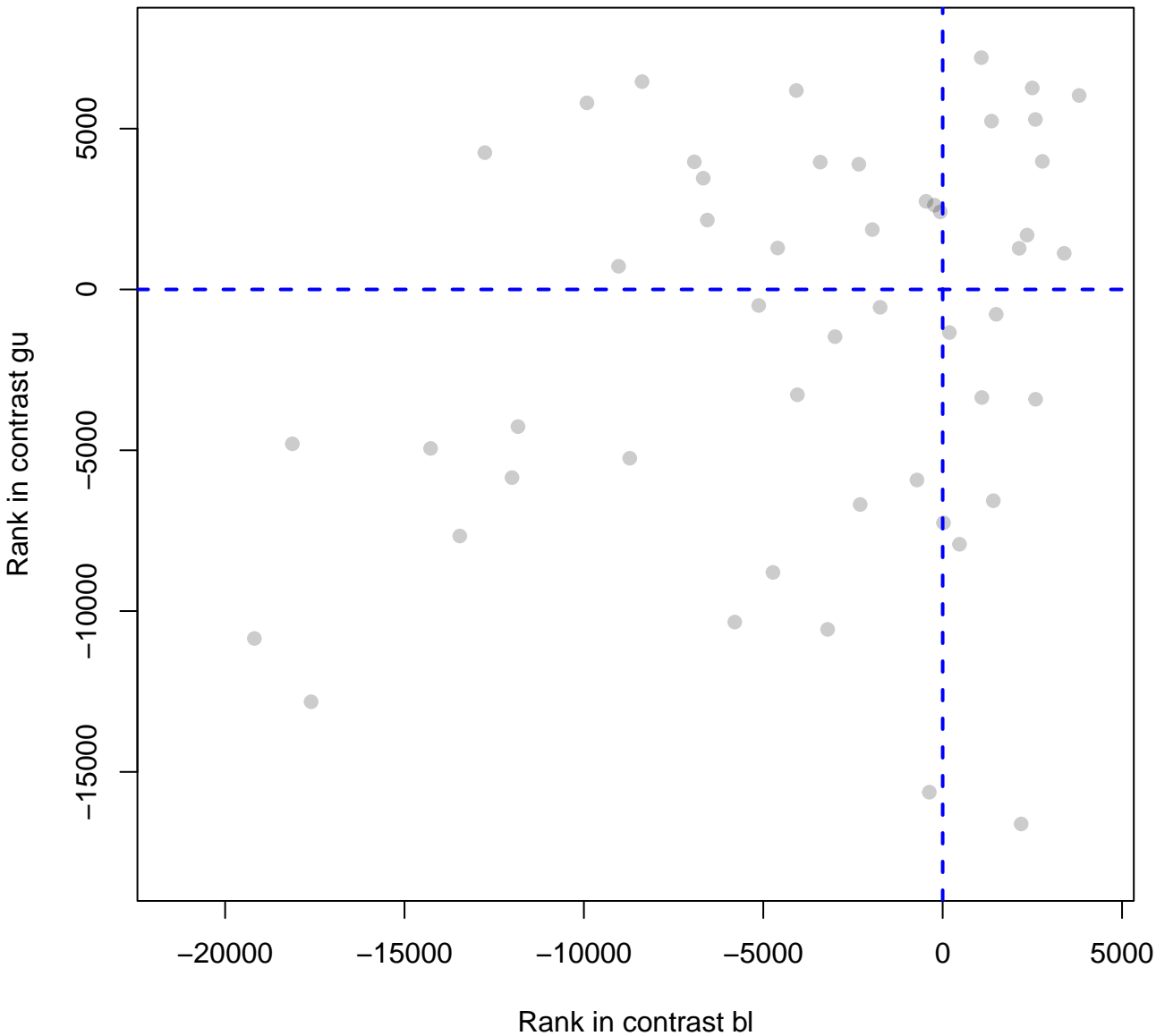
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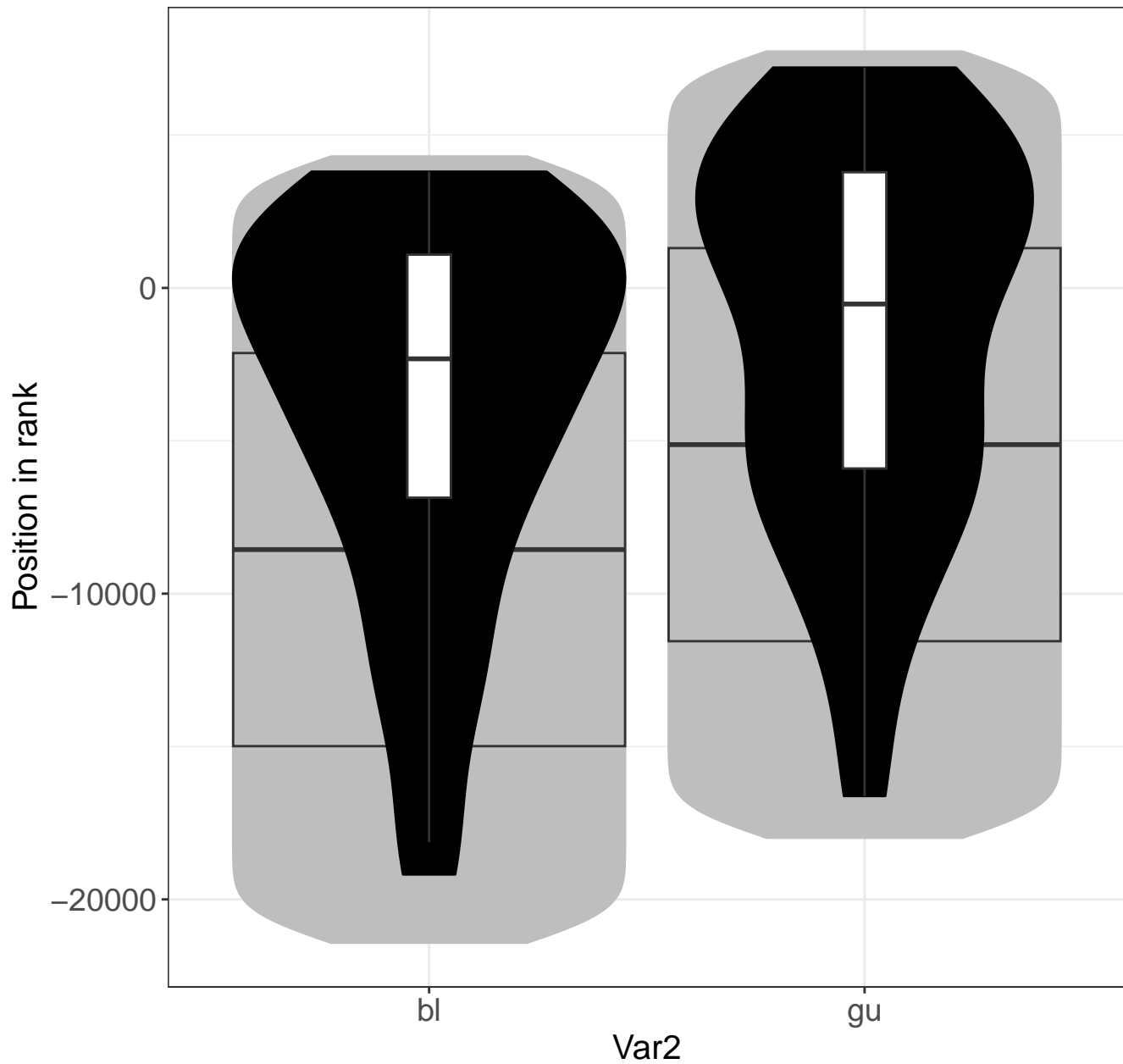
Autodegradation of the E3 ubiquitin ligase COP1



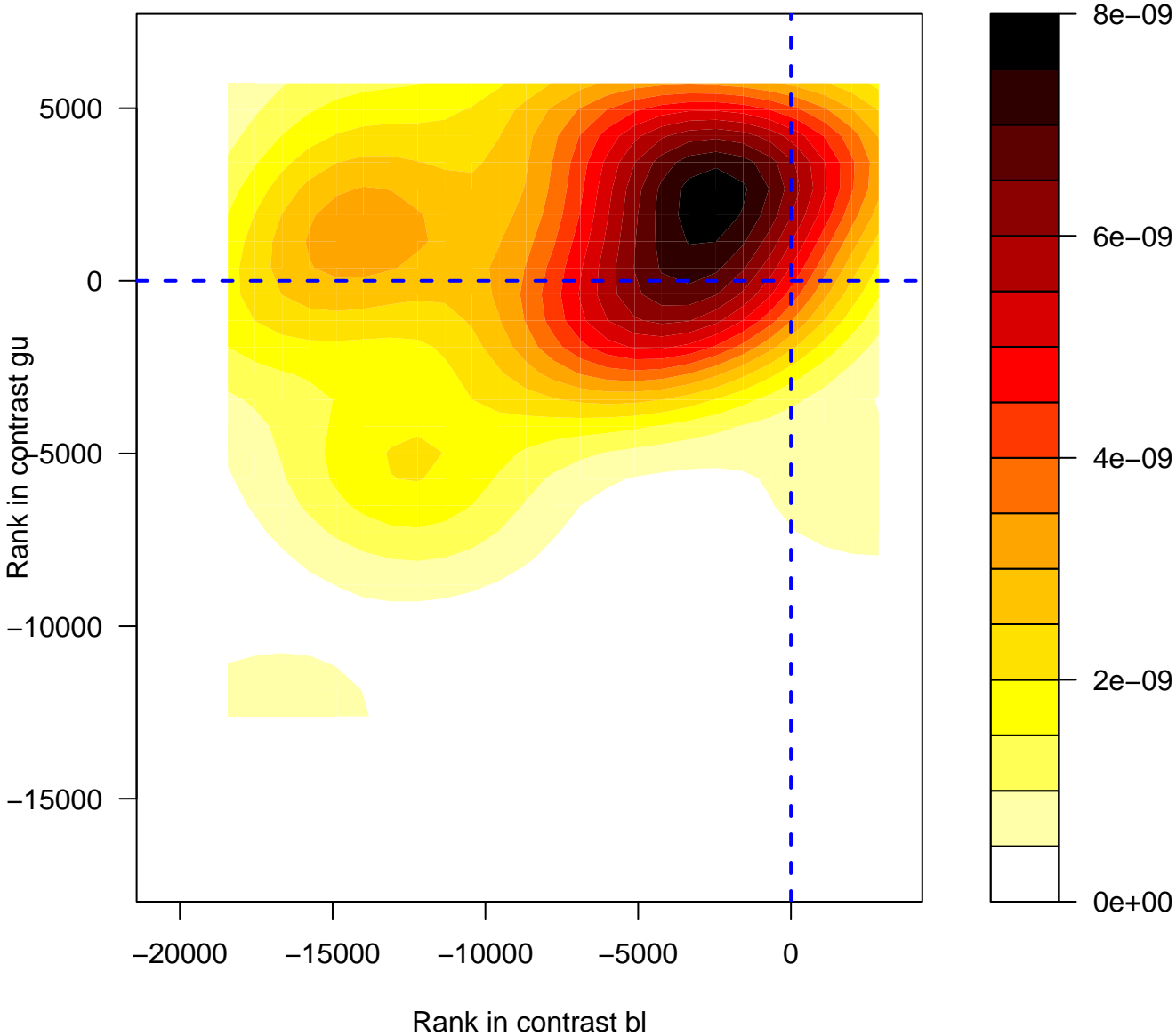
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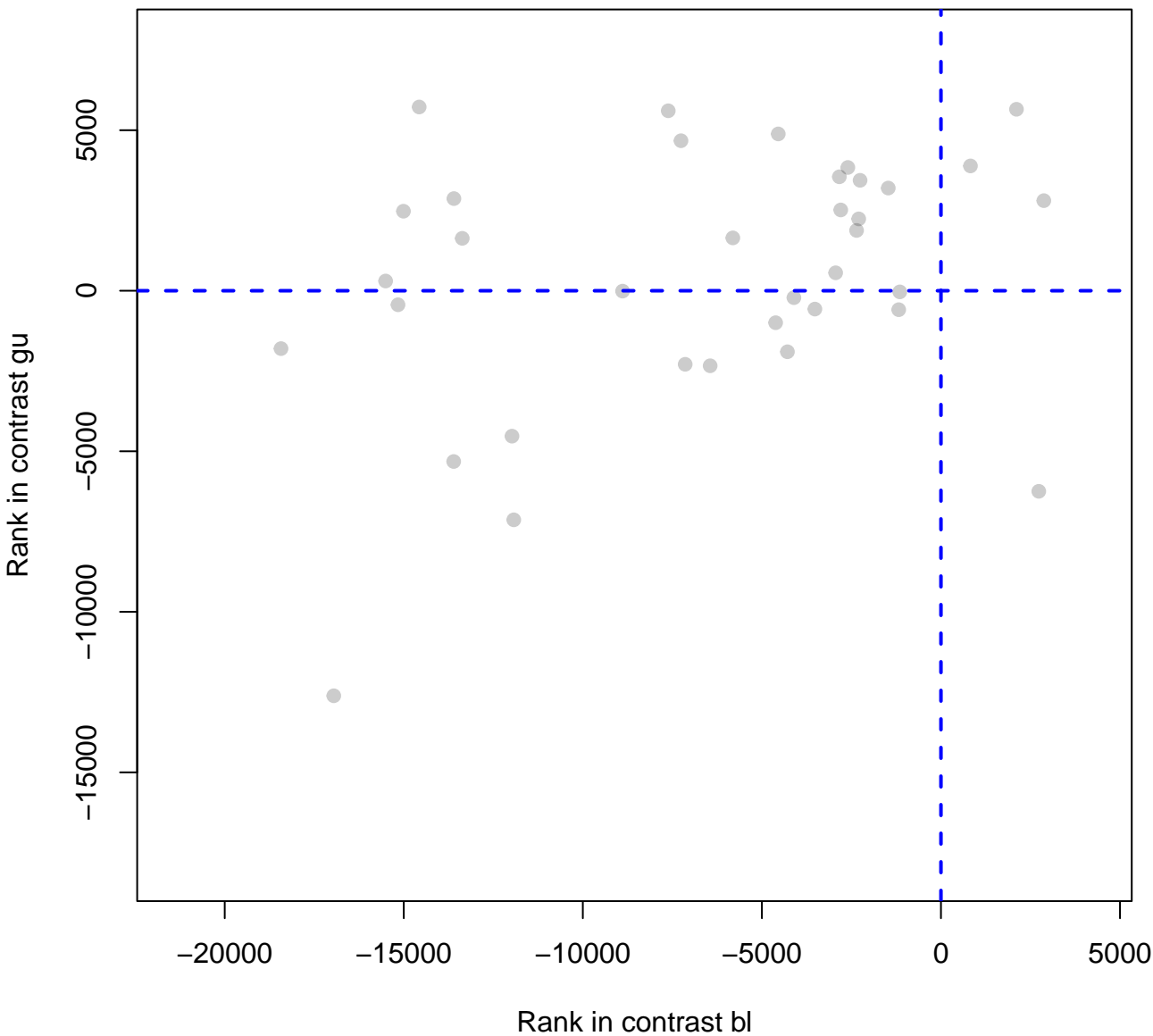
Autodegradation of the E3 ubiquitin ligase COP1



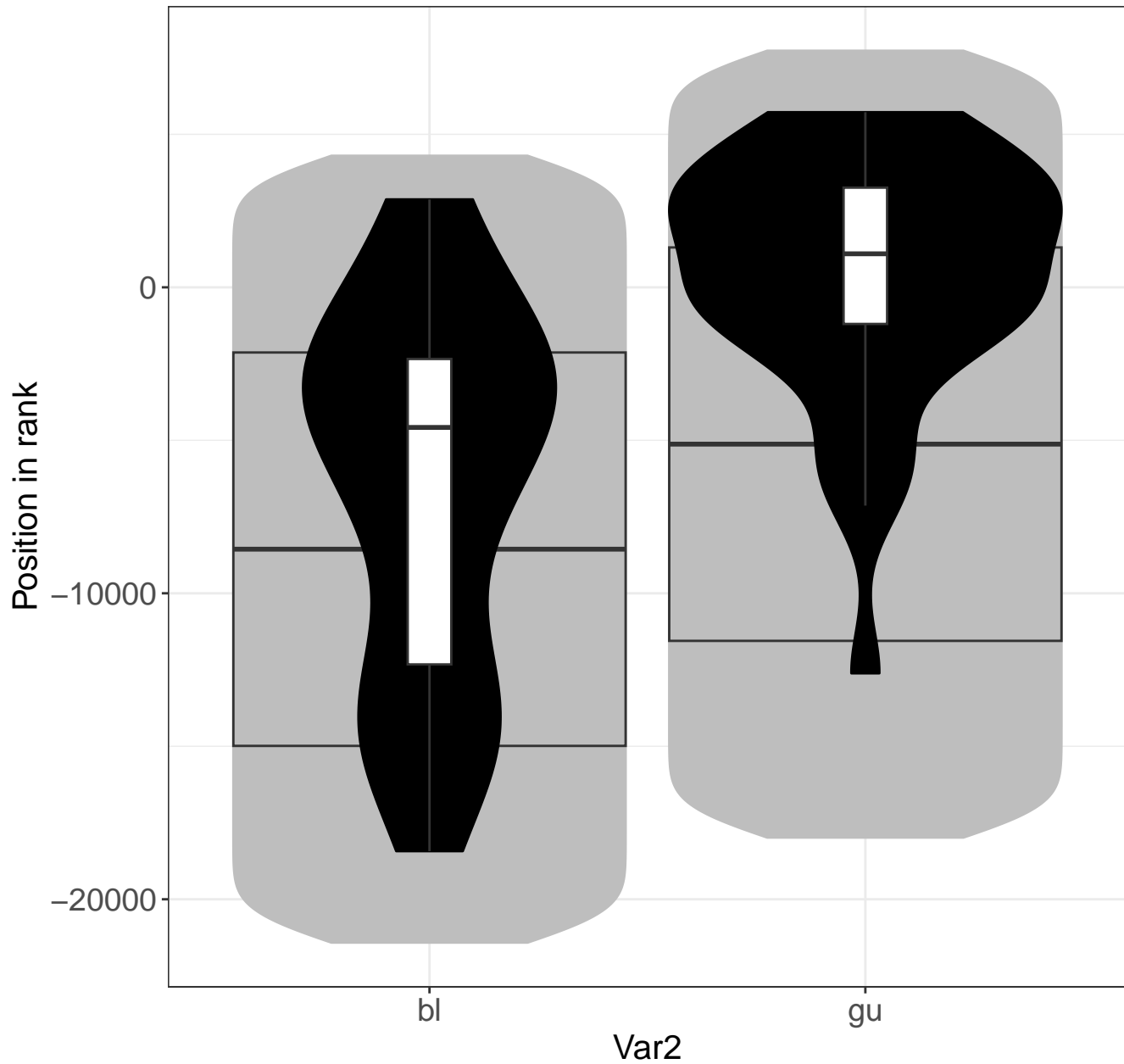
Aberrant regulation of mitotic cell cycle due to RB1 defec



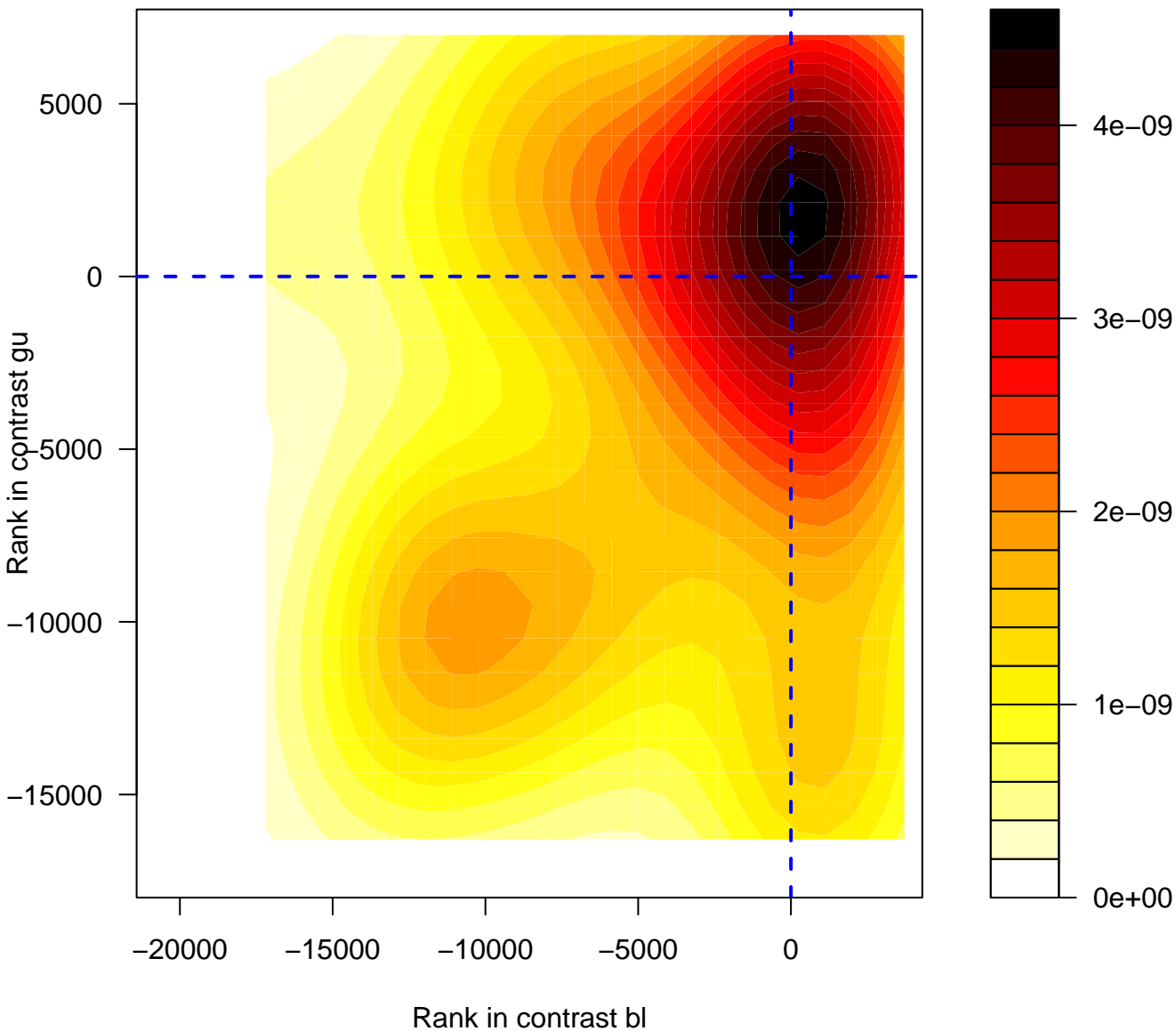
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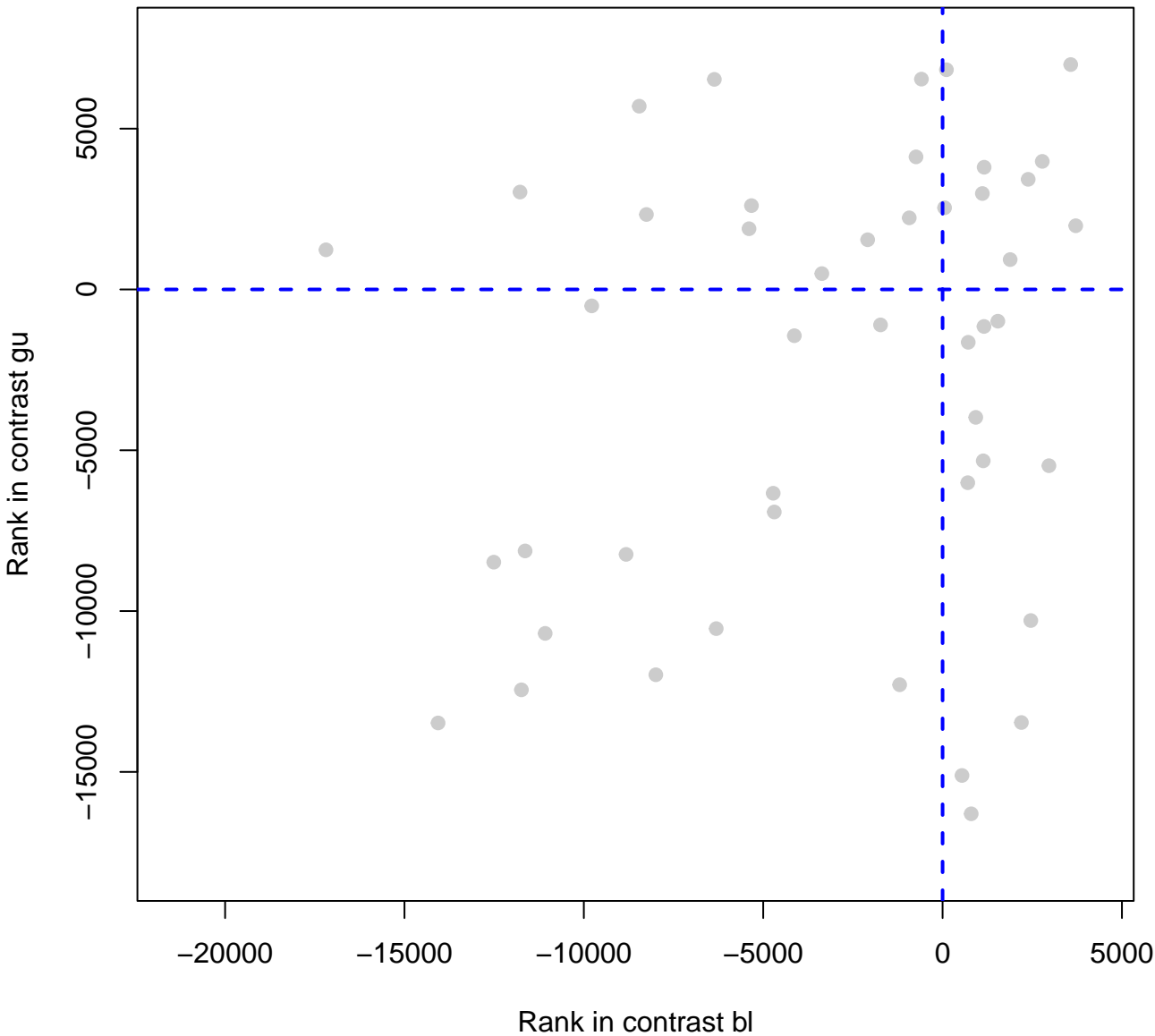
Aberrant regulation of mitotic cell cycle due to RE



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Formation of the ternary complex, and subsequently, the 43S complex



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