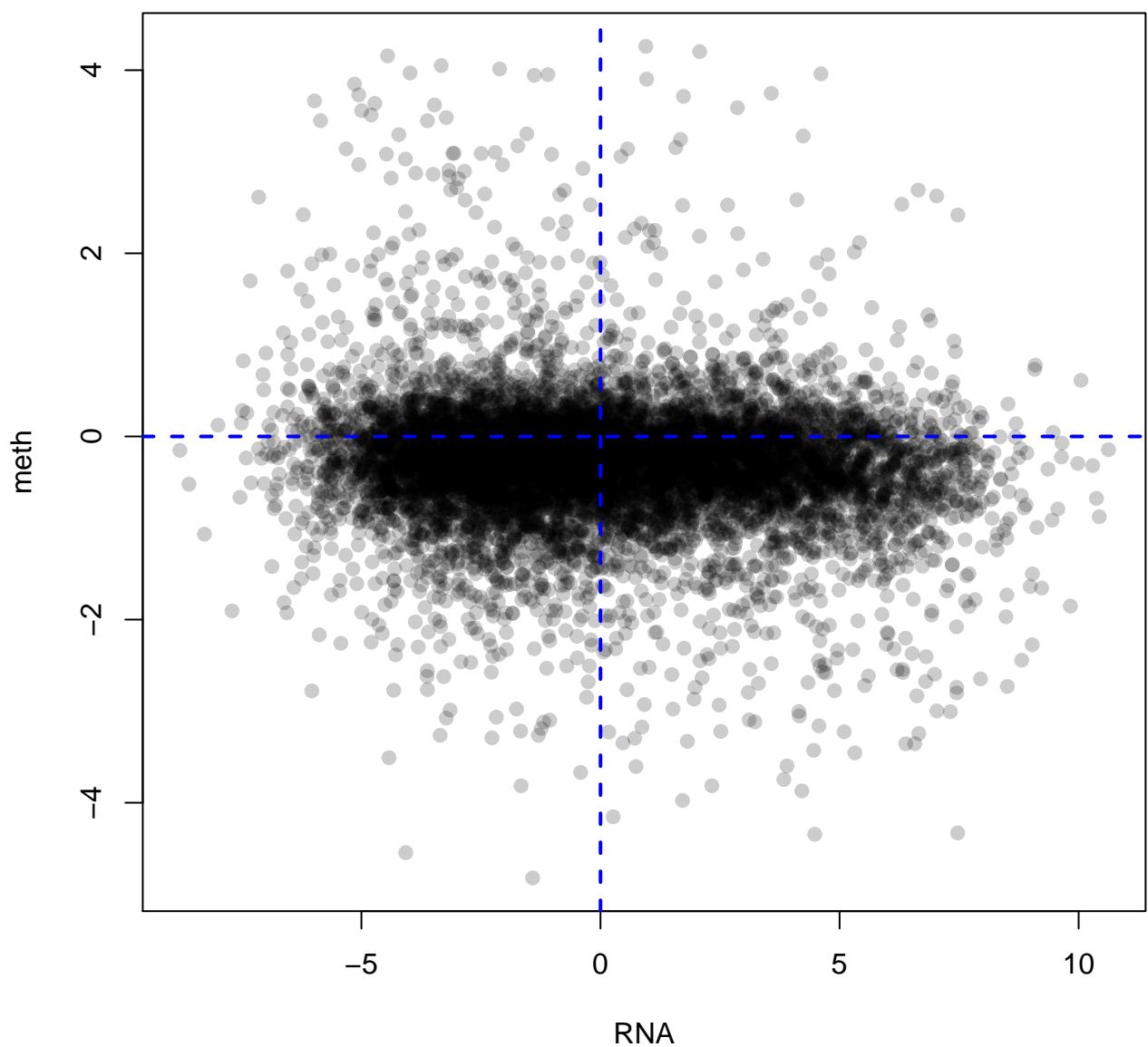
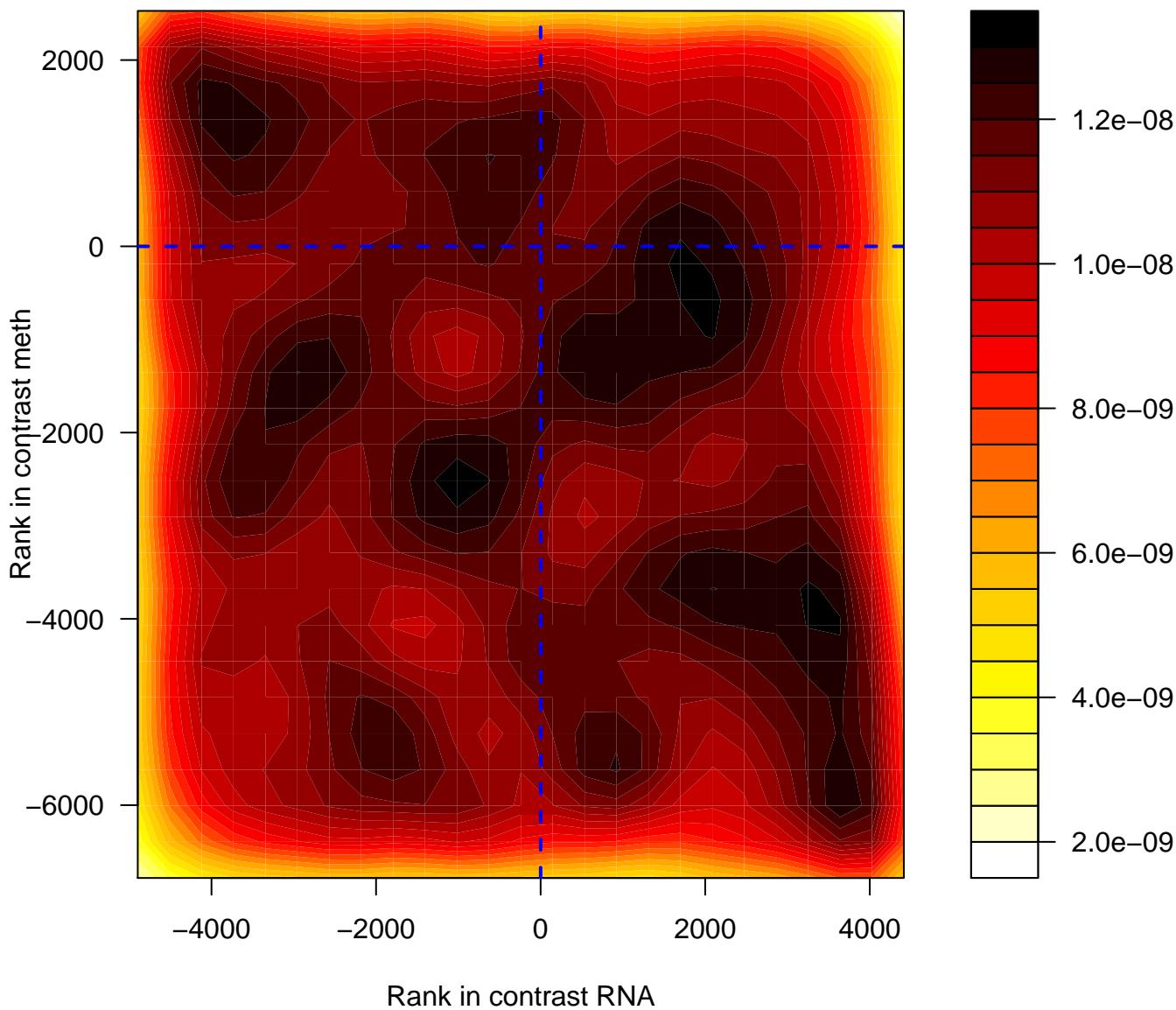


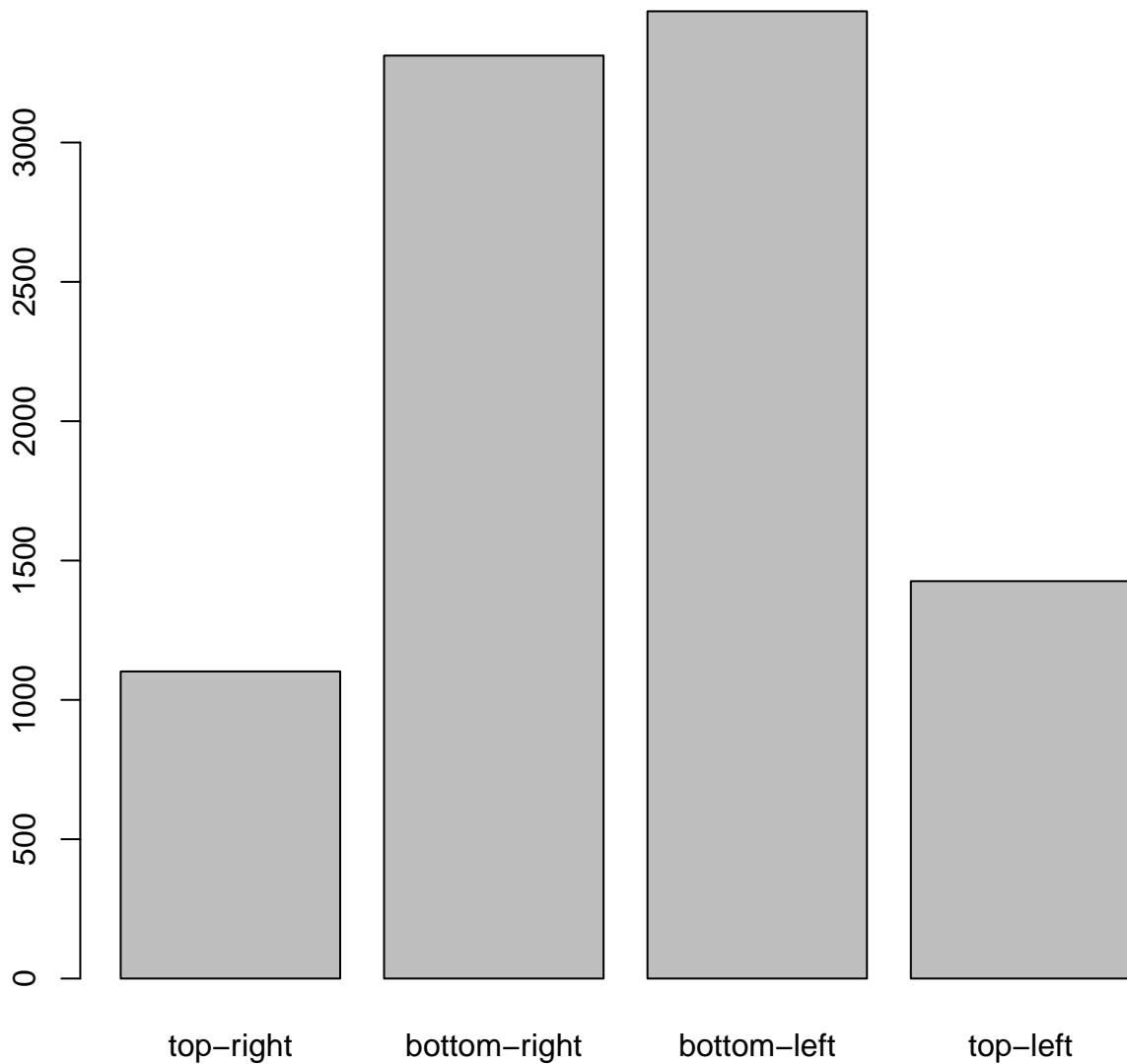
Scatterplot of all genes



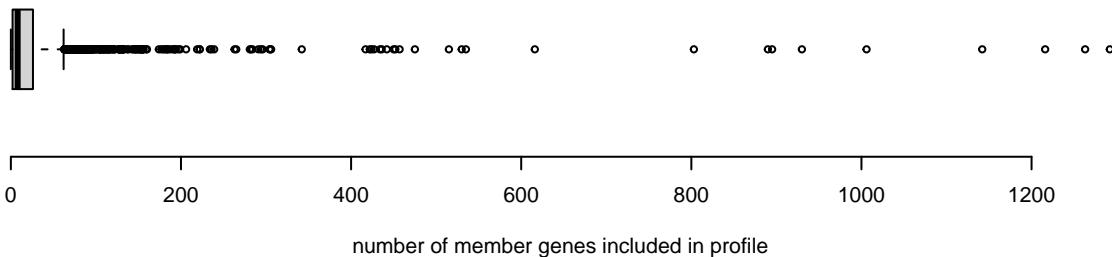
Rank–rank plot of all genes



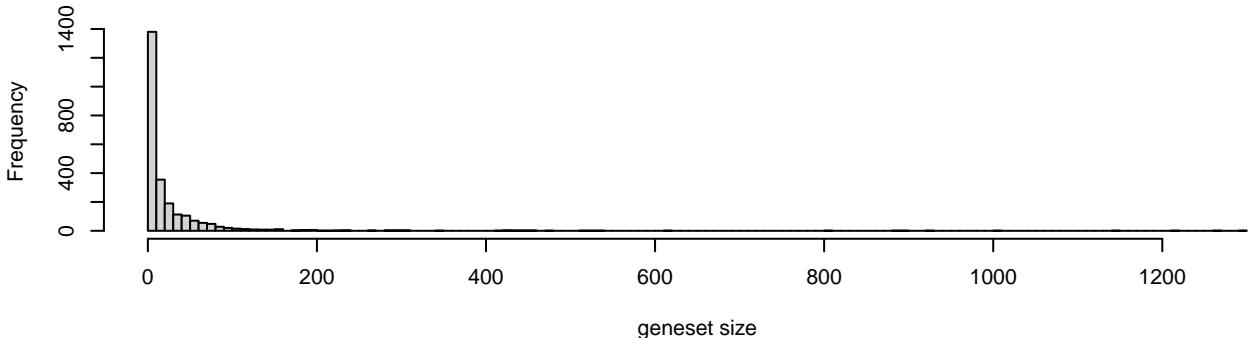
number of genes in each quadrant



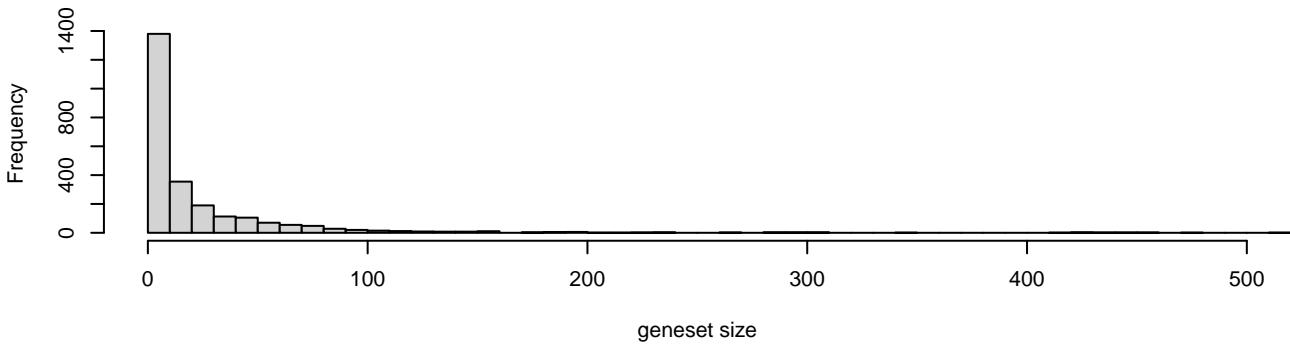
Gene set size



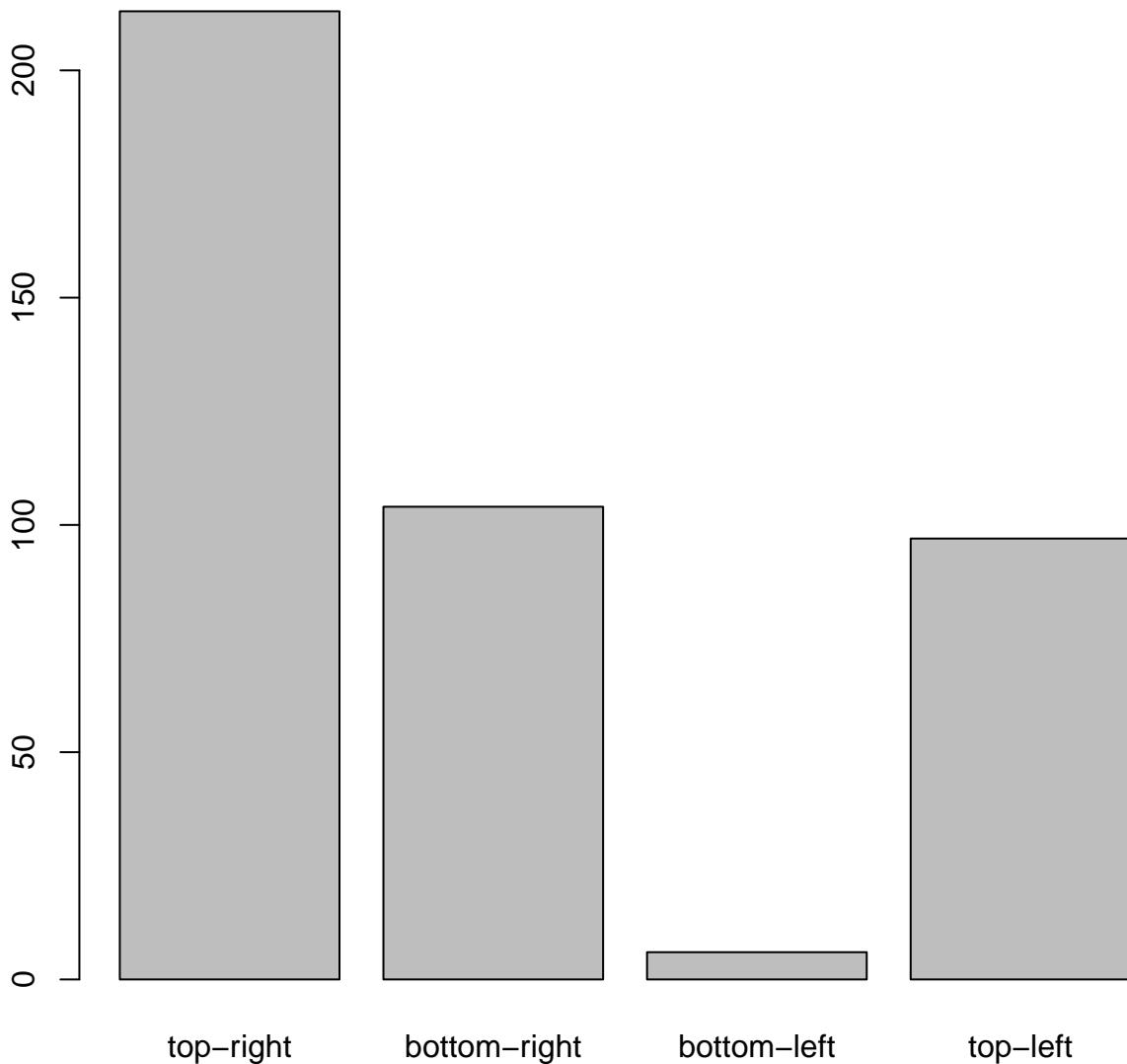
Histogram of geneset size



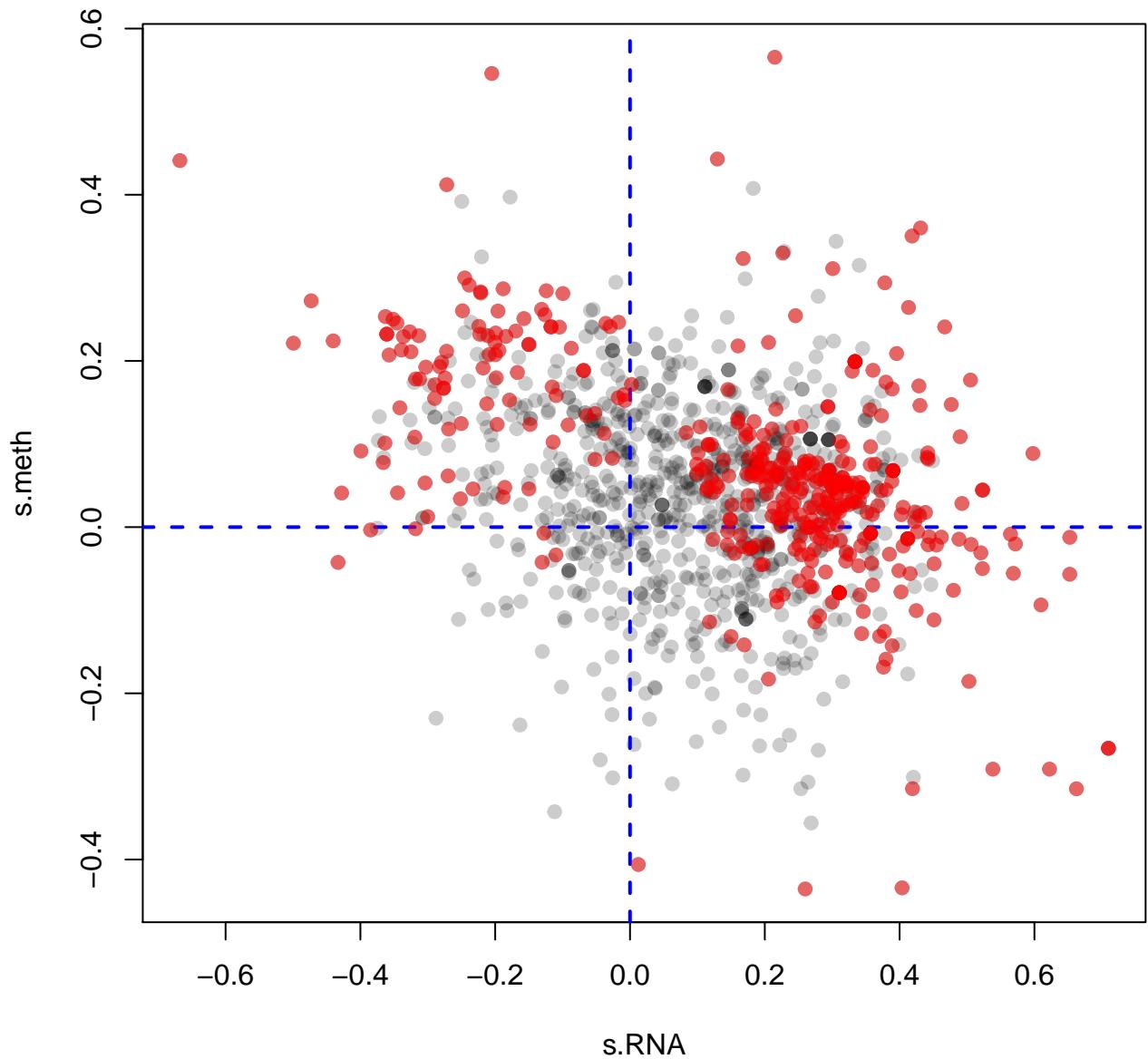
Trimmed histogram of geneset size



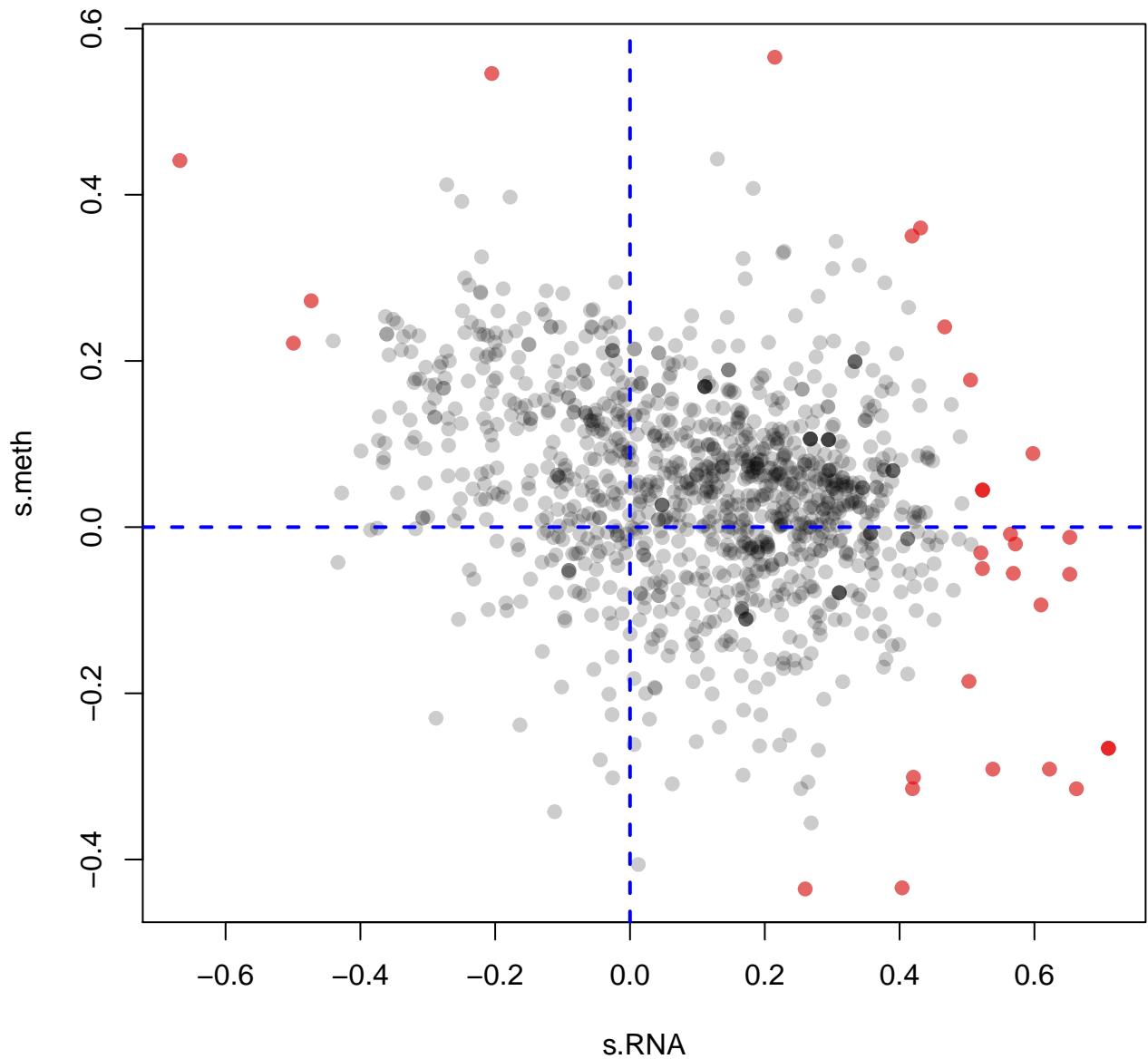
number of genesets FDR<0.05



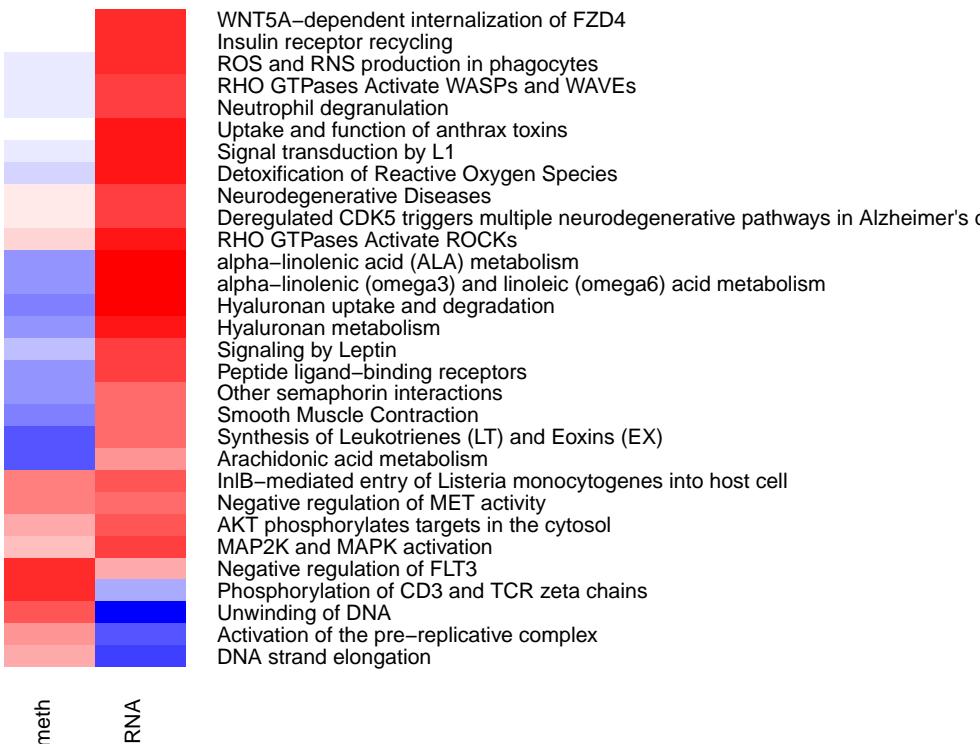
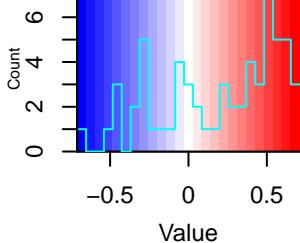
Scatterplot of all gene sets; FDR<0.05 in red



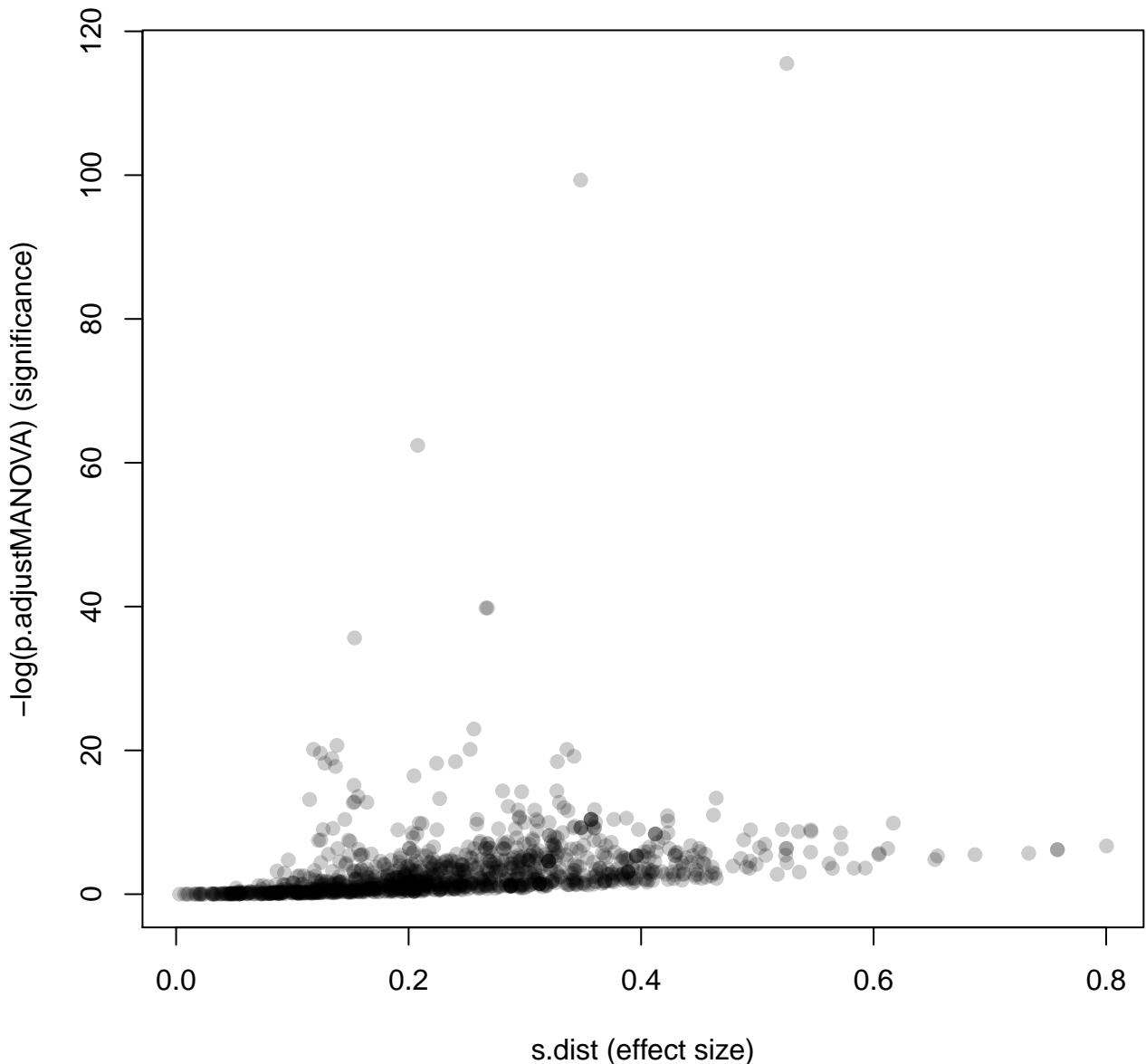
Scatterplot of all gene sets; top 30 in red



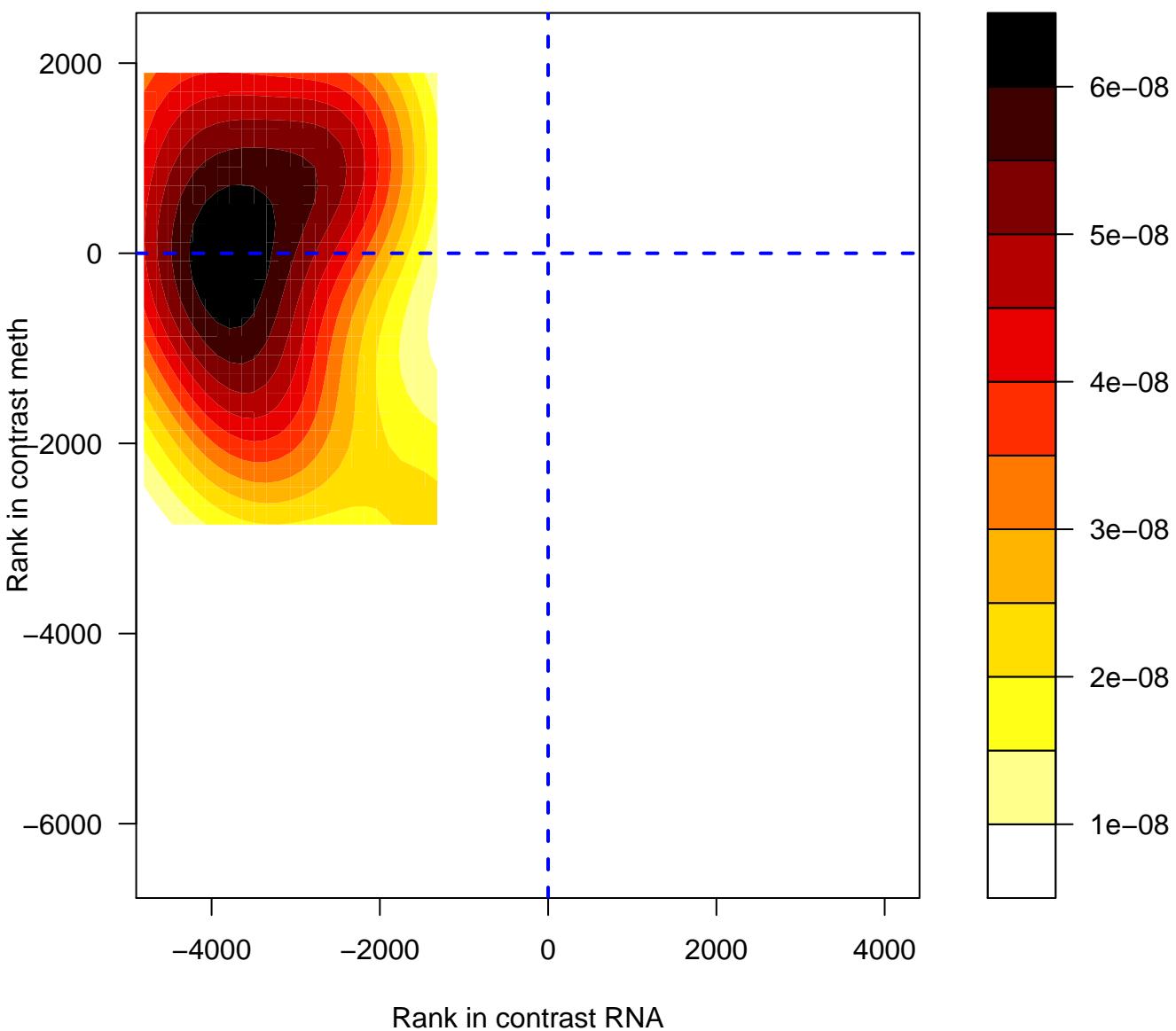
Color Key and Histogram



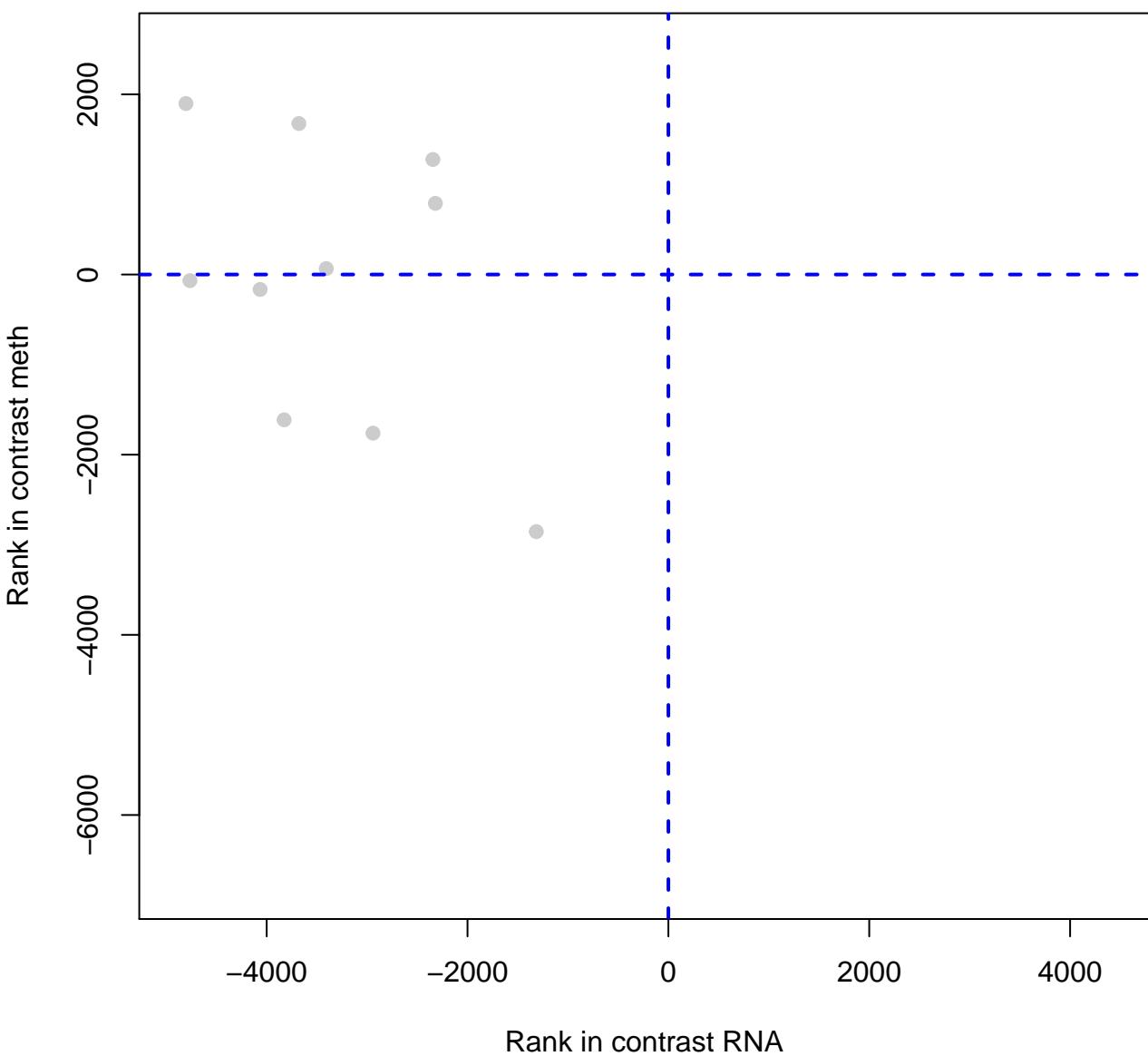
effect size versus statistical significance



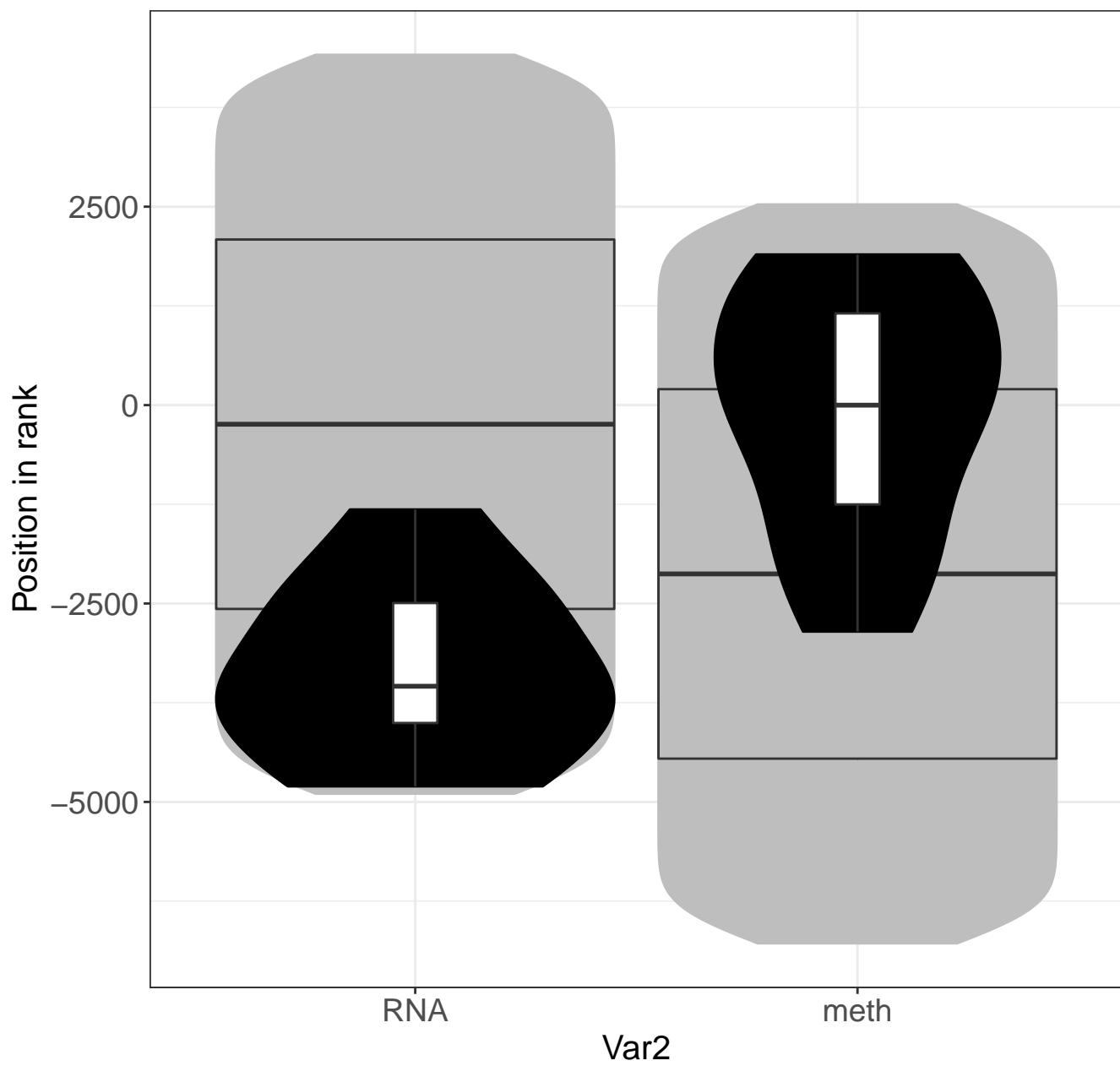
Unwinding of DNA



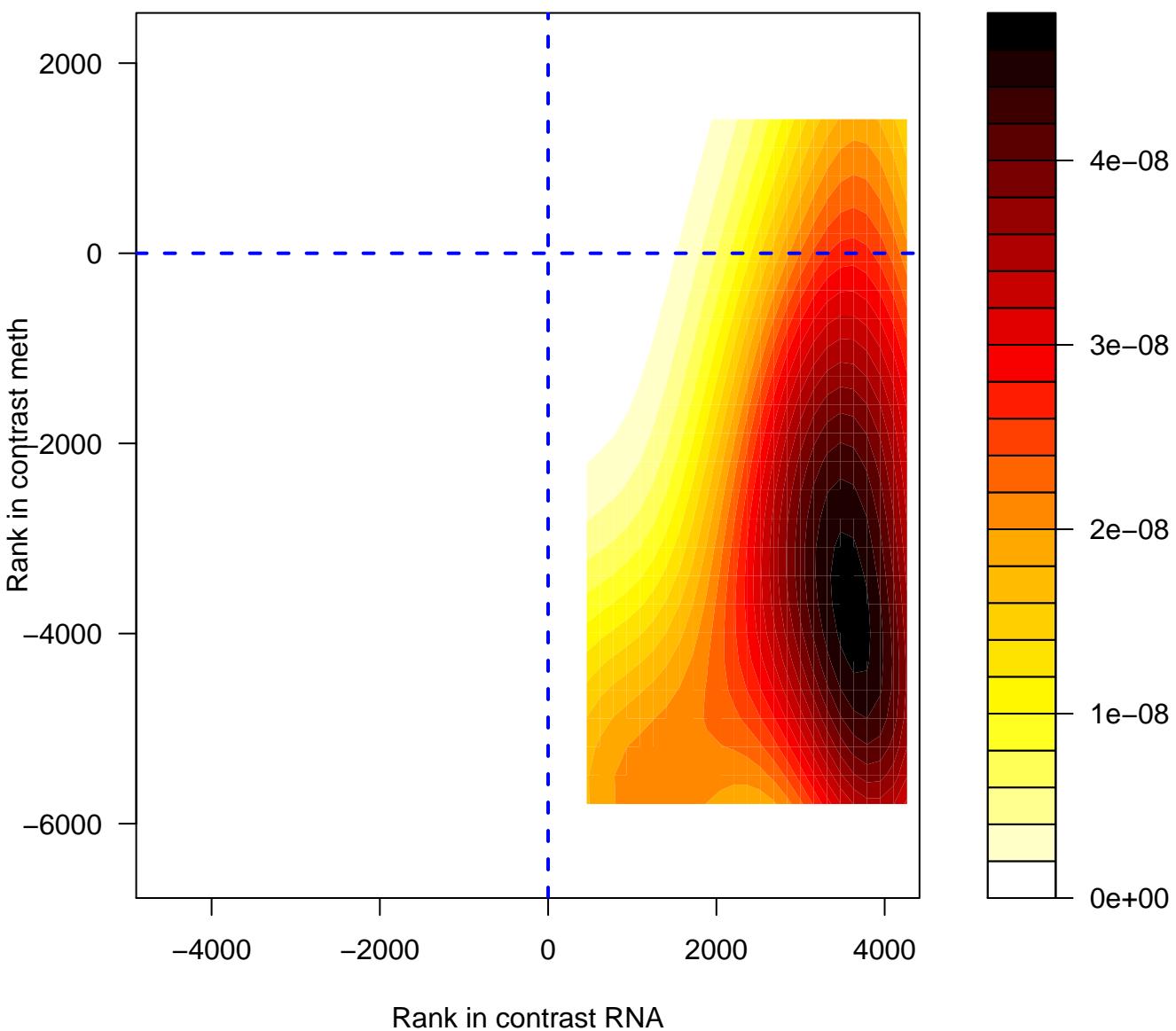
Unwinding of DNA



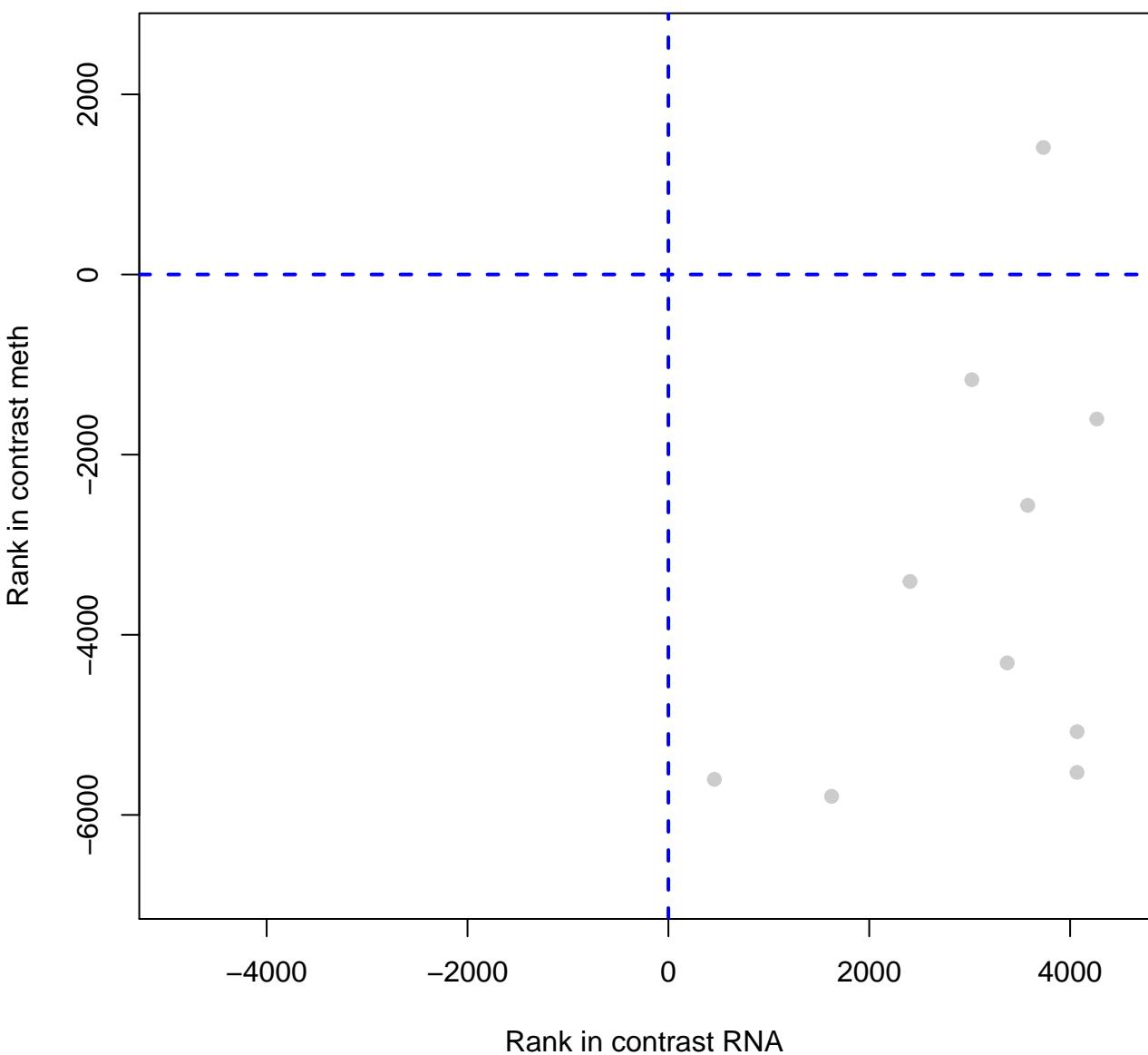
Unwinding of DNA



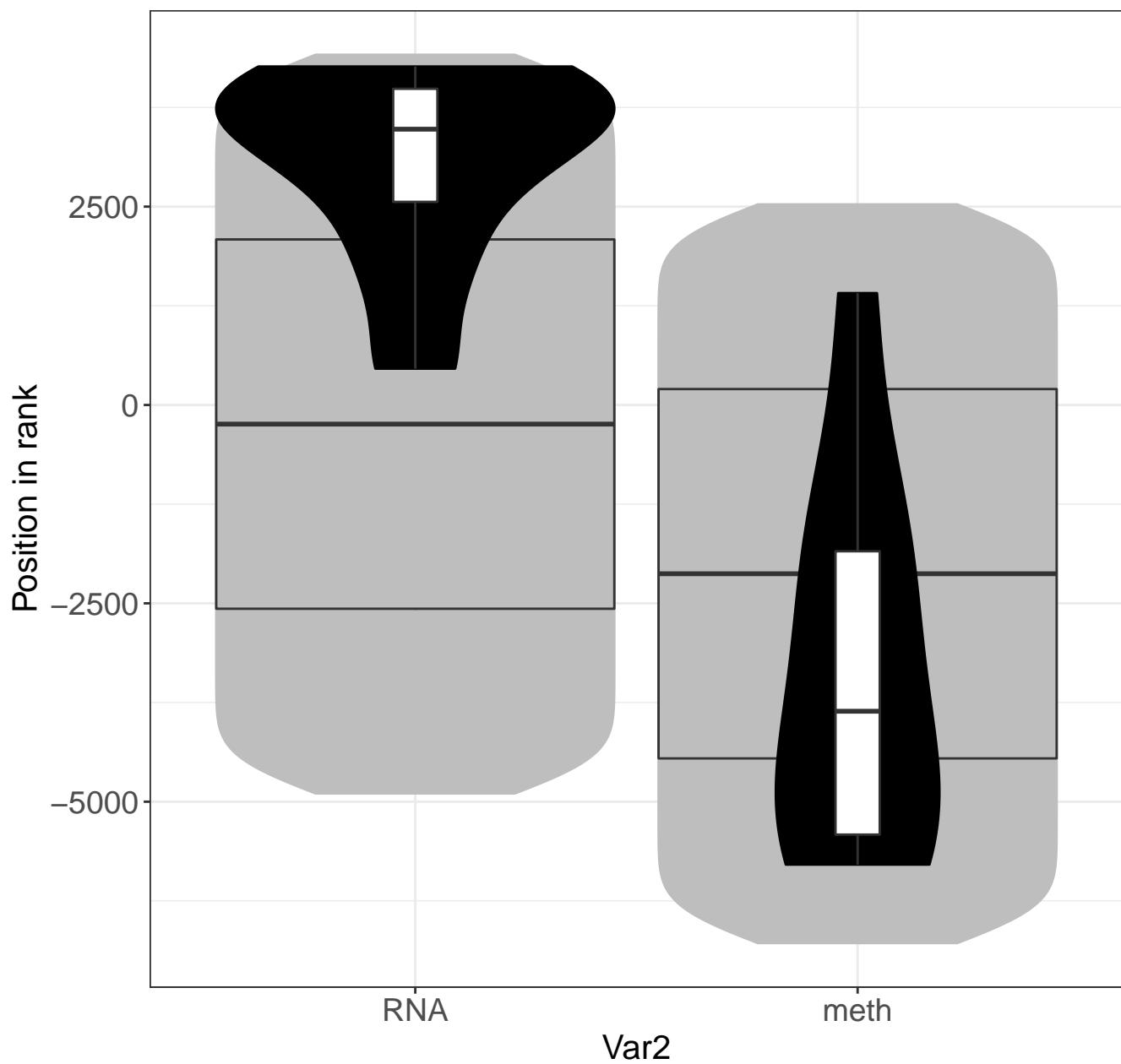
alpha-linolenic (omega3) and linoleic (omega6) acid metabolism



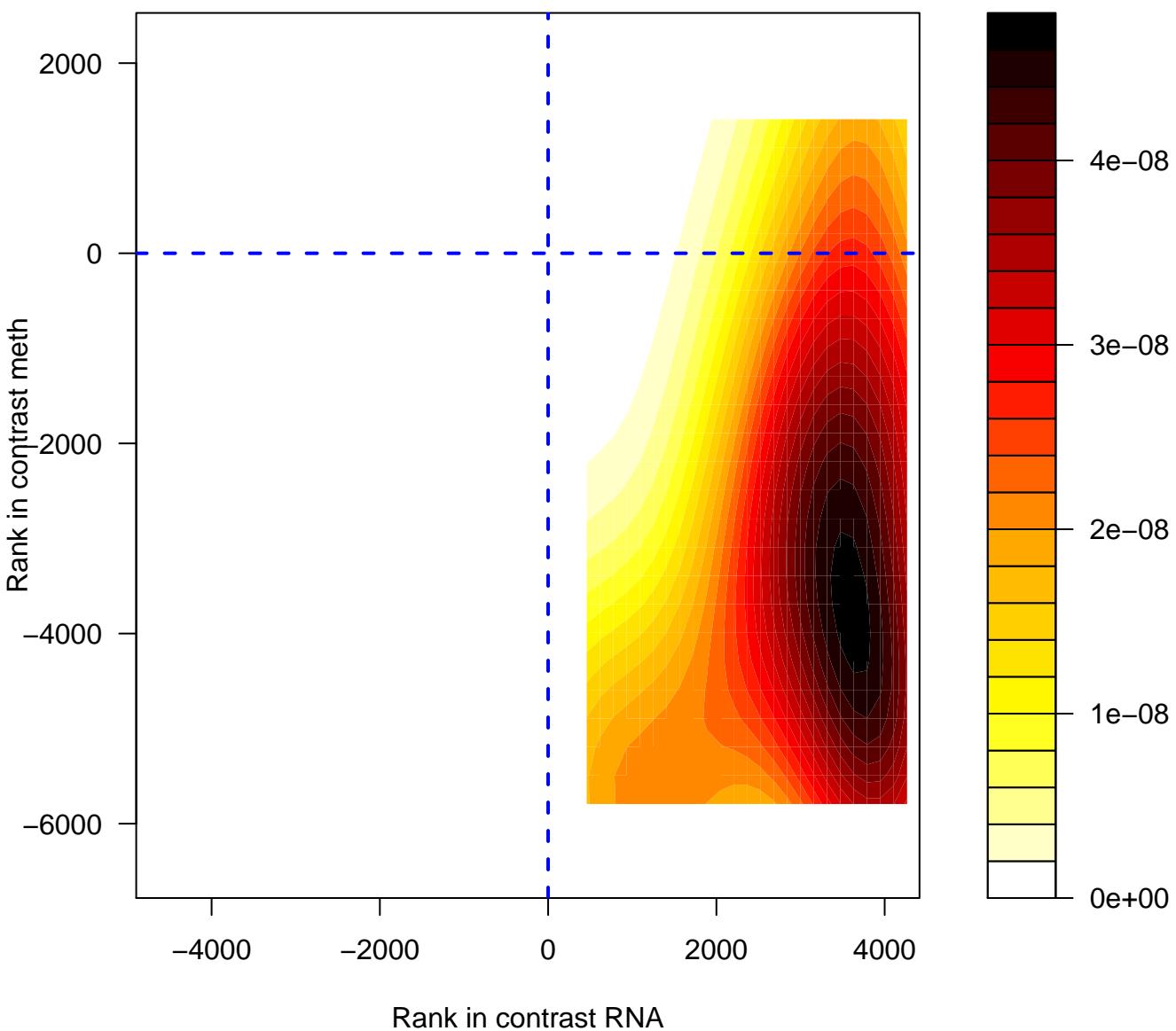
alpha-linolenic (omega3) and linoleic (omega6) acid metabolism



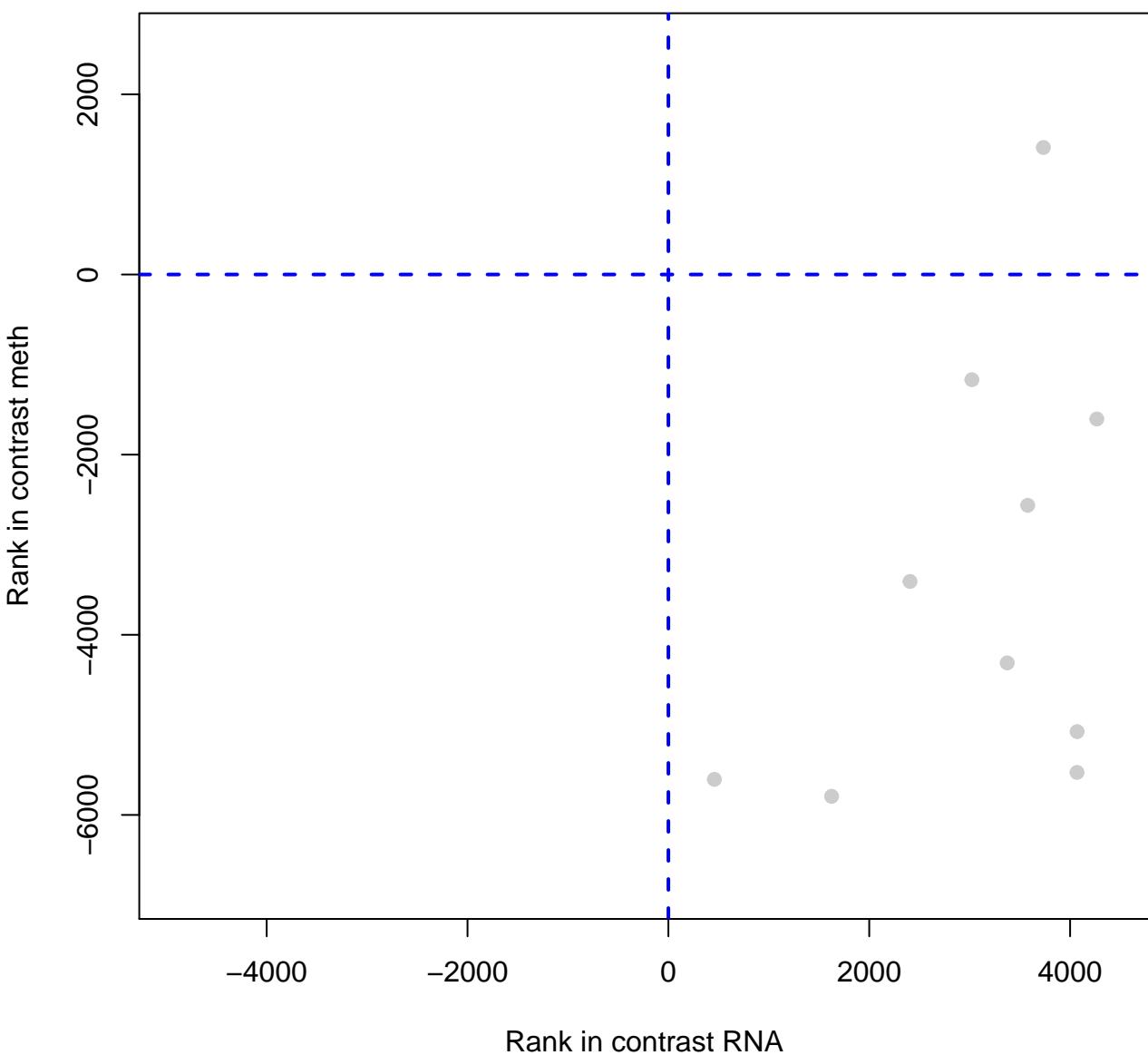
alpha-linolenic (omega3) and linoleic (omega6) a



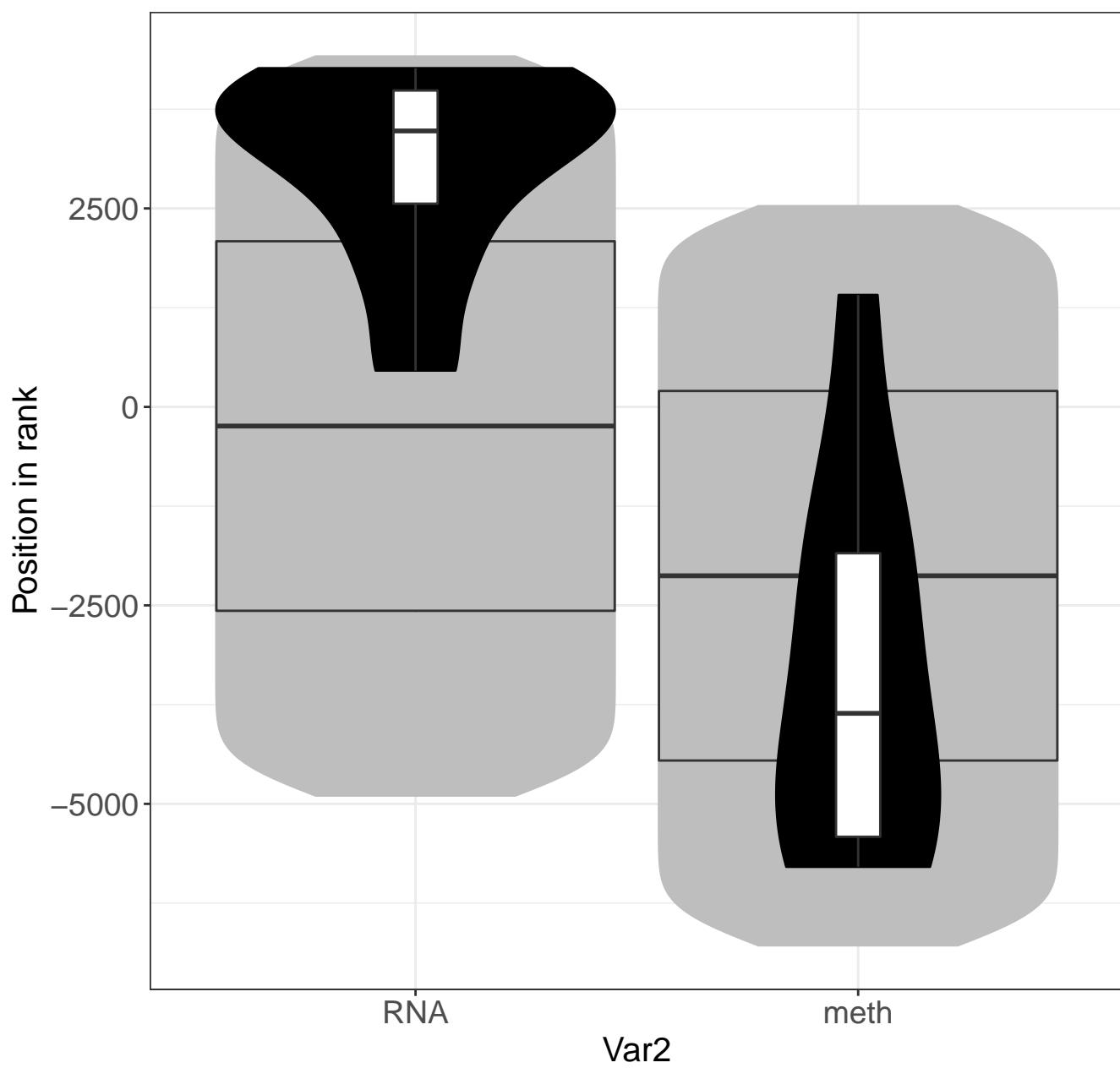
alpha-linolenic acid (ALA) metabolism



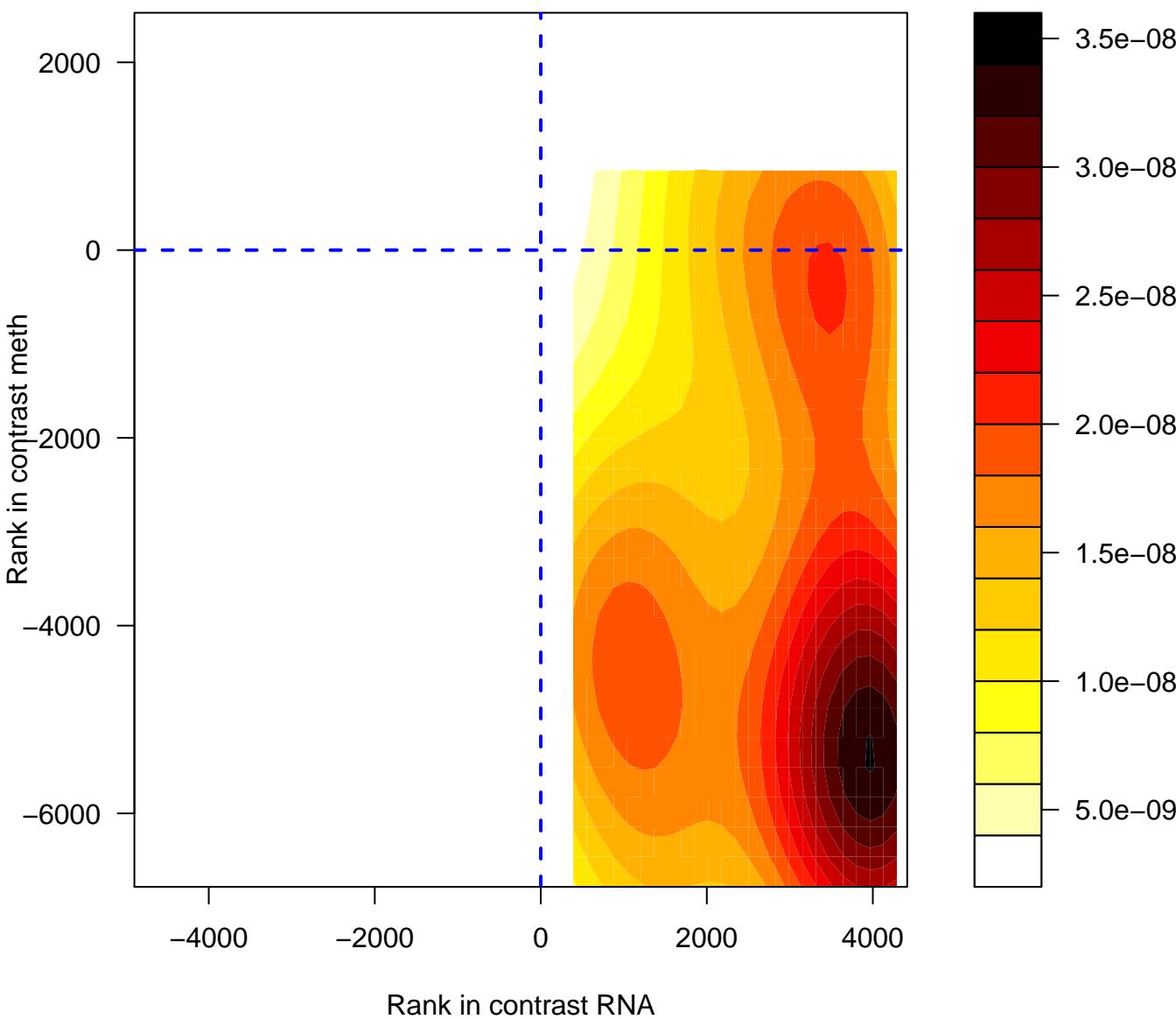
alpha-linolenic acid (ALA) metabolism



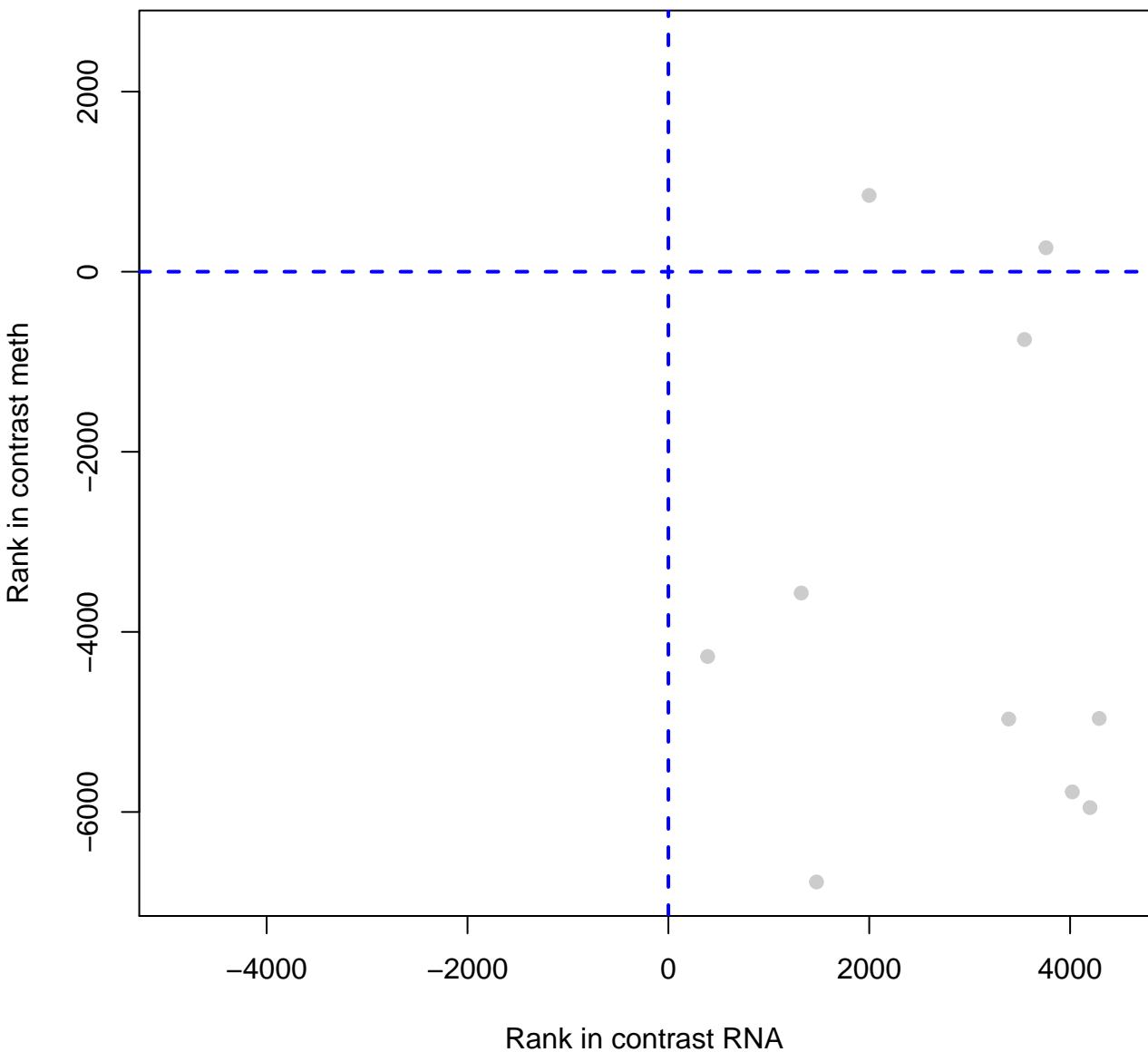
alpha-linolenic acid (ALA) metabolism



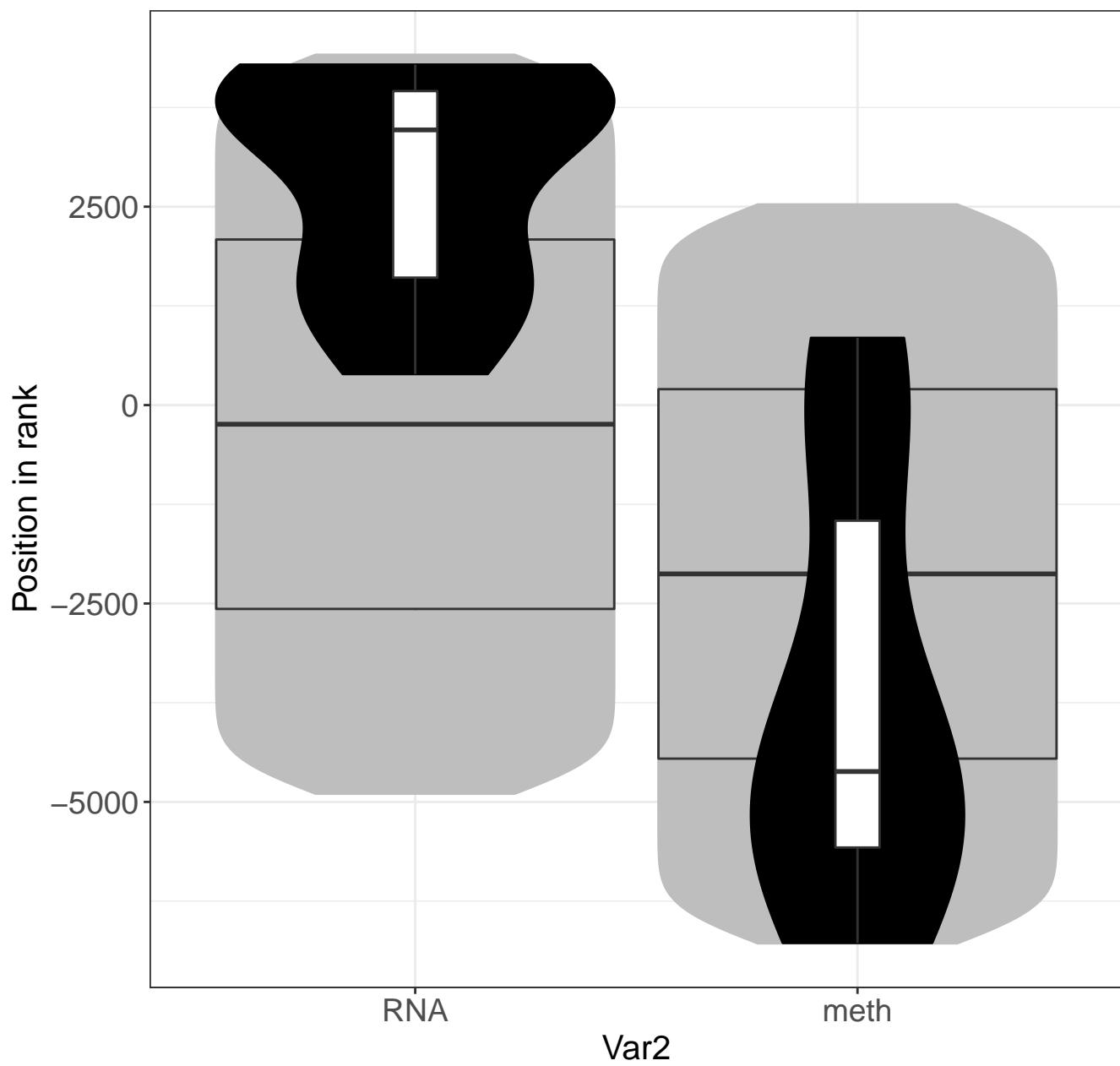
Hyaluronan uptake and degradation



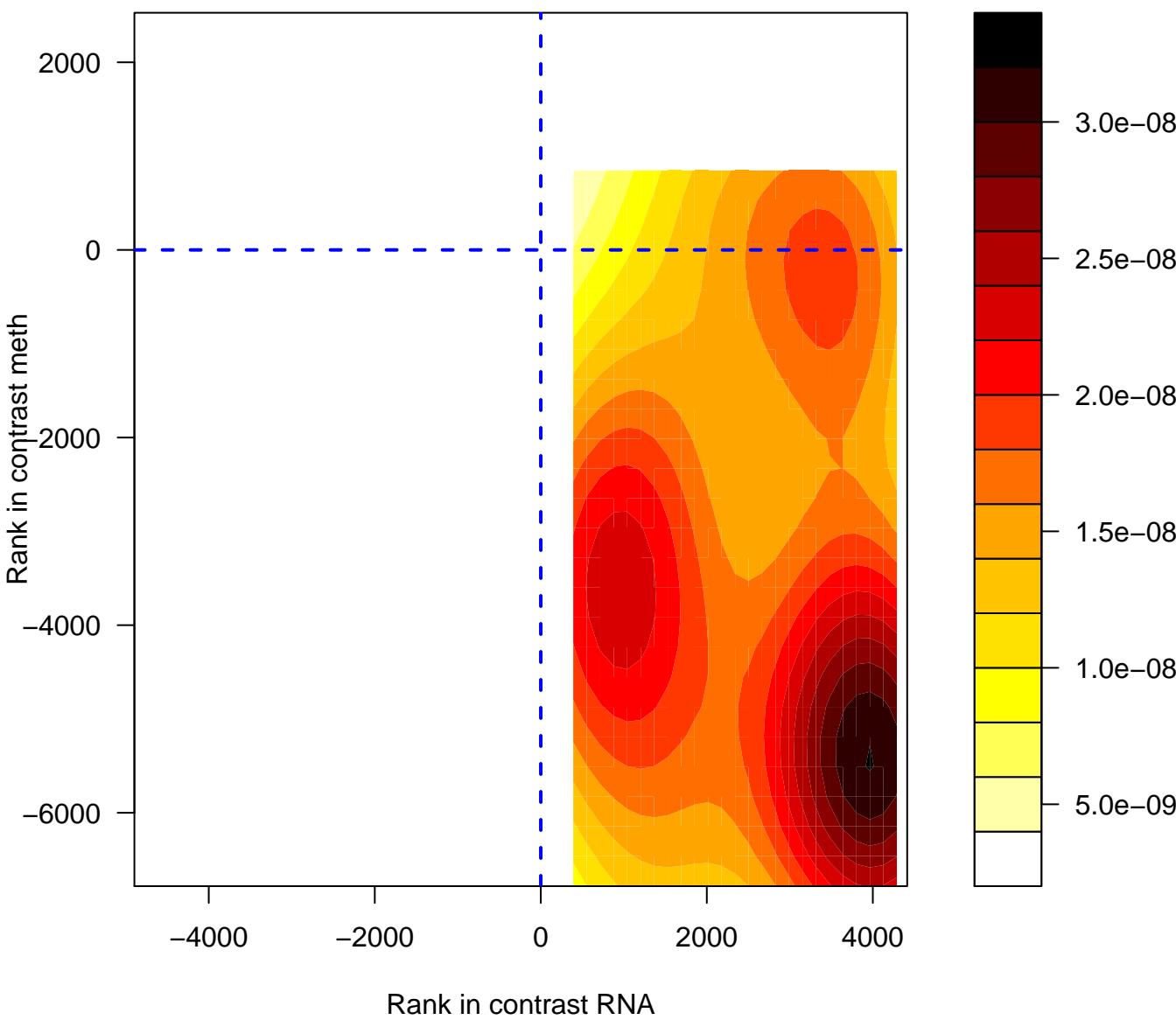
Hyaluronan uptake and degradation



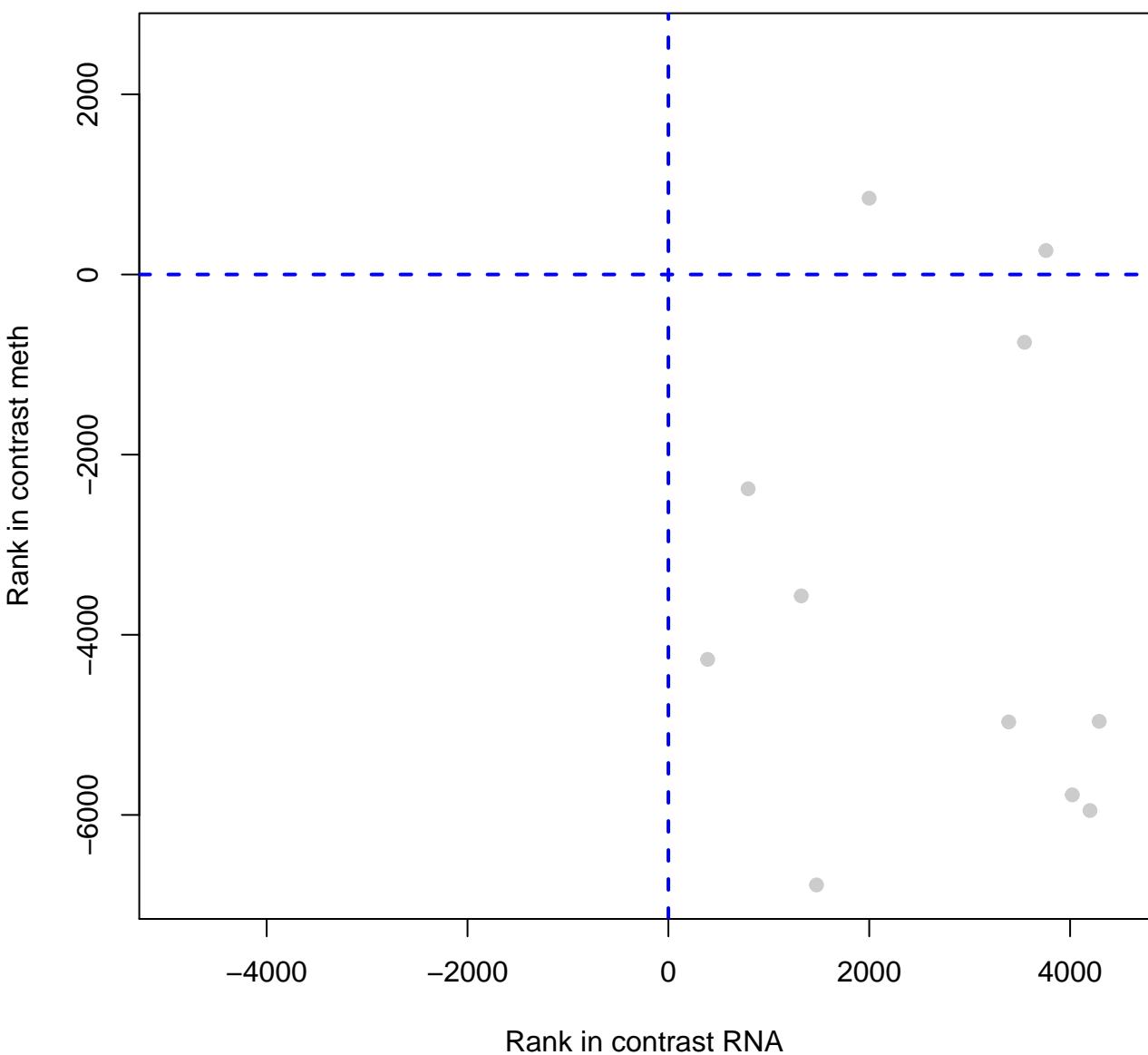
Hyaluronan uptake and degradation



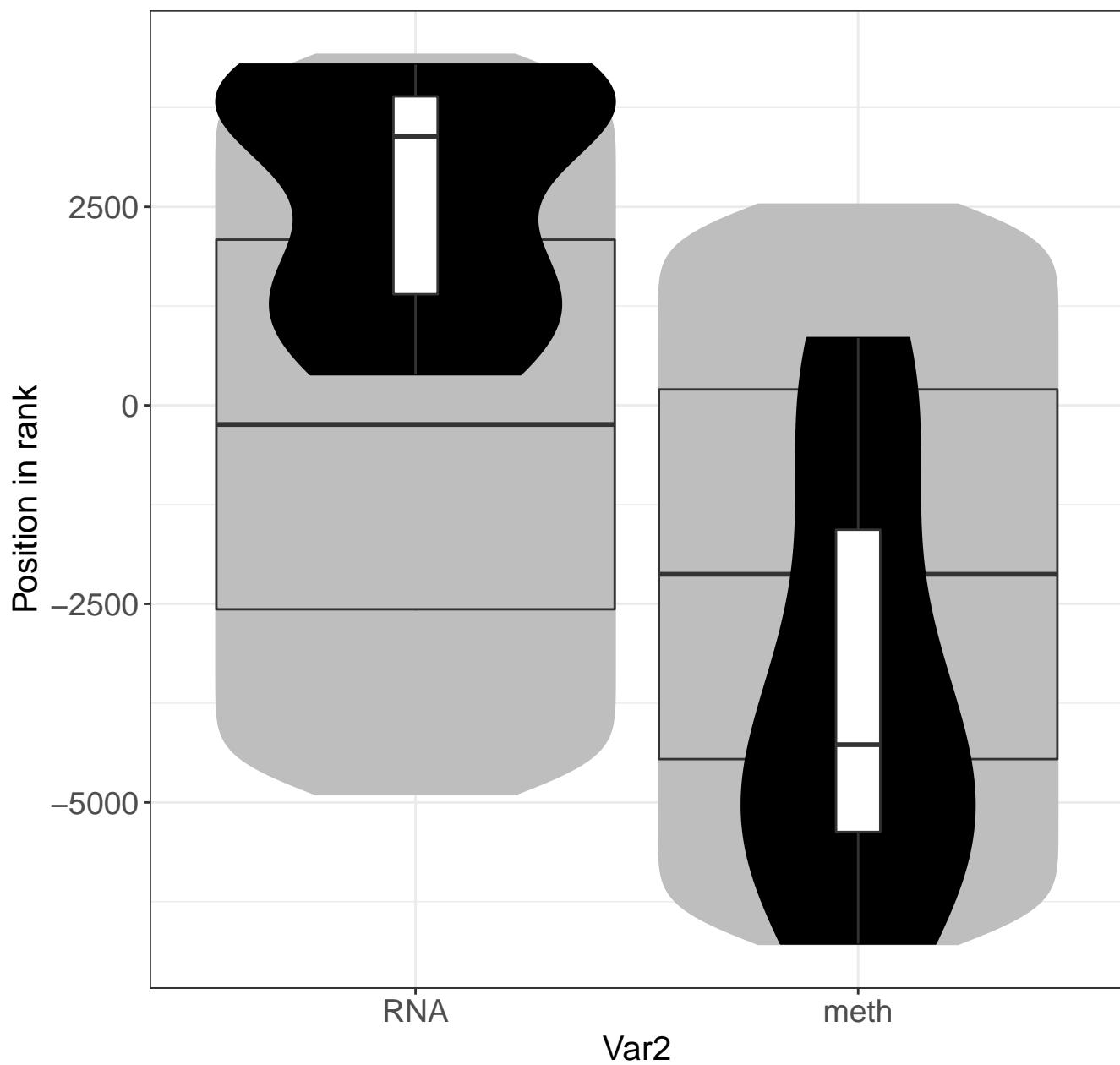
Hyaluronan metabolism



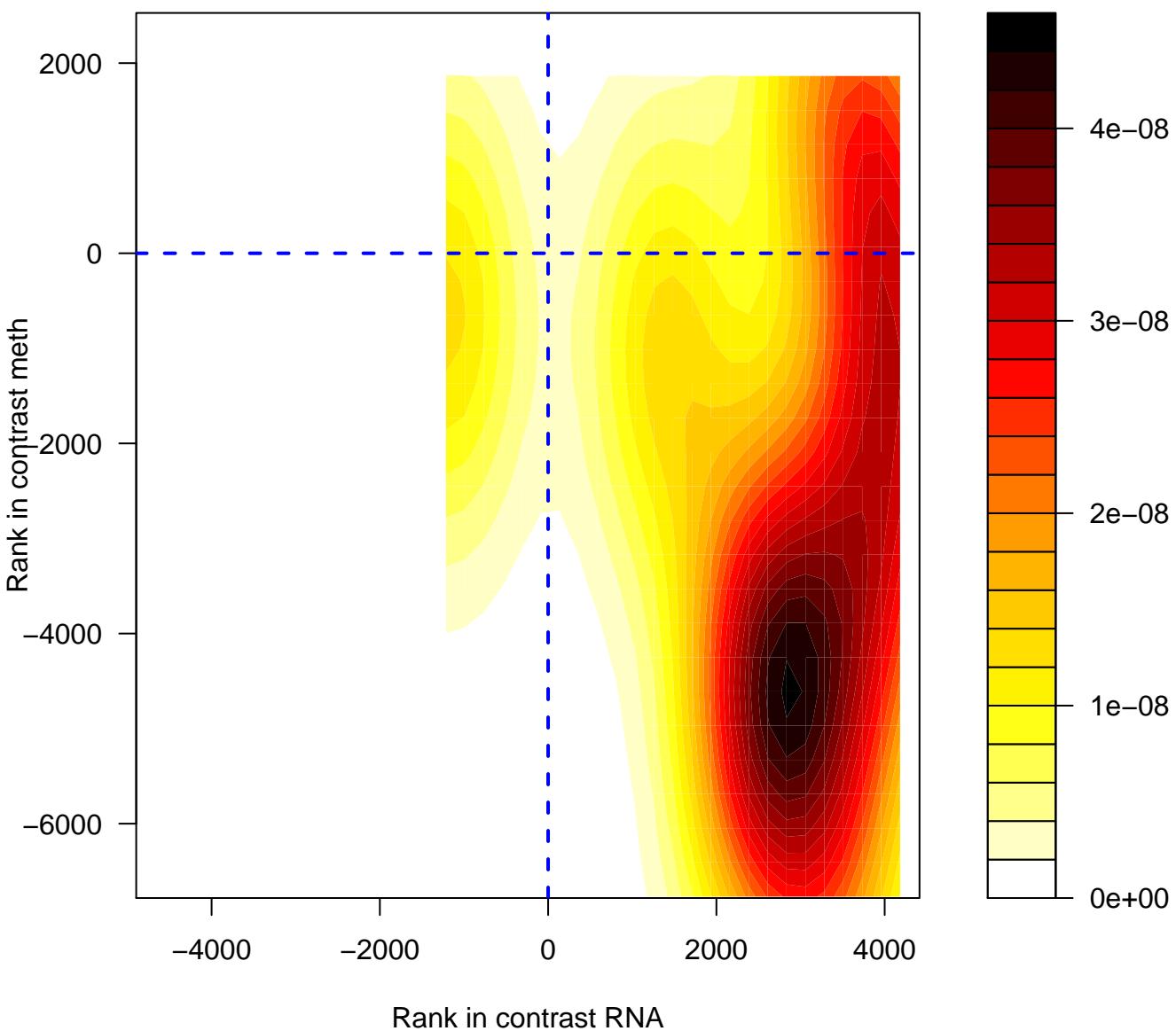
Hyaluronan metabolism



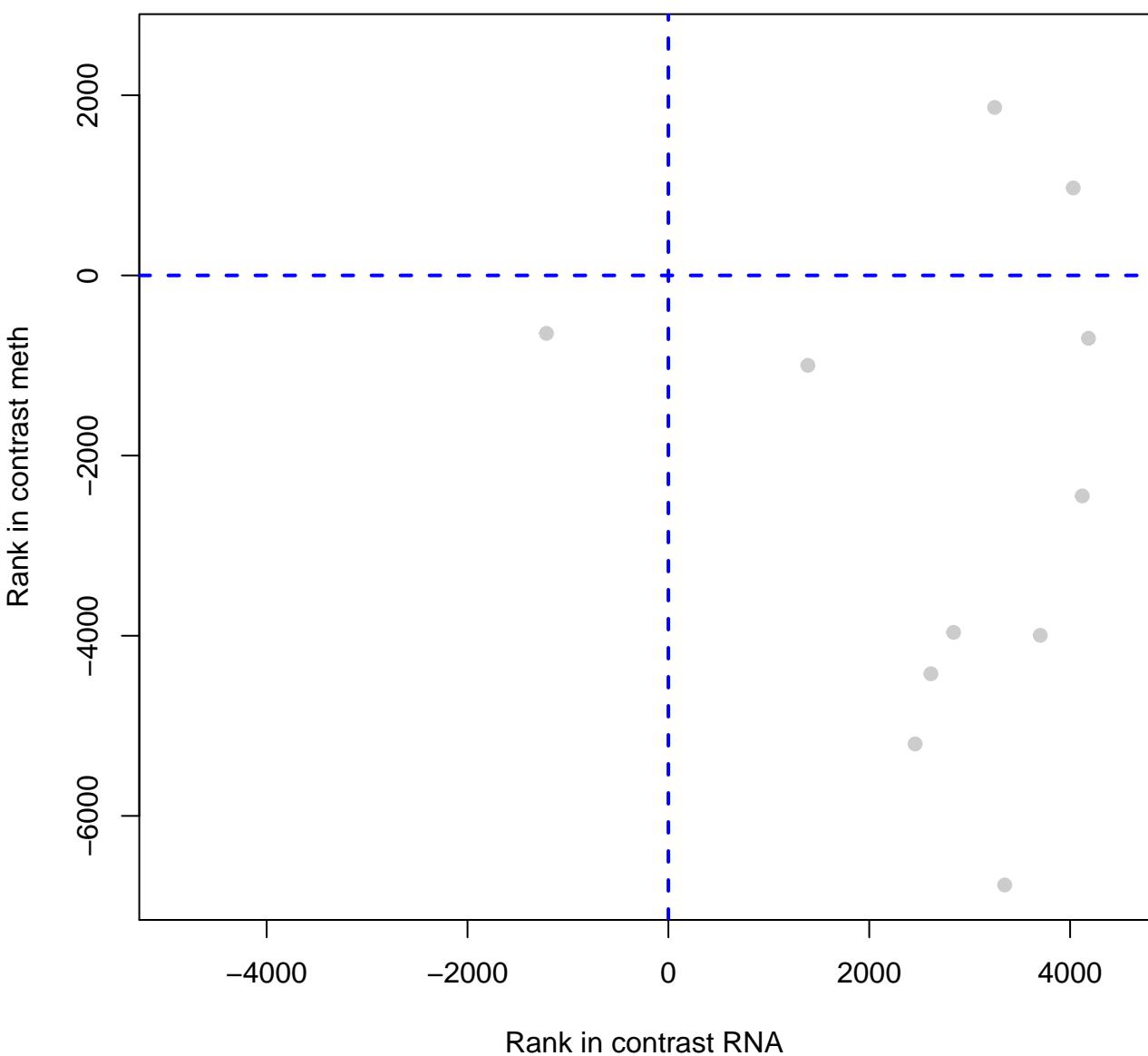
Hyaluronan metabolism



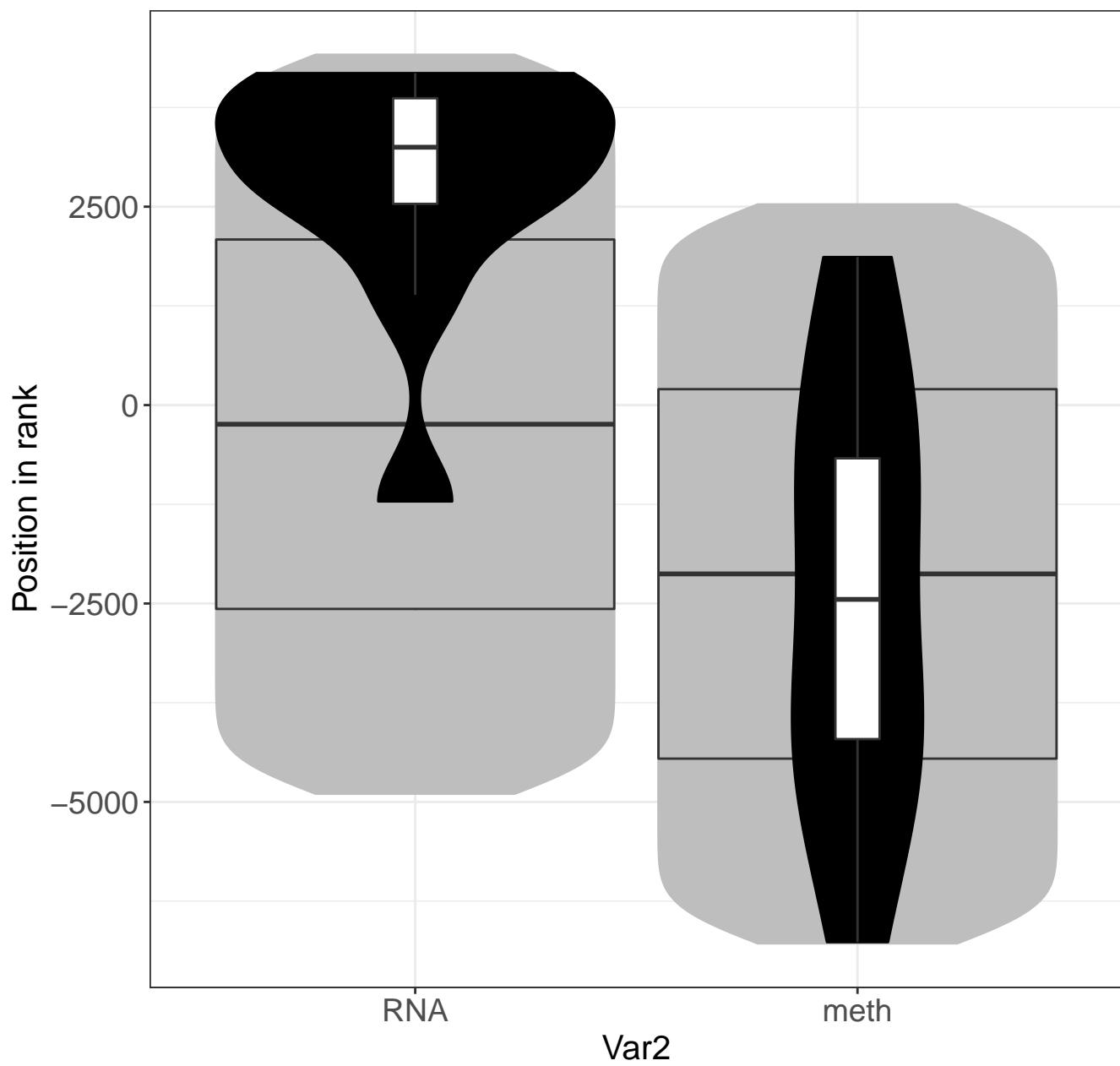
Signal transduction by L1



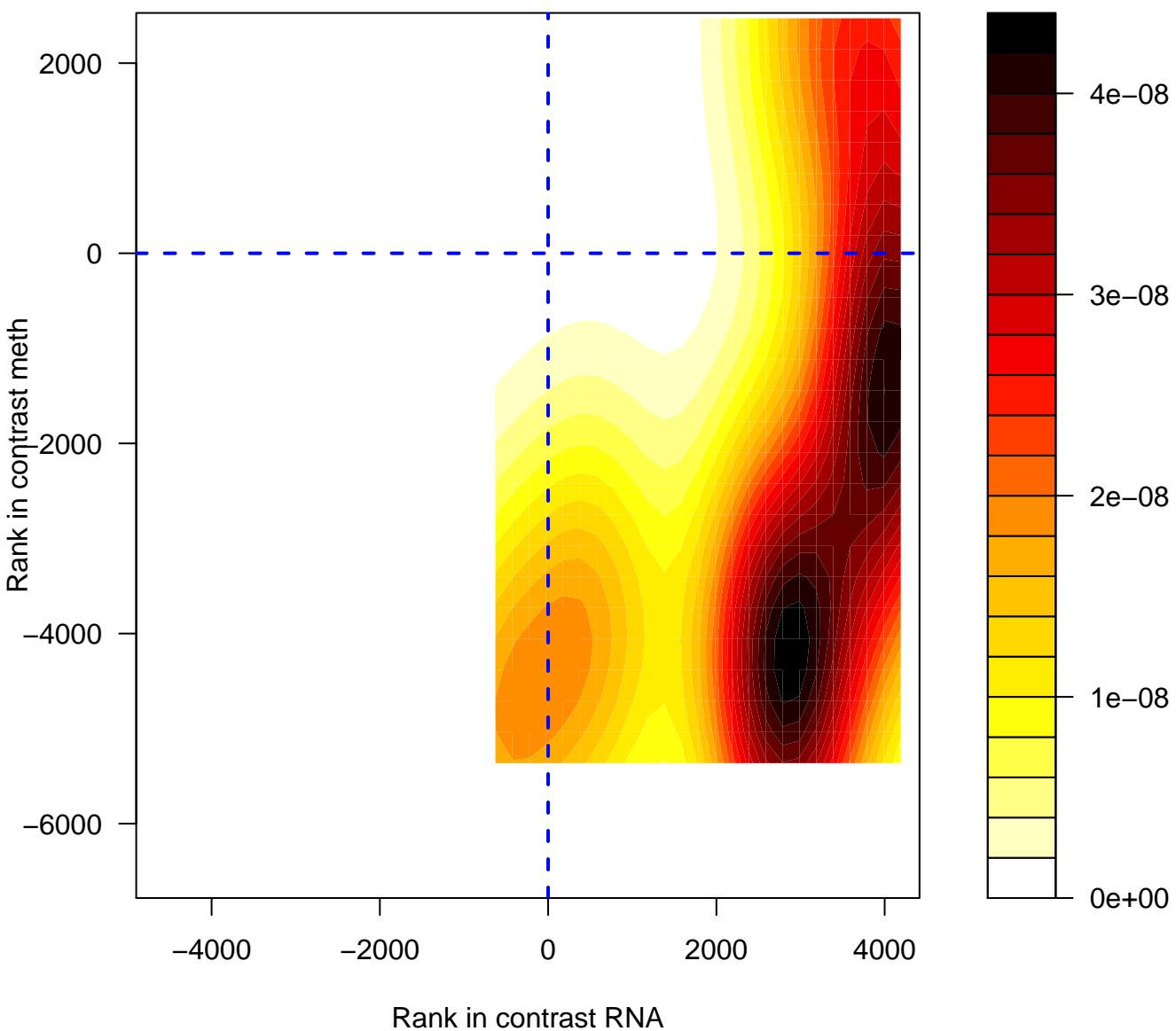
Signal transduction by L1



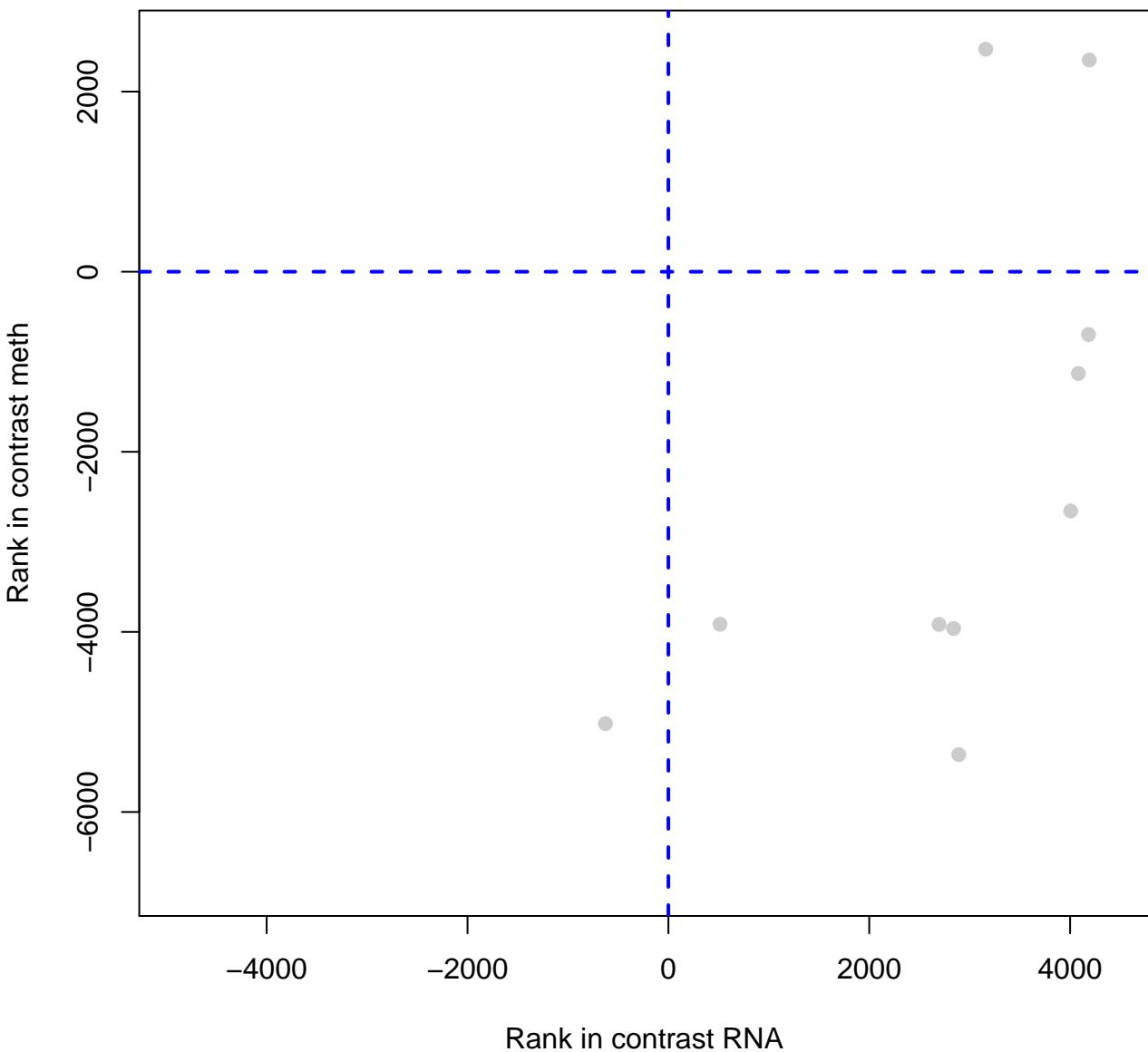
Signal transduction by L1



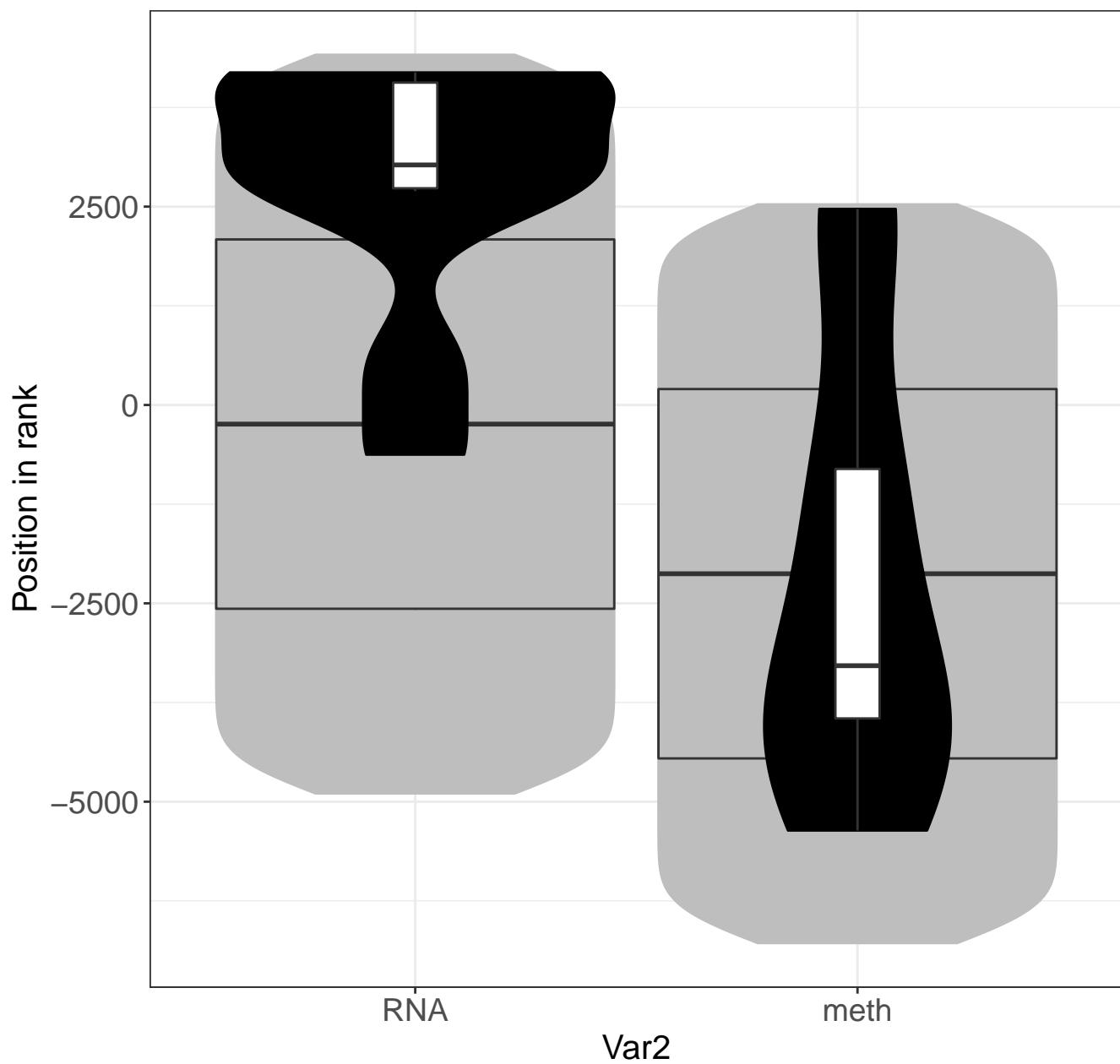
Uptake and function of anthrax toxins



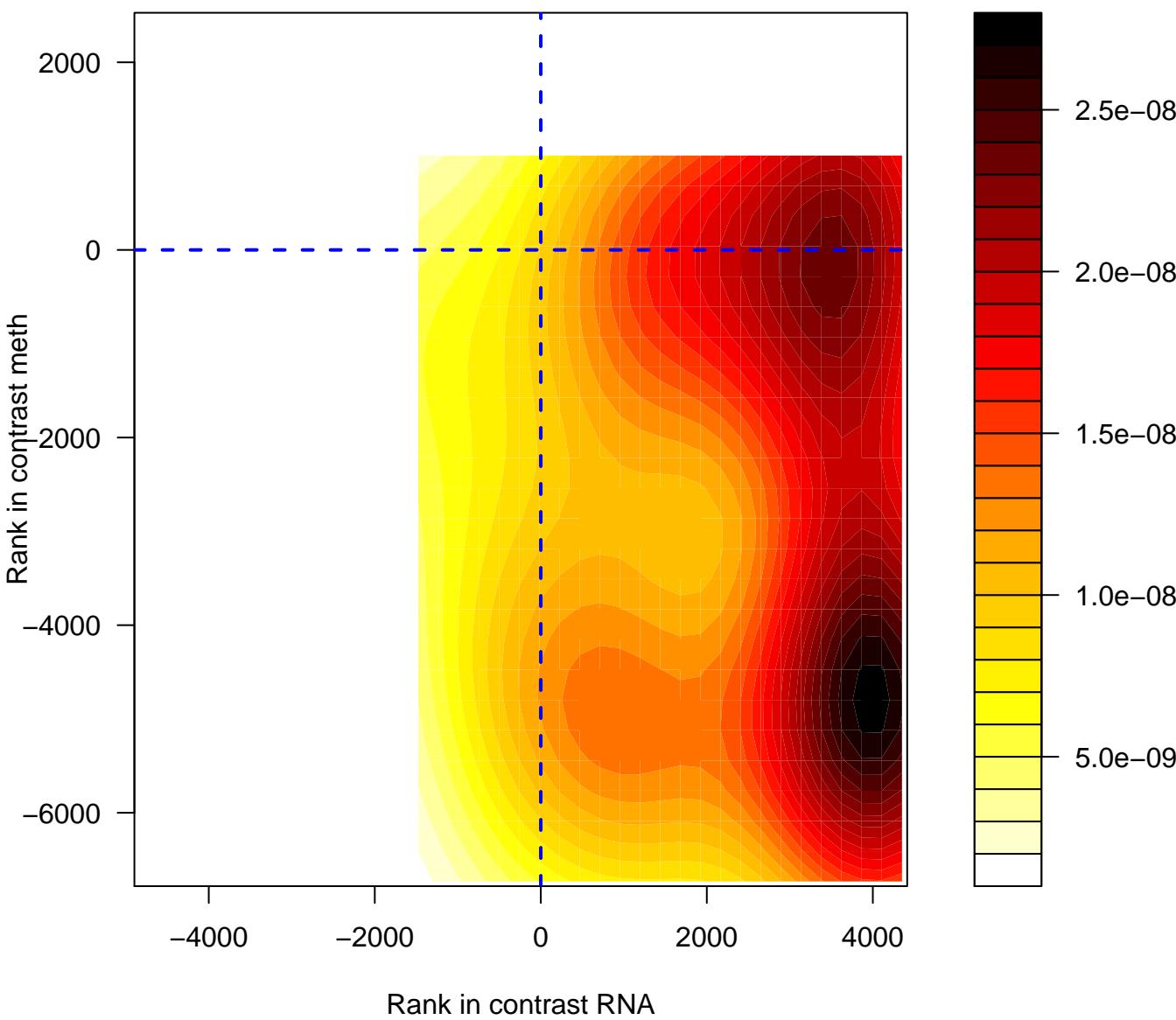
Uptake and function of anthrax toxins



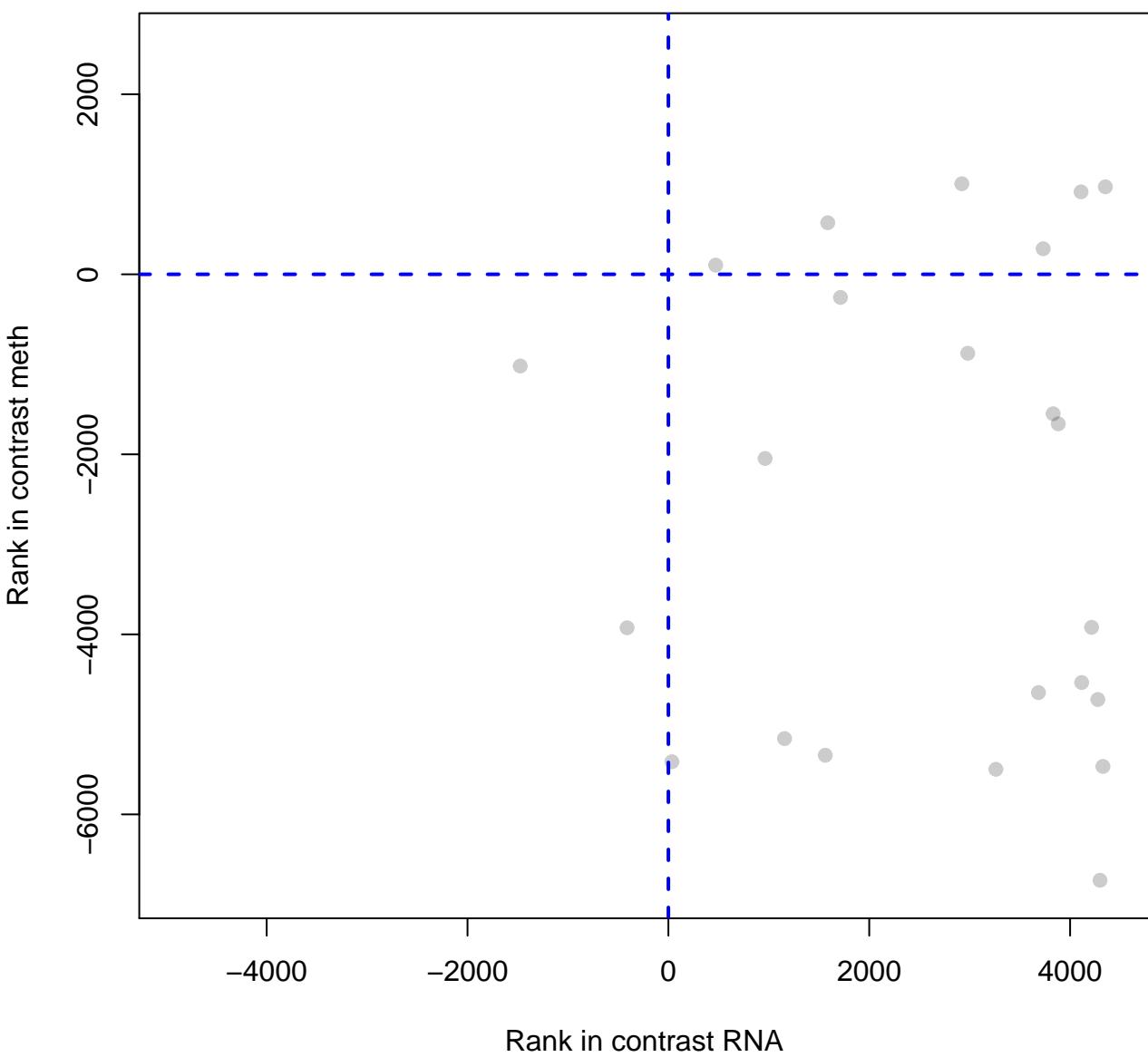
Uptake and function of anthrax toxins



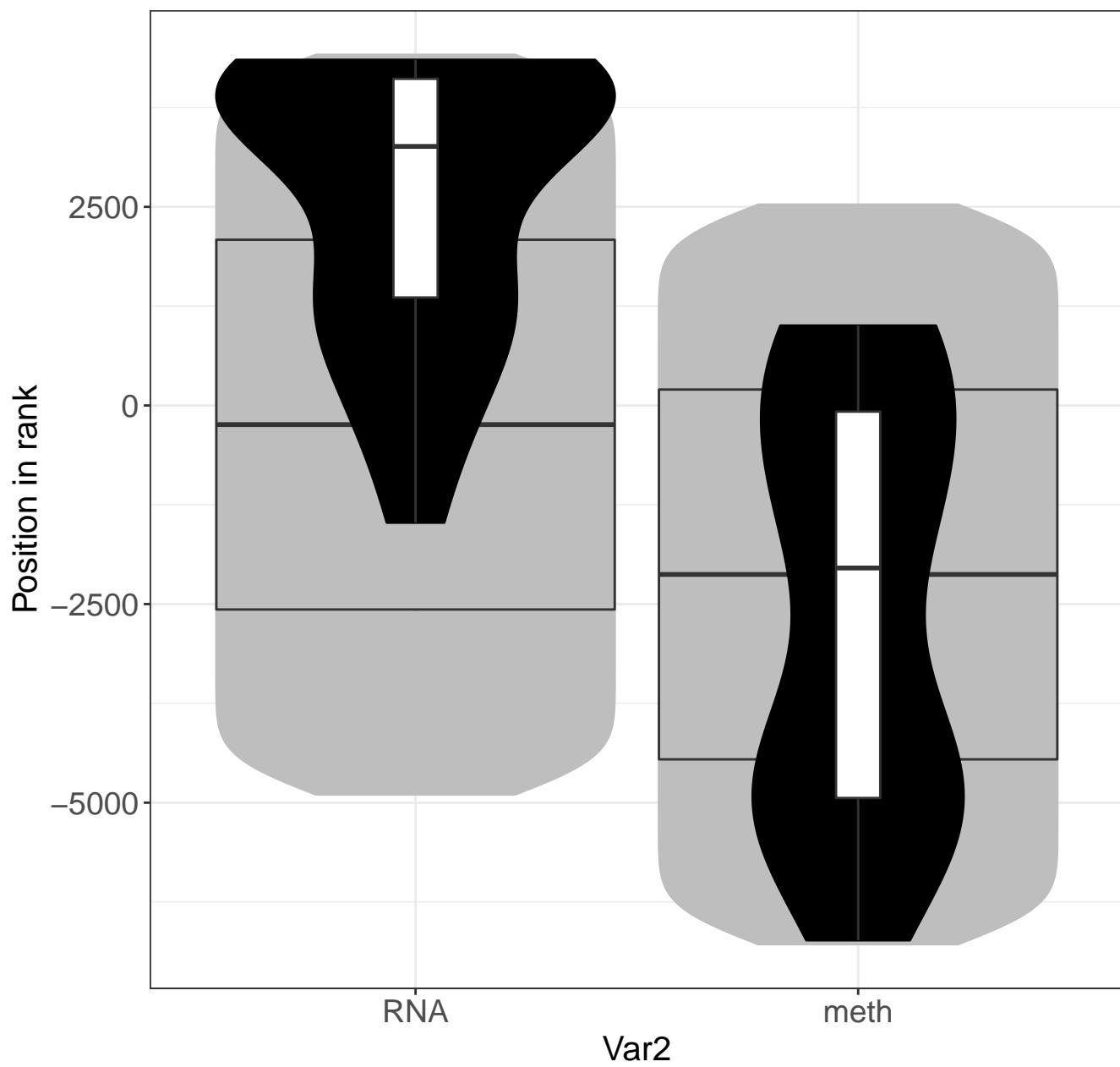
Detoxification of Reactive Oxygen Species



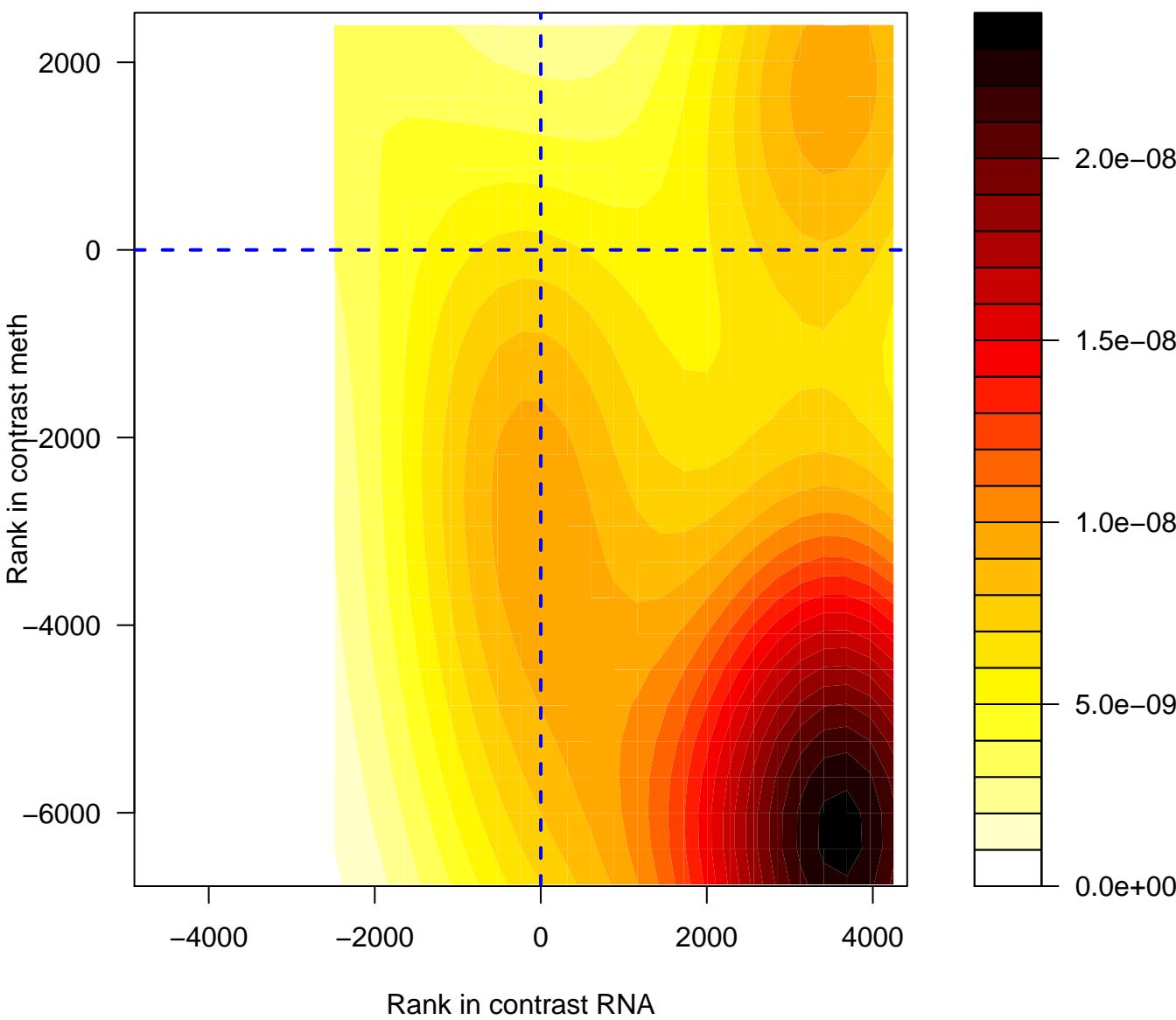
Detoxification of Reactive Oxygen Species



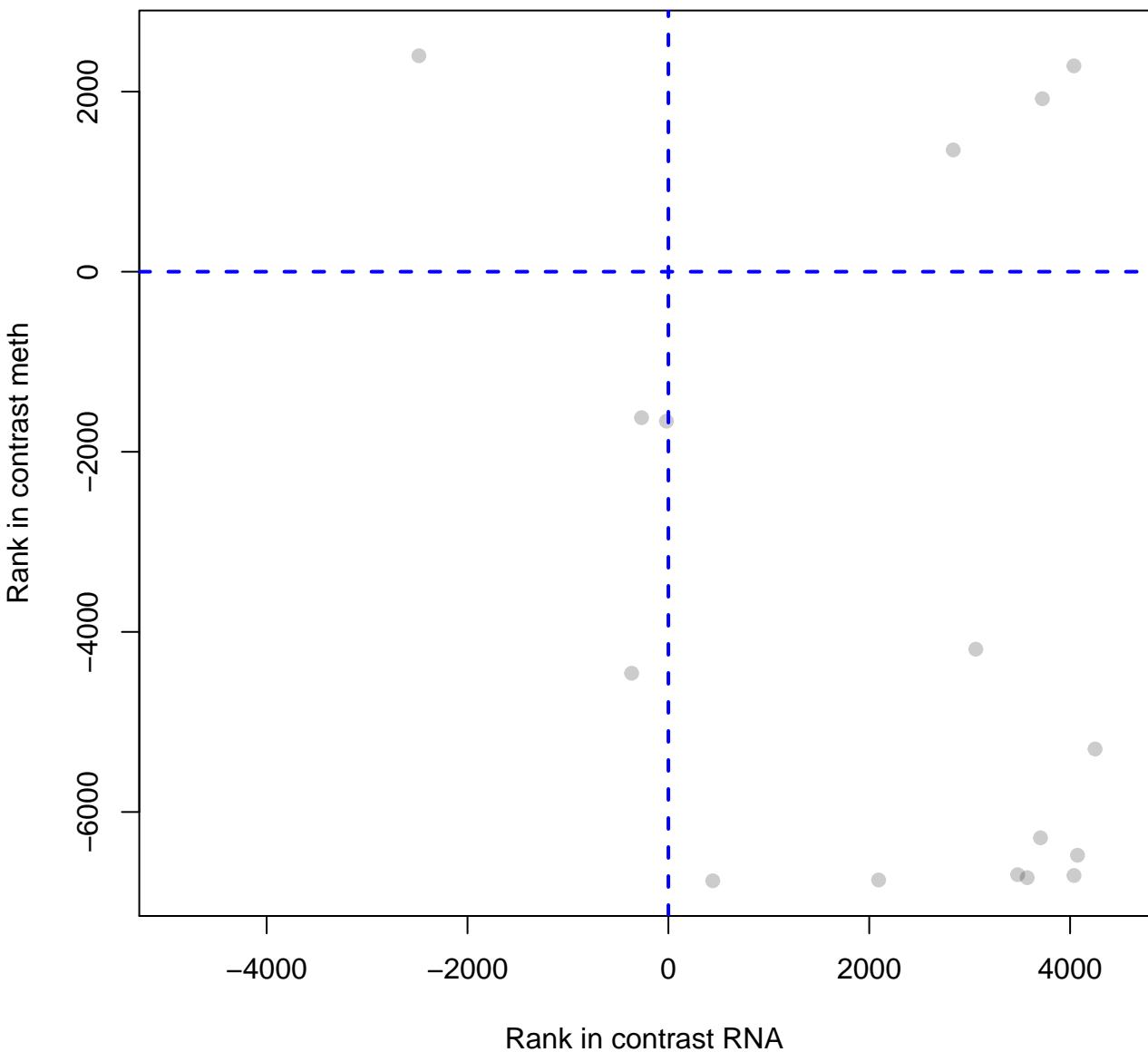
Detoxification of Reactive Oxygen Species



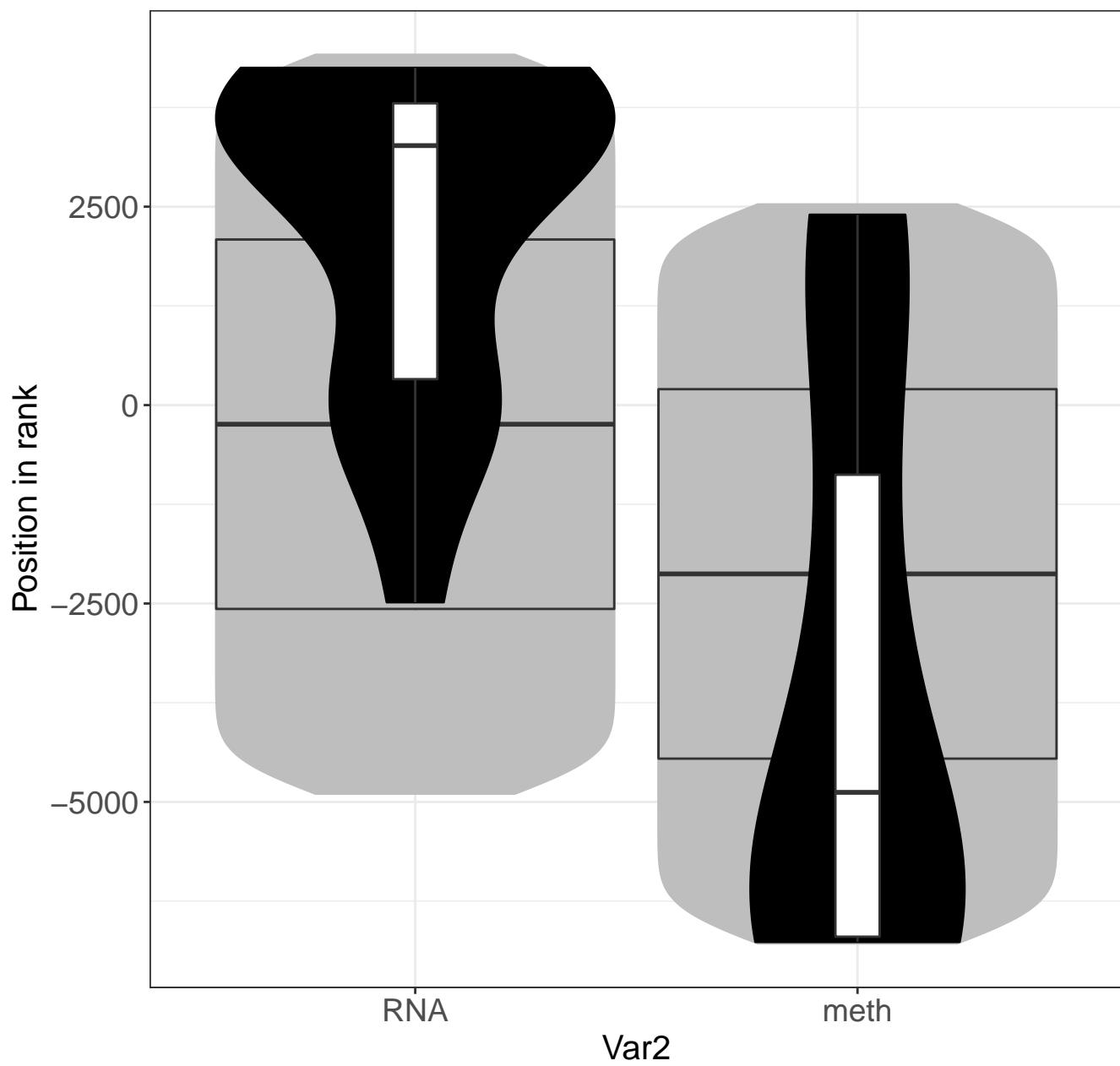
Peptide ligand–binding receptors



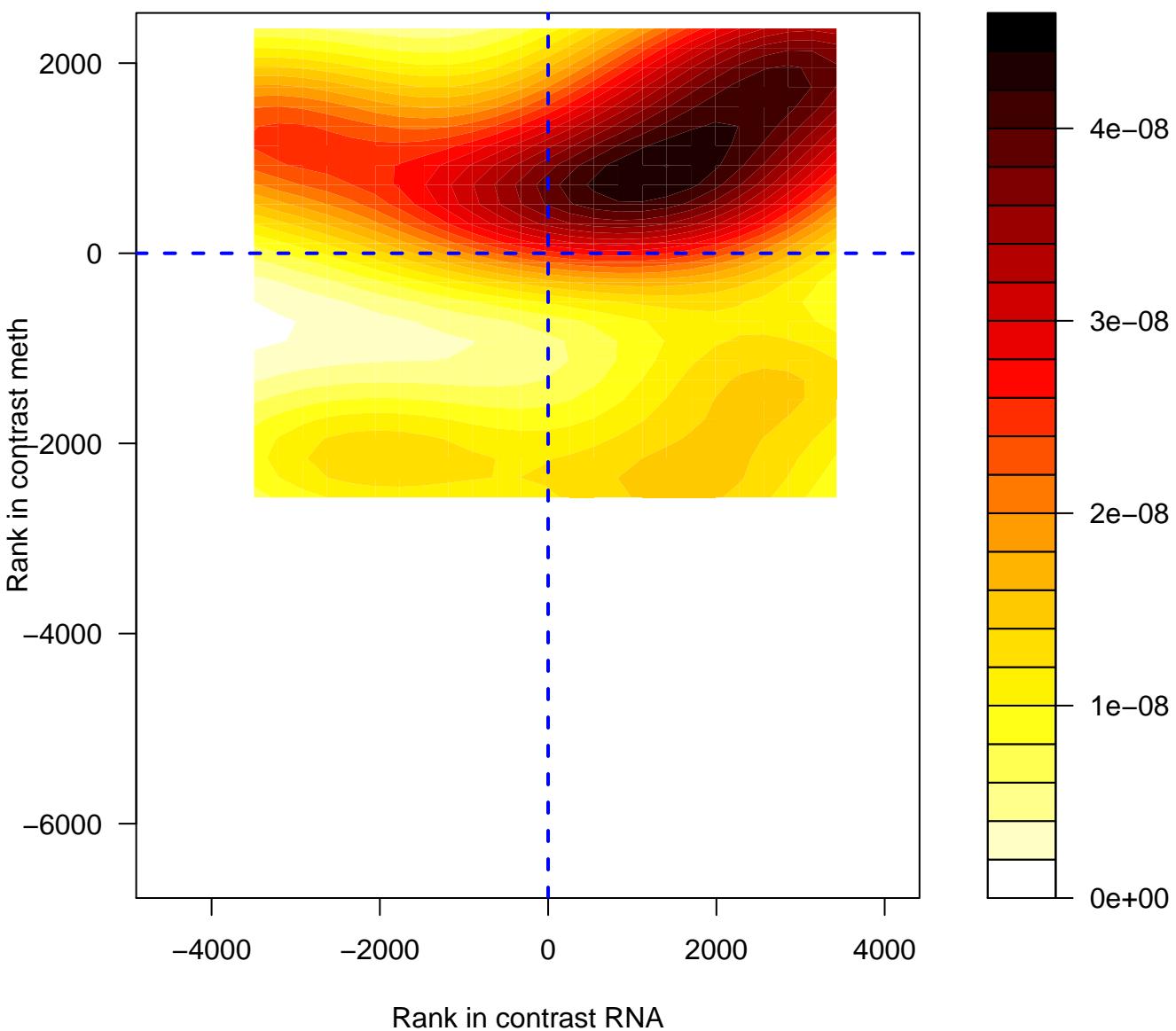
Peptide ligand–binding receptors



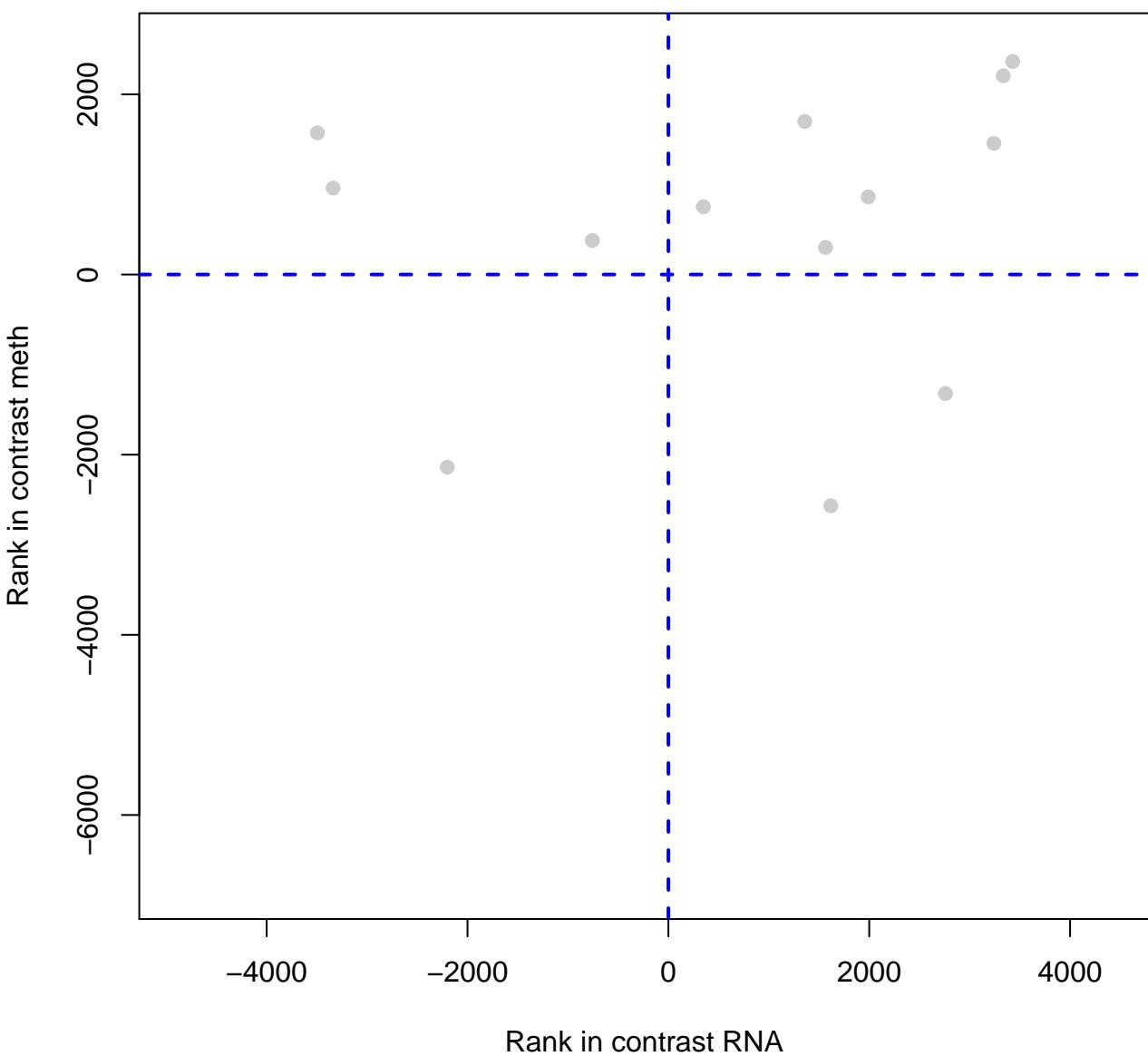
Peptide ligand–binding receptors



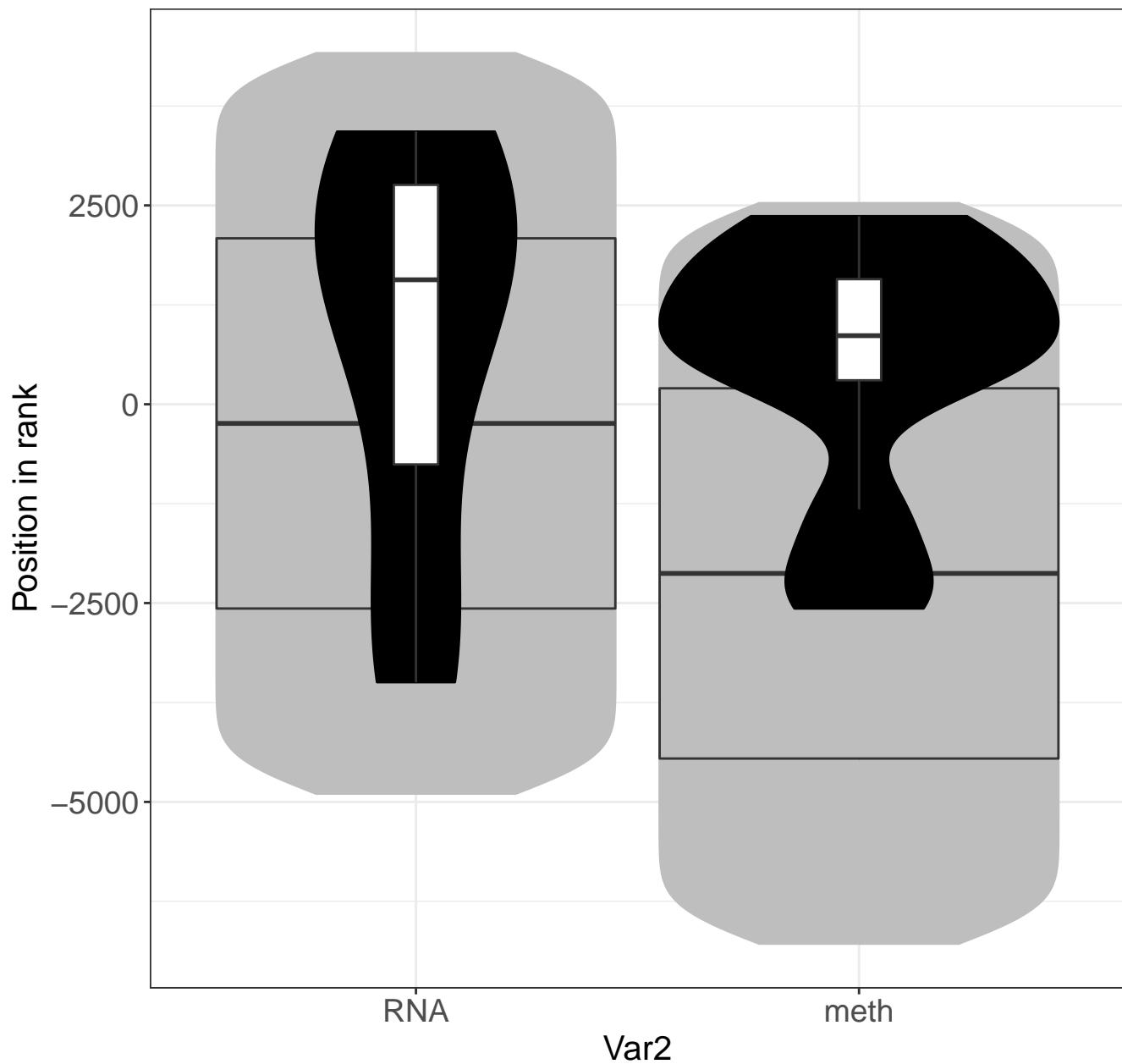
Negative regulation of FLT3



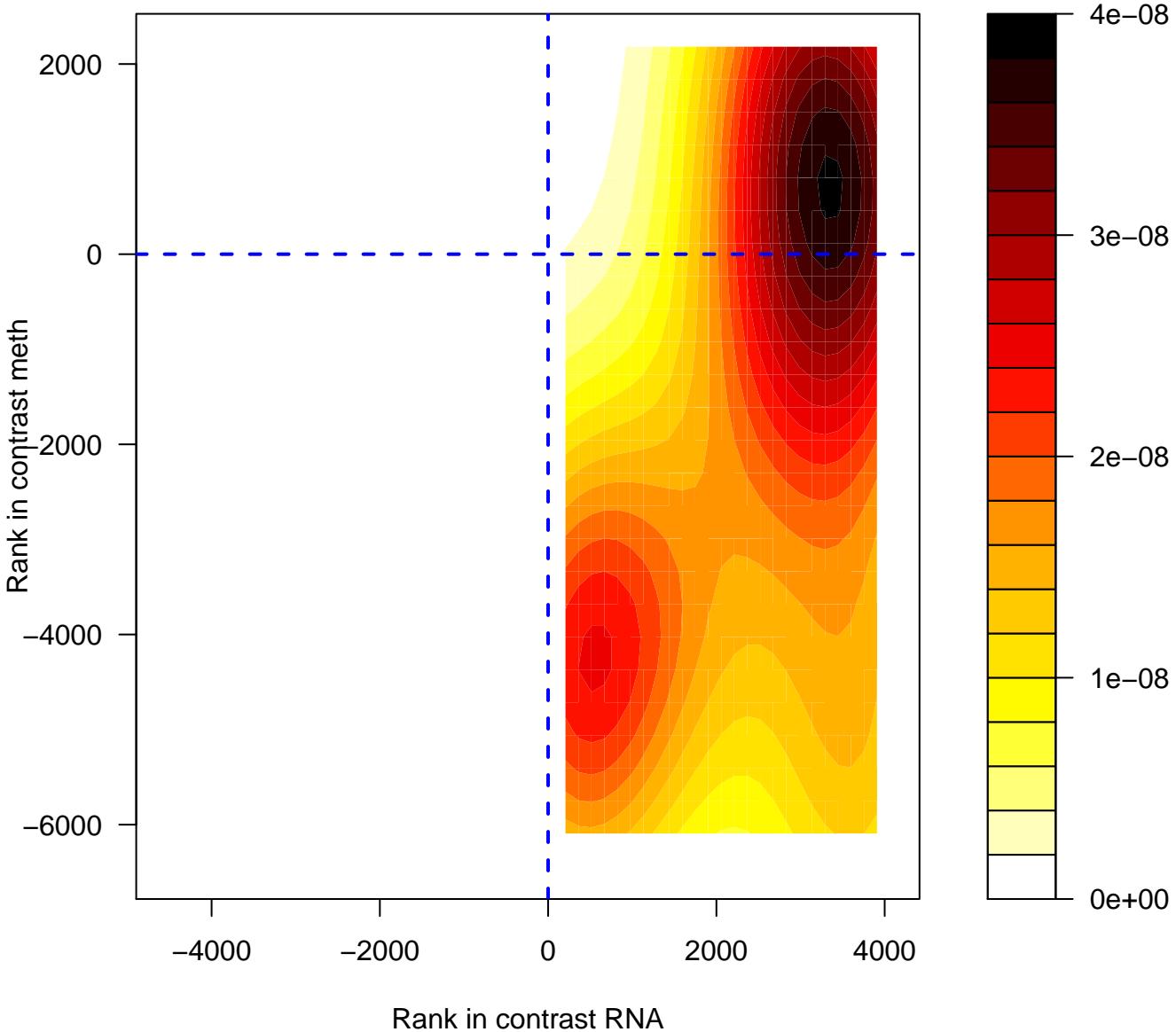
Negative regulation of FLT3



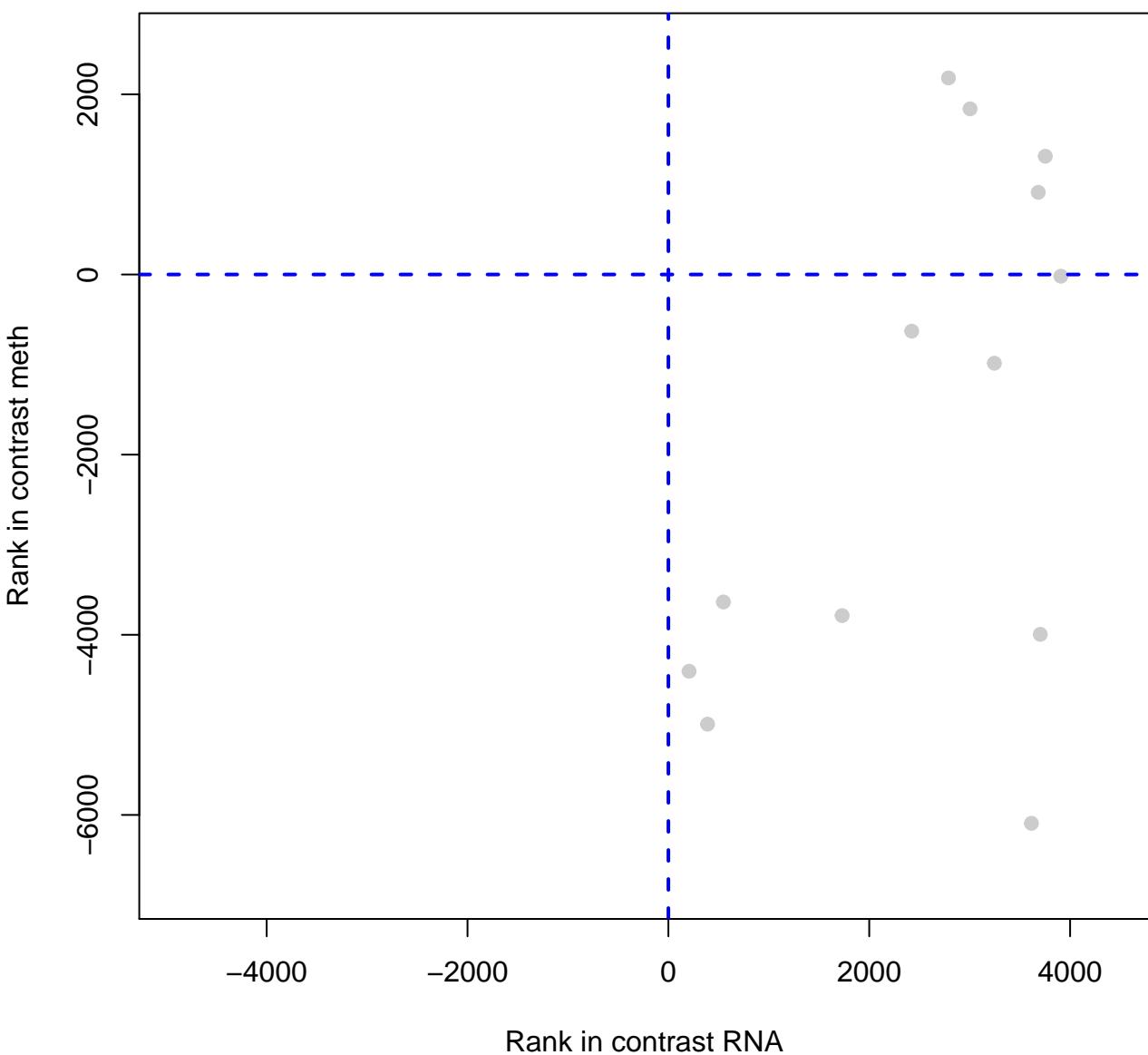
Negative regulation of FLT3



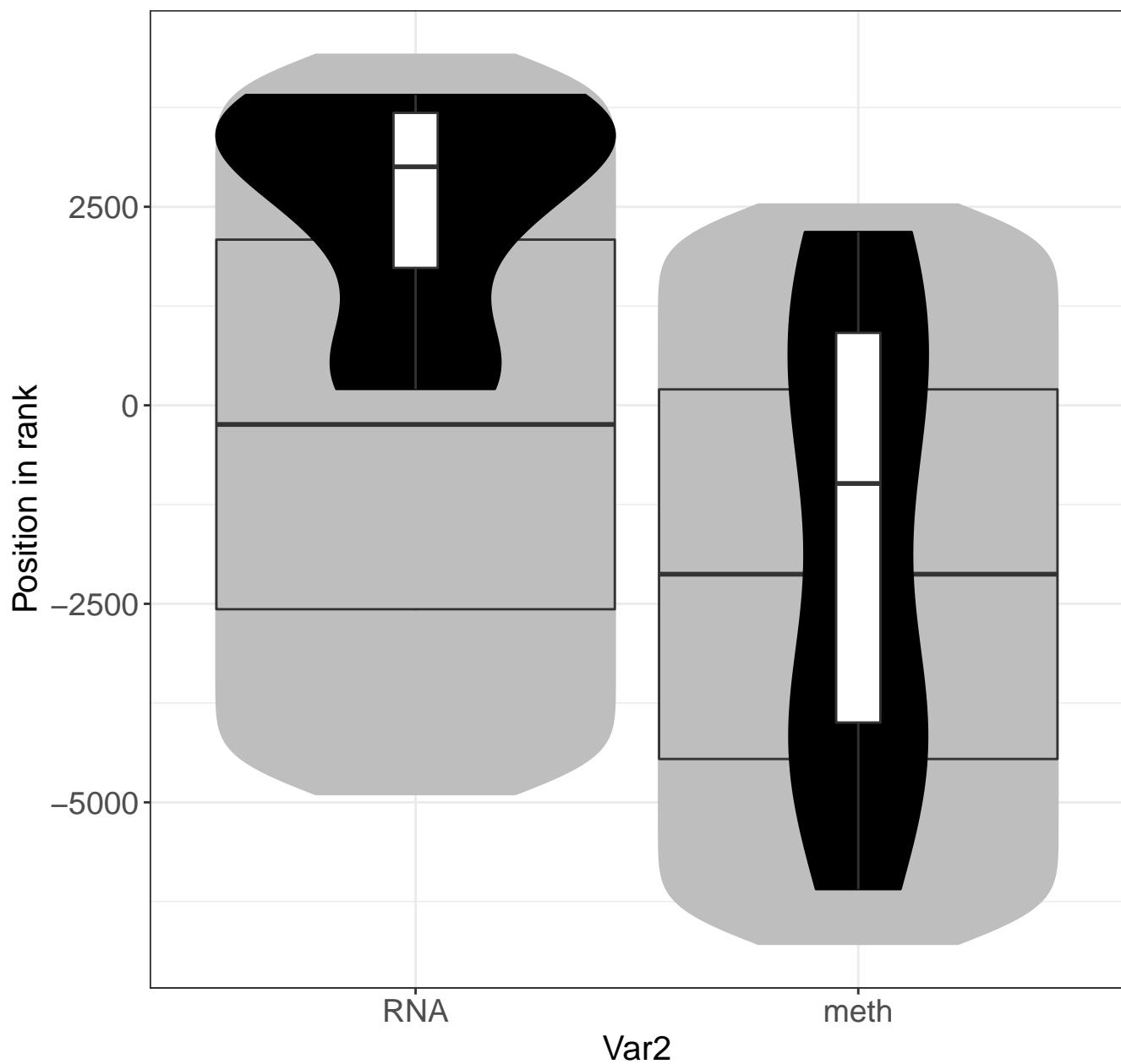
RHO GTPases Activate ROCKs



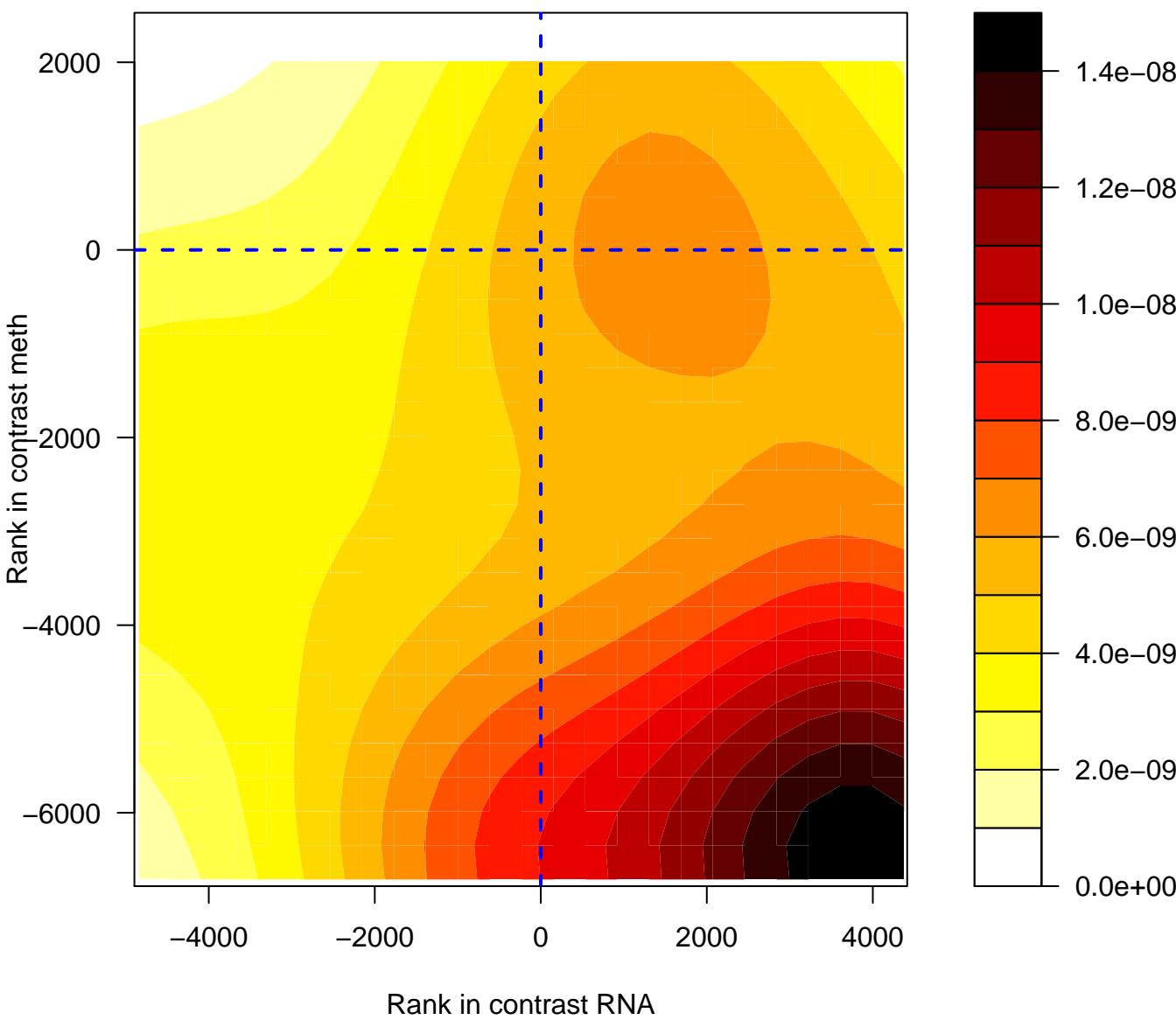
RHO GTPases Activate ROCKs



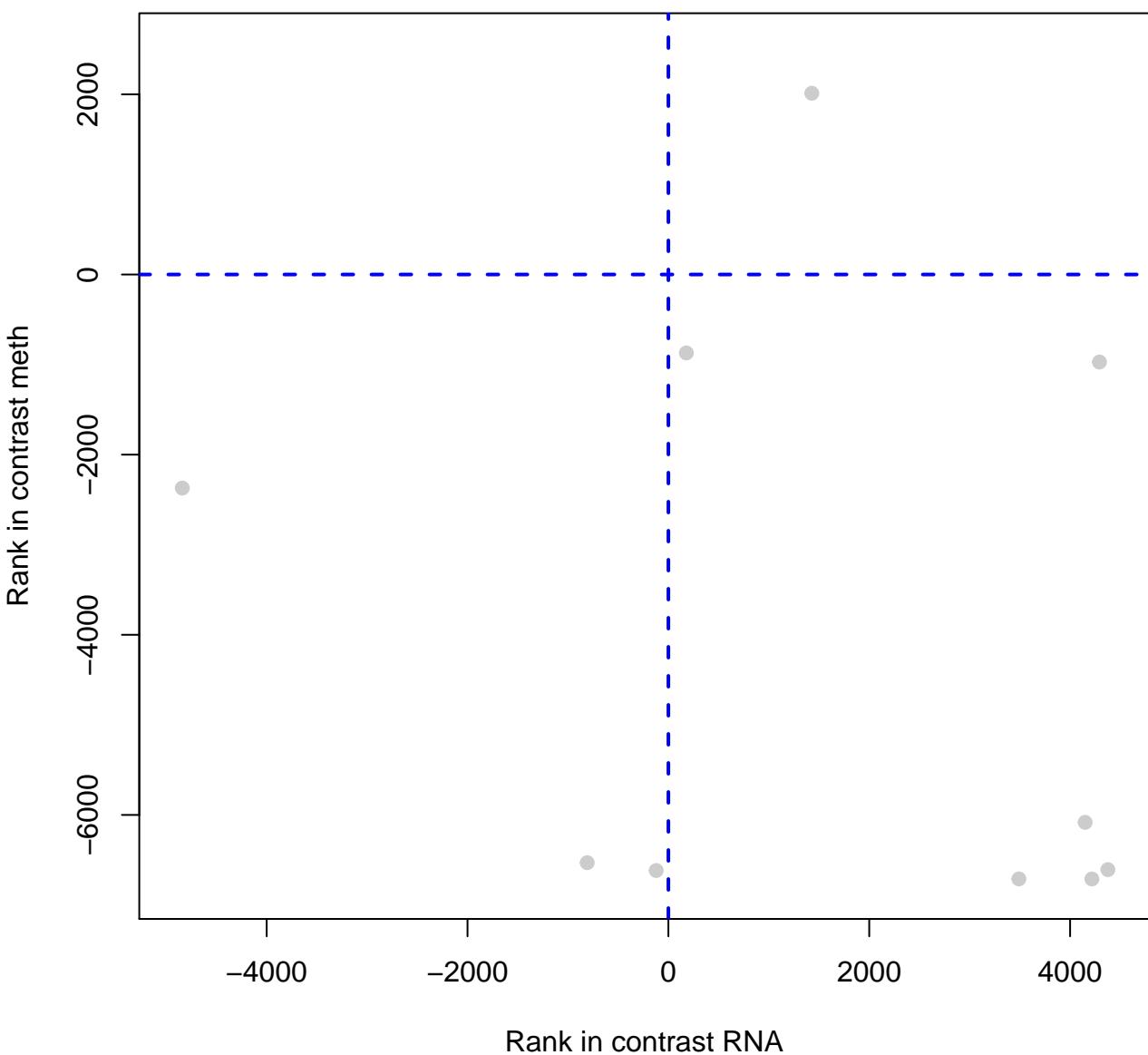
RHO GTPases Activate ROCKs



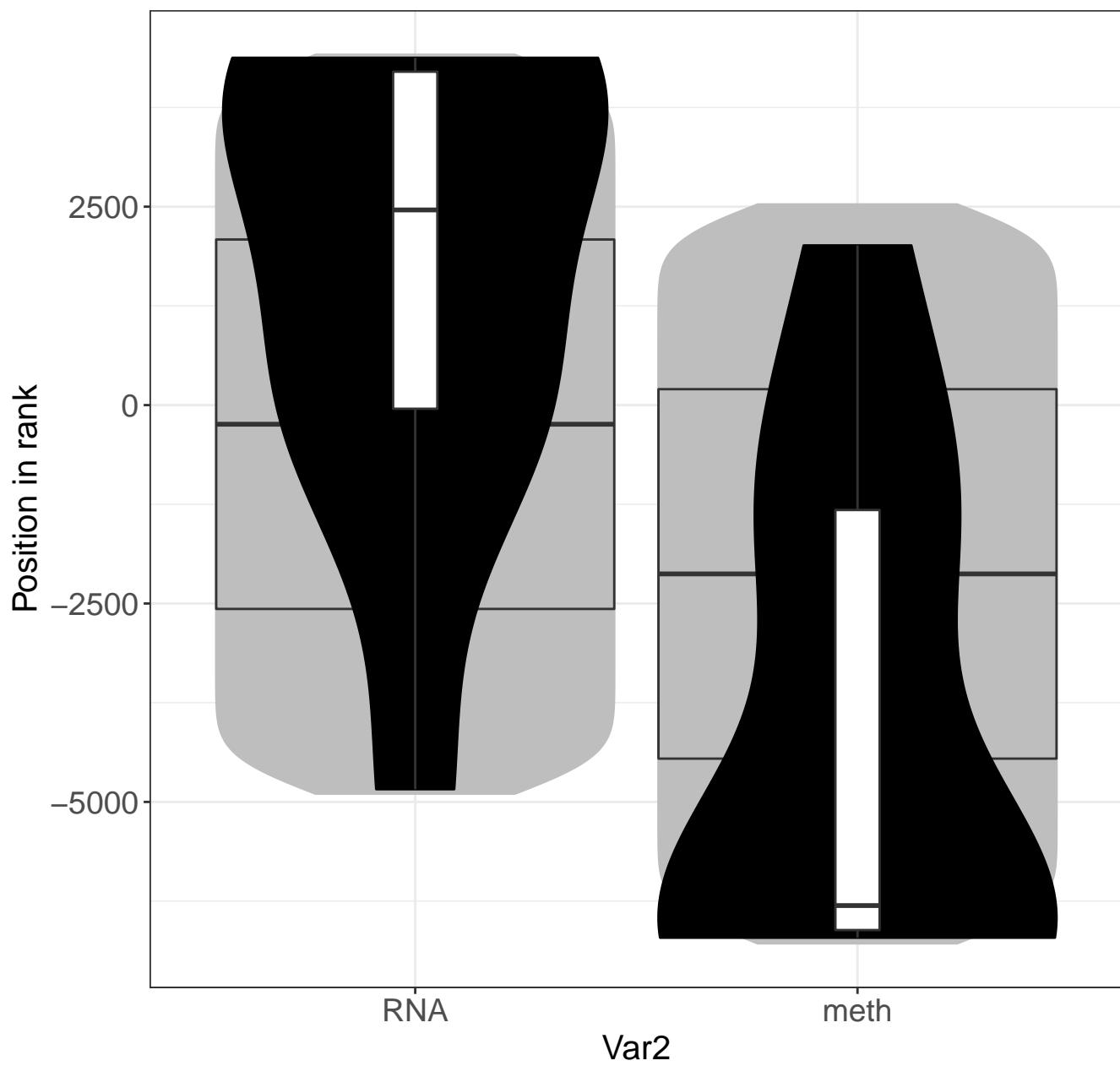
Synthesis of Leukotrienes (LT) and Exoxins (EX)



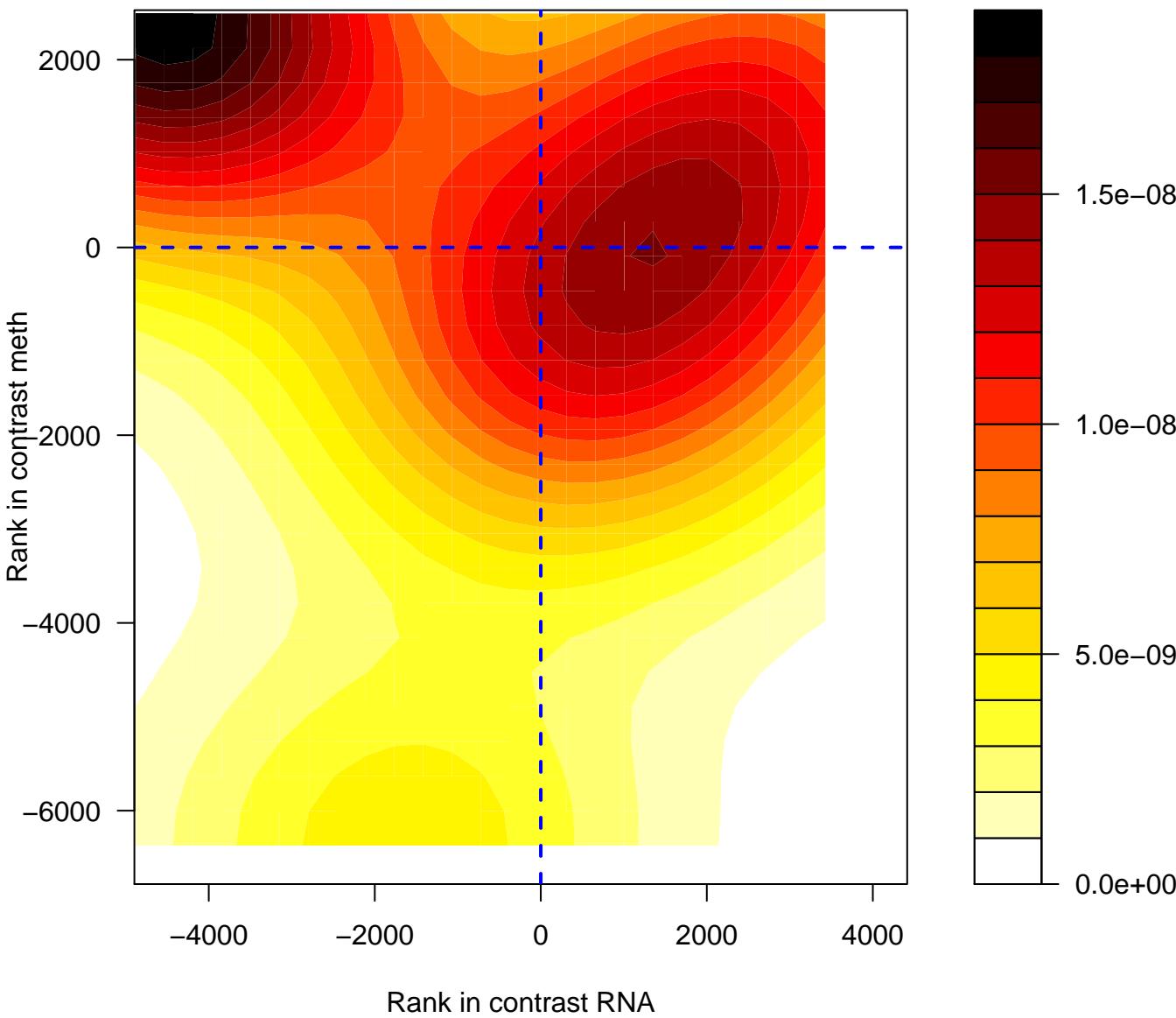
Synthesis of Leukotrienes (LT) and Eoxins (EX)



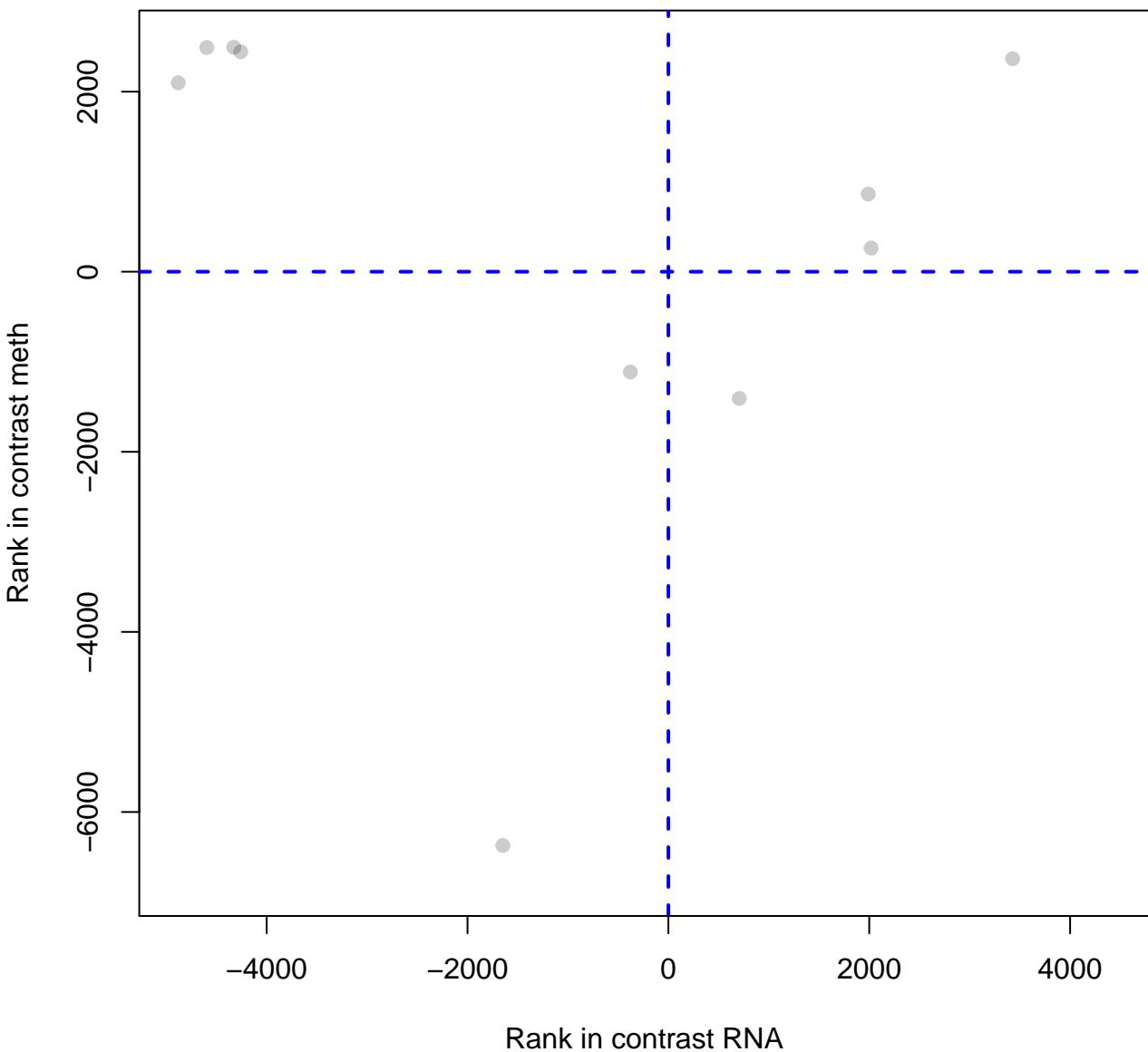
Synthesis of Leukotrienes (LT) and Eoxins (EX)



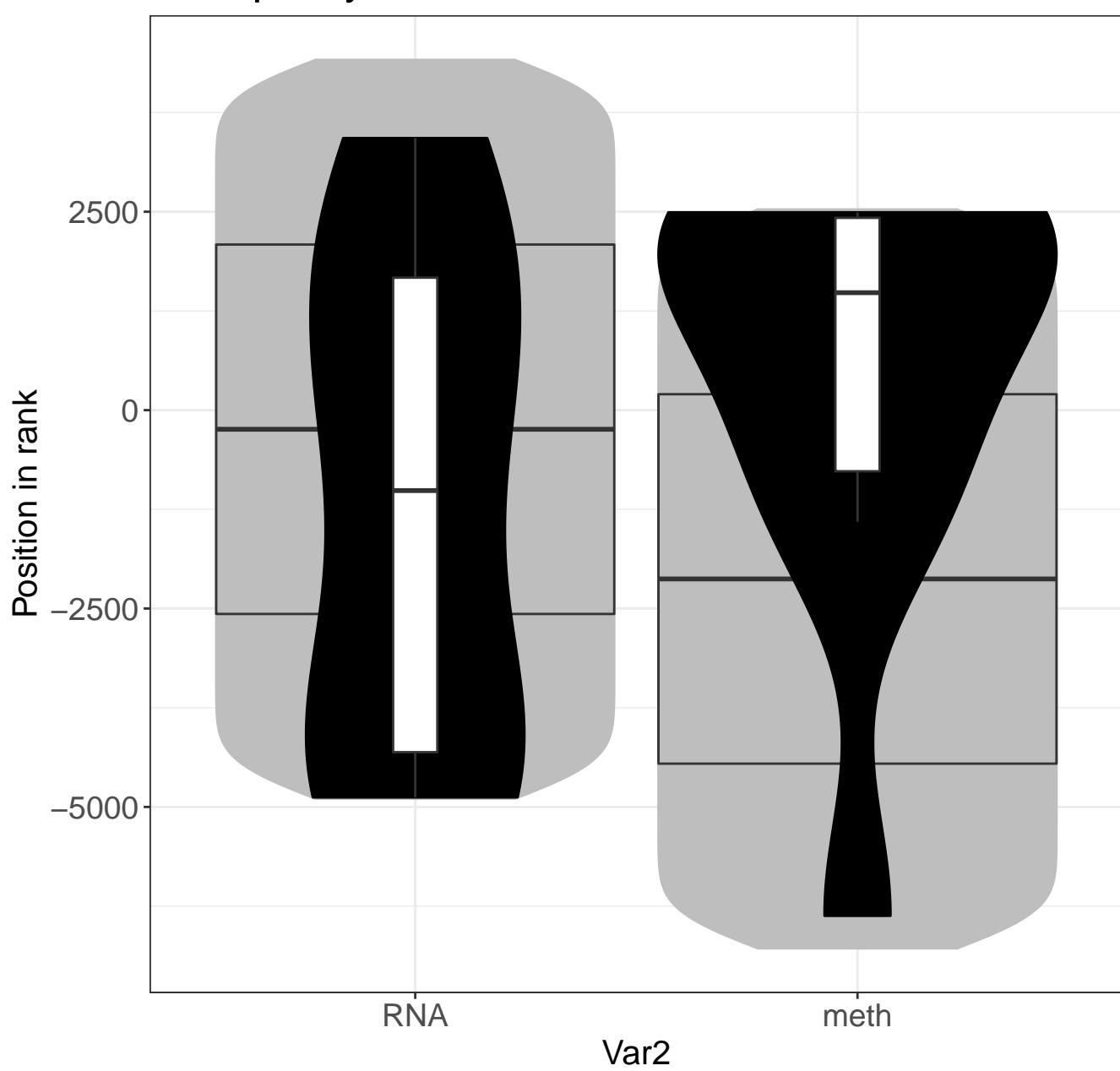
Phosphorylation of CD3 and TCR zeta chains



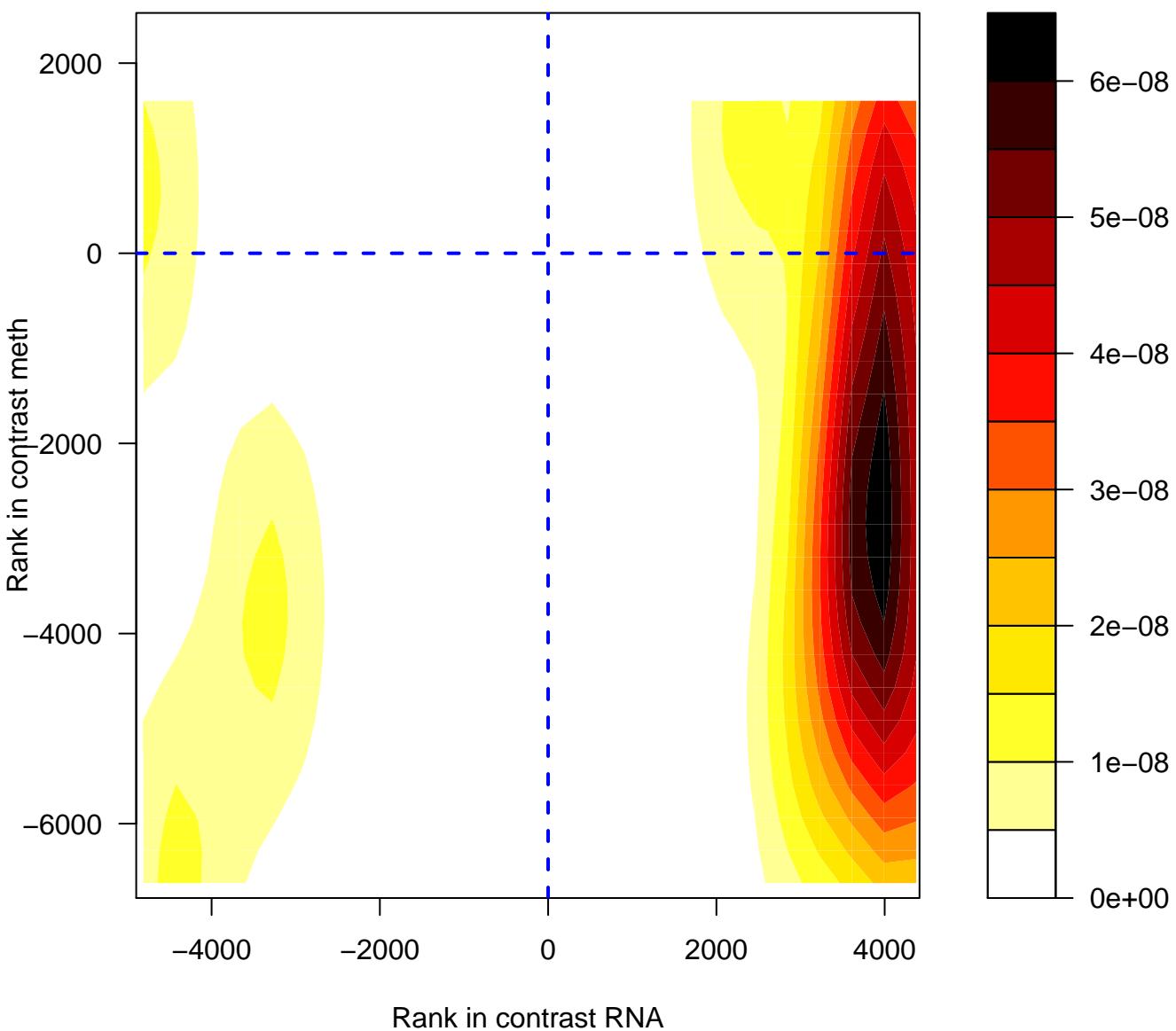
Phosphorylation of CD3 and TCR zeta chains



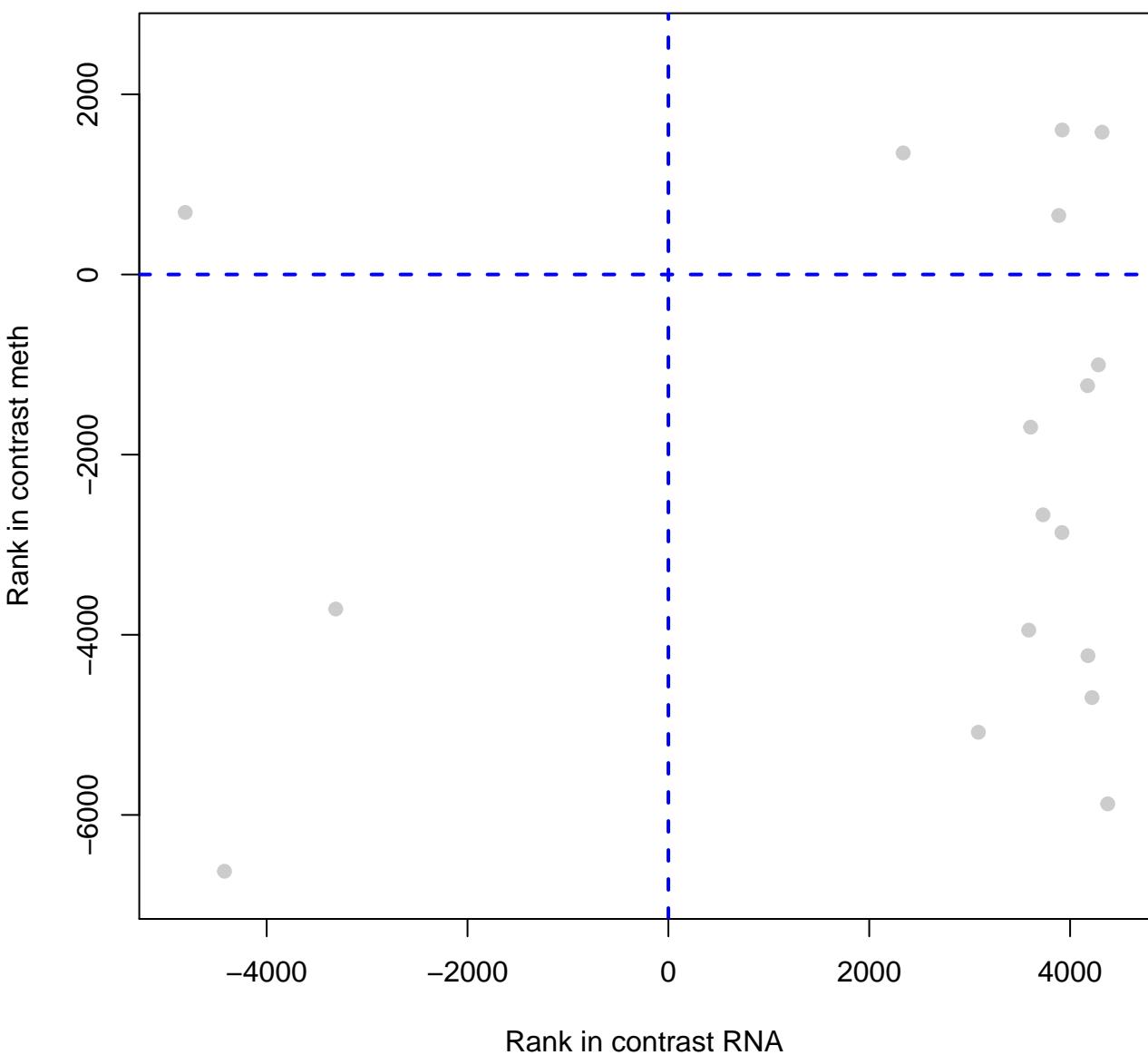
Phosphorylation of CD3 and TCR zeta chains



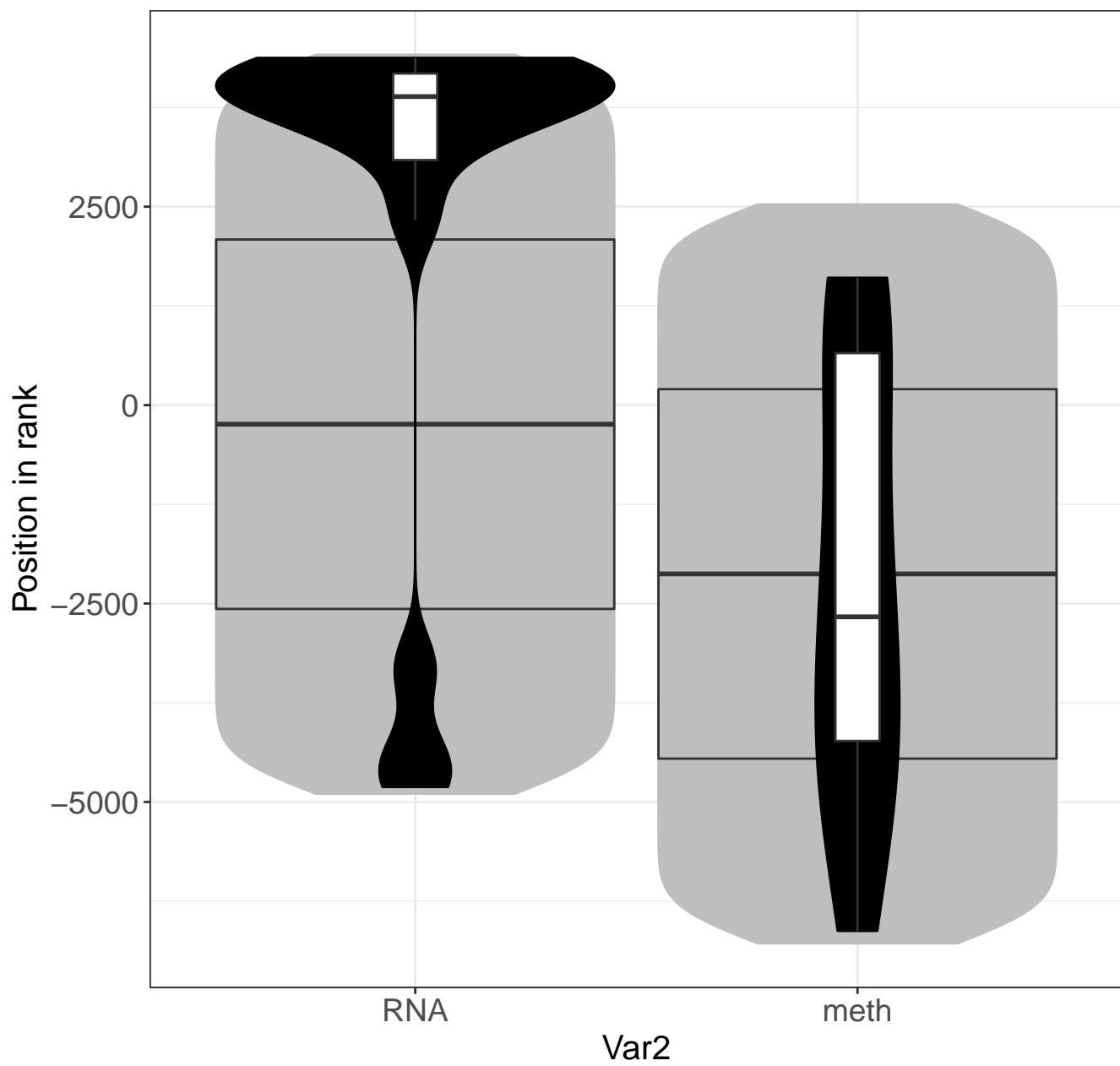
Insulin receptor recycling



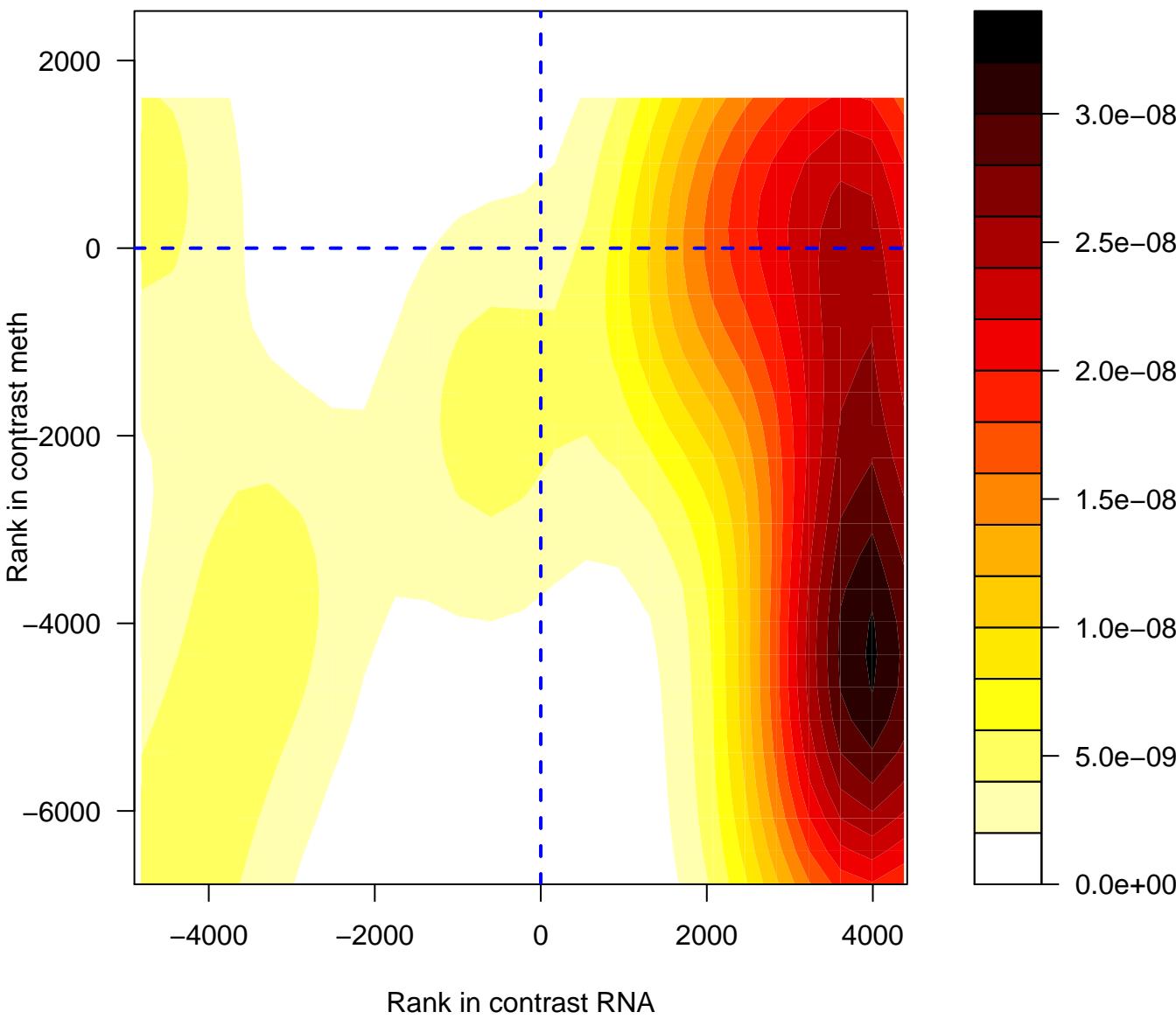
Insulin receptor recycling



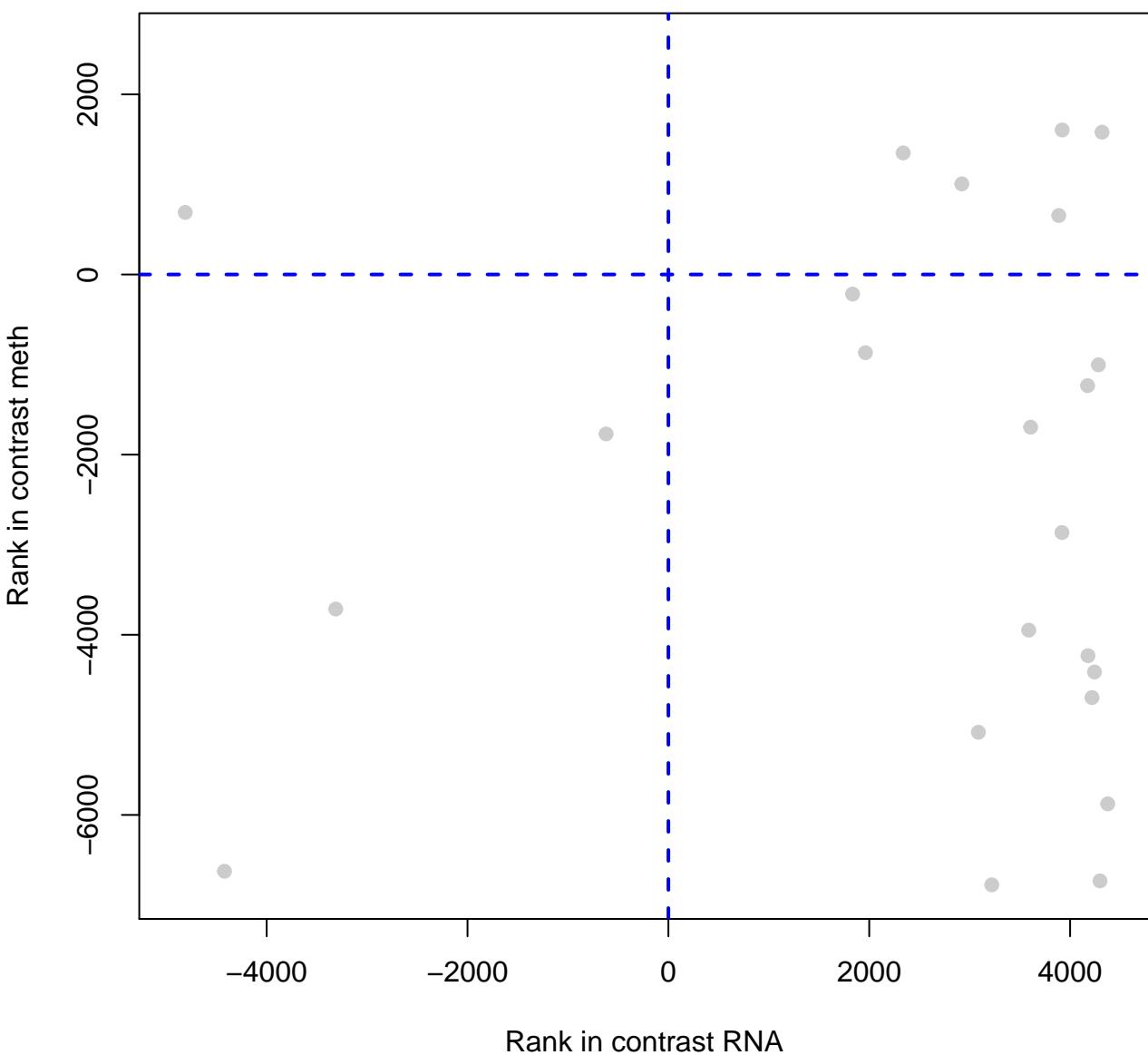
Insulin receptor recycling



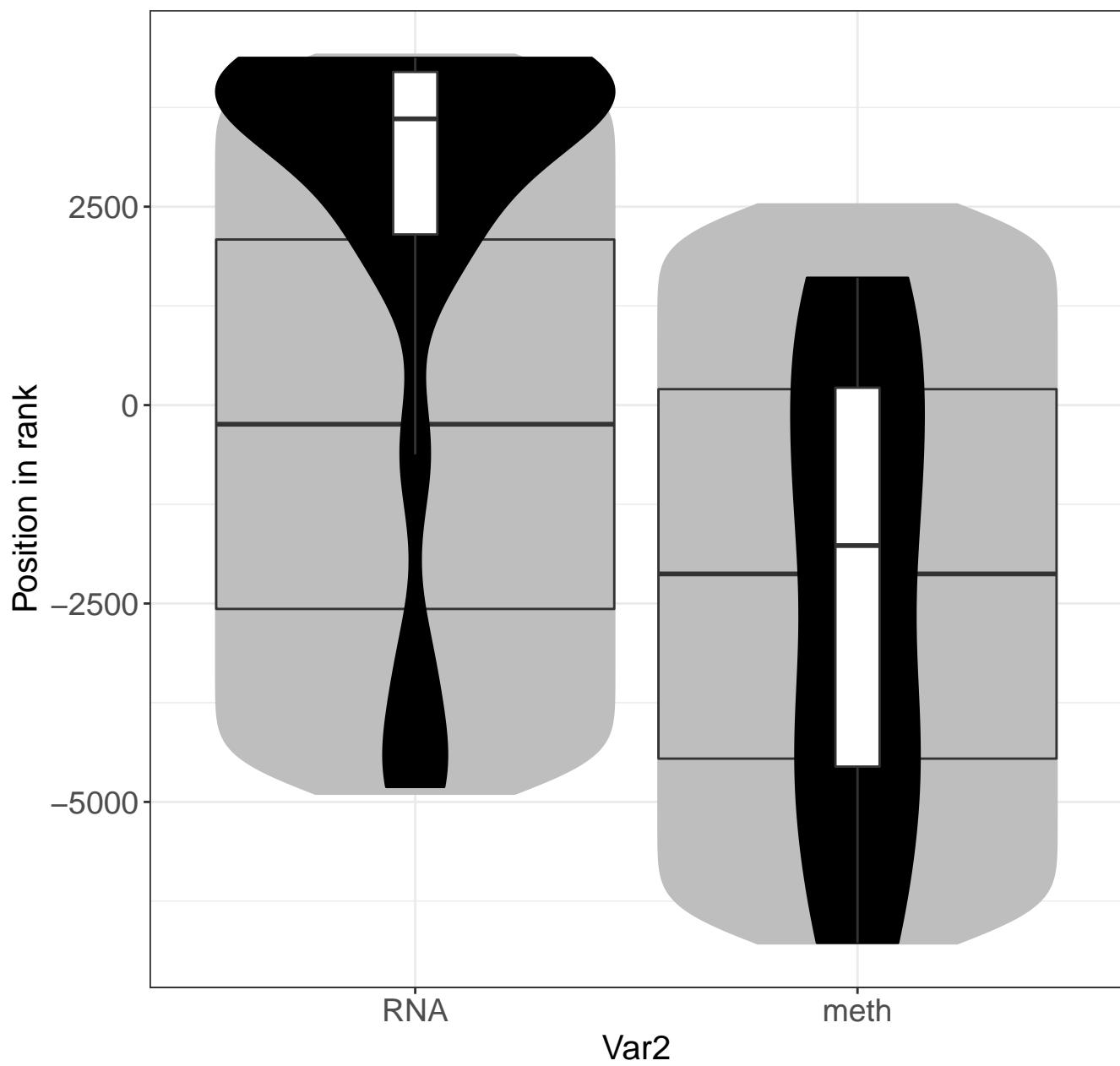
ROS and RNS production in phagocytes



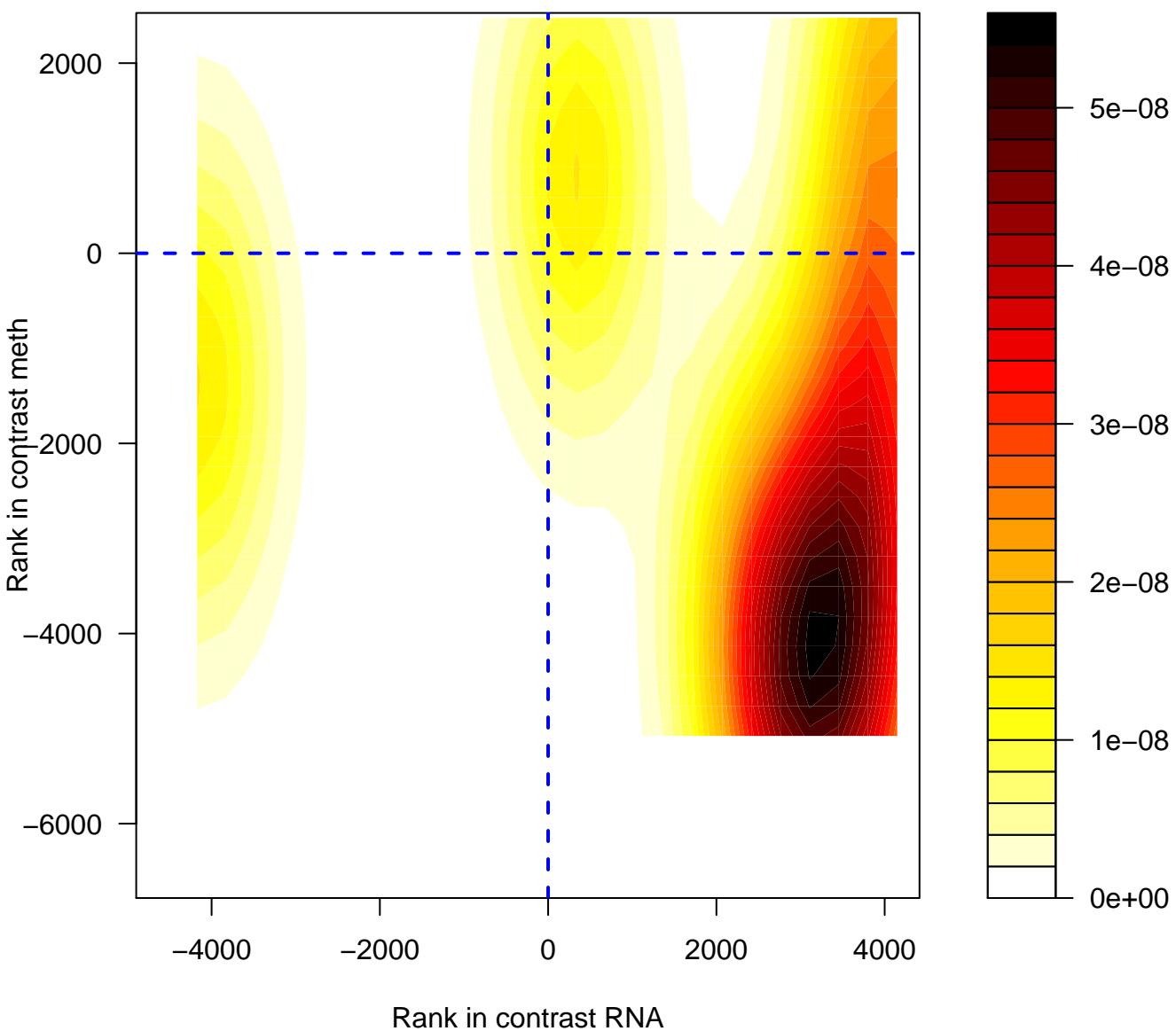
ROS and RNS production in phagocytes



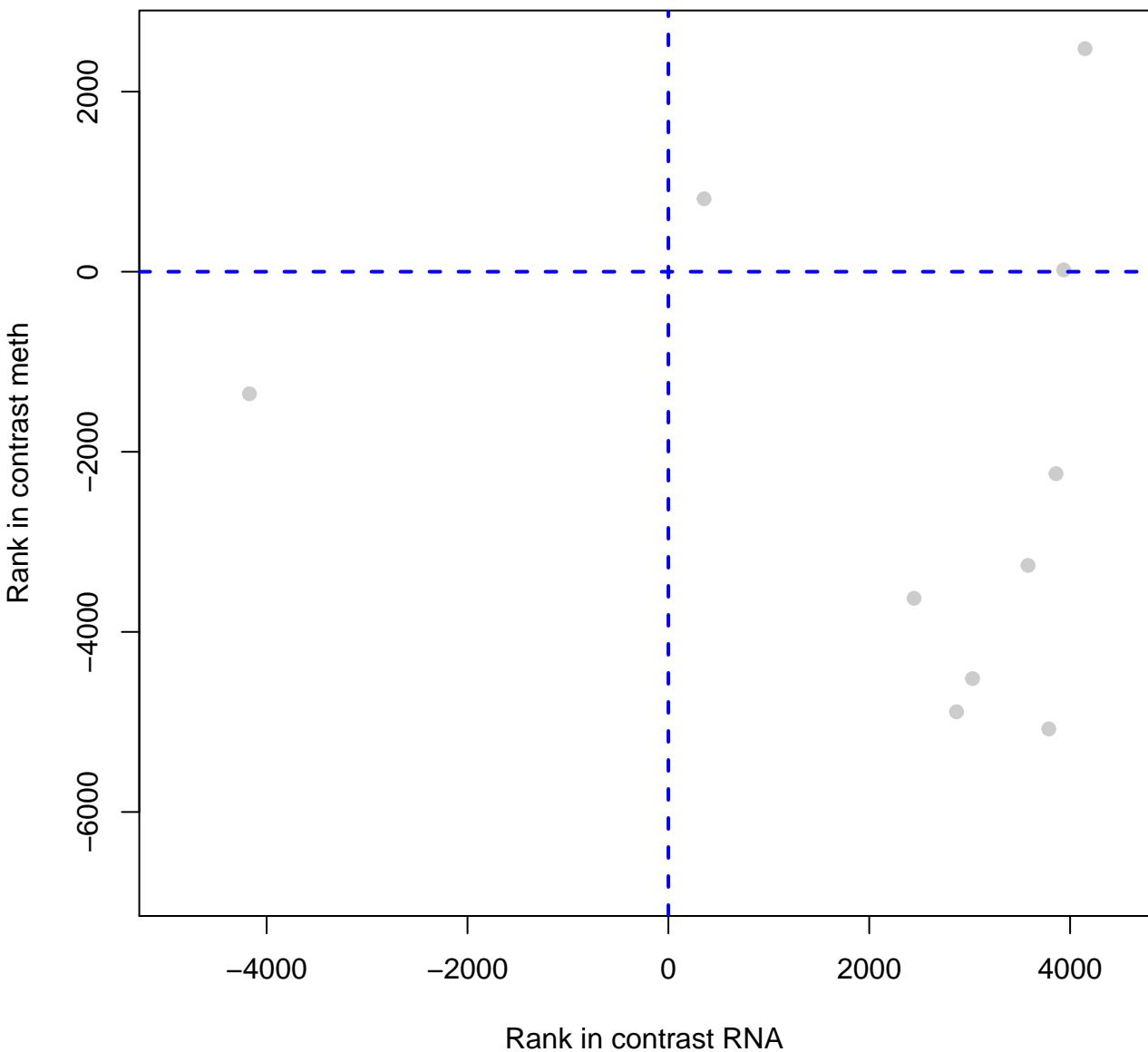
ROS and RNS production in phagocytes



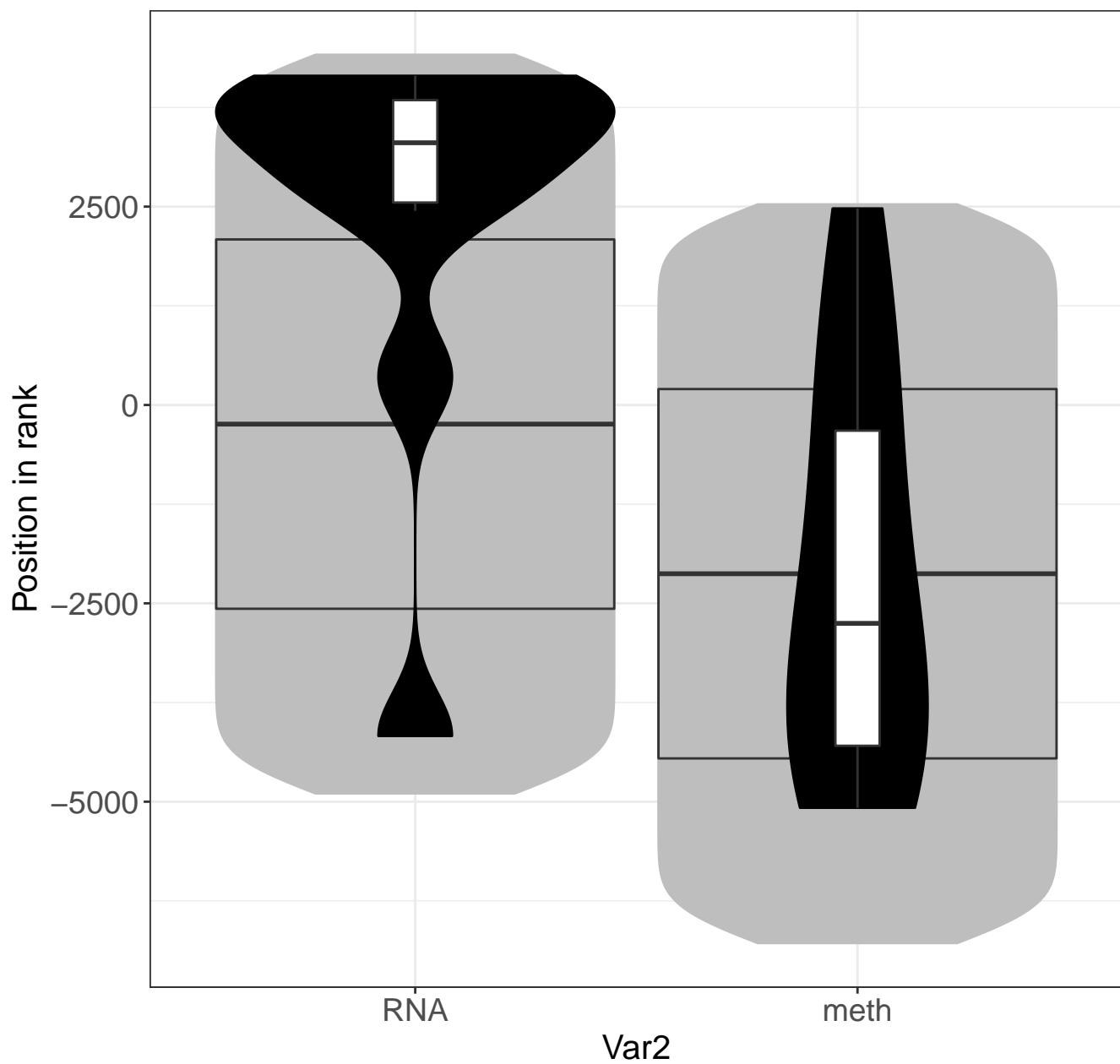
WNT5A-dependent internalization of FZD4



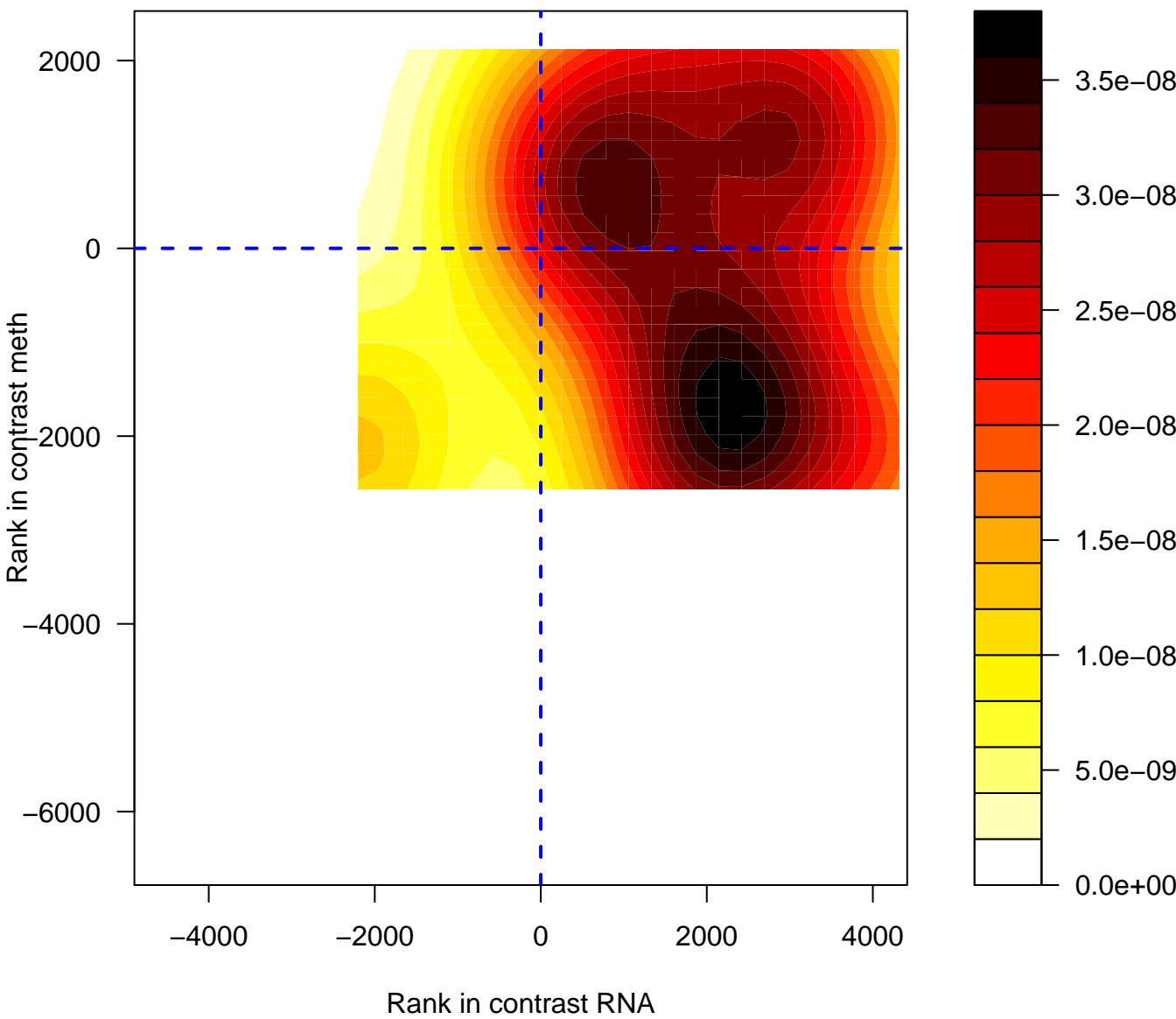
WNT5A-dependent internalization of FZD4



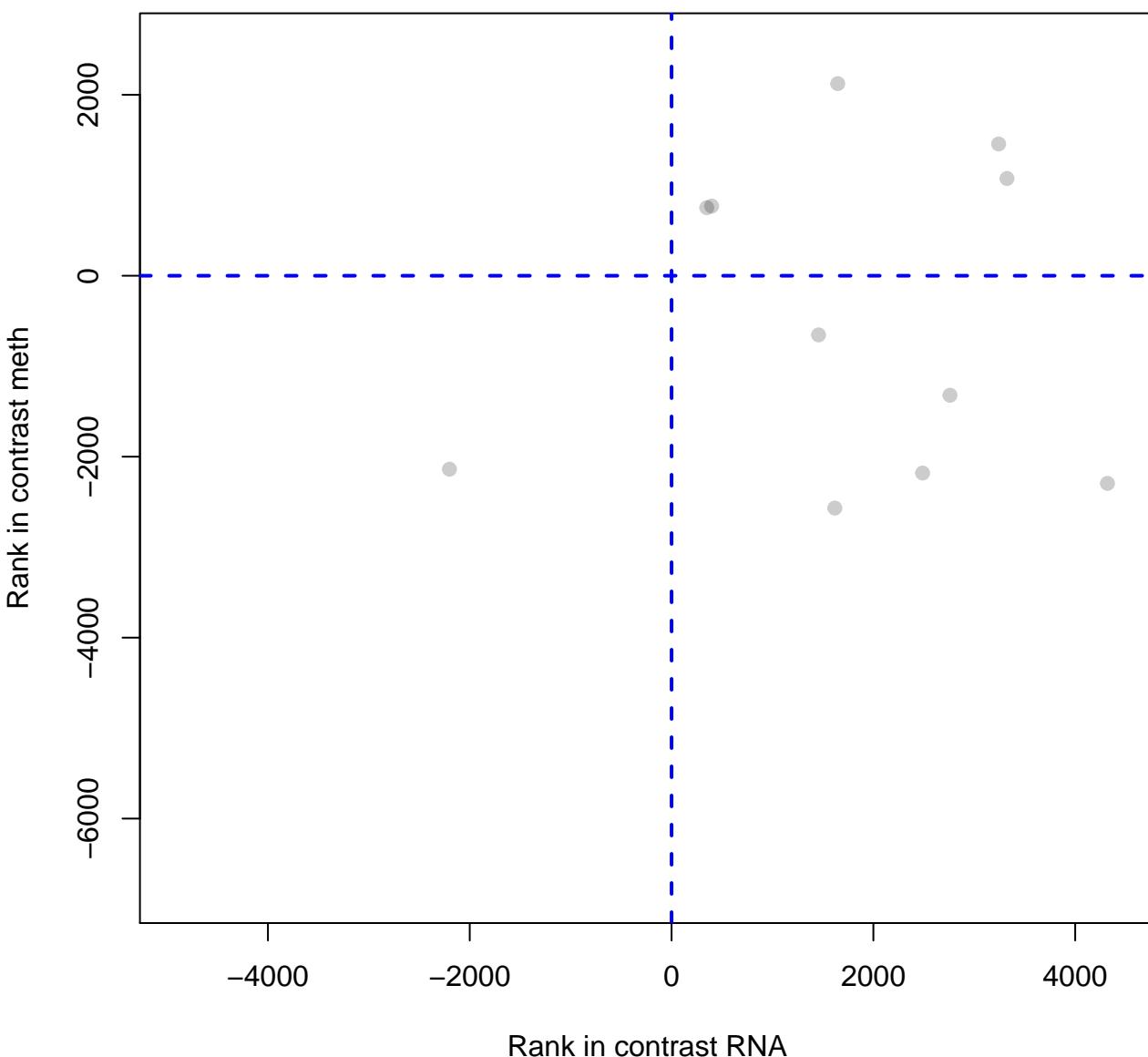
WNT5A-dependent internalization of FZD4



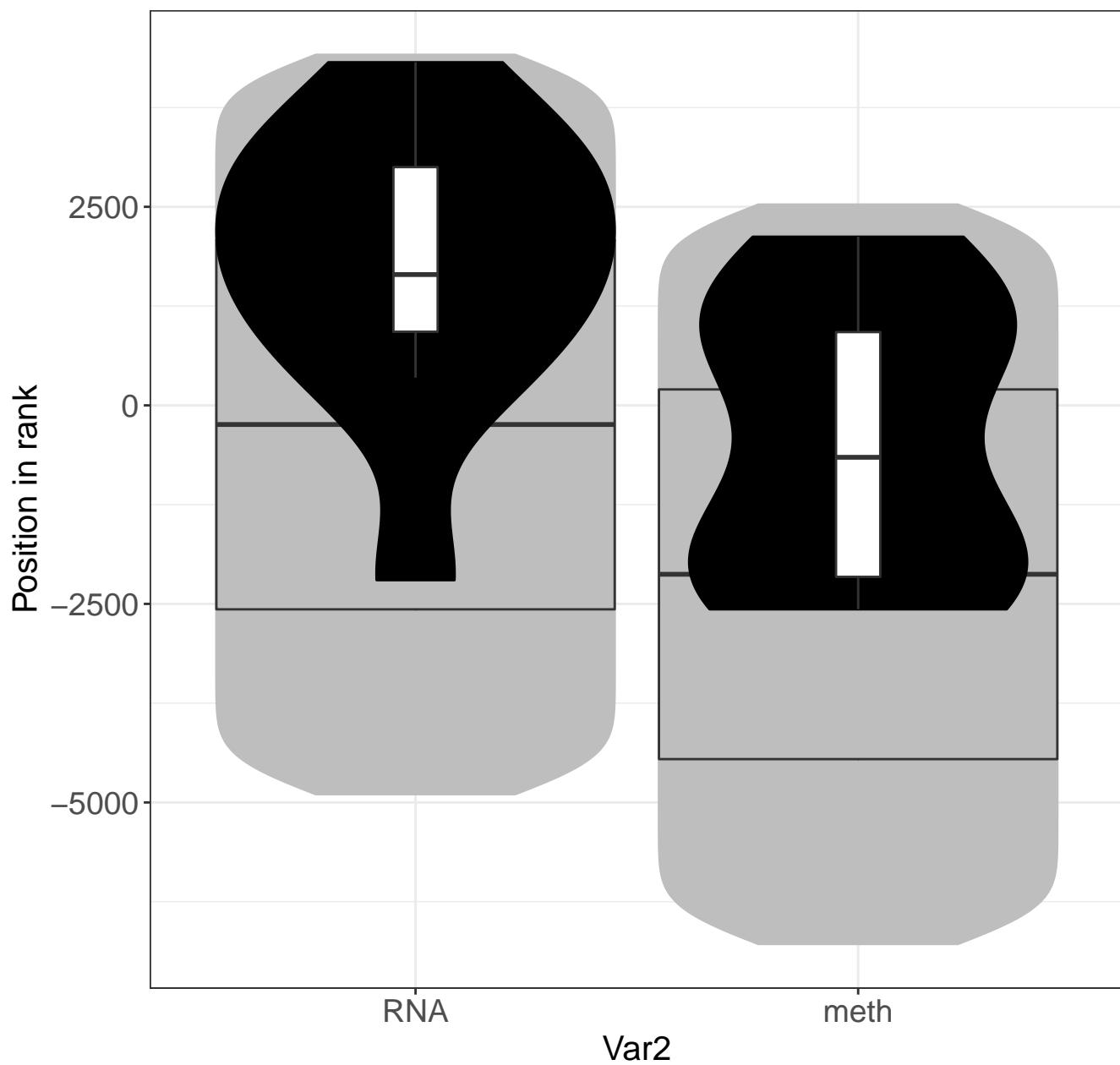
InIB-mediated entry of *Listeria monocytogenes* into host



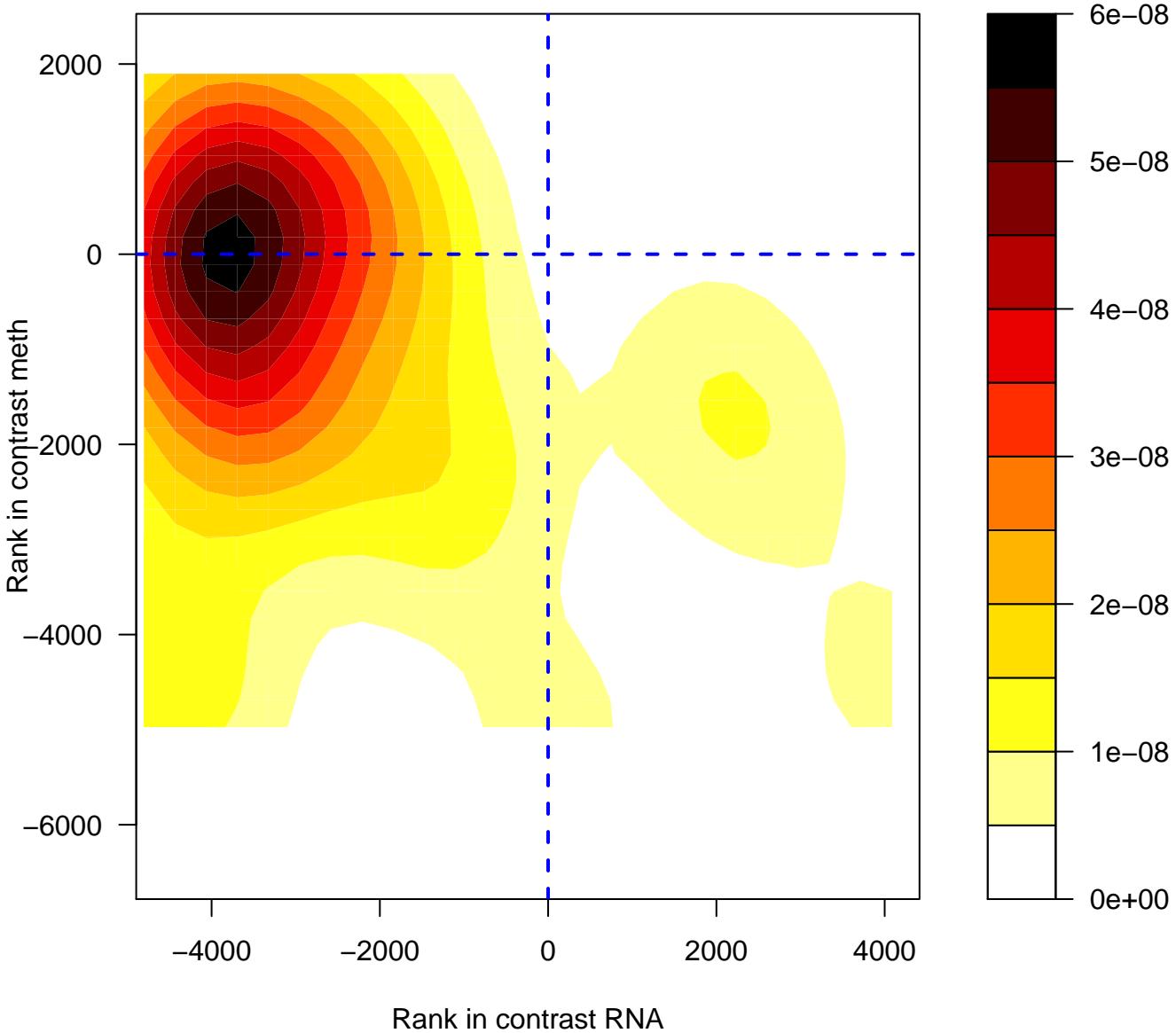
InIB-mediated entry of Listeria monocytogenes into host cell



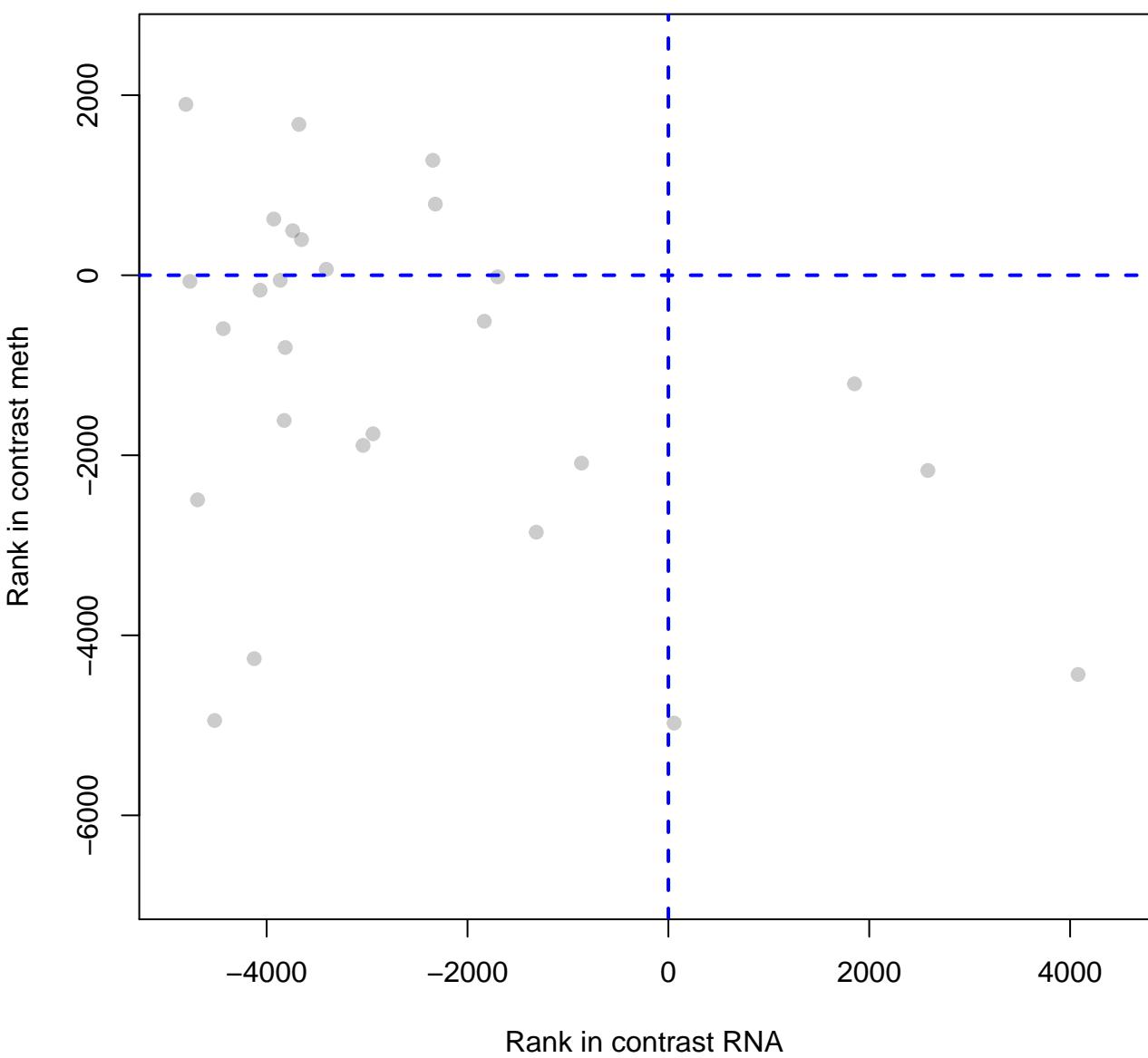
InlB-mediated entry of *Listeria monocytogenes* in



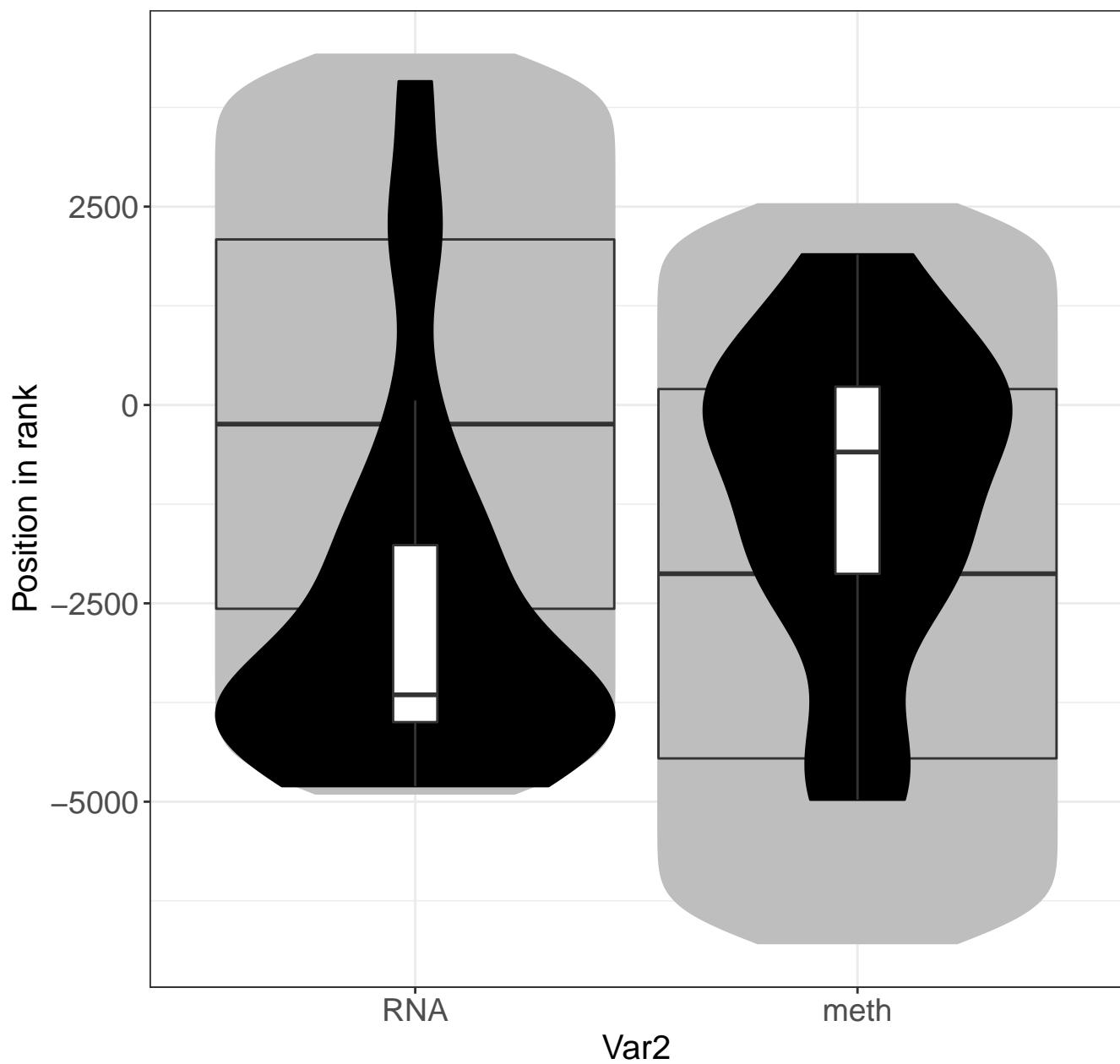
DNA strand elongation



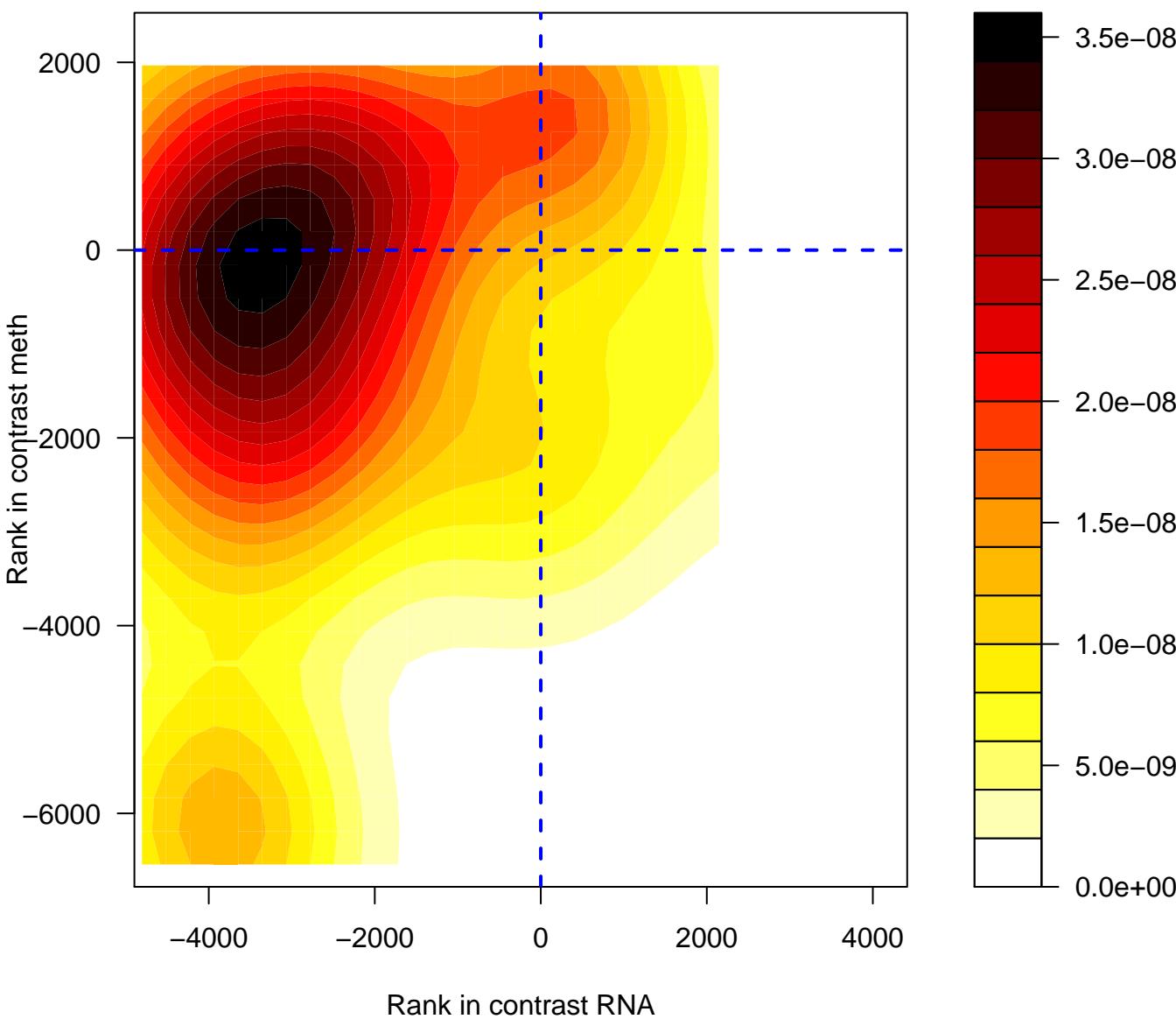
DNA strand elongation



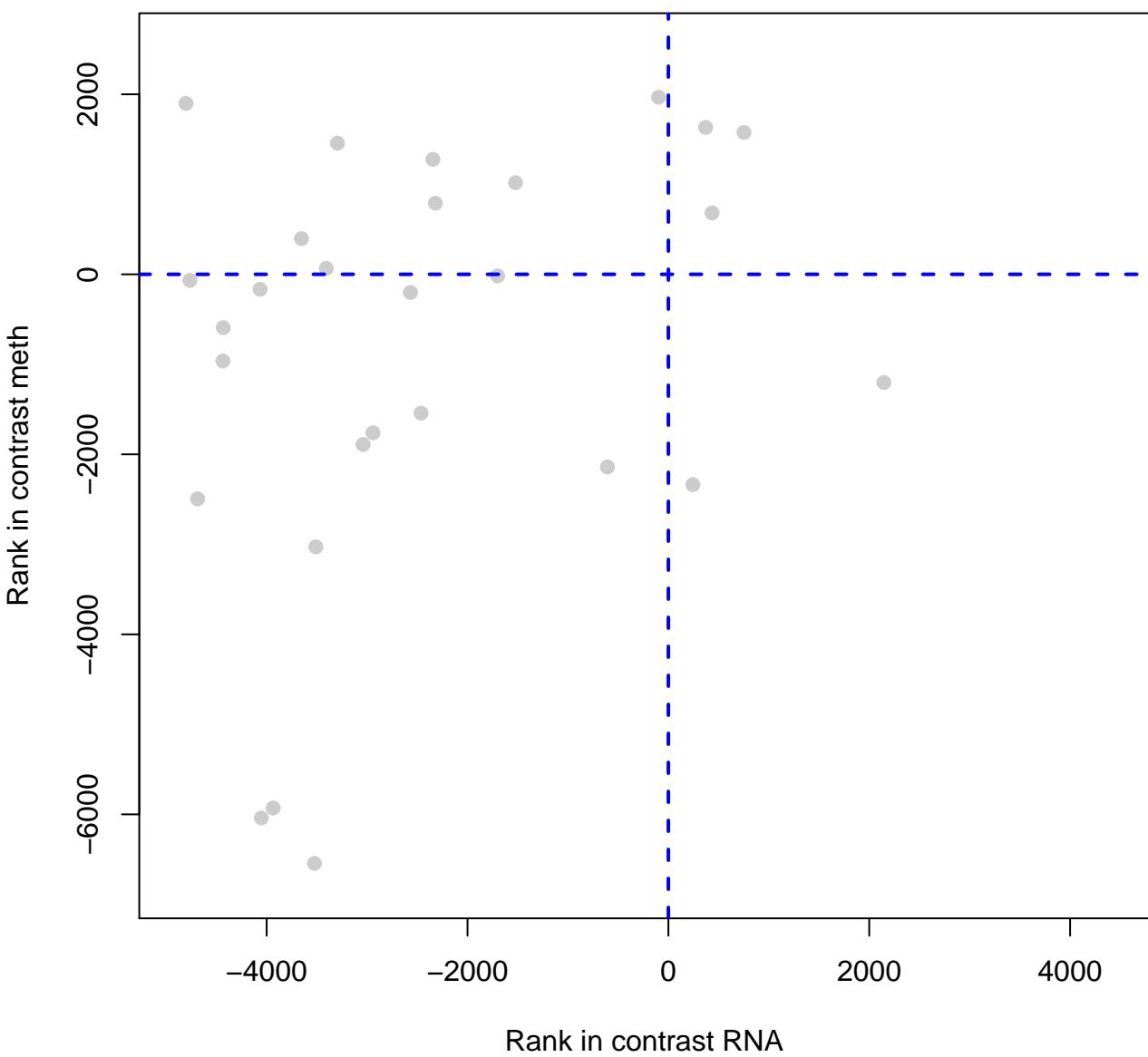
DNA strand elongation



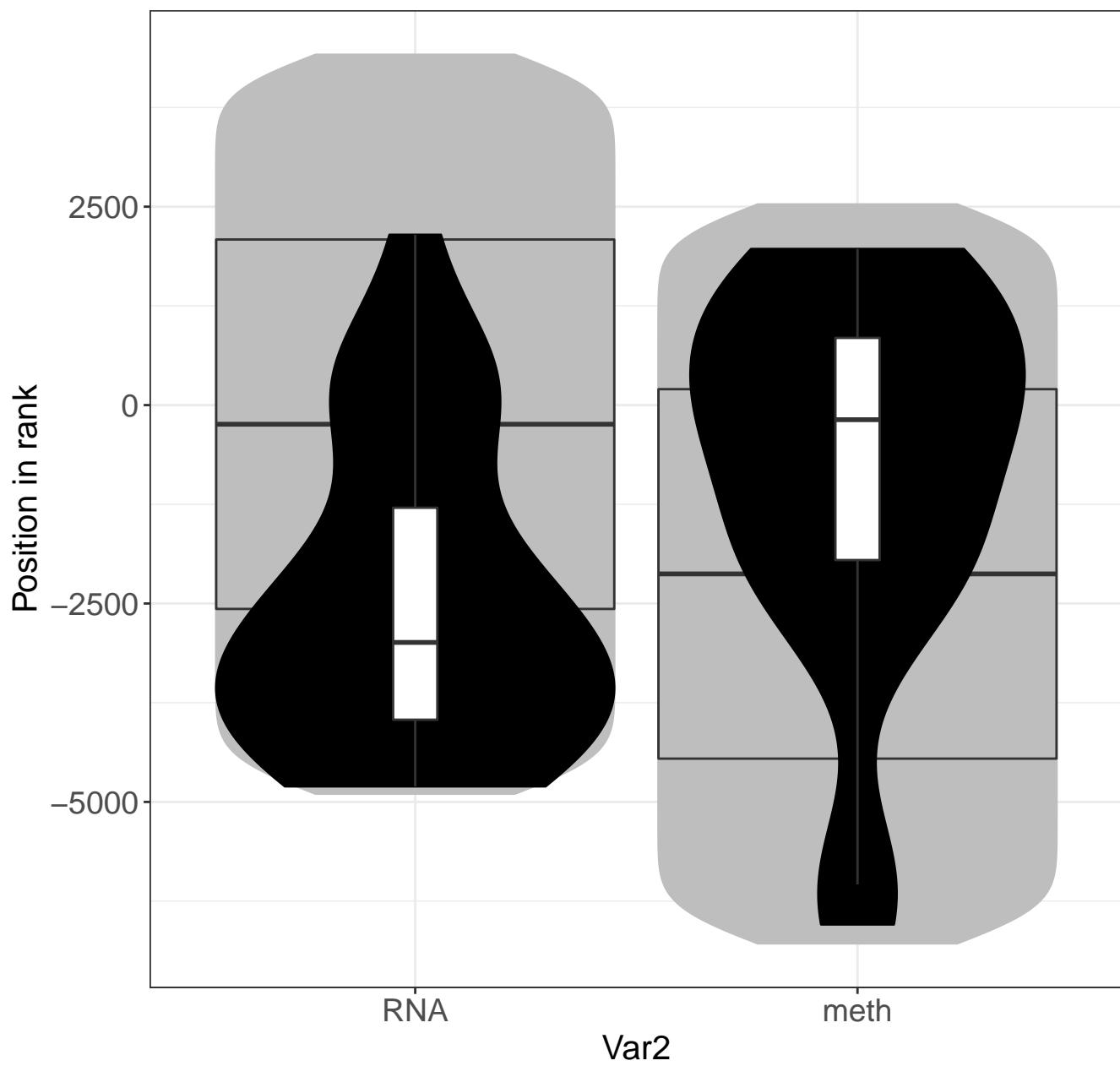
Activation of the pre-replicative complex



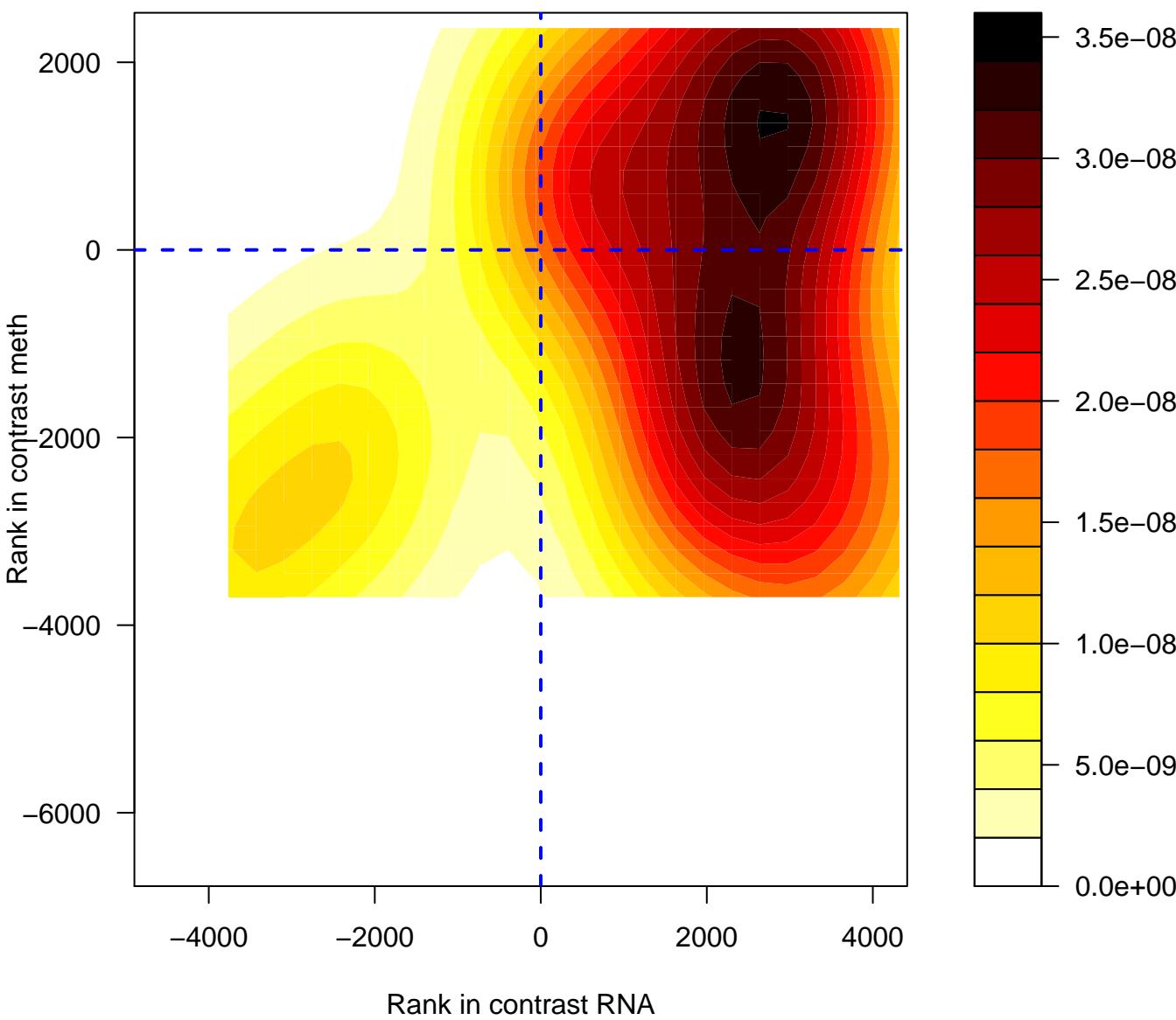
Activation of the pre-replicative complex



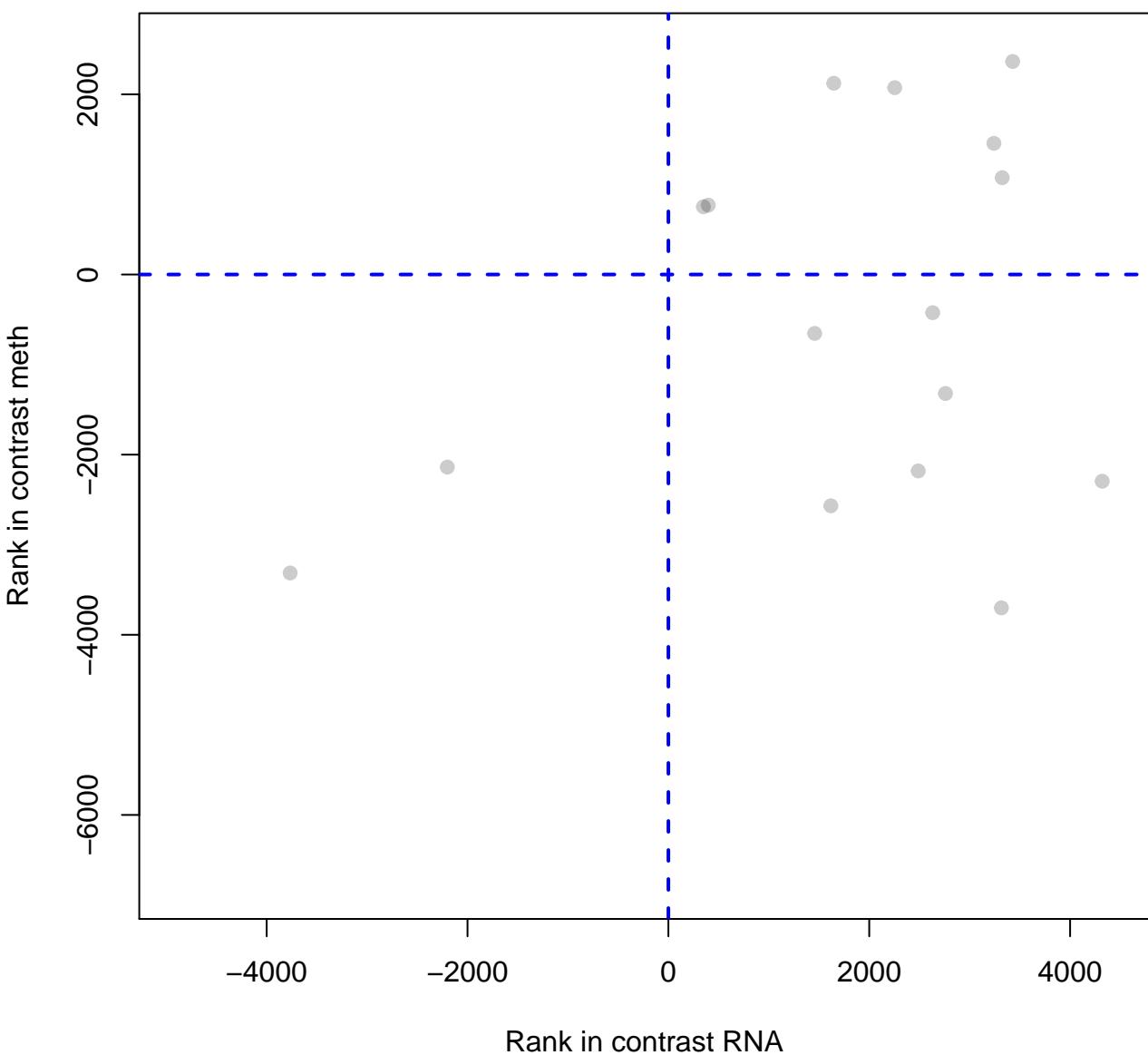
Activation of the pre-replicative complex



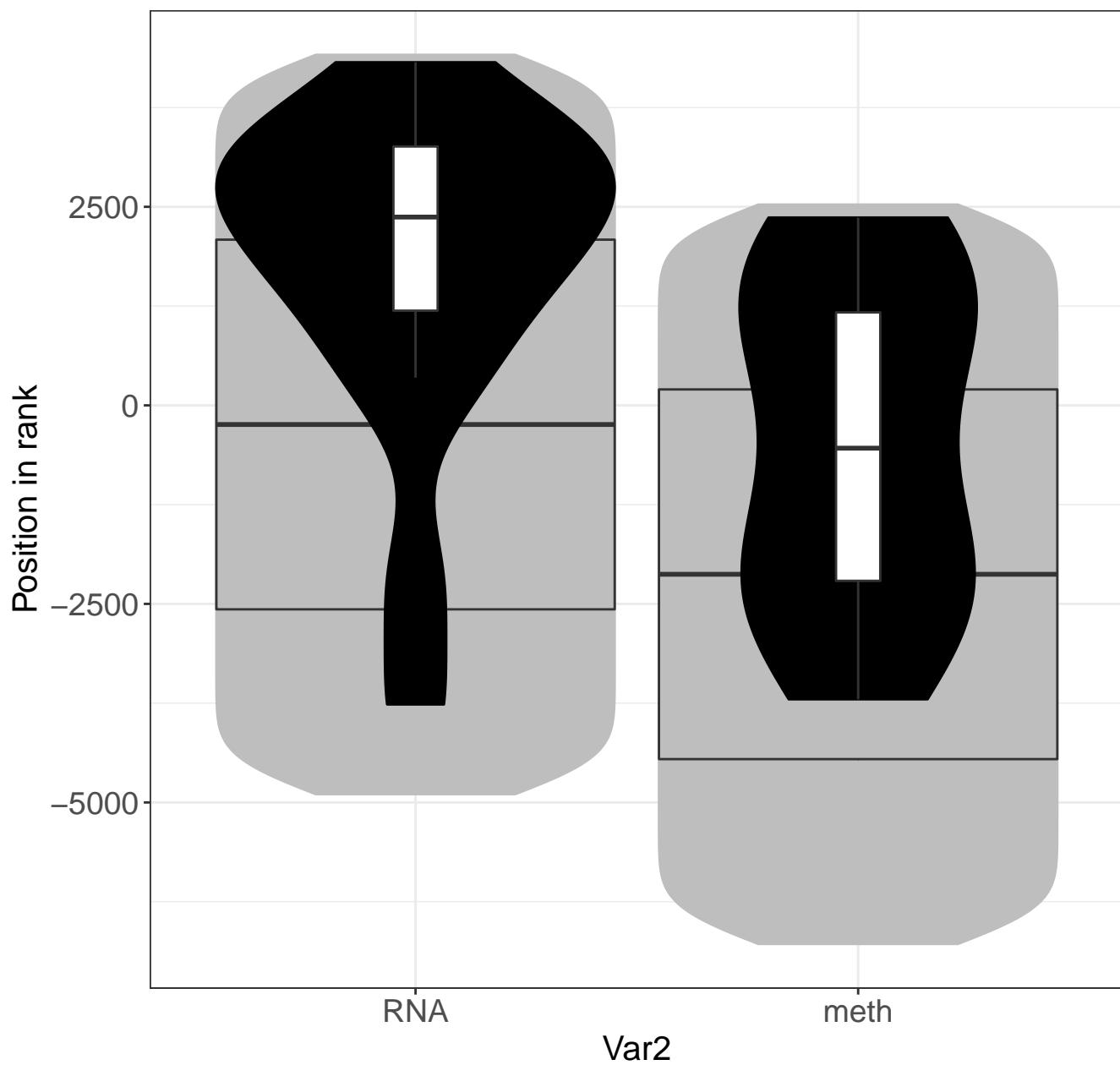
Negative regulation of MET activity



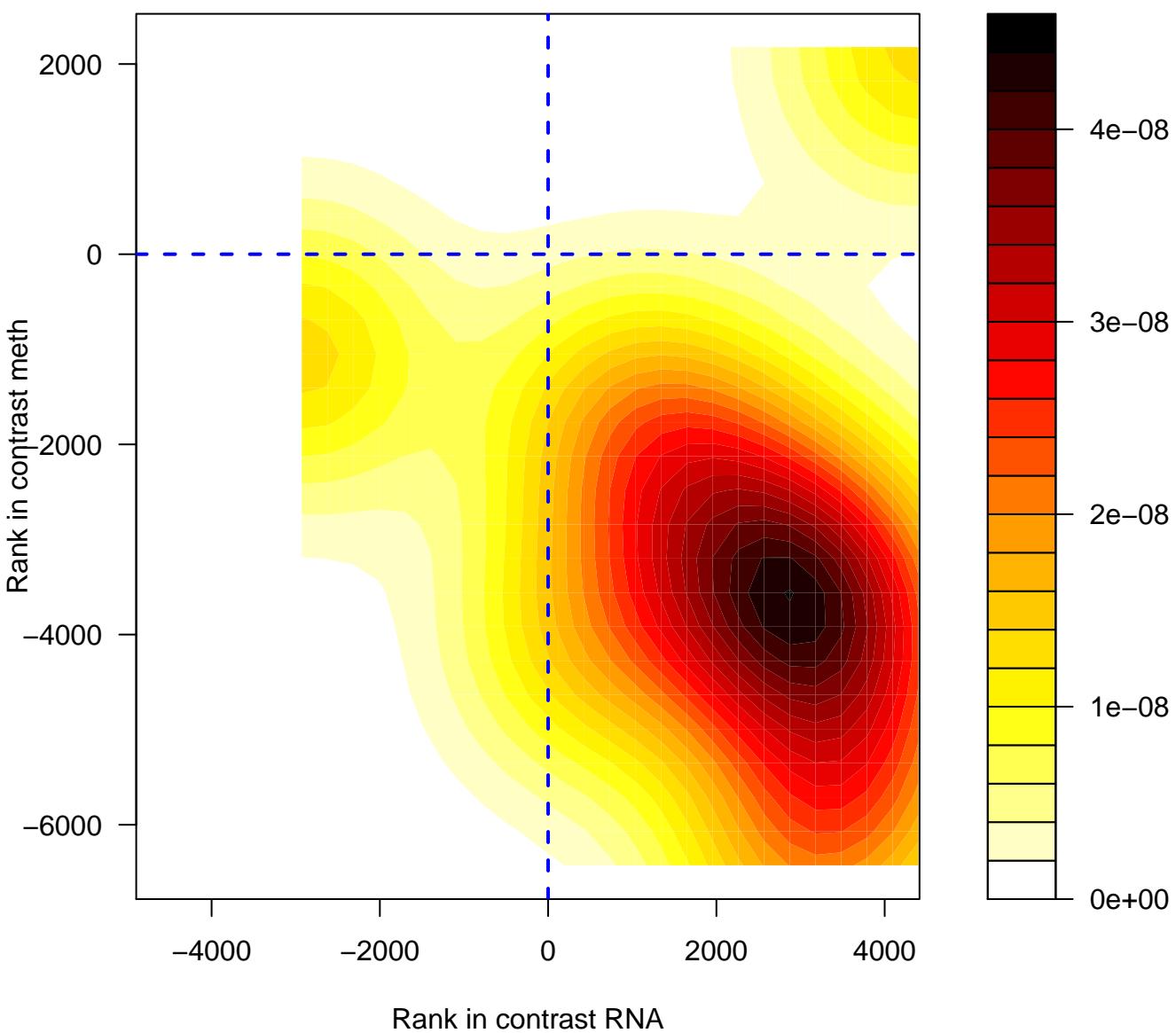
Negative regulation of MET activity



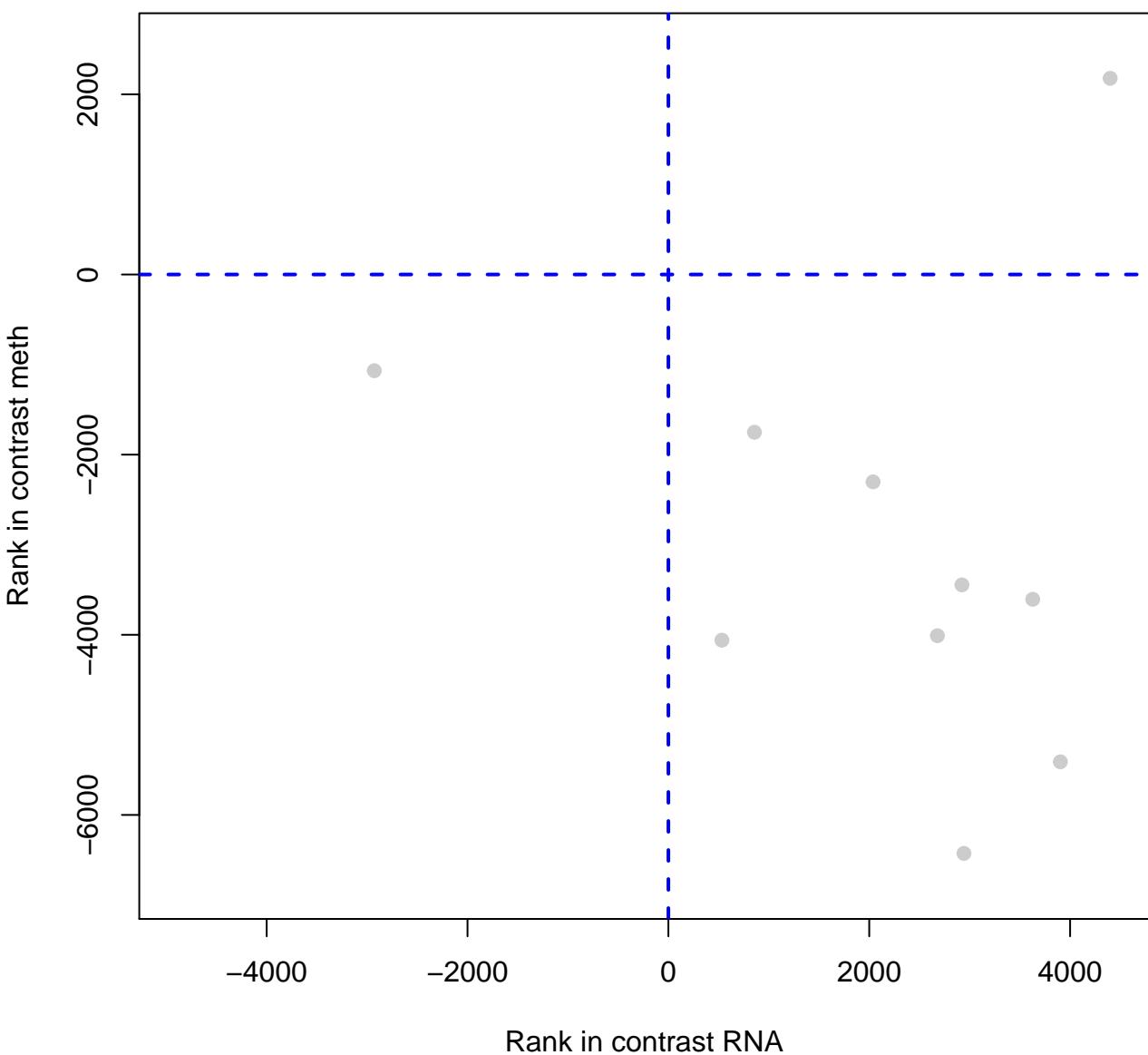
Negative regulation of MET activity



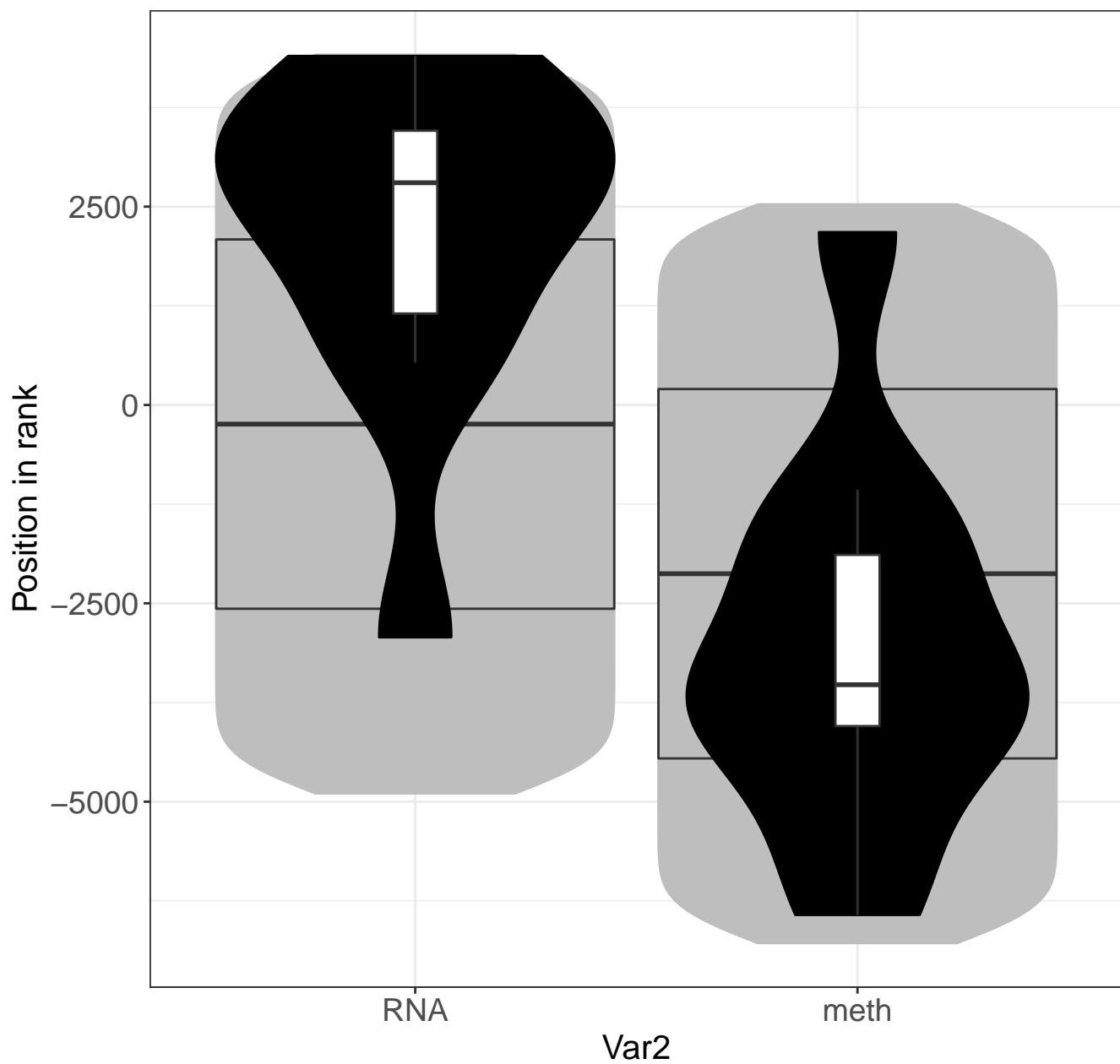
Signaling by Leptin



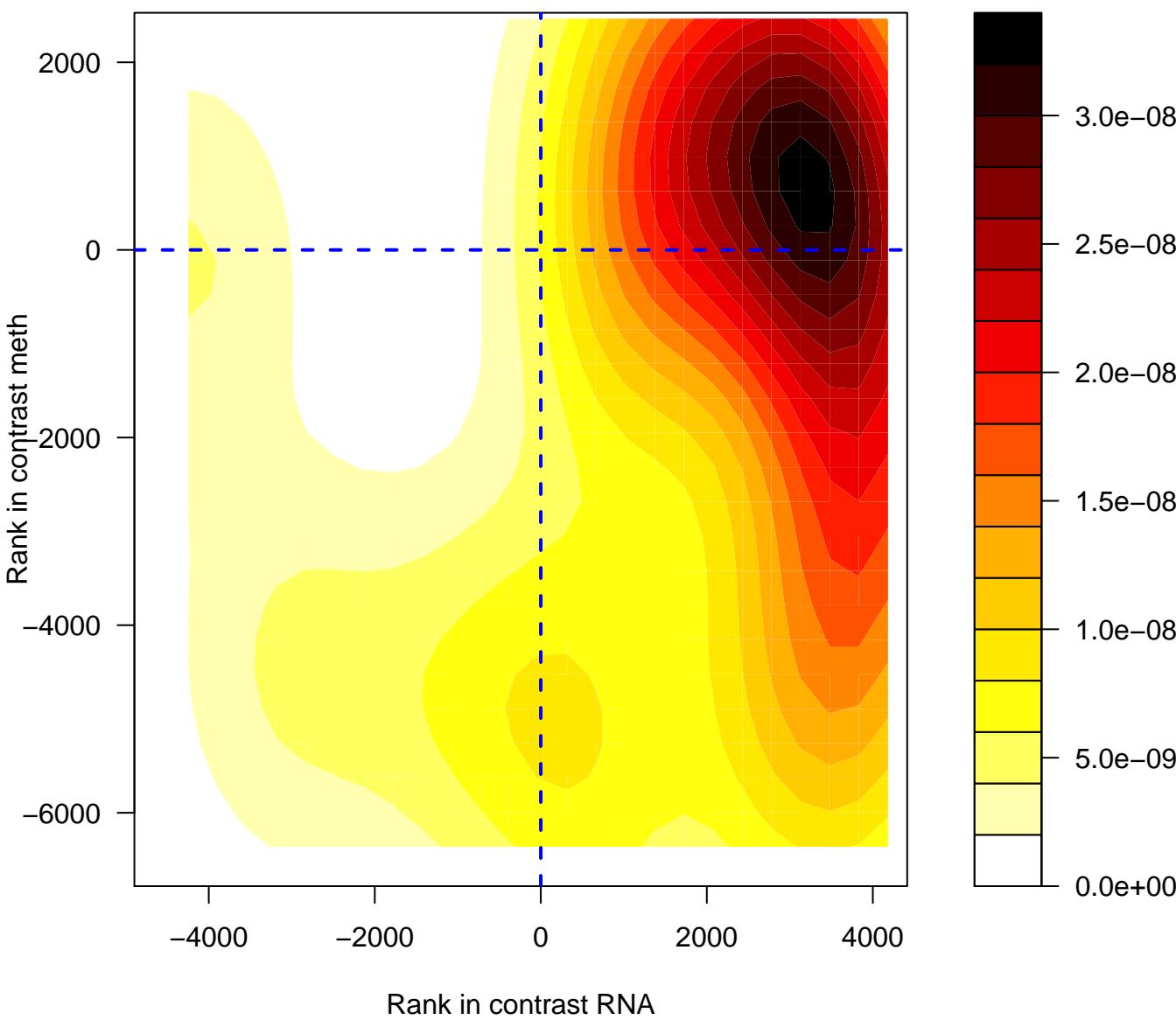
Signaling by Leptin



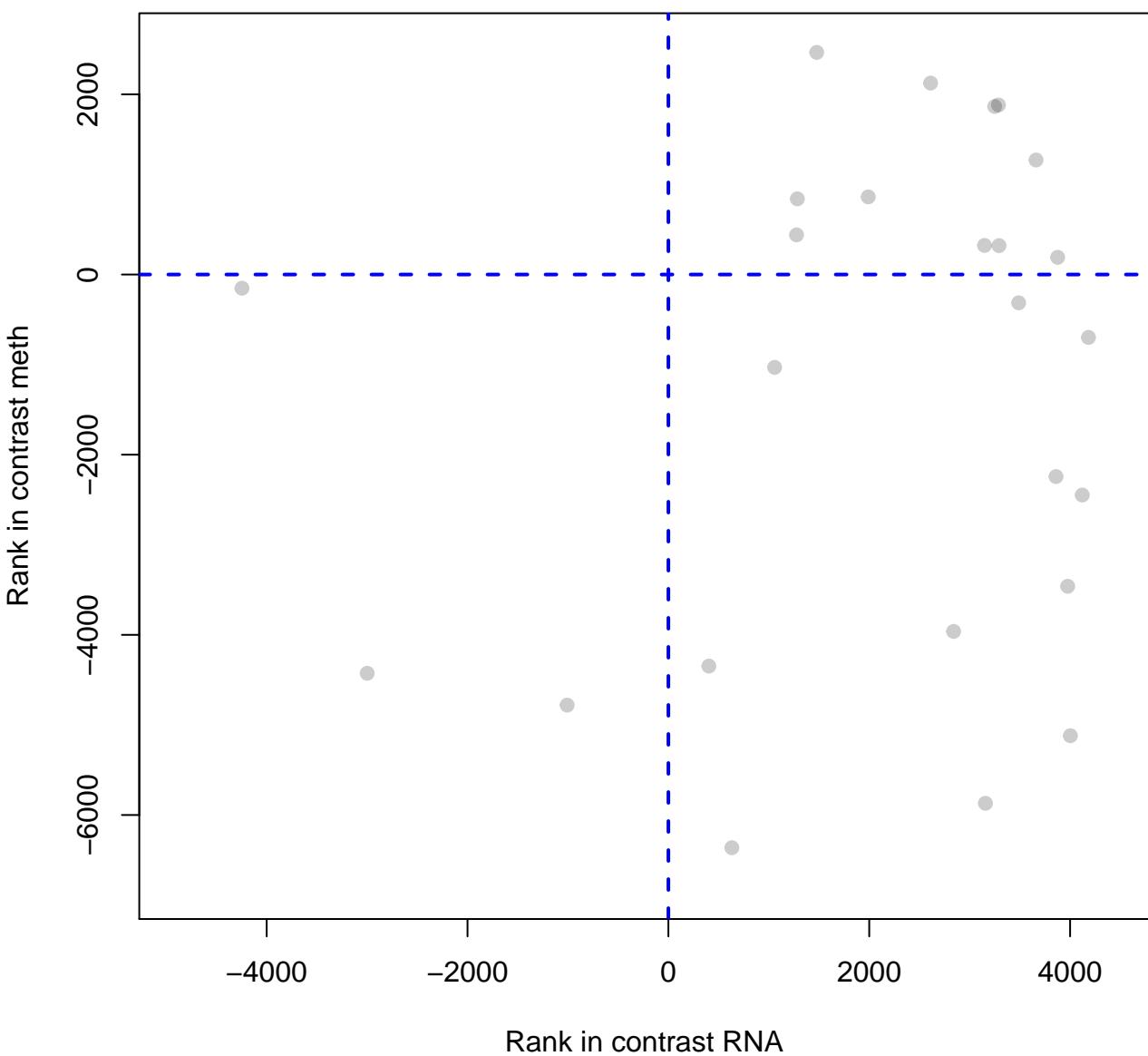
Signaling by Leptin



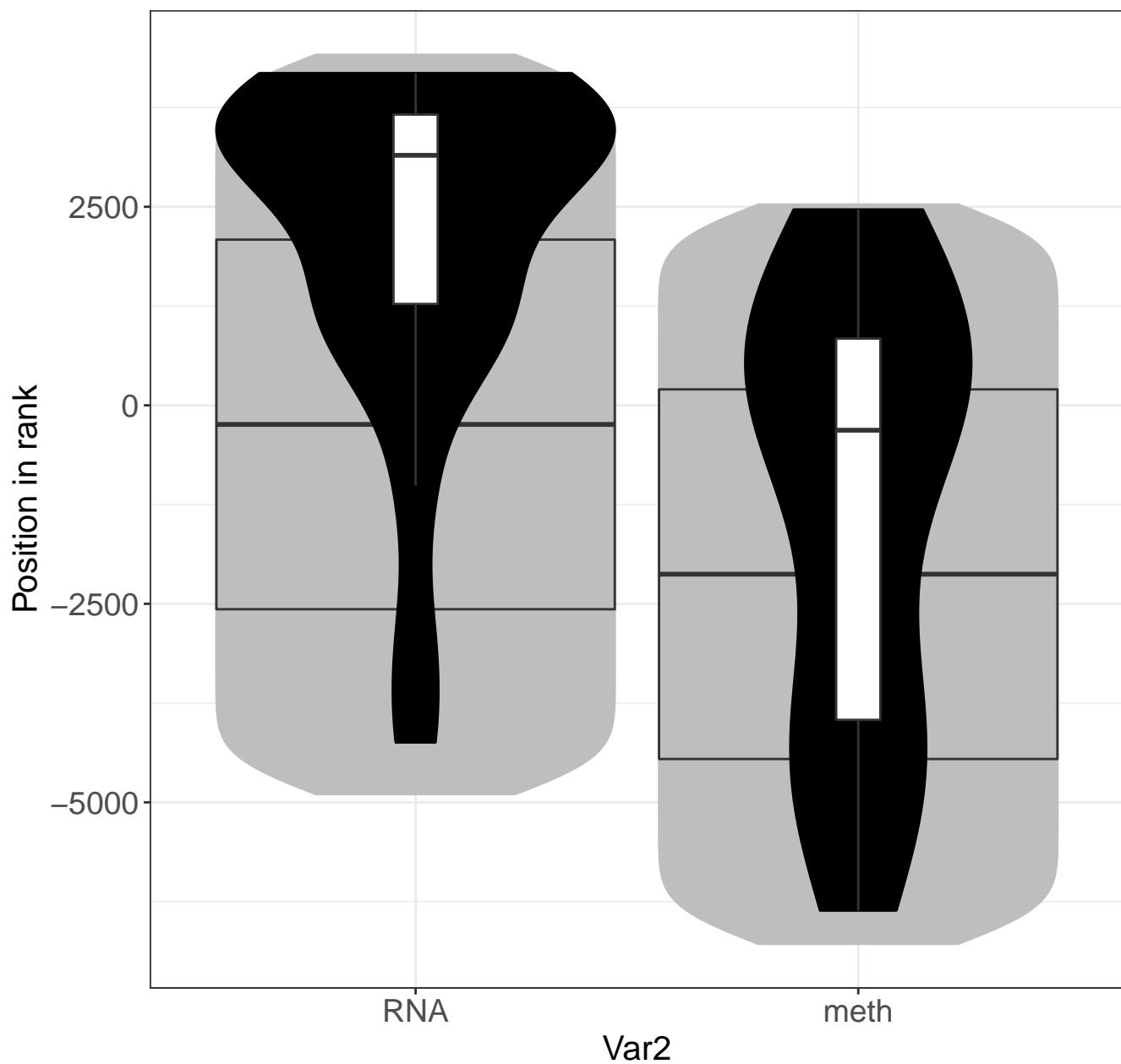
MAP2K and MAPK activation



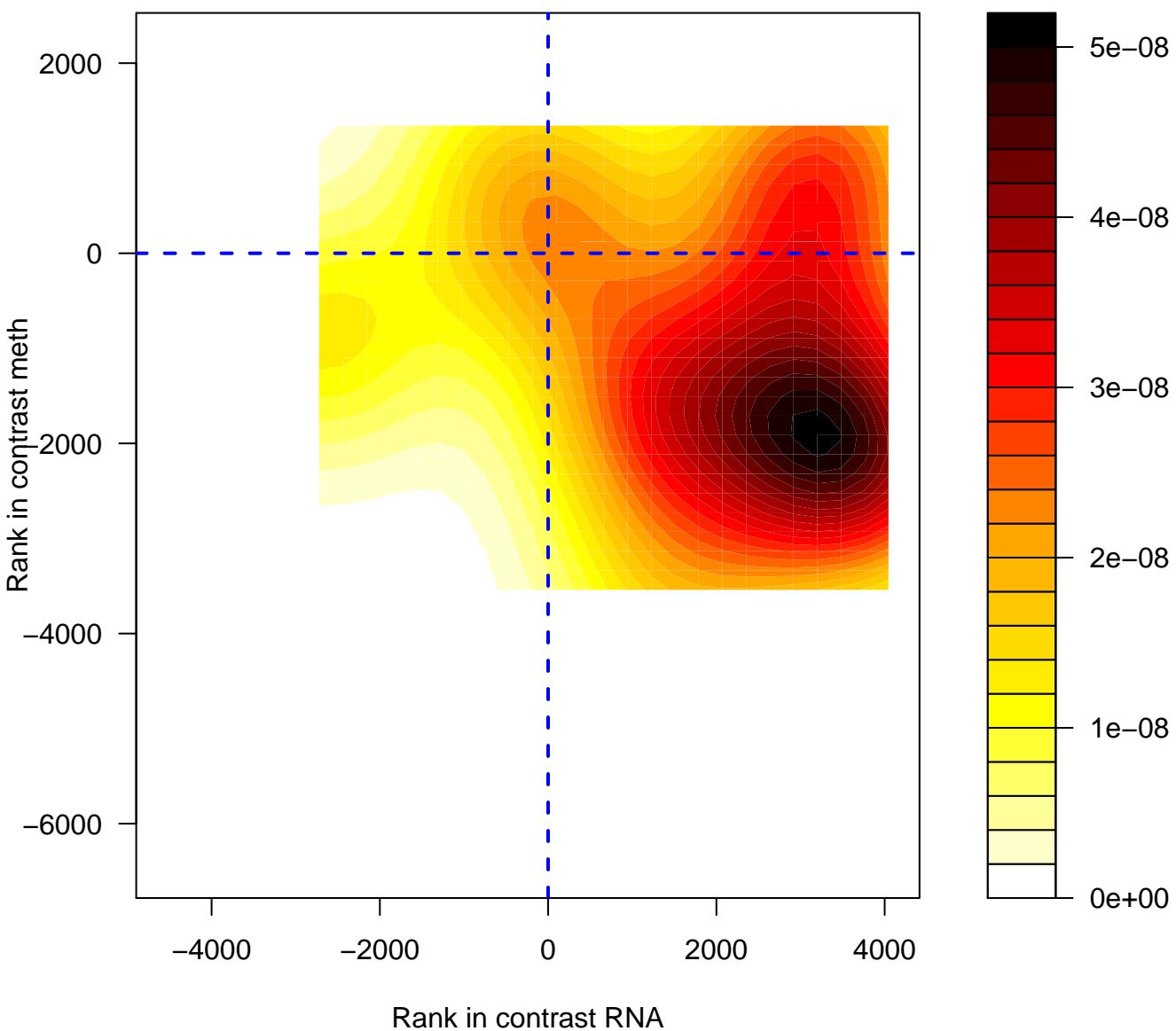
MAP2K and MAPK activation



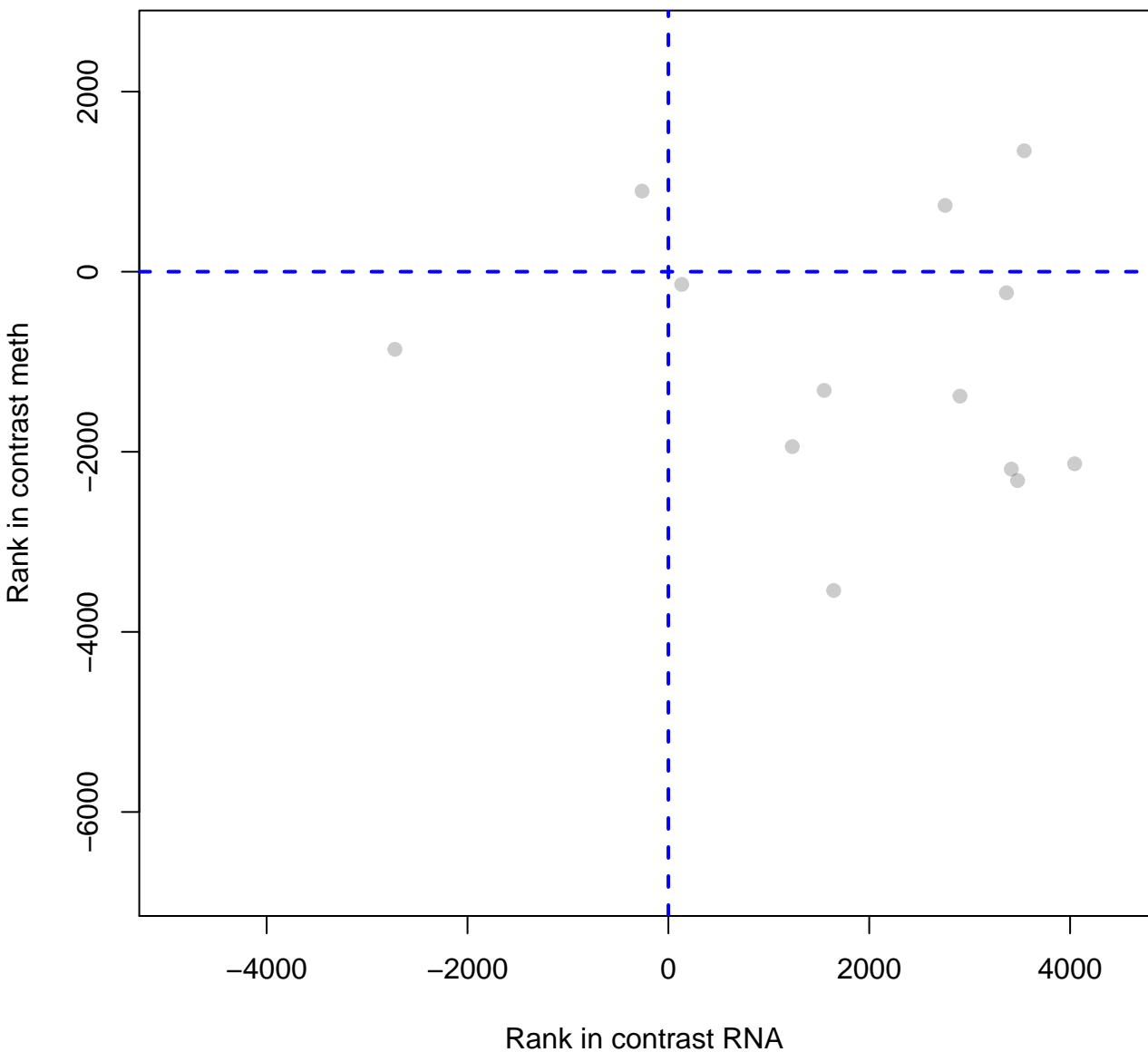
MAP2K and MAPK activation



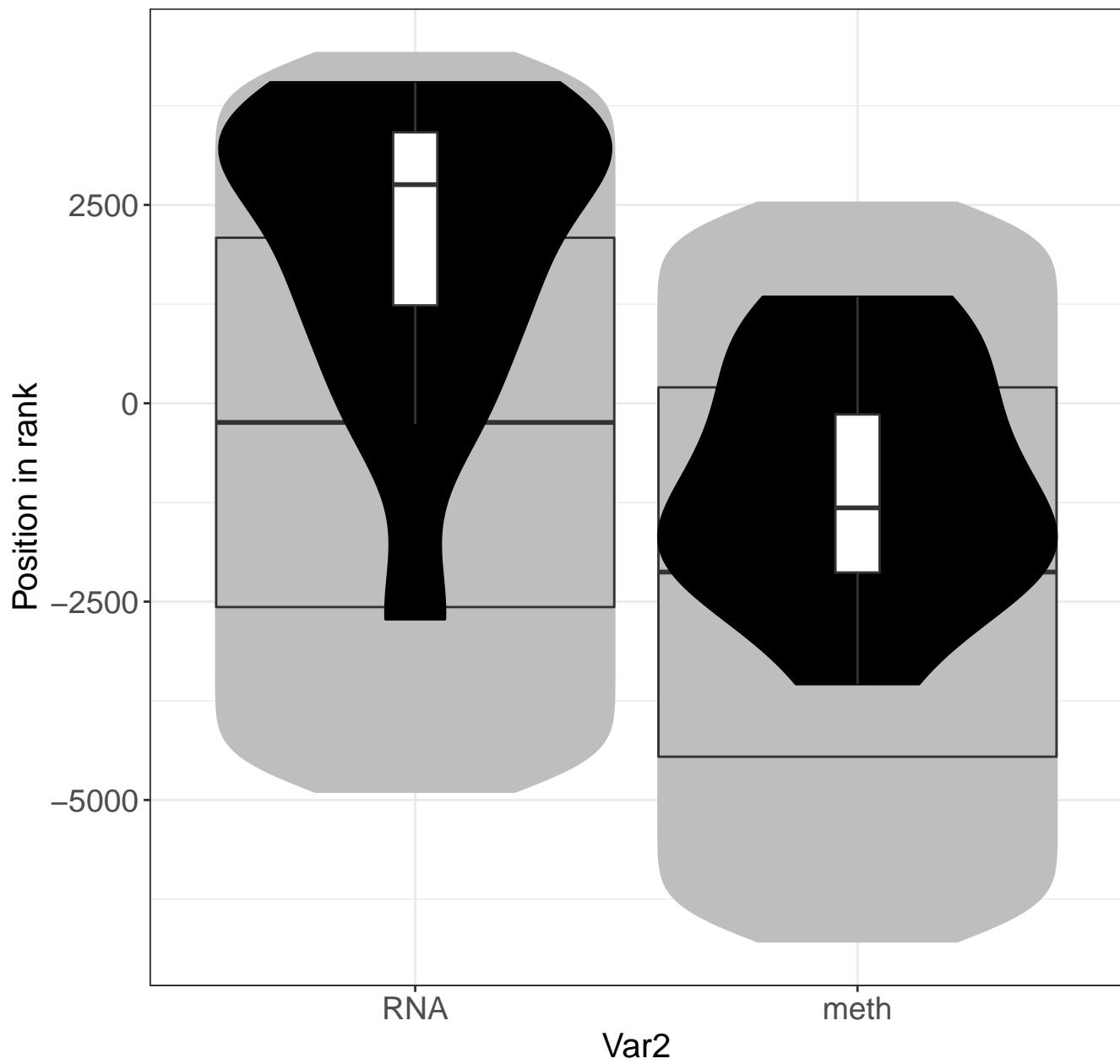
AKT phosphorylates targets in the cytosol



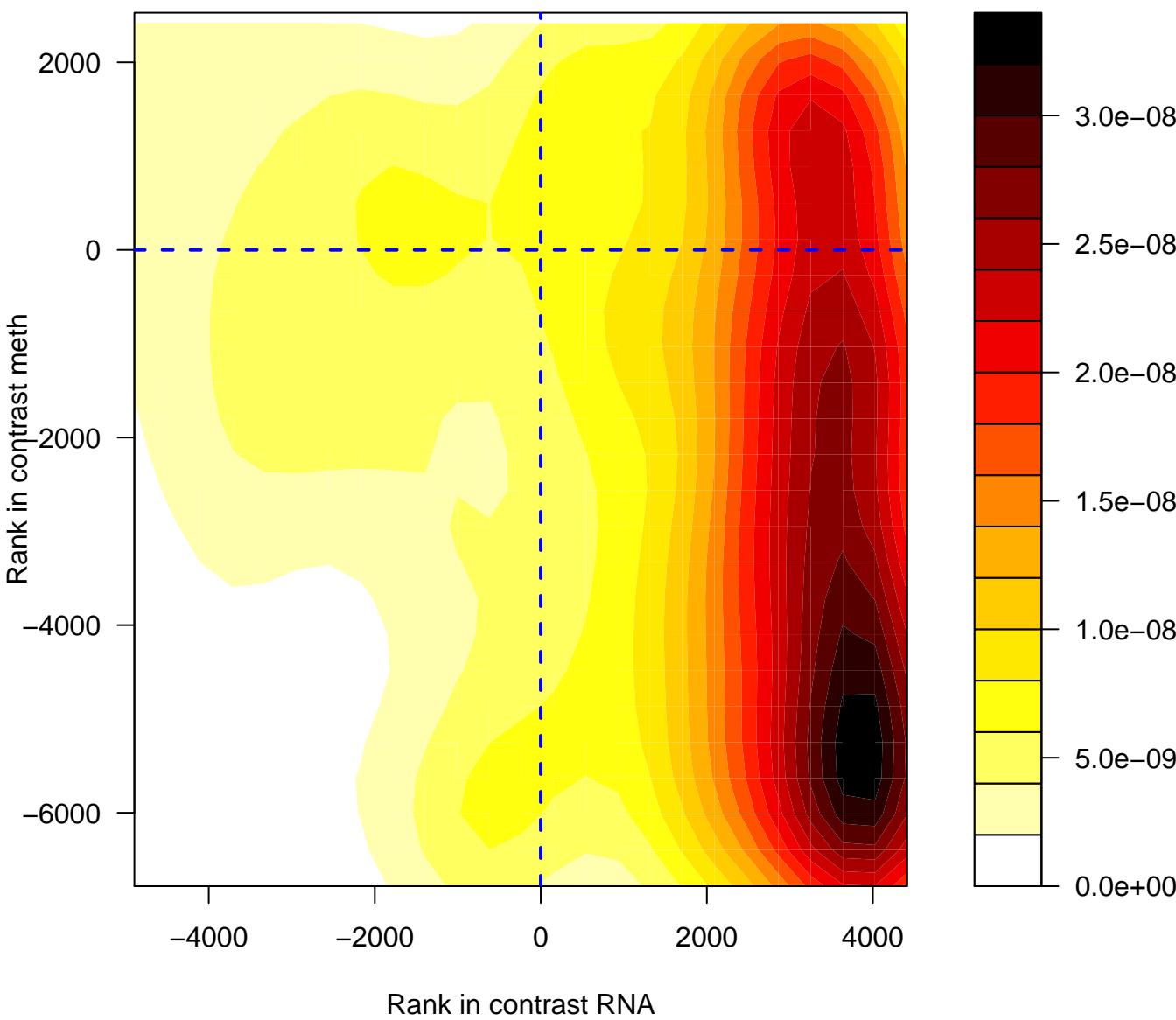
AKT phosphorylates targets in the cytosol



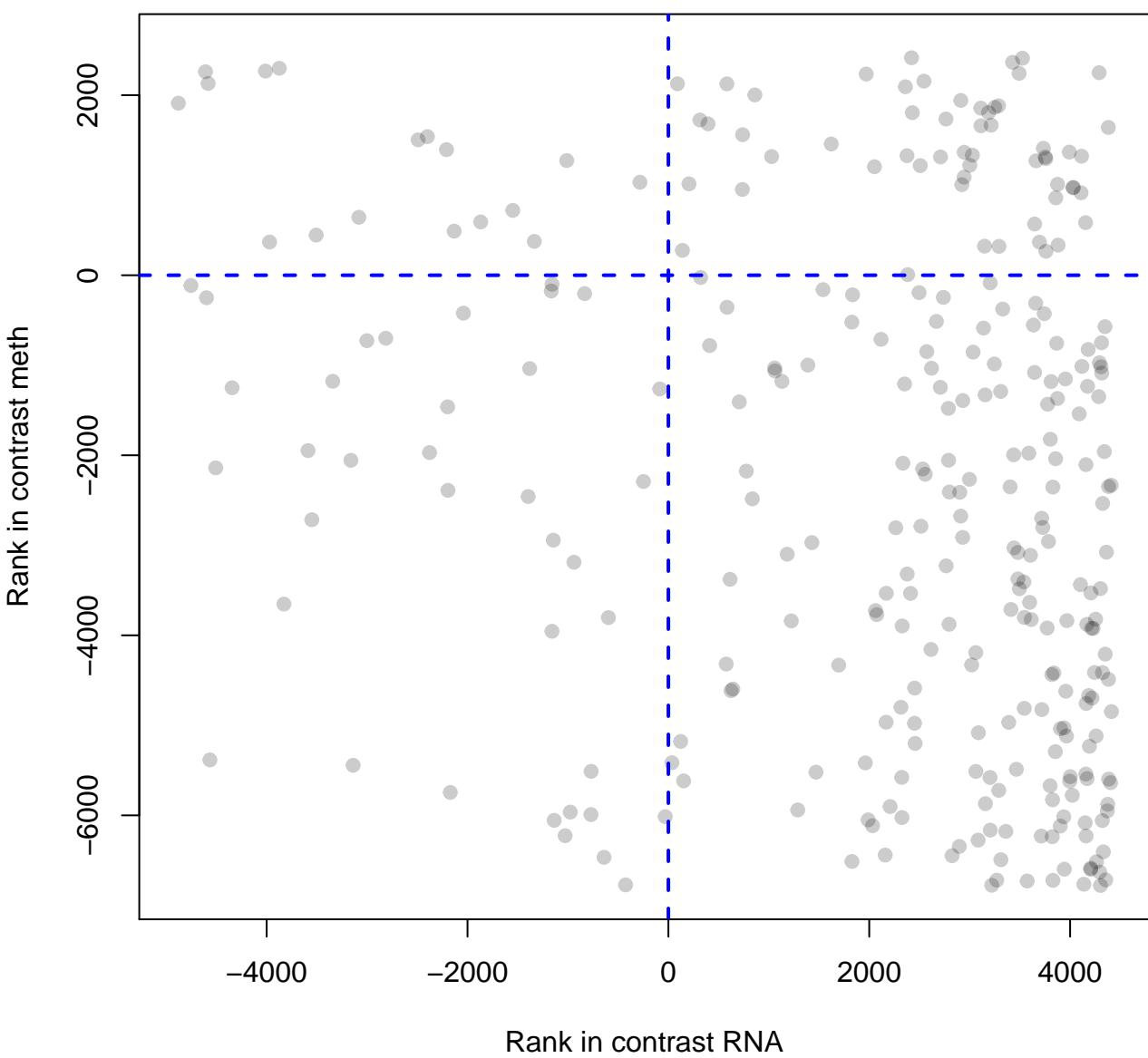
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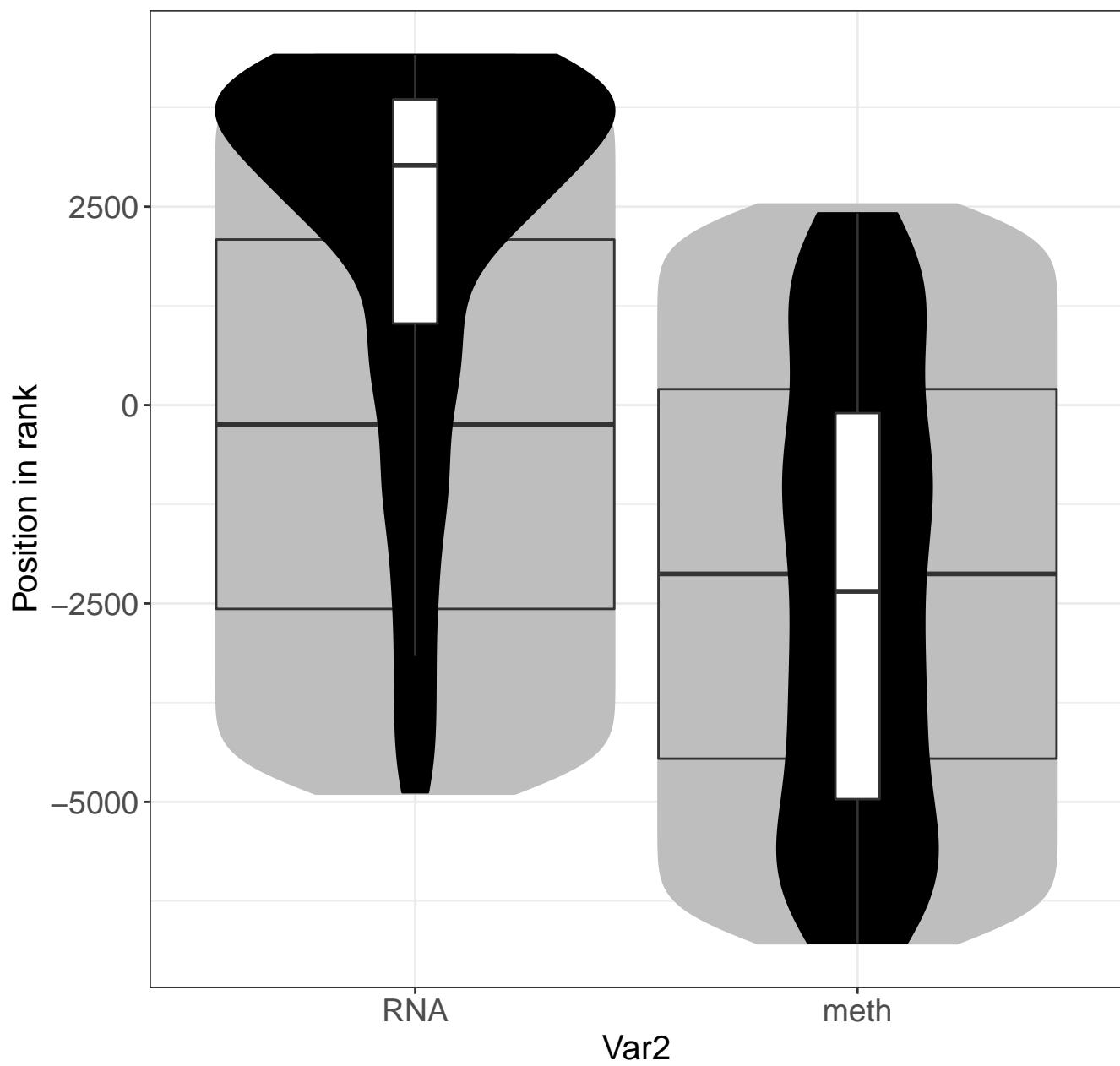
Neutrophil degranulation



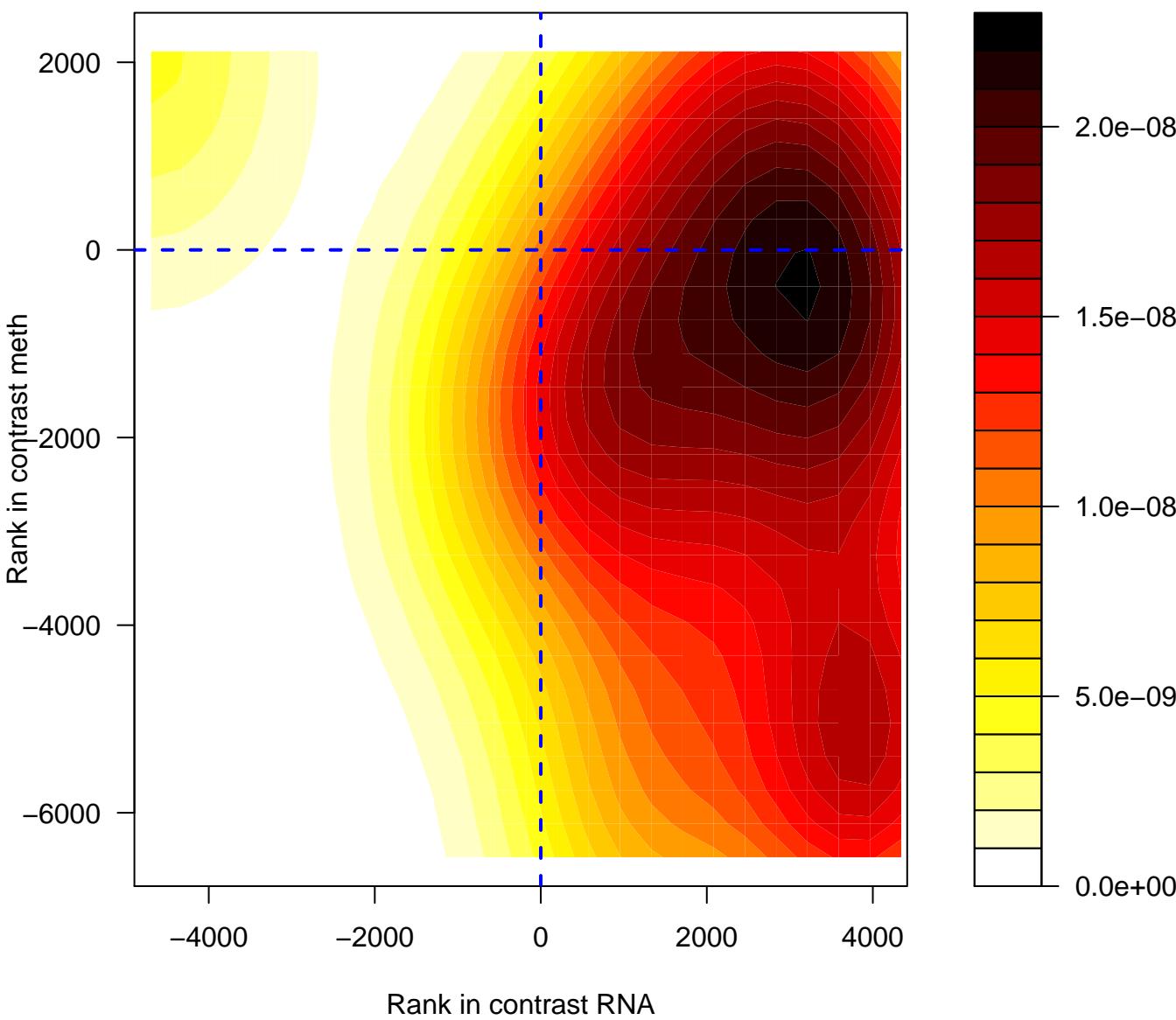
Neutrophil degranulation



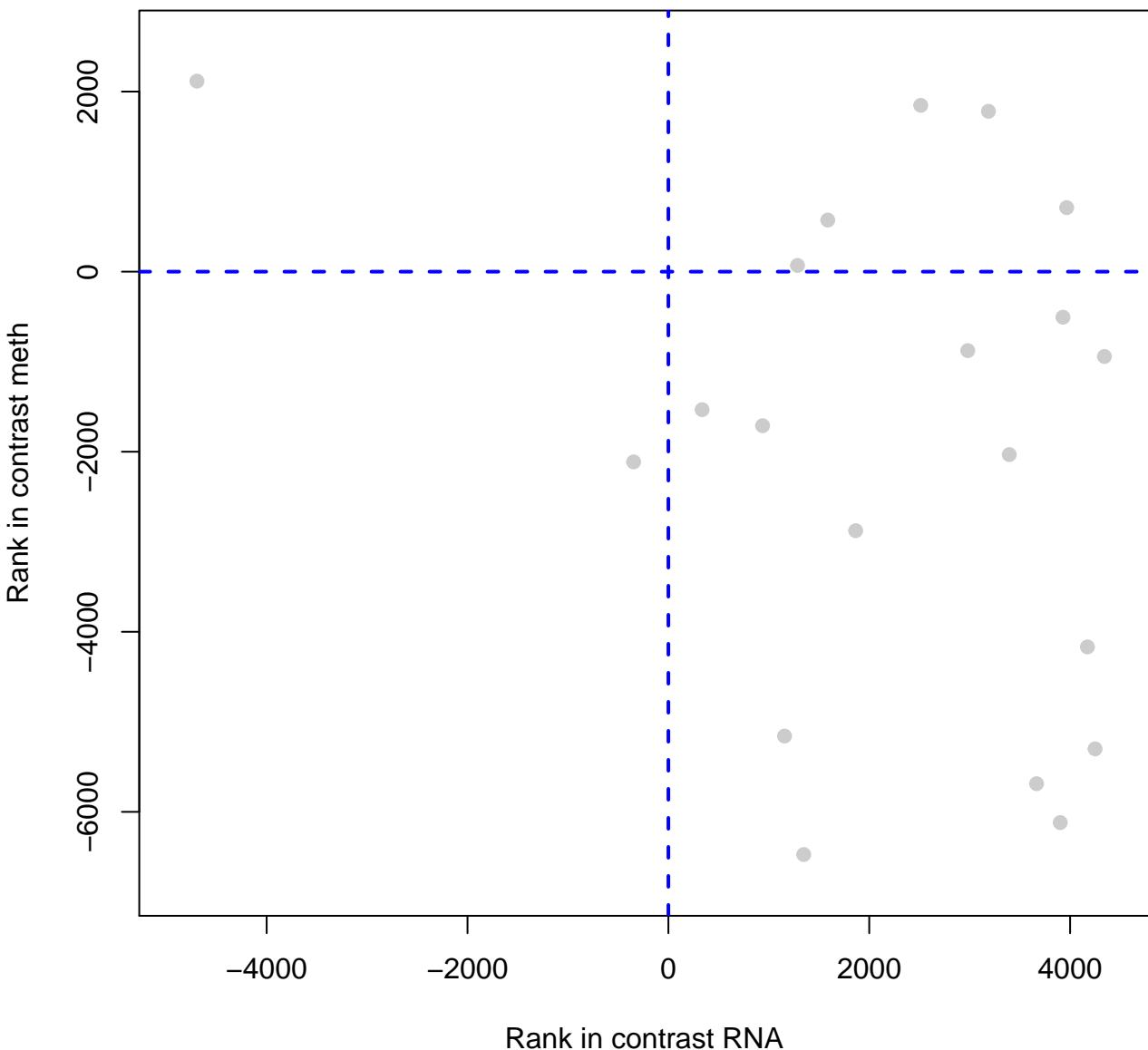
Neutrophil degranulation



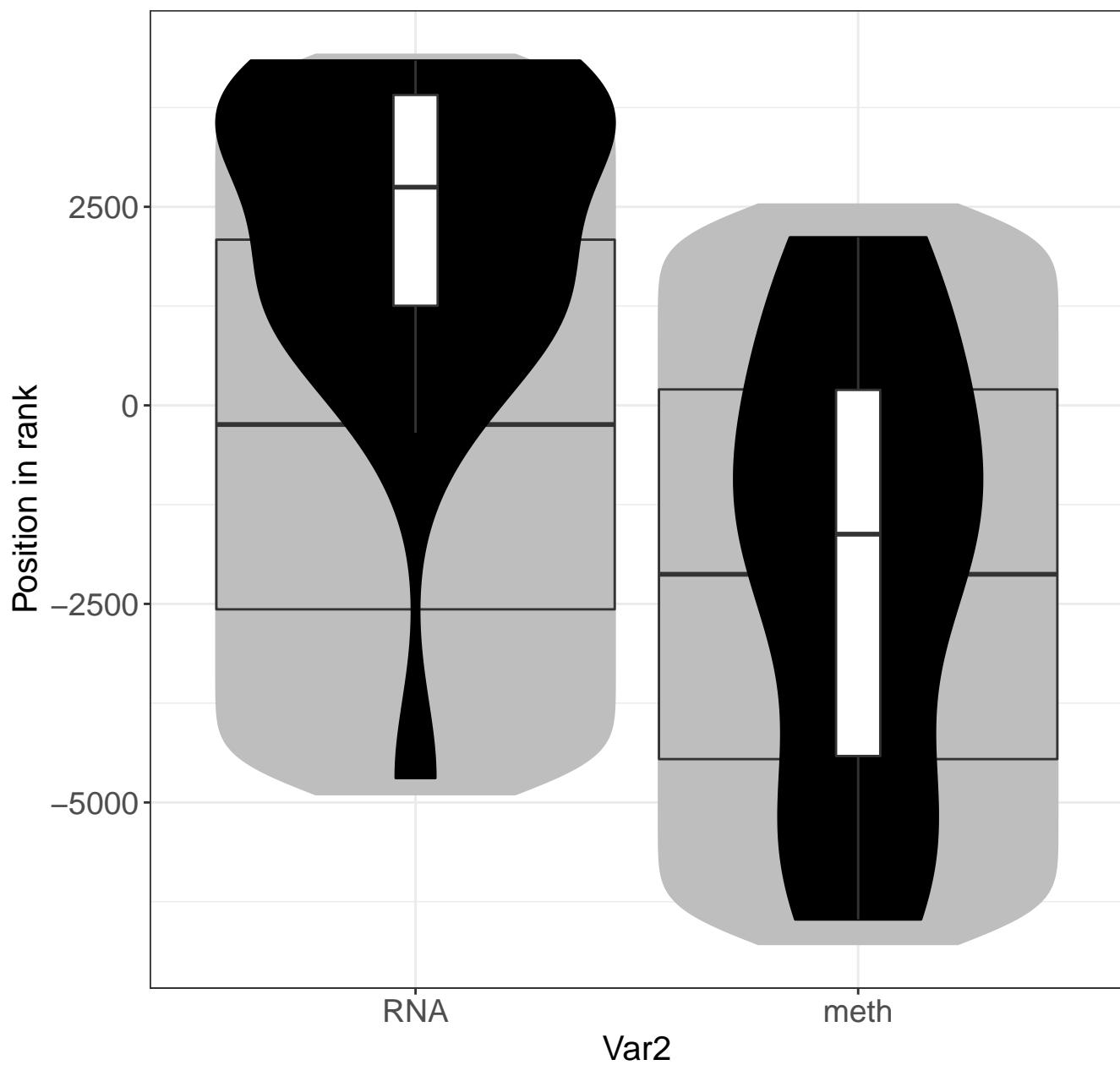
DK5 triggers multiple neurodegenerative pathways in Alzheimer



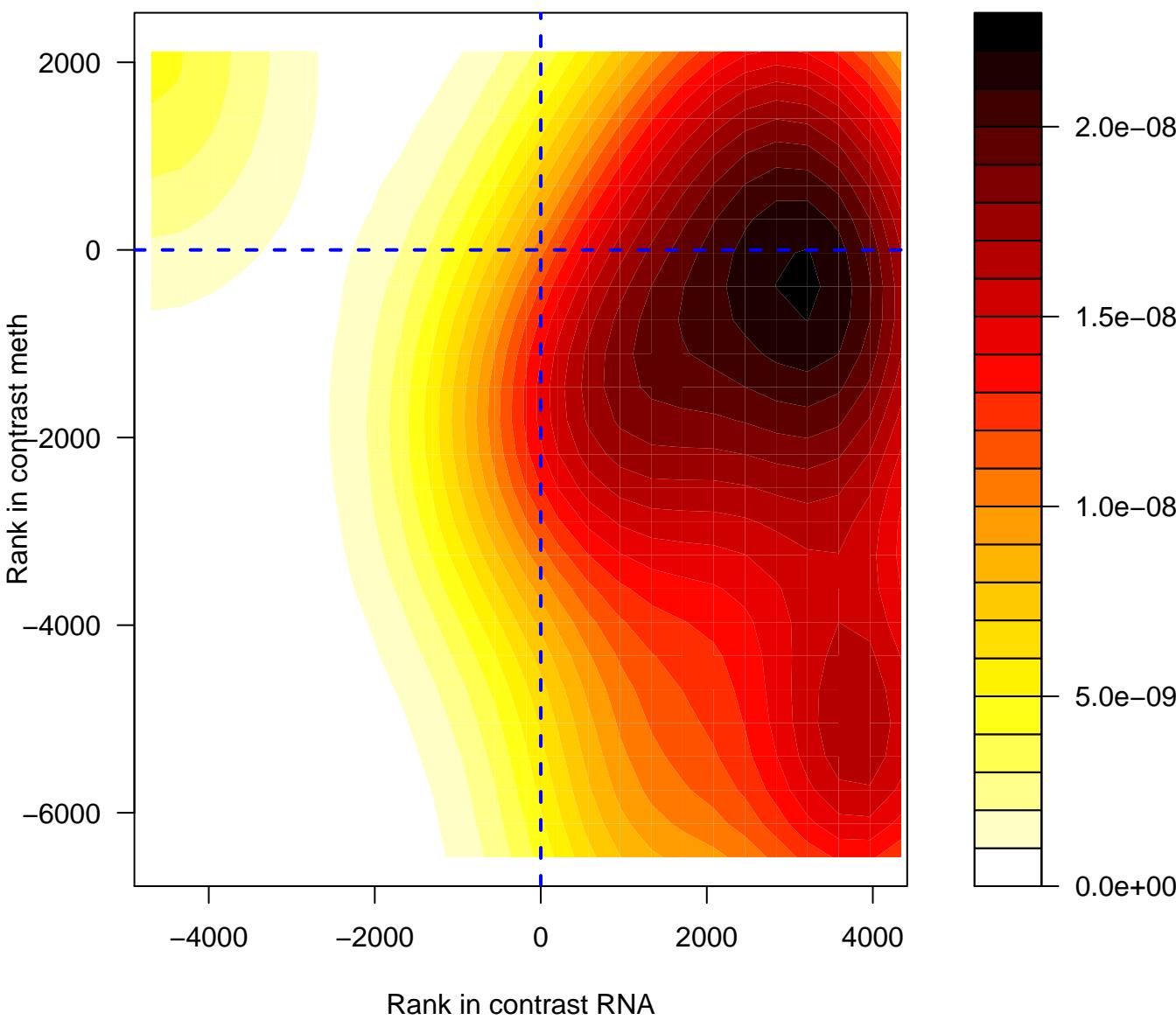
Regulated CDK5 triggers multiple neurodegenerative pathways in Alzheimer's disease



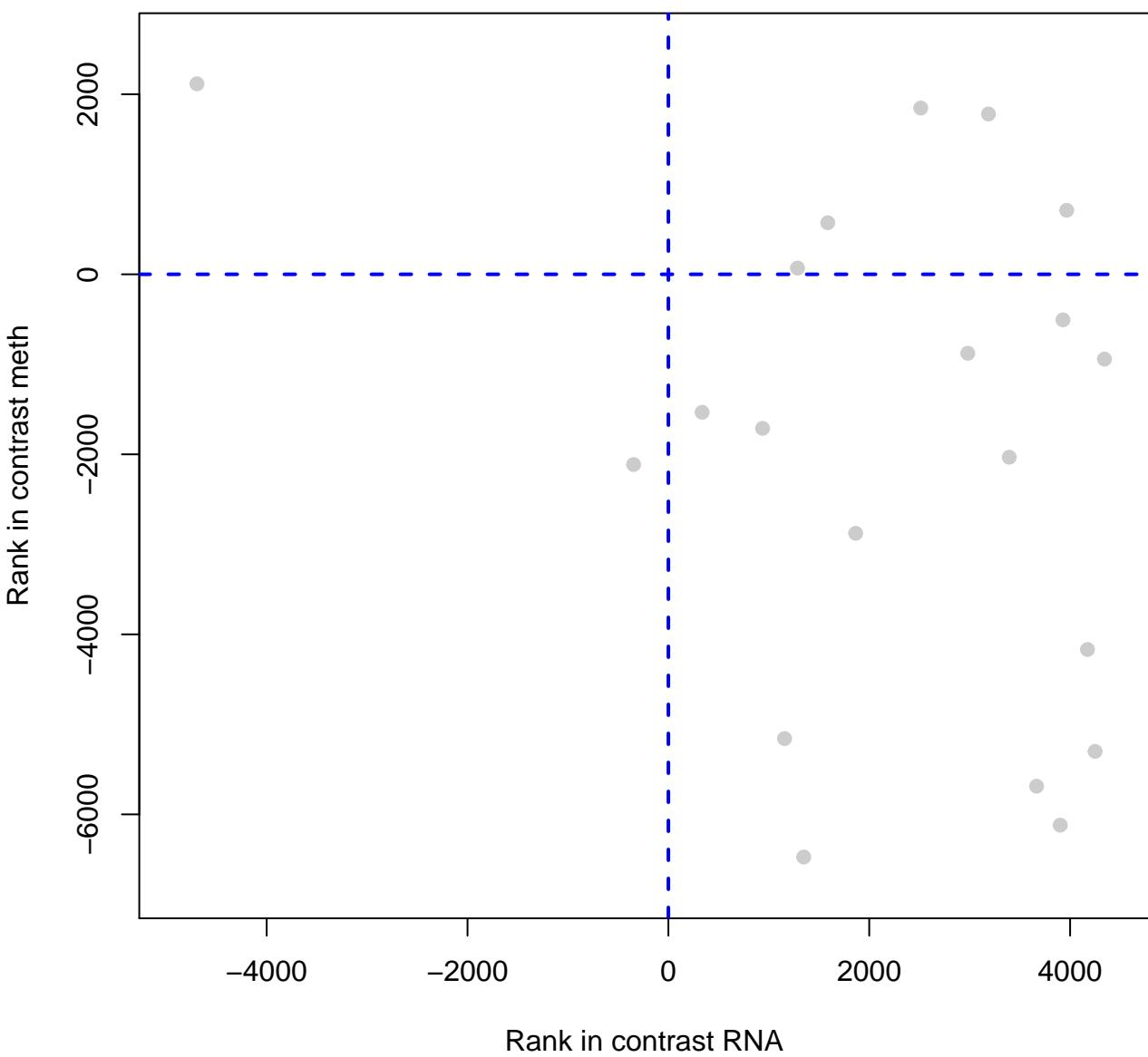
Deregulated CDK5 triggers multiple neurodegener



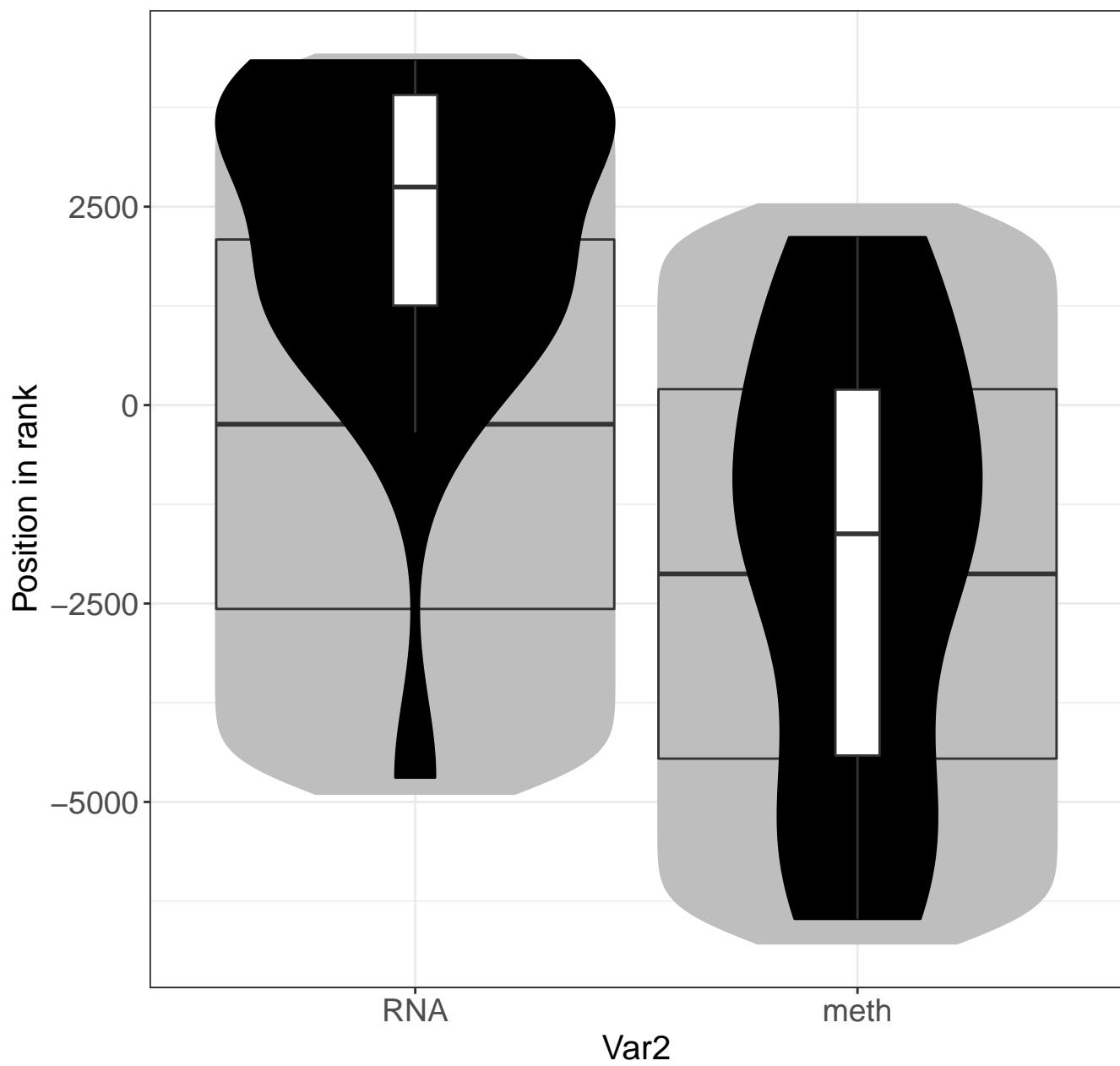
Neurodegenerative Diseases



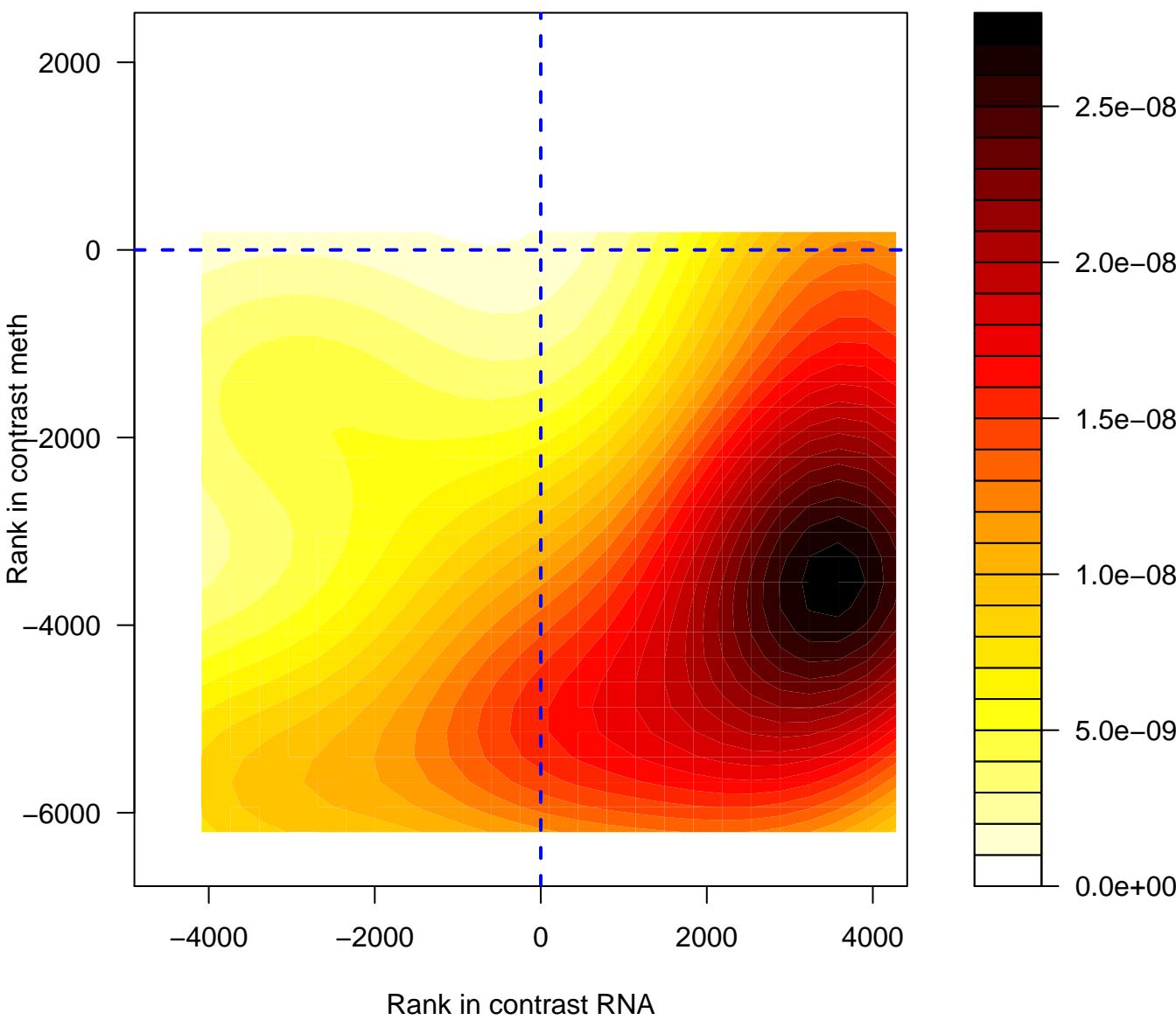
Neurodegenerative Diseases



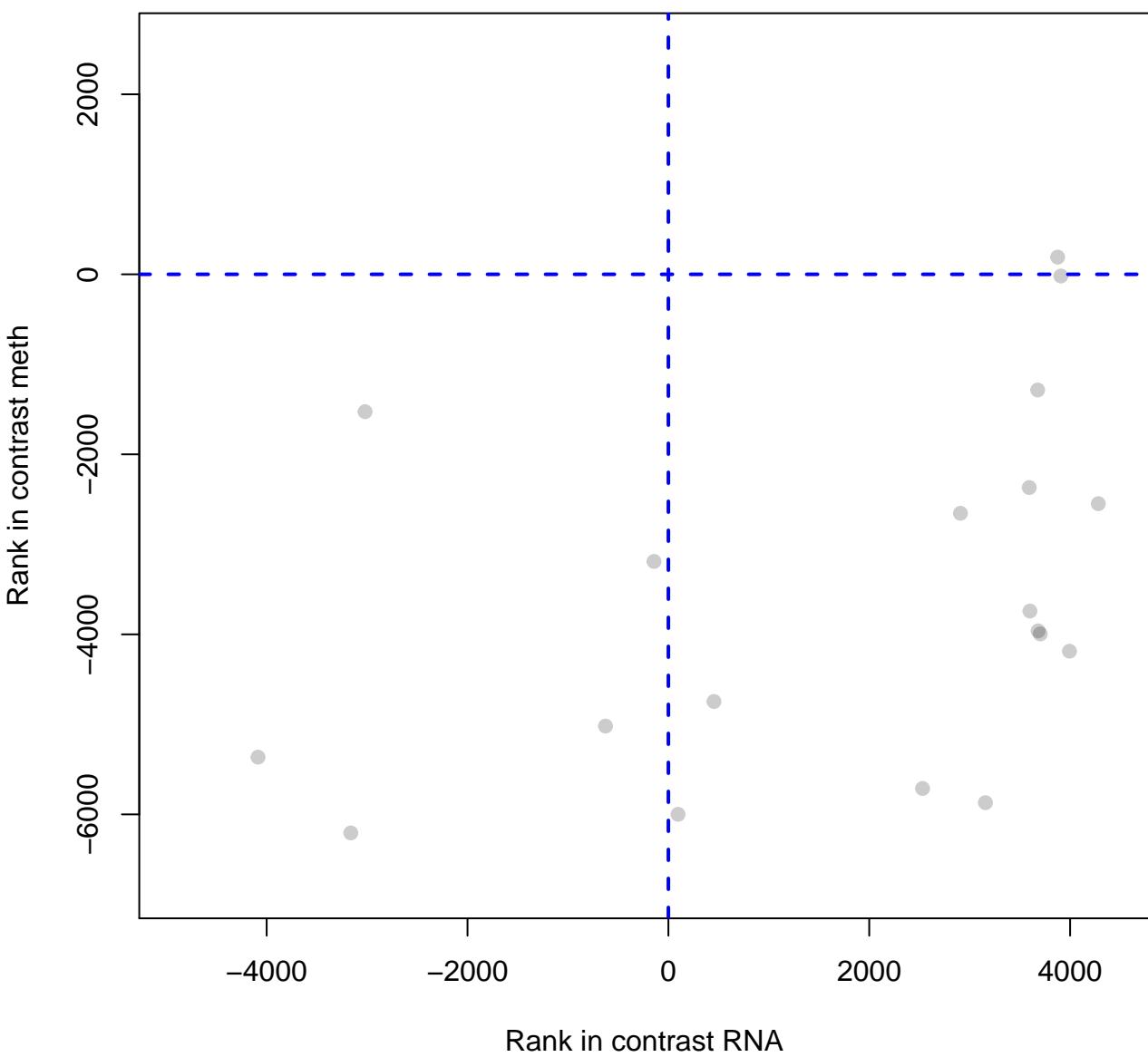
Neurodegenerative Diseases



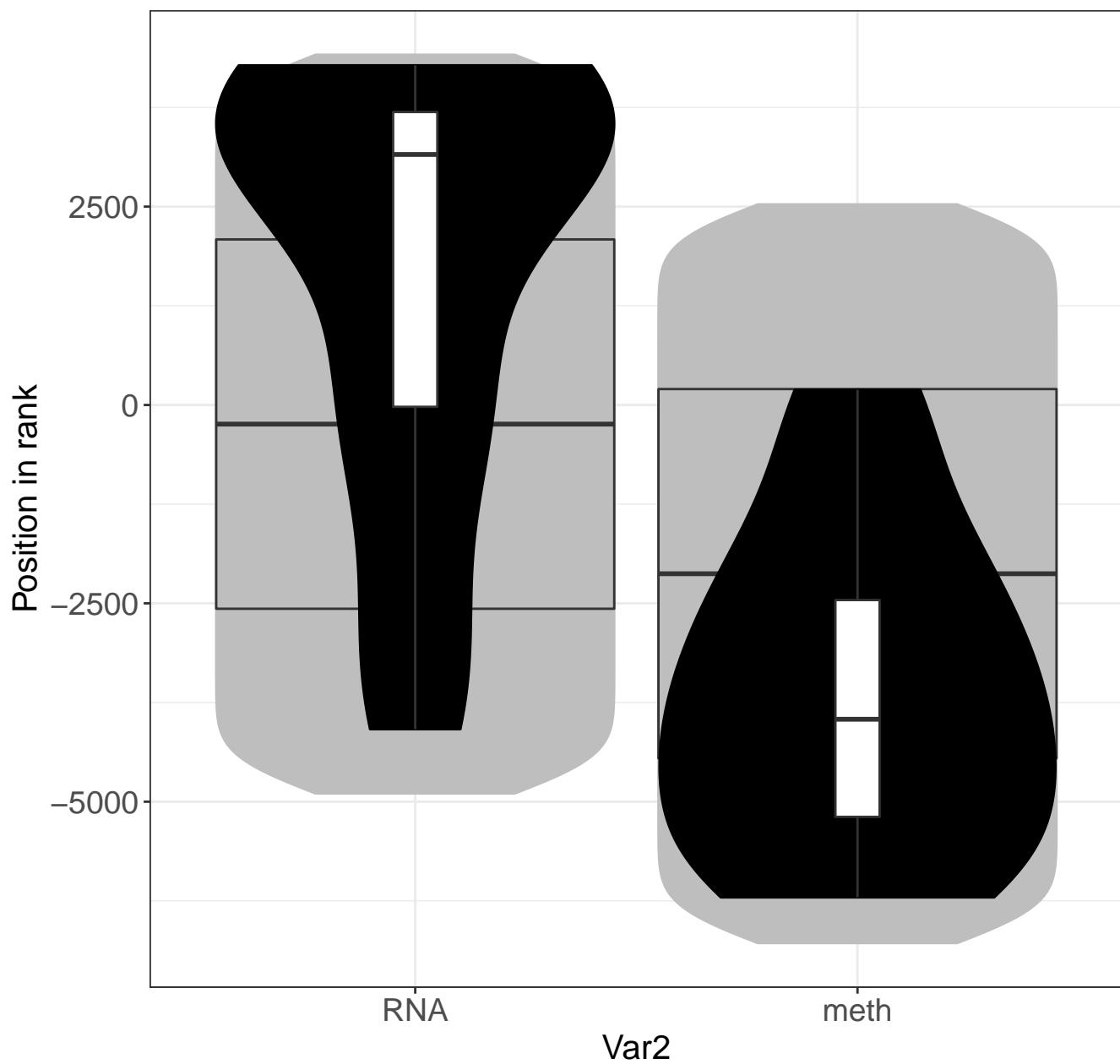
Smooth Muscle Contraction



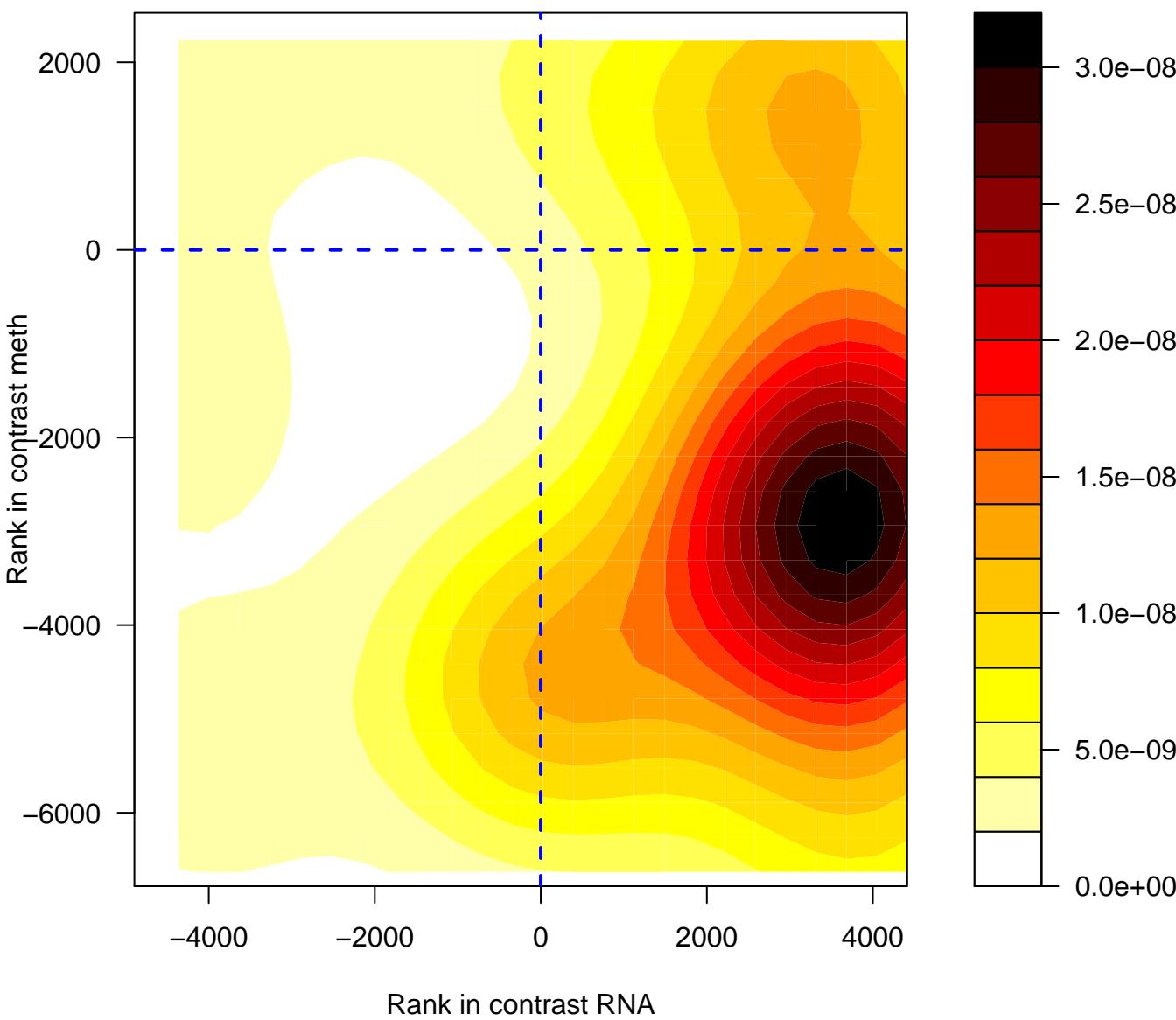
Smooth Muscle Contraction



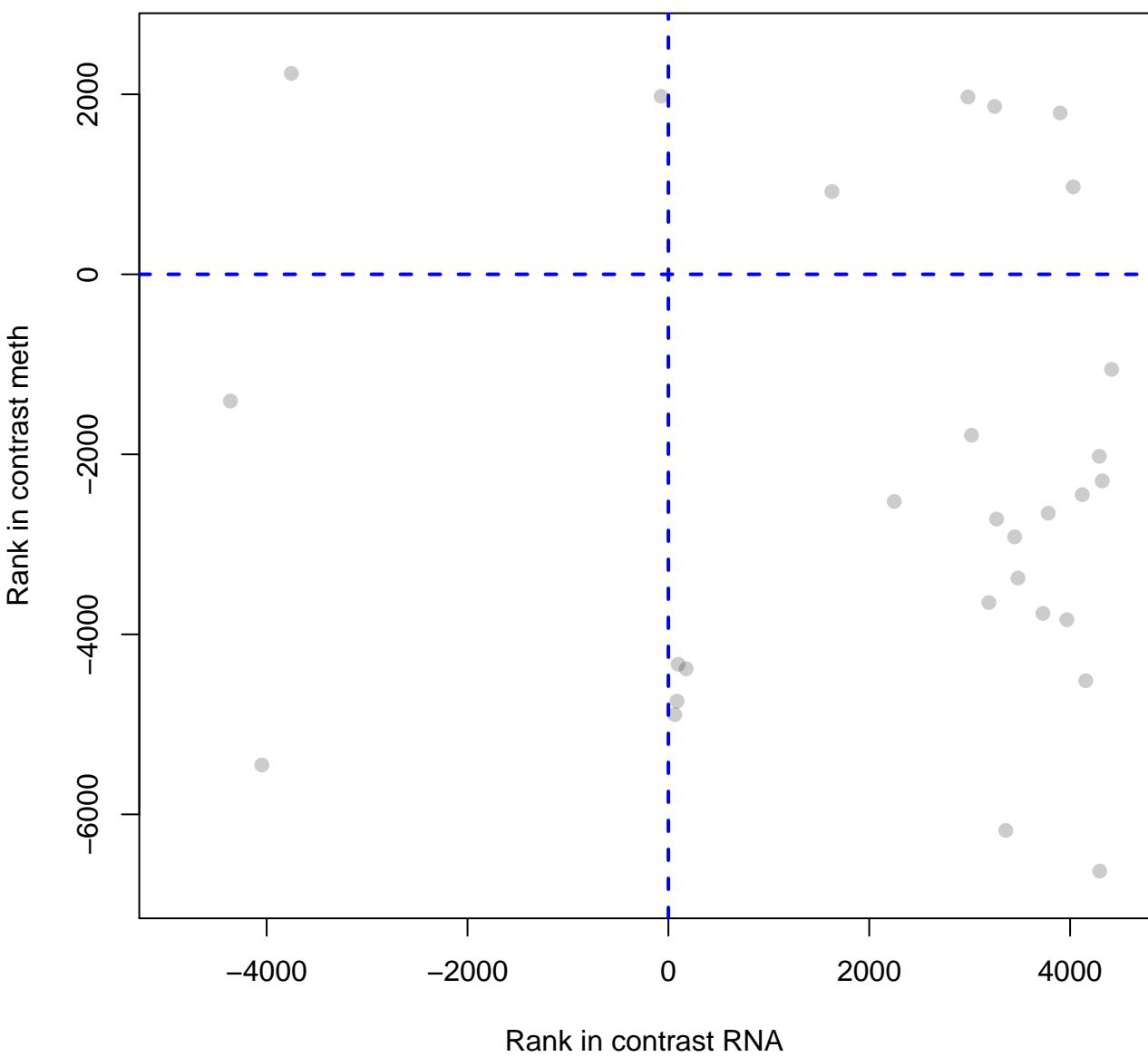
Smooth Muscle Contraction



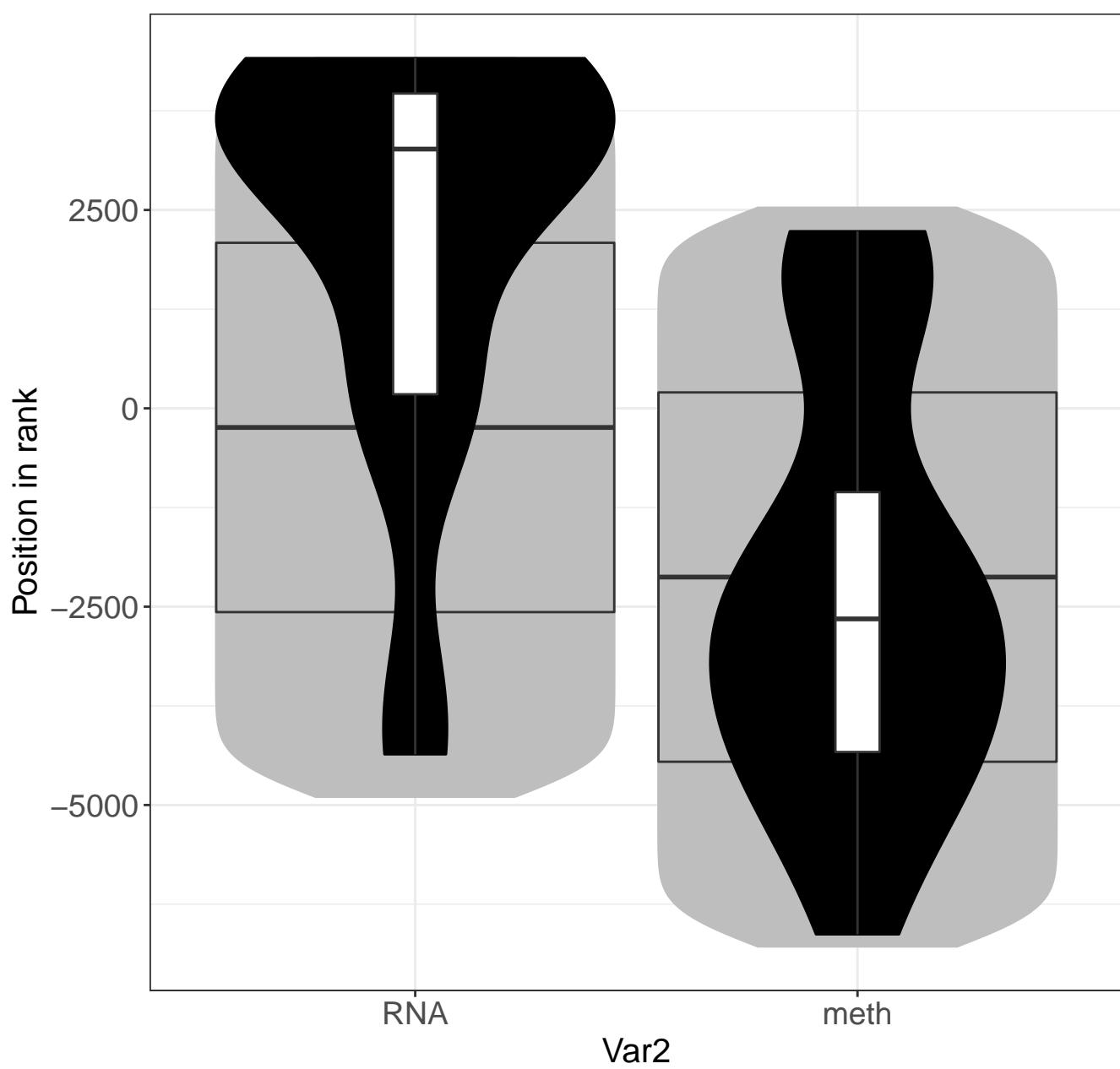
RHO GTPases Activate WASPs and WAVEs



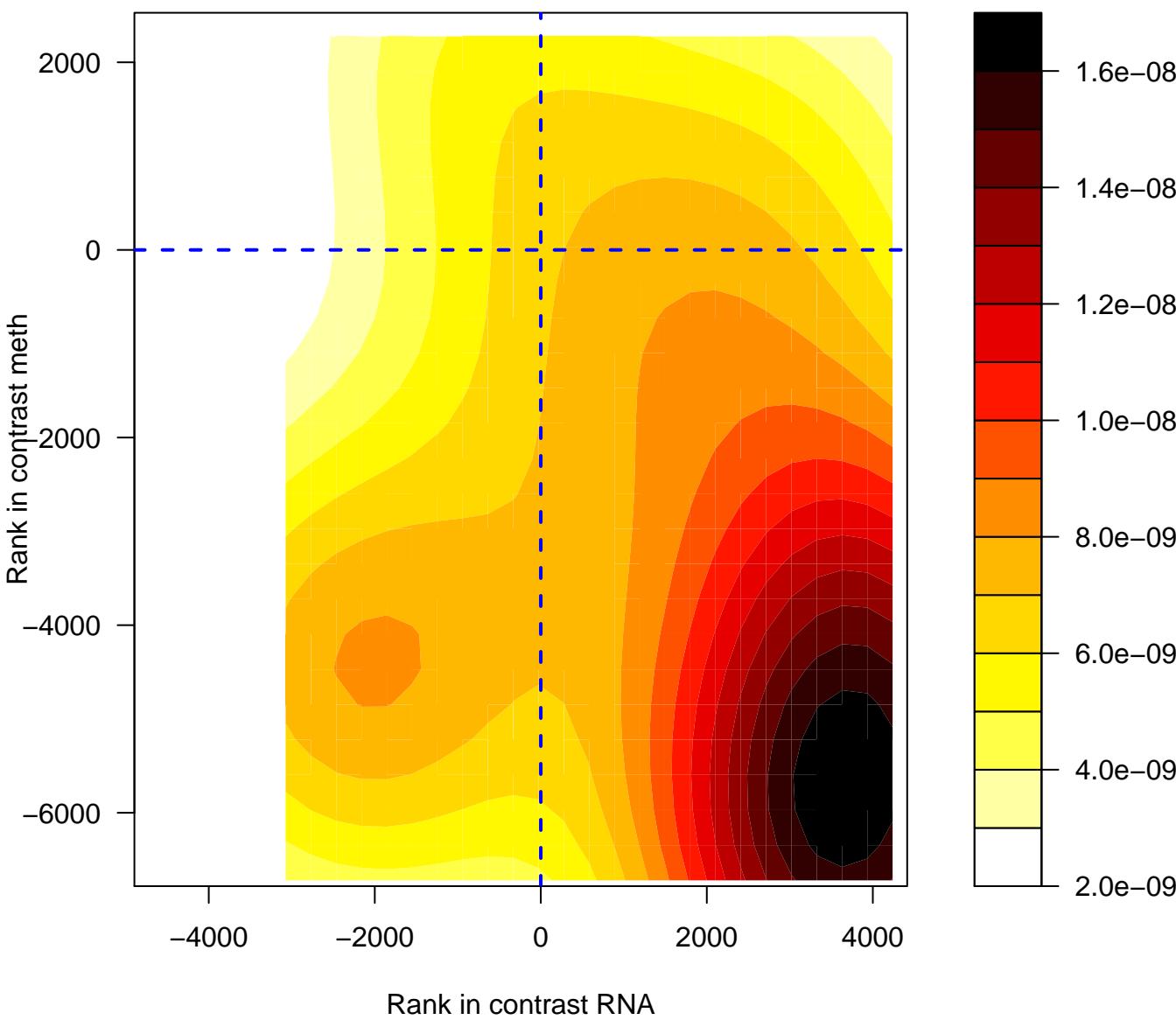
RHO GTPases Activate WASPs and WAVEs



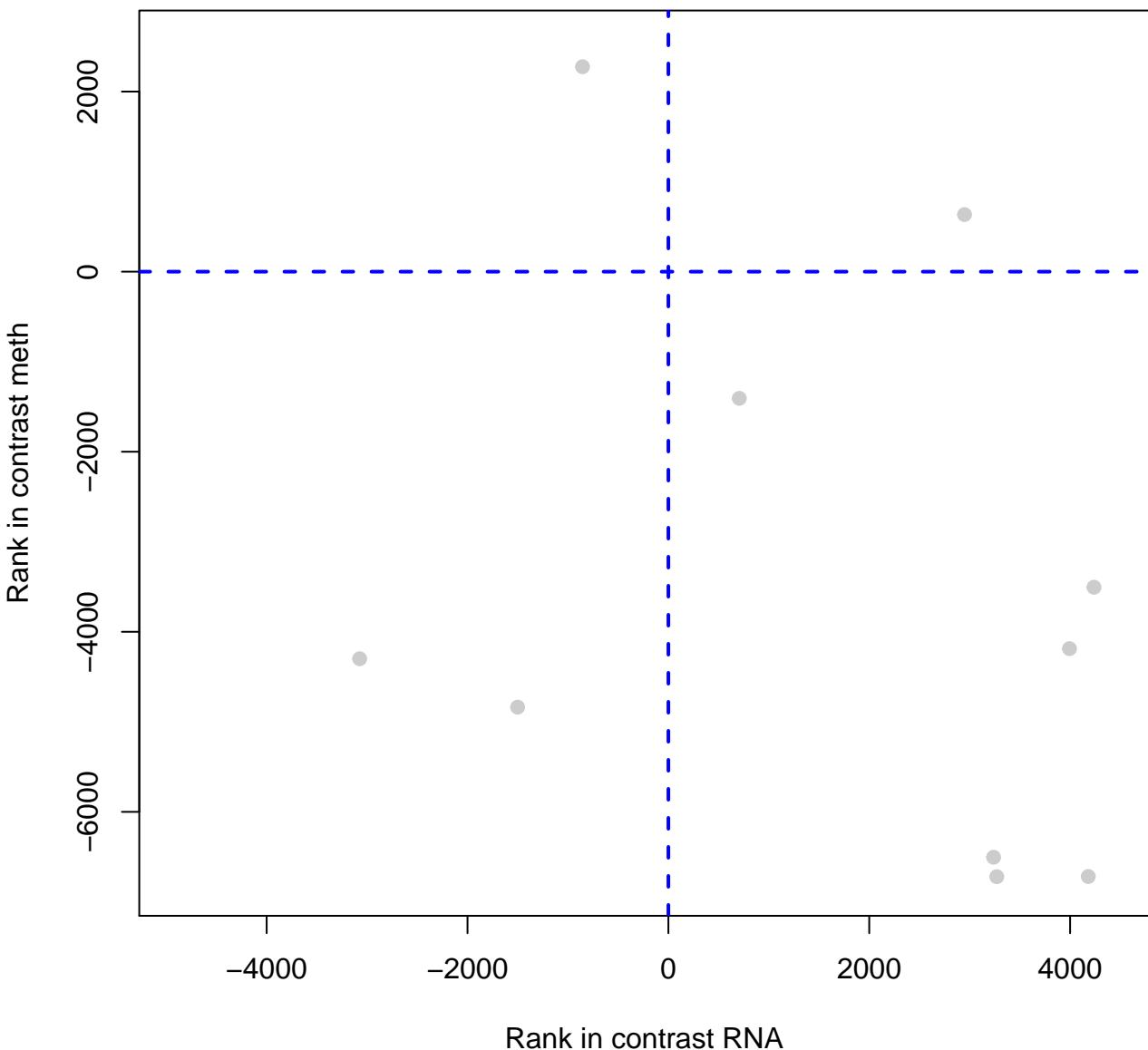
RHO GTPases Activate WASPs and WAVEs



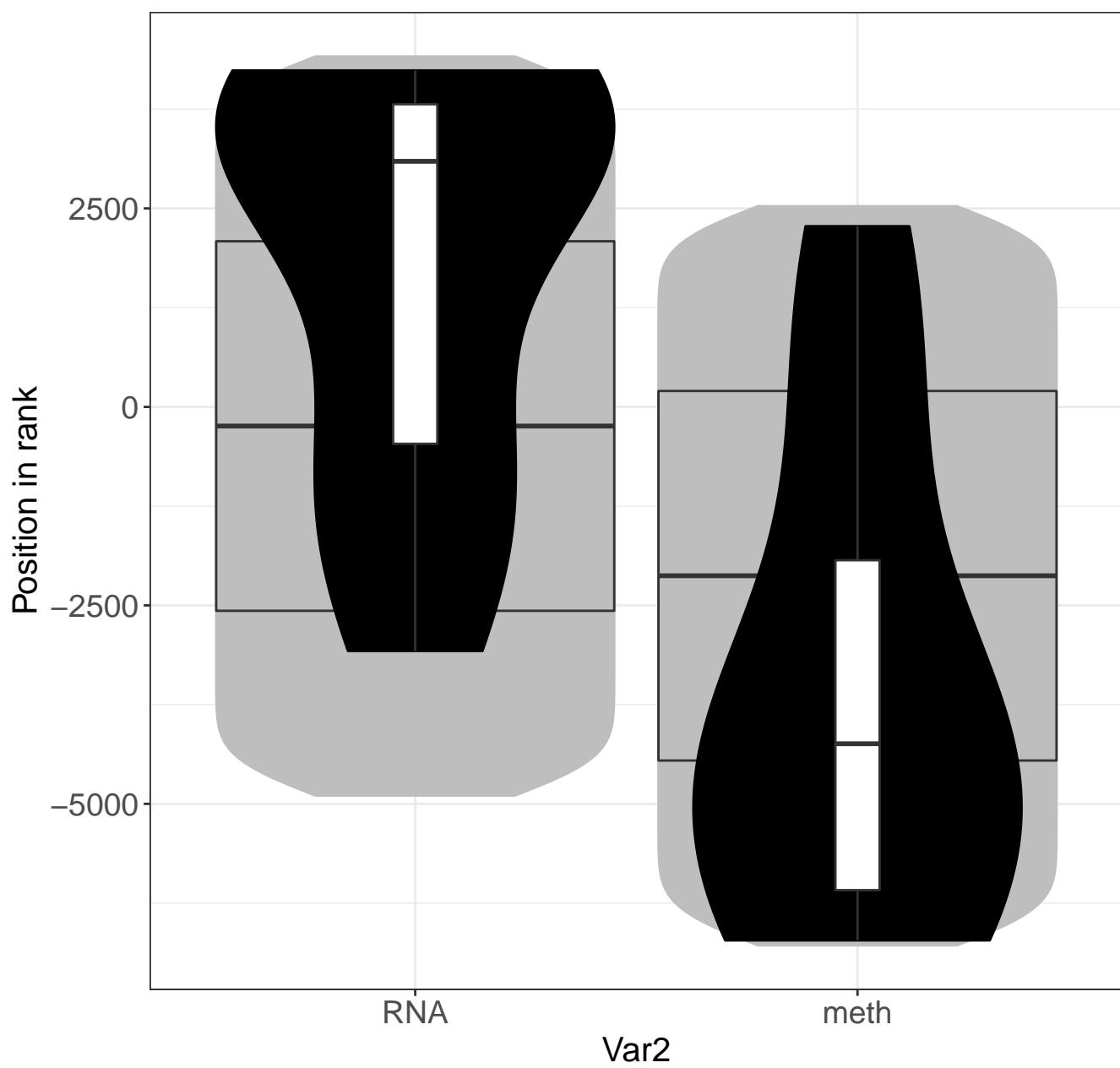
Other semaphorin interactions



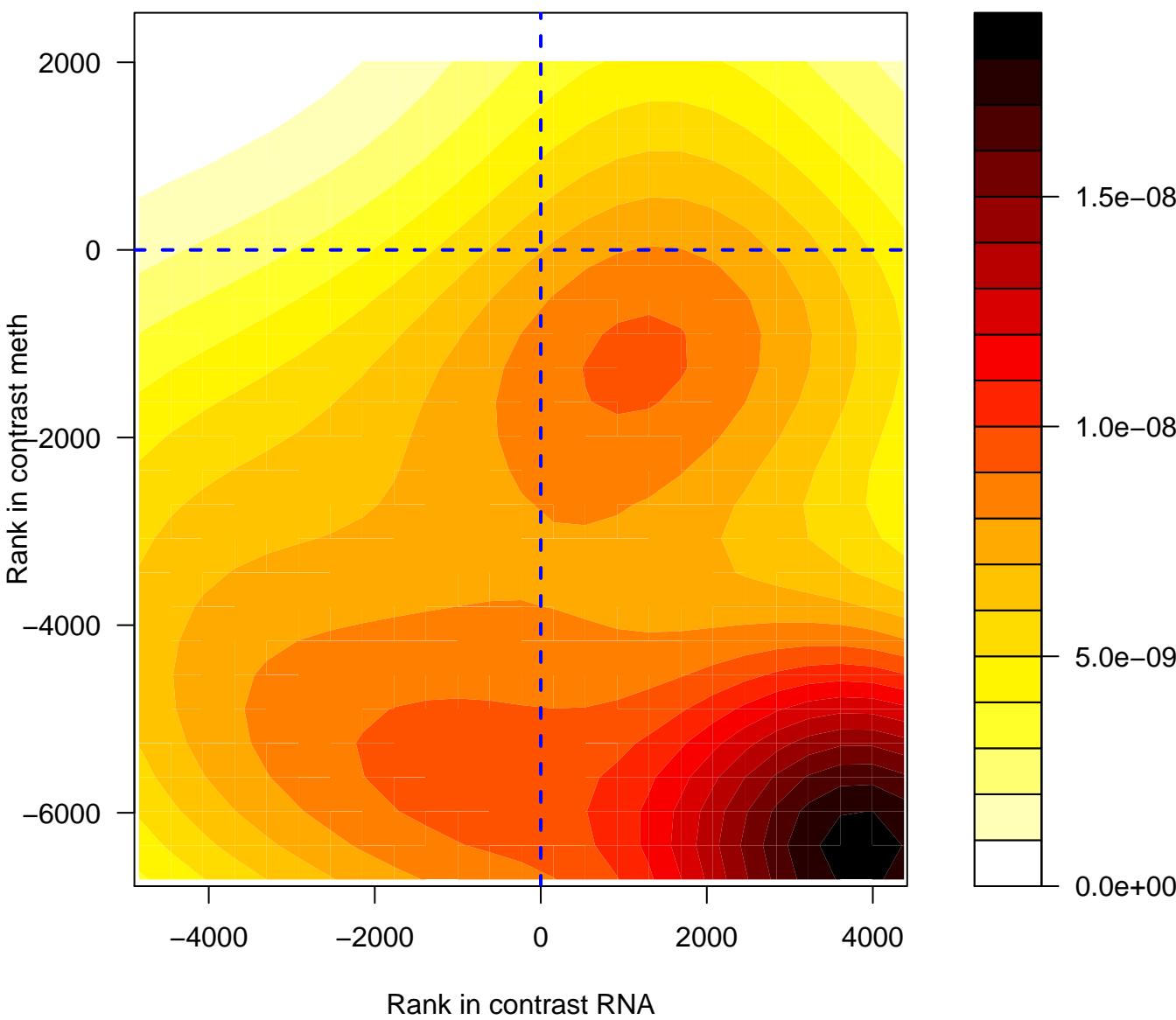
Other semaphorin interactions



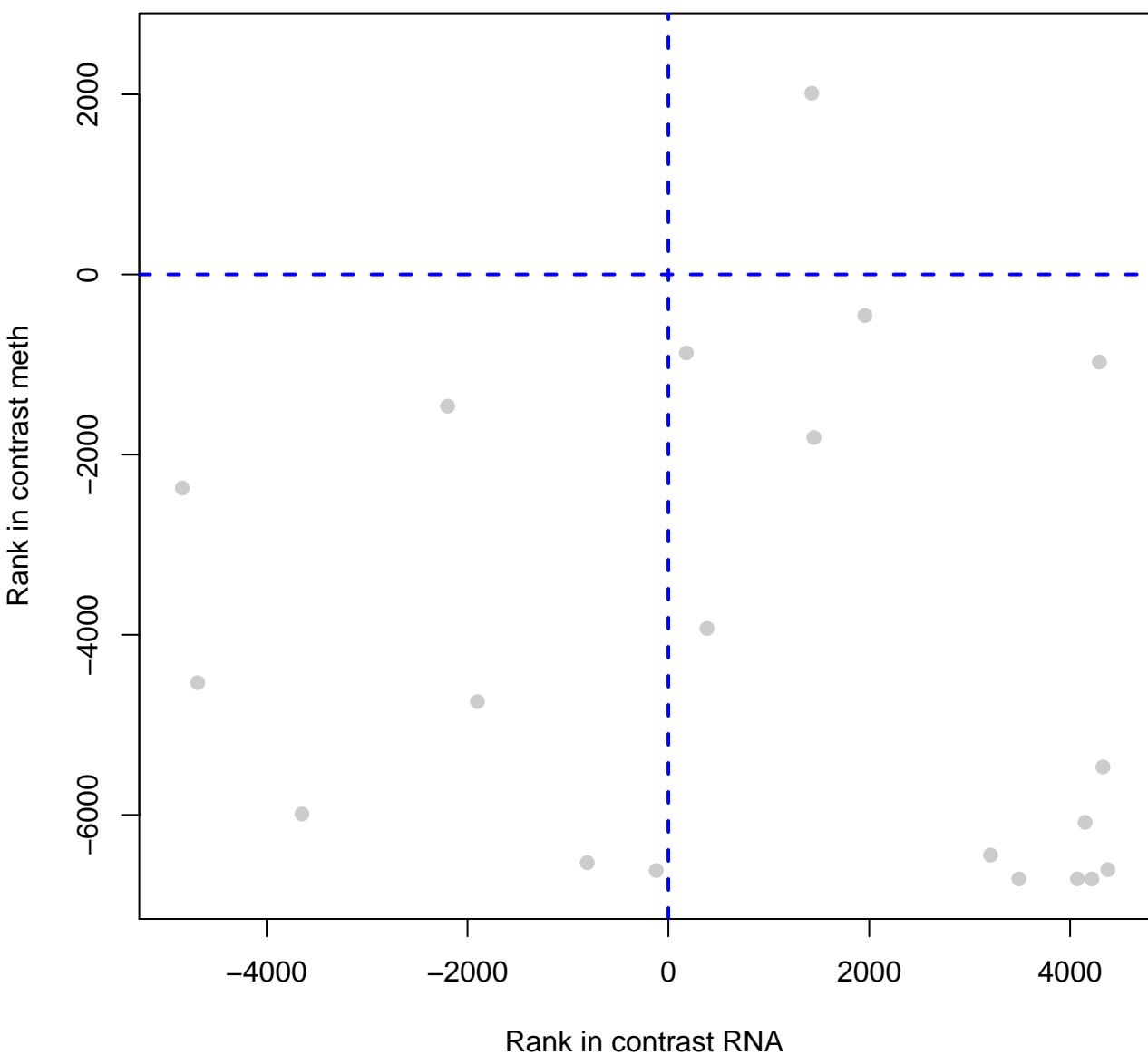
Other semaphorin interactions



Arachidonic acid metabolism



Arachidonic acid metabolism



Arachidonic acid metabolism

