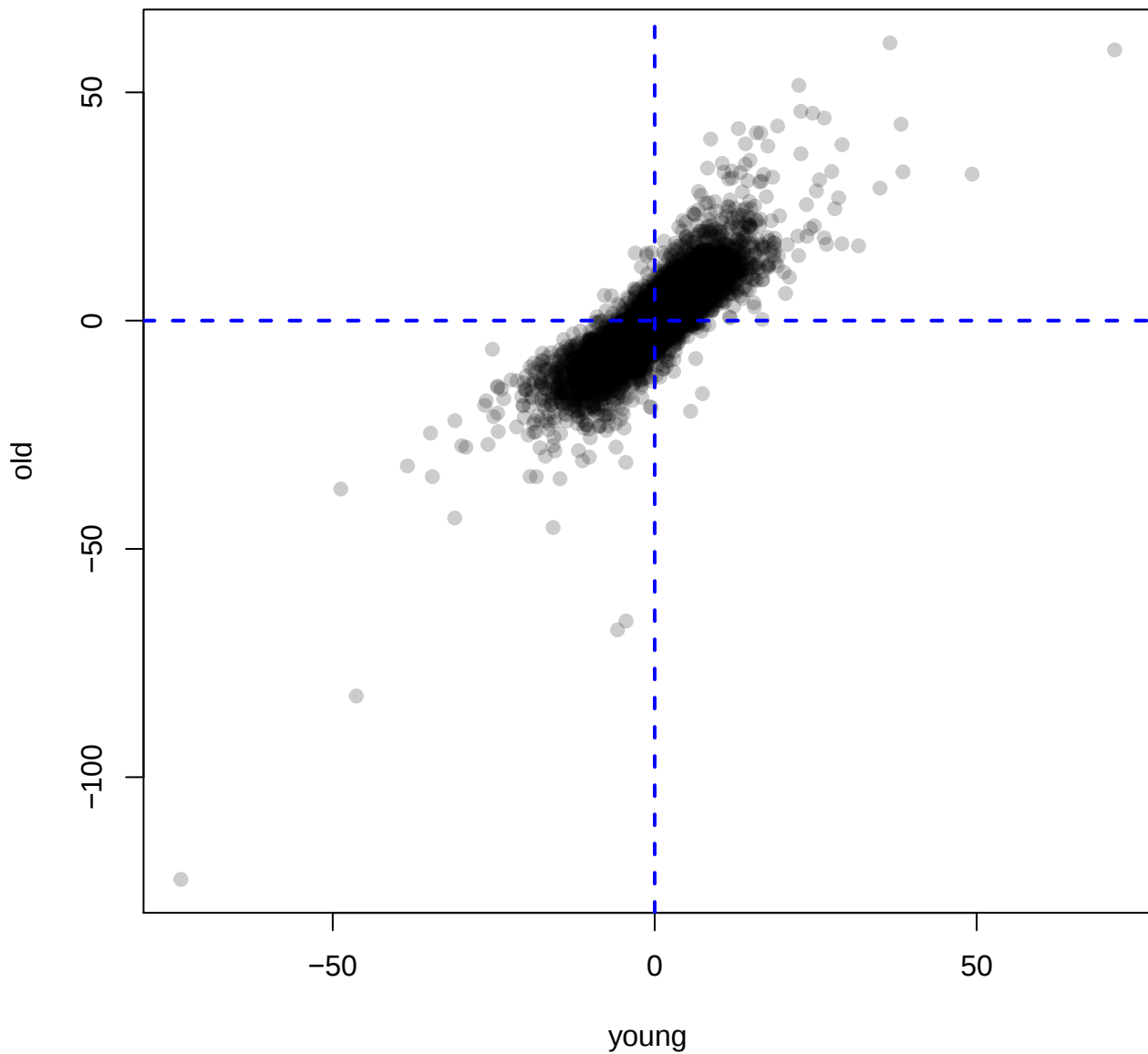
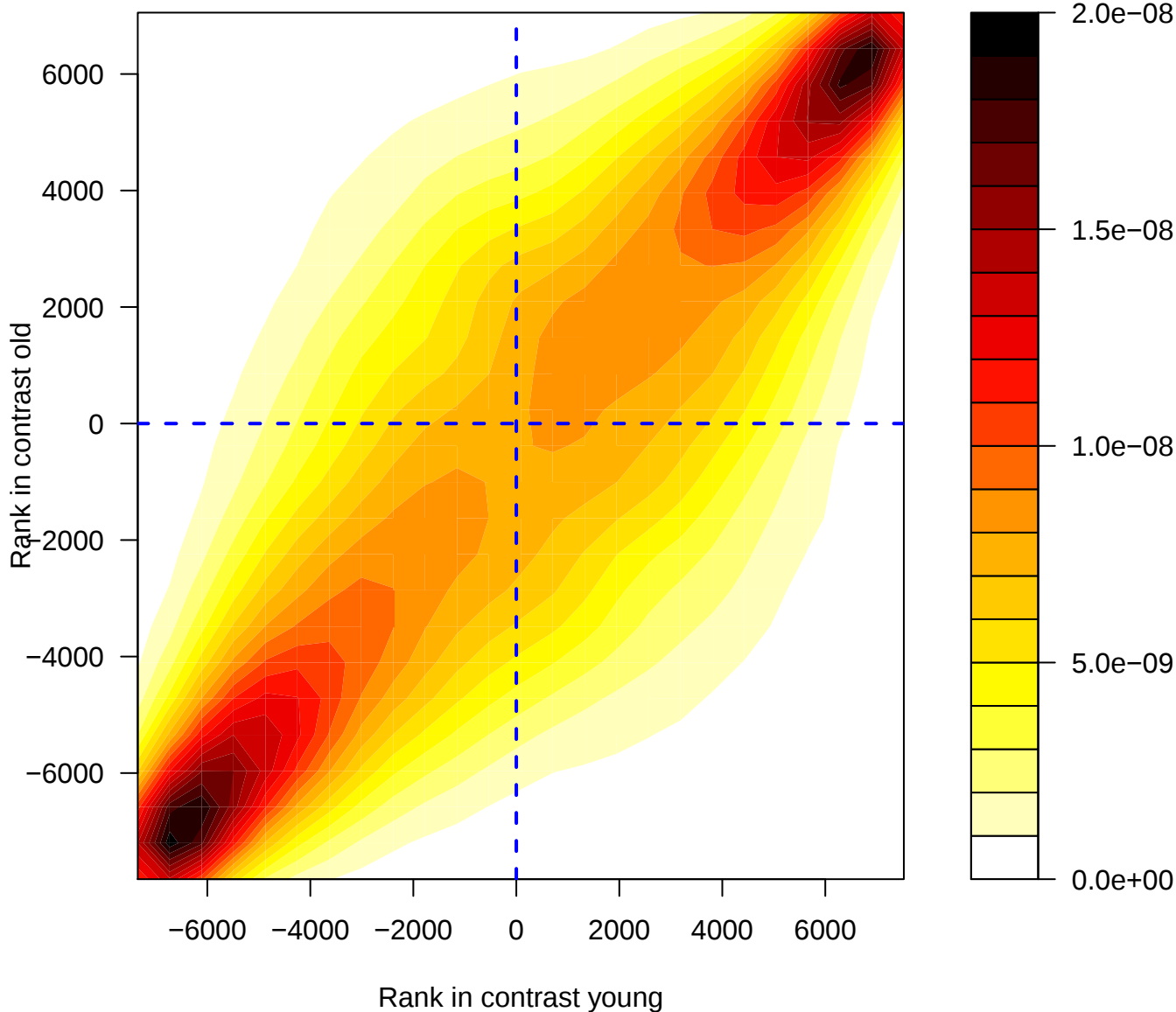


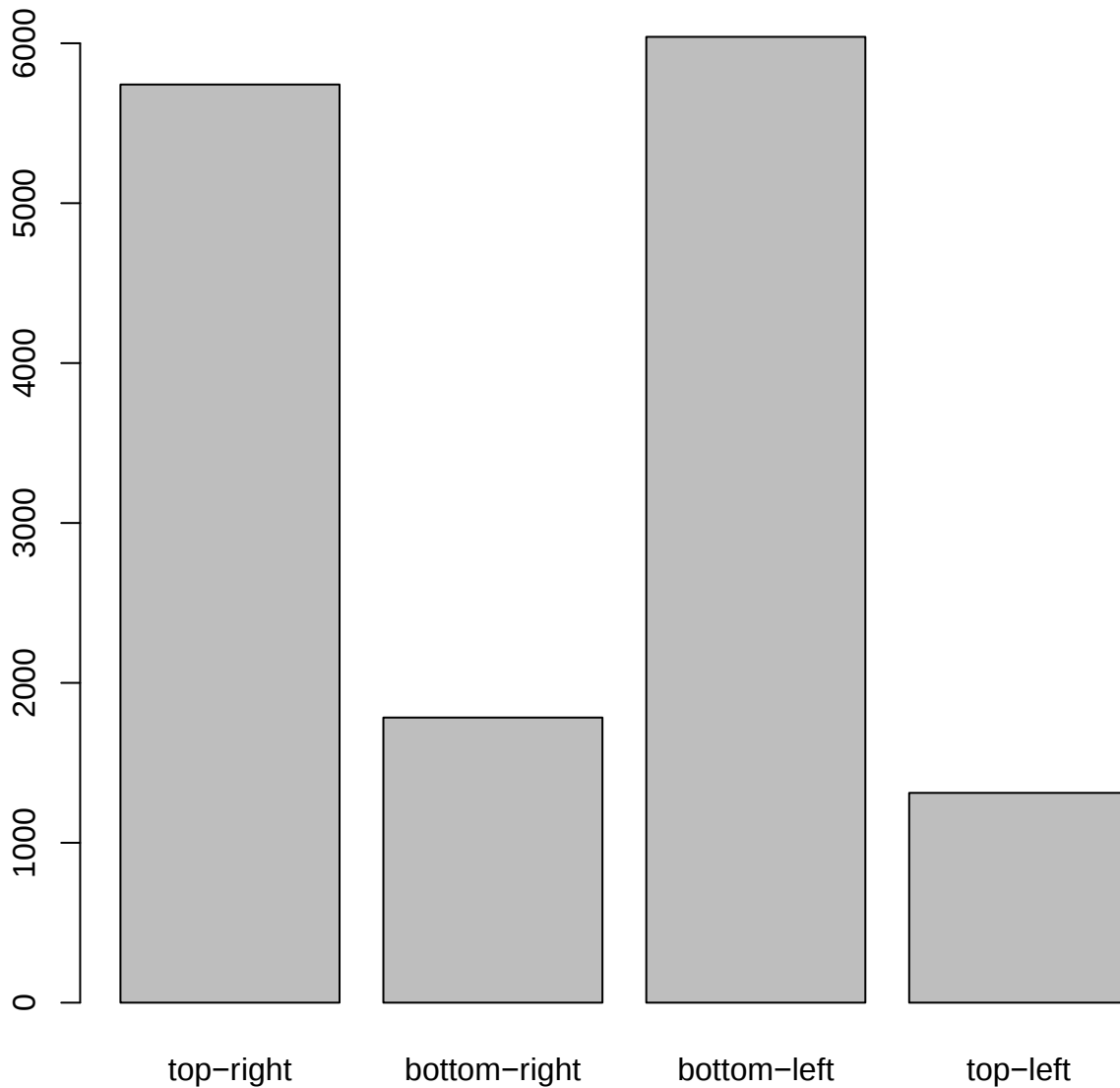
Scatterplot of all genes



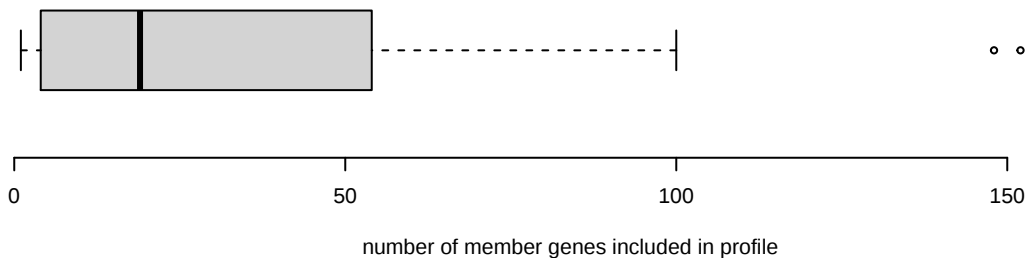
Rank–rank plot of all genes



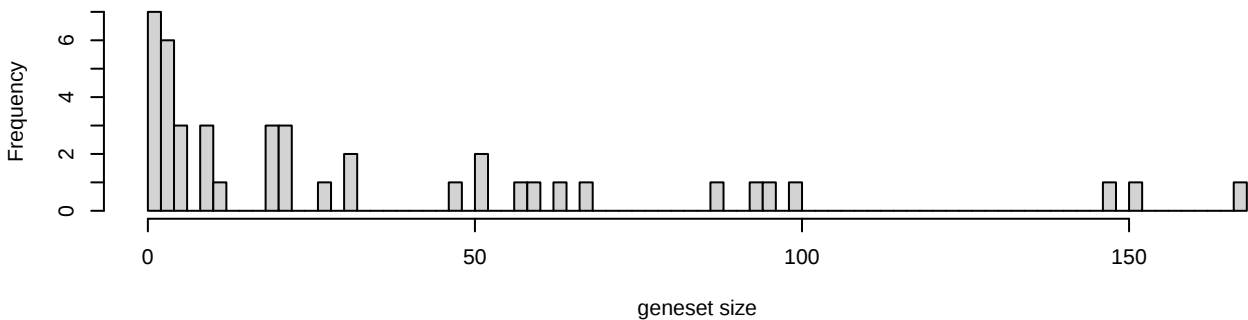
number of genes in each quadrant



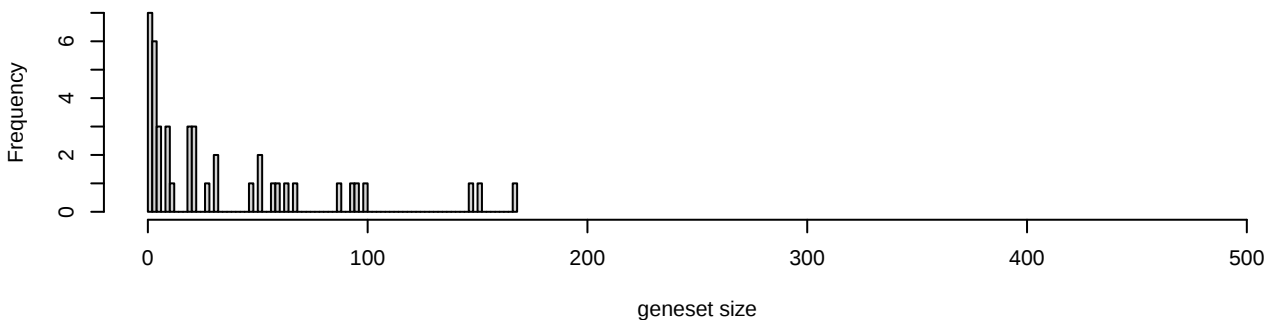
Gene set size



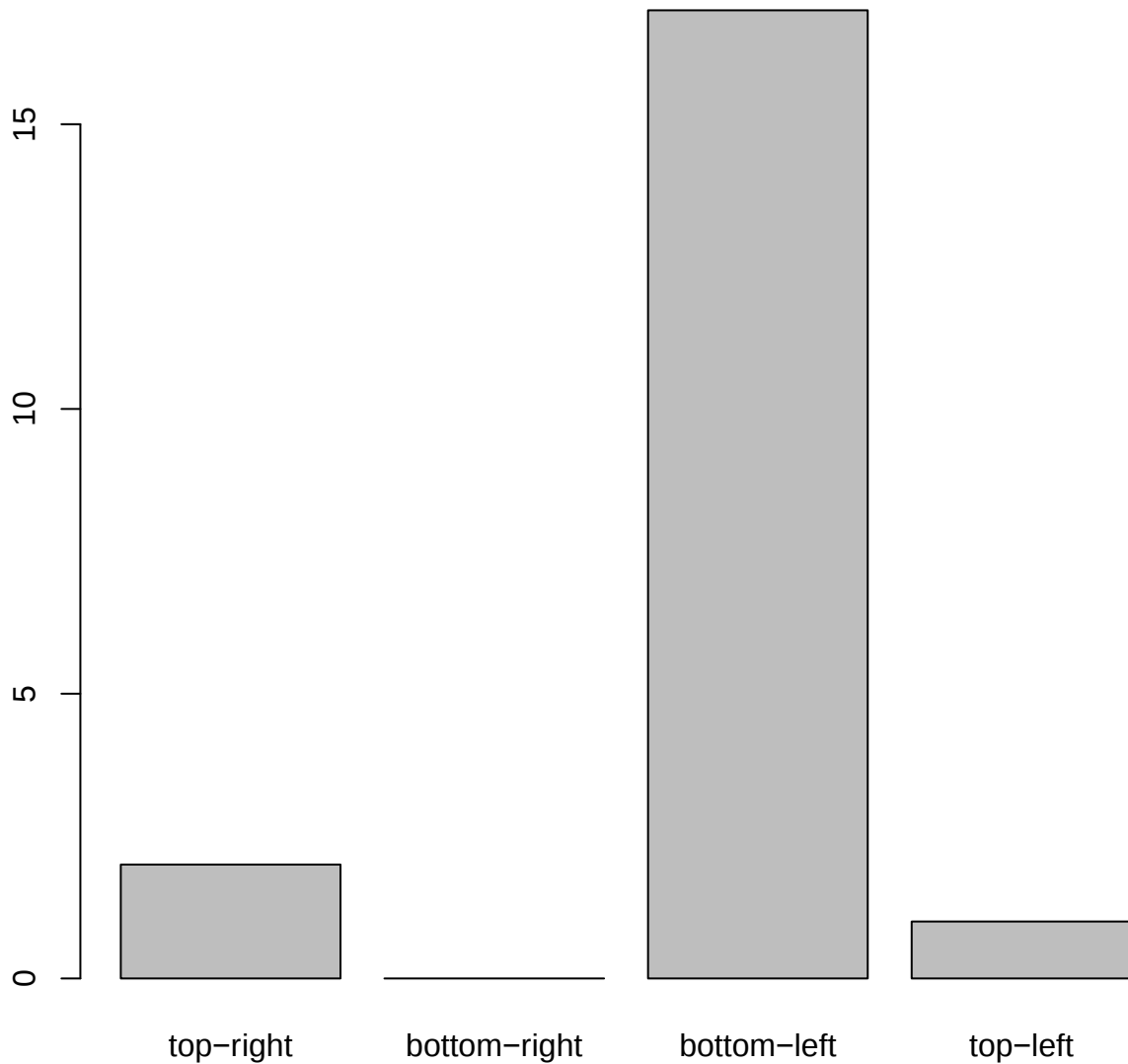
Histogram of geneset size



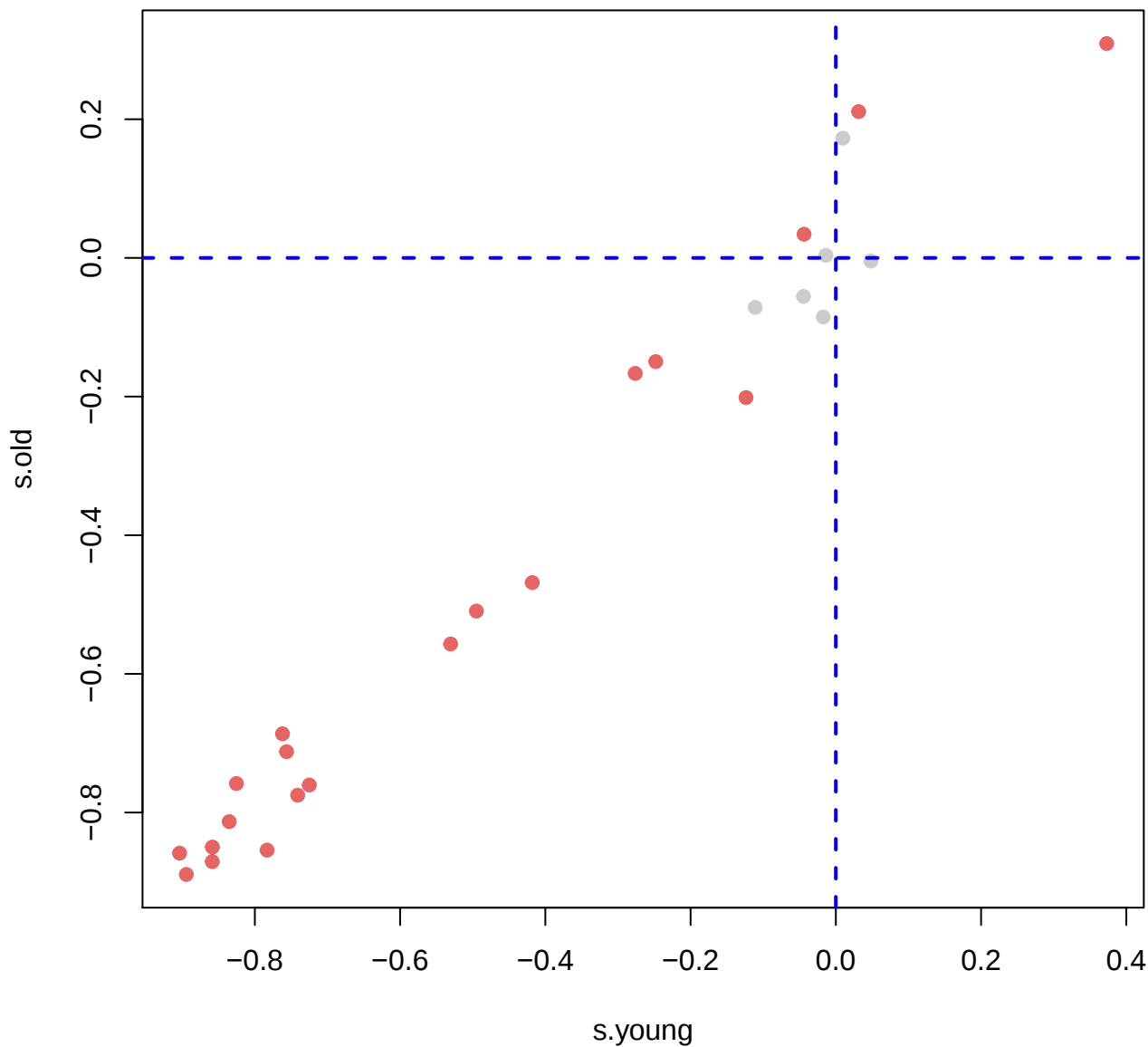
Trimmed histogram of geneset size



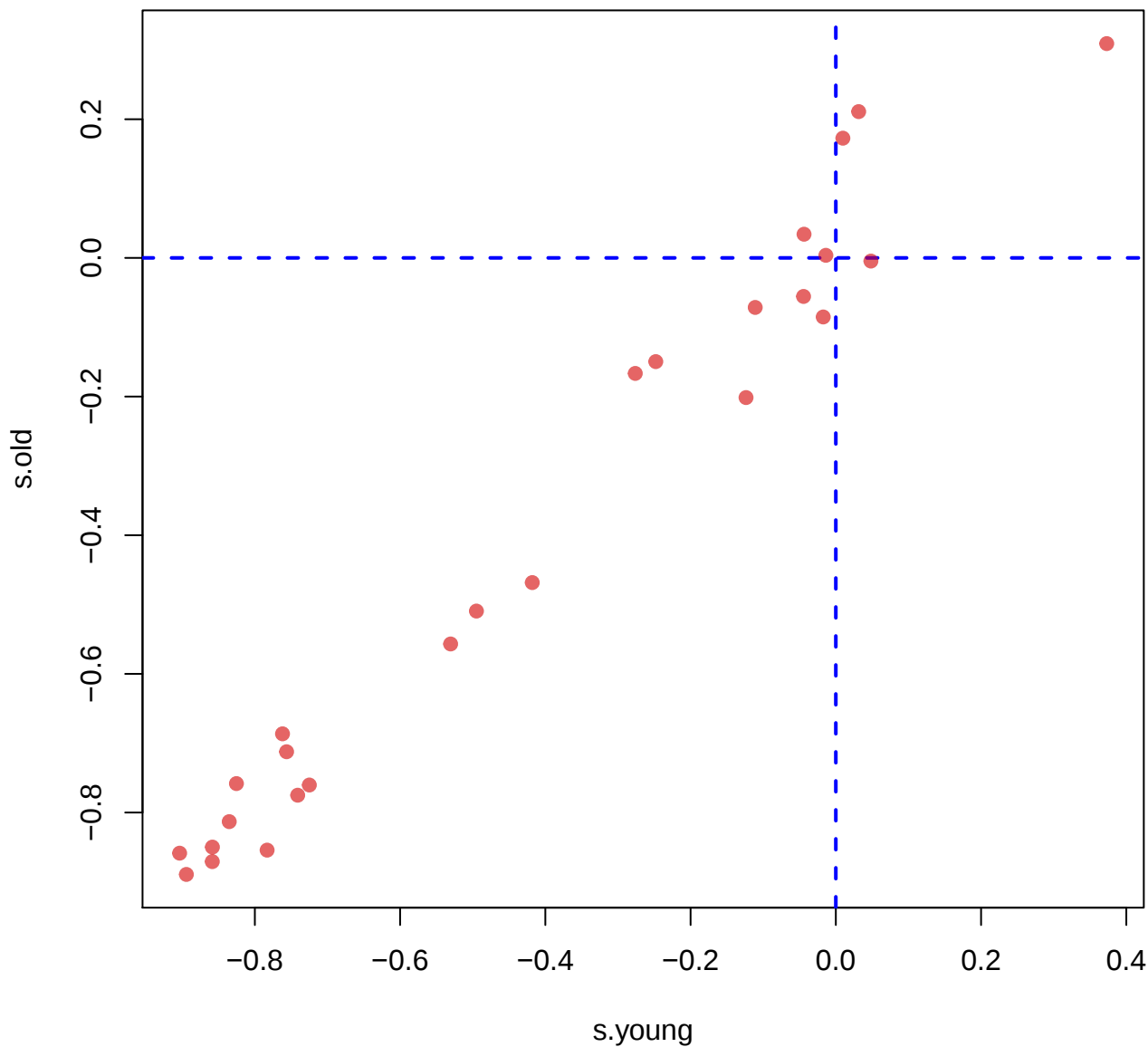
number of genesets FDR<0.05



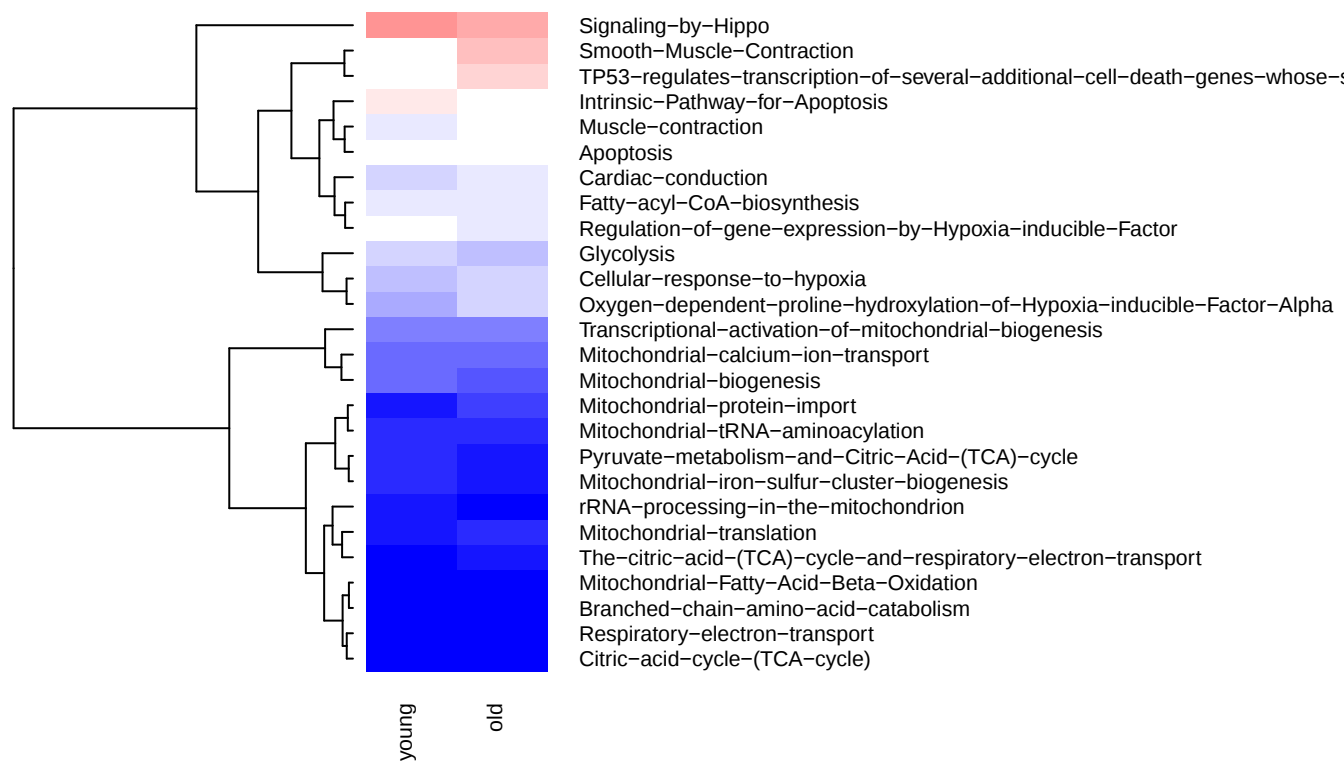
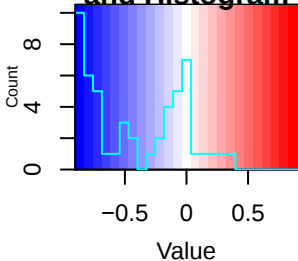
Scatterplot of all gene sets; FDR<0.05 in red



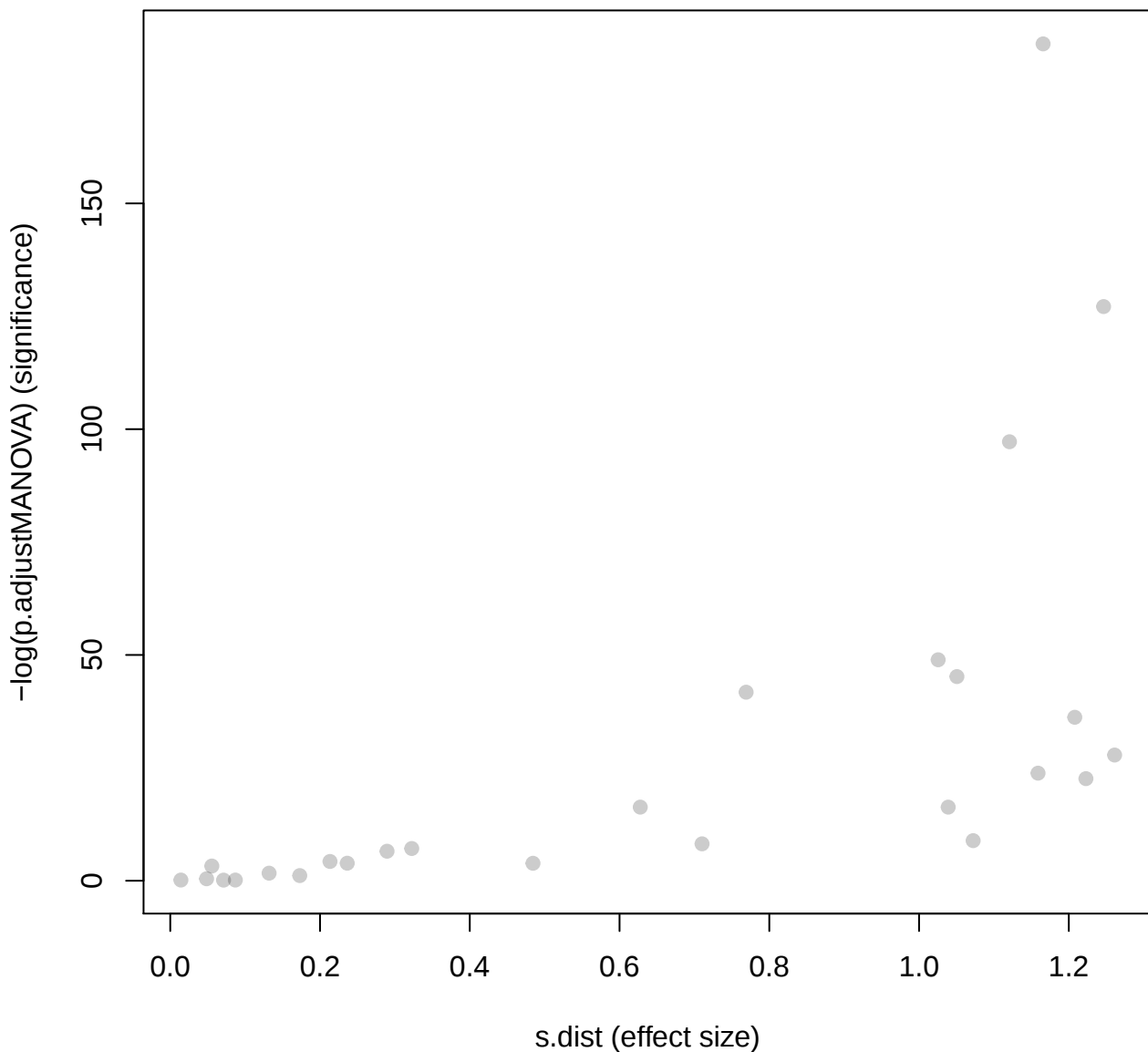
Scatterplot of all gene sets; top 26 in red



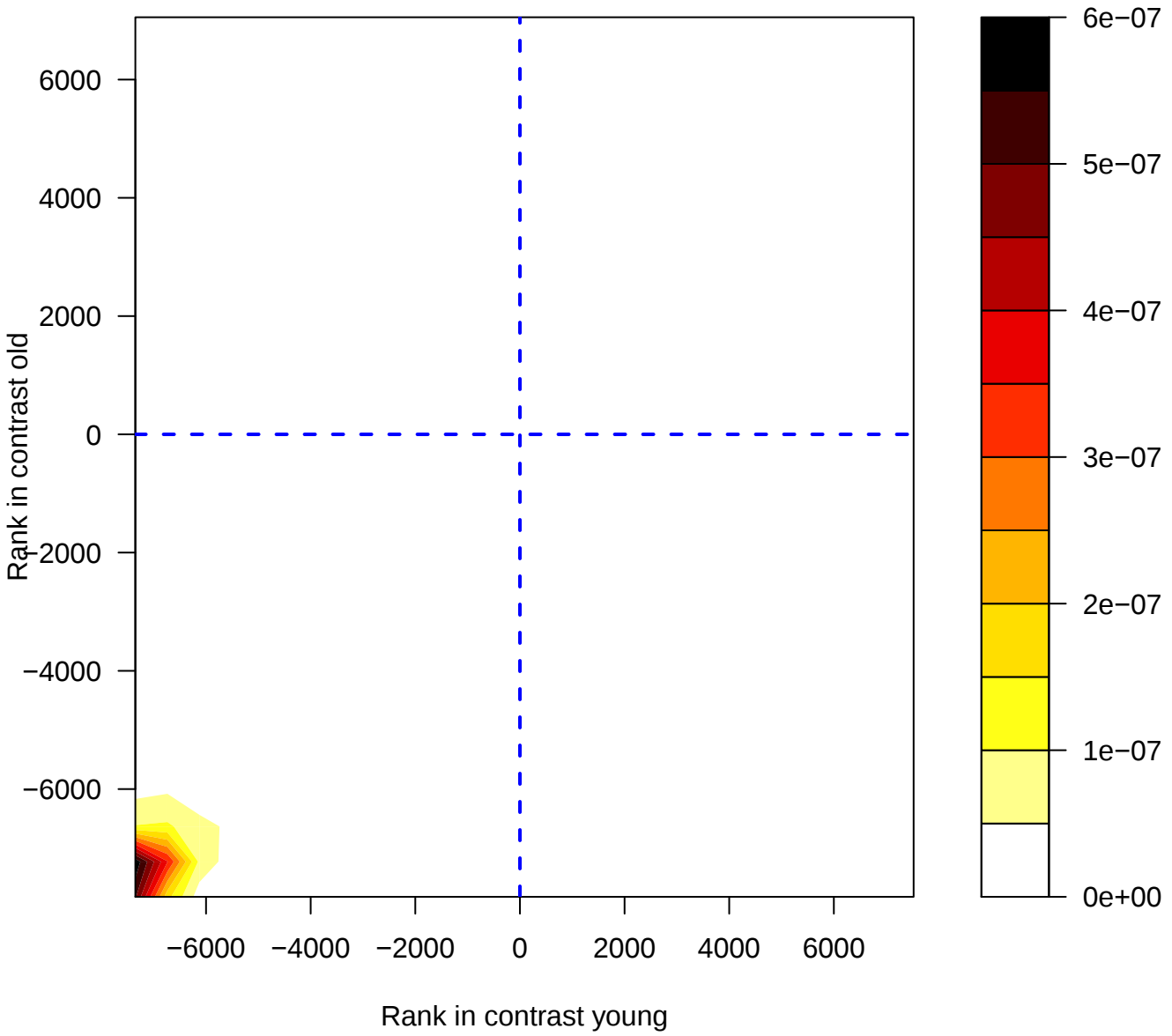
Color Key
and Histogram



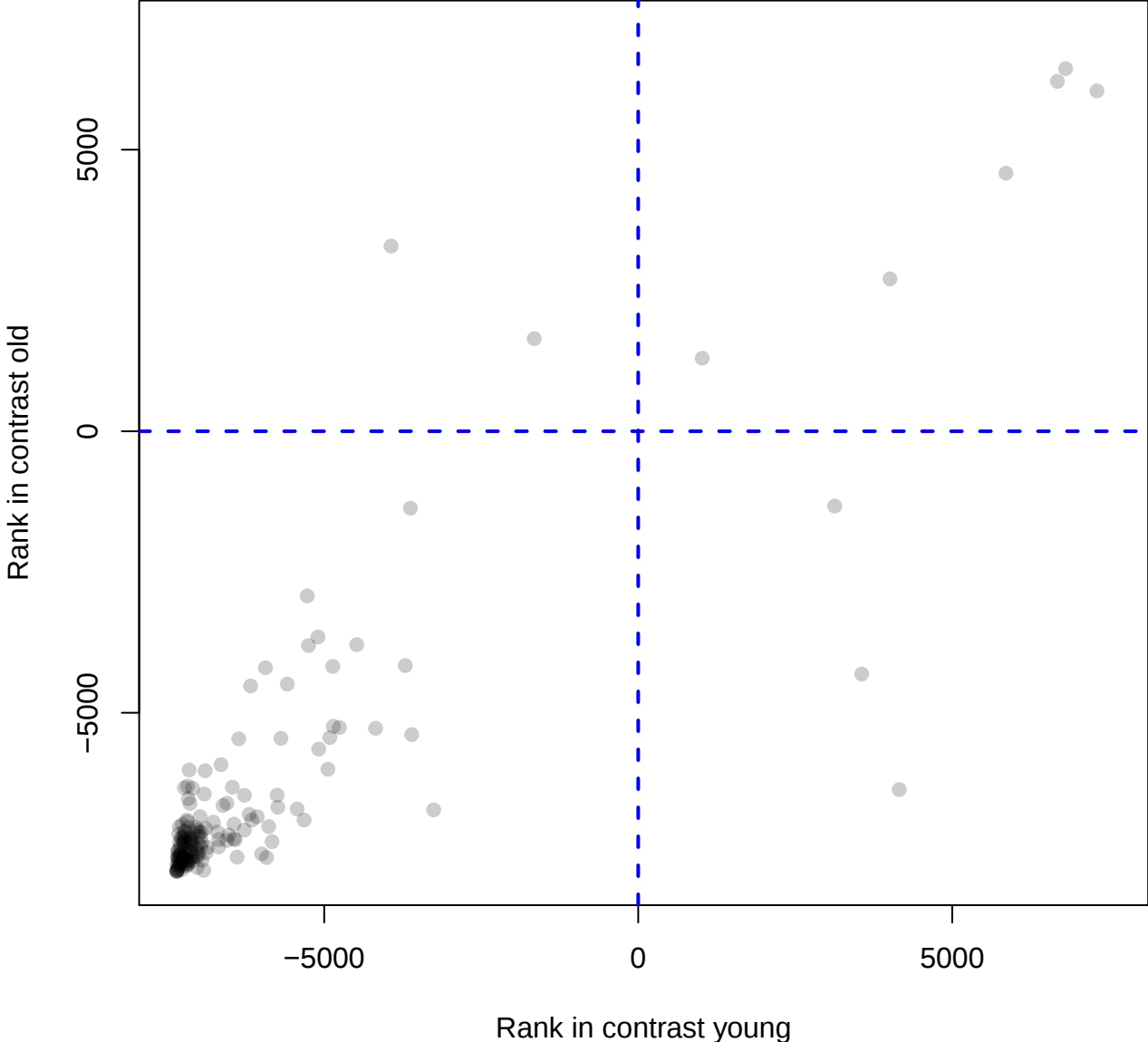
effect size versus statistical significance



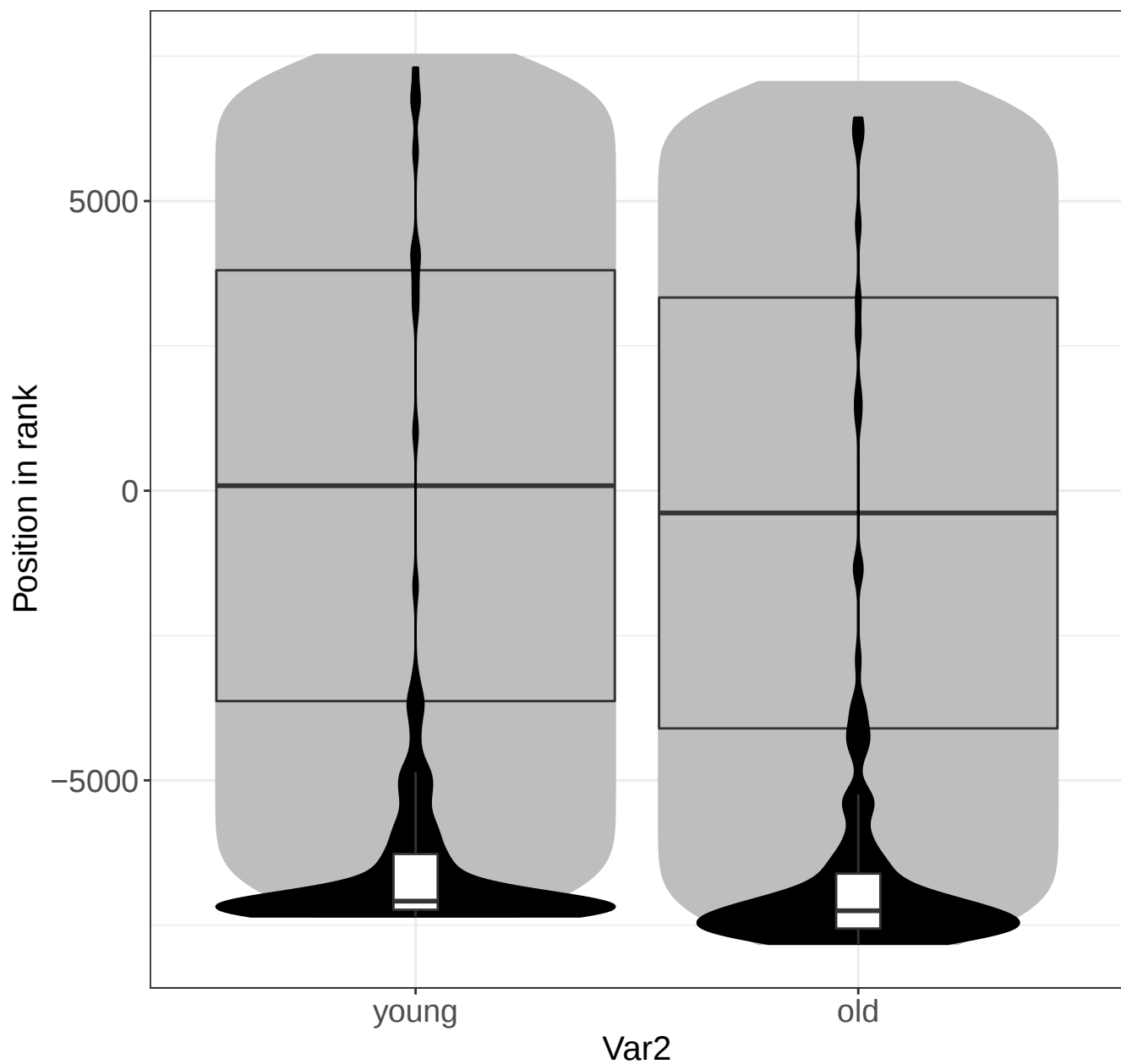
The-citric-acid-(TCA)-cycle-and-respiratory-electron-trar



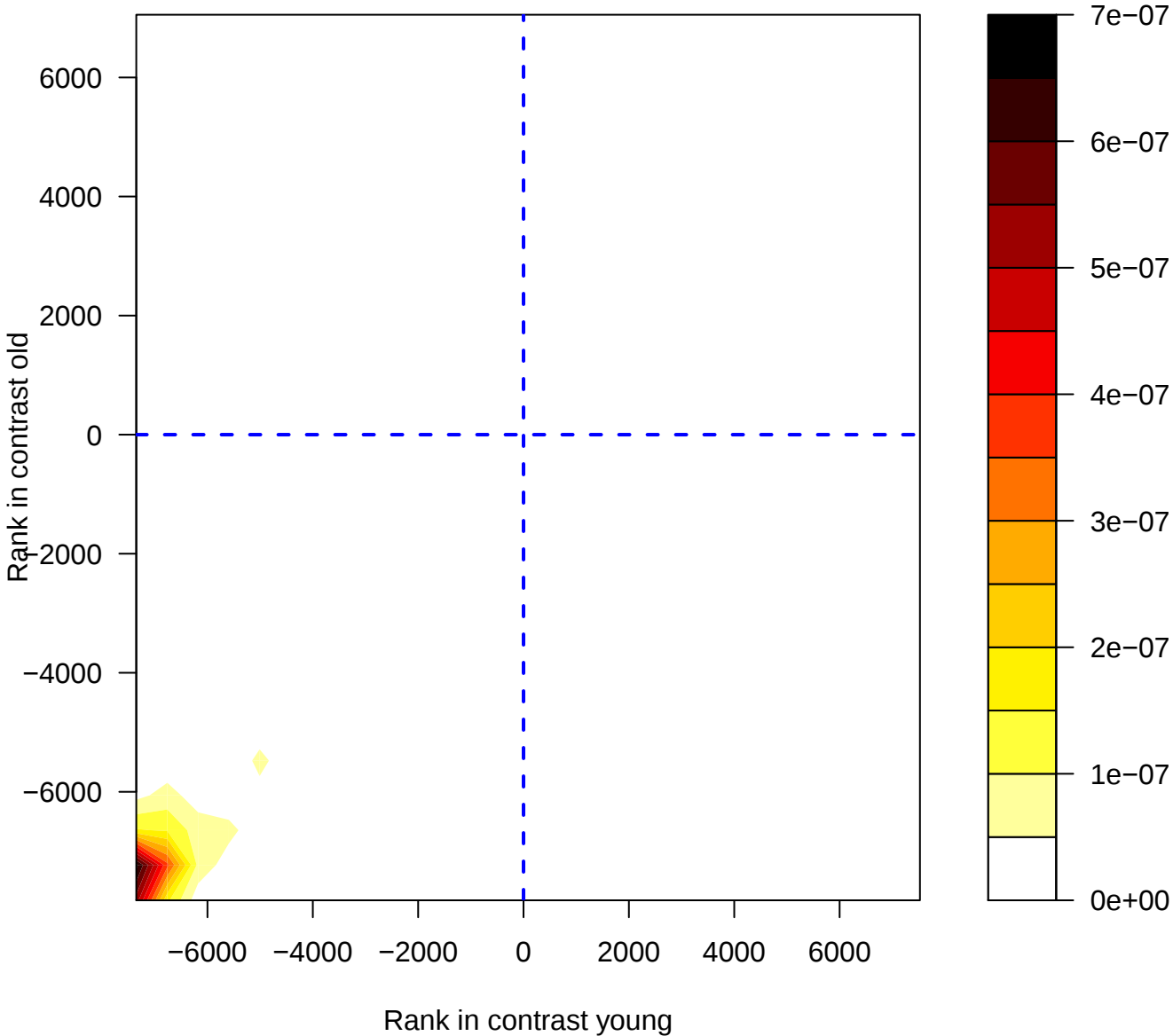
The-citric-acid-(TCA)-cycle-and-respiratory-electron-transport



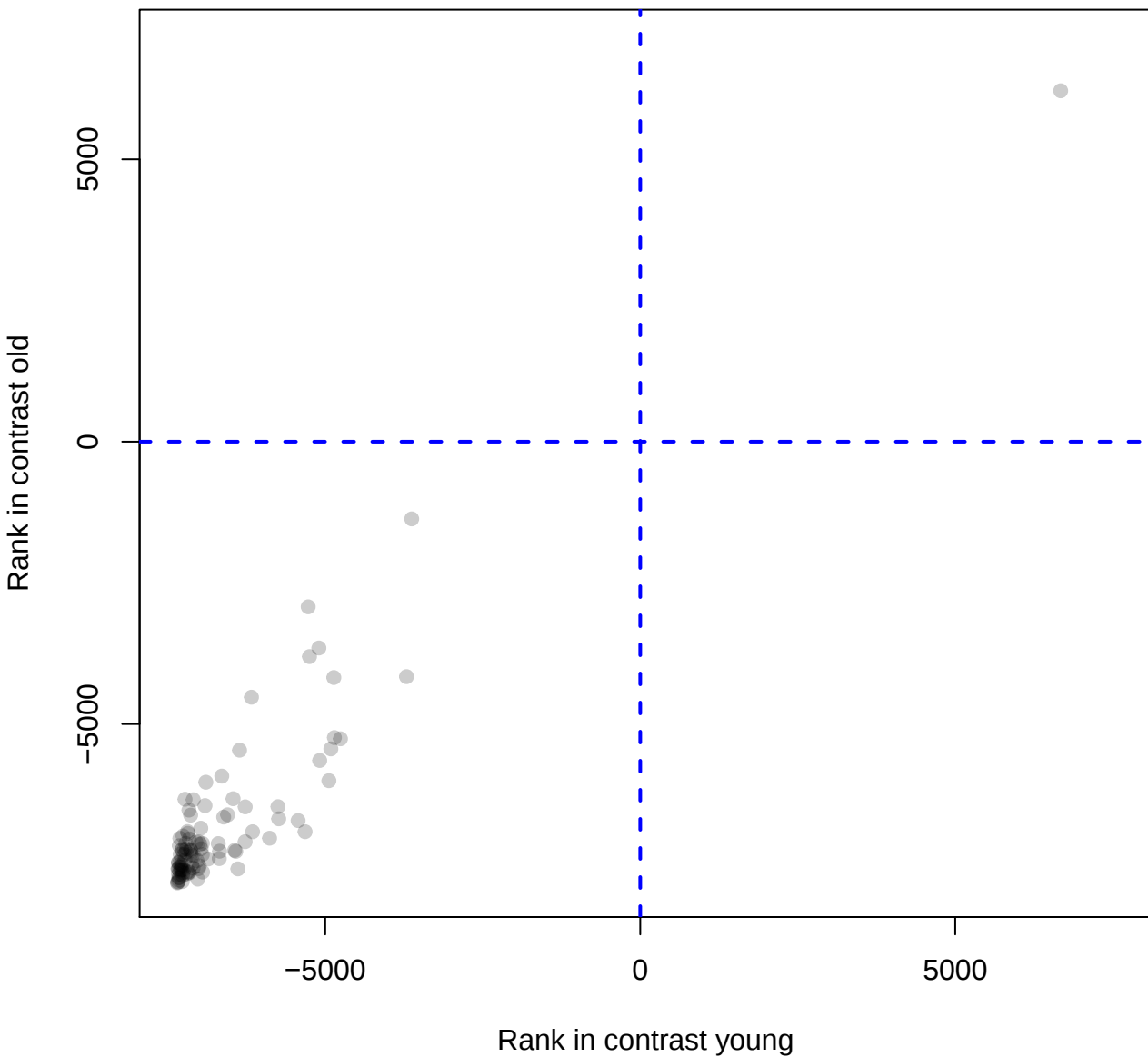
The-citric-acid-(TCA)-cycle-and-respiratory-el



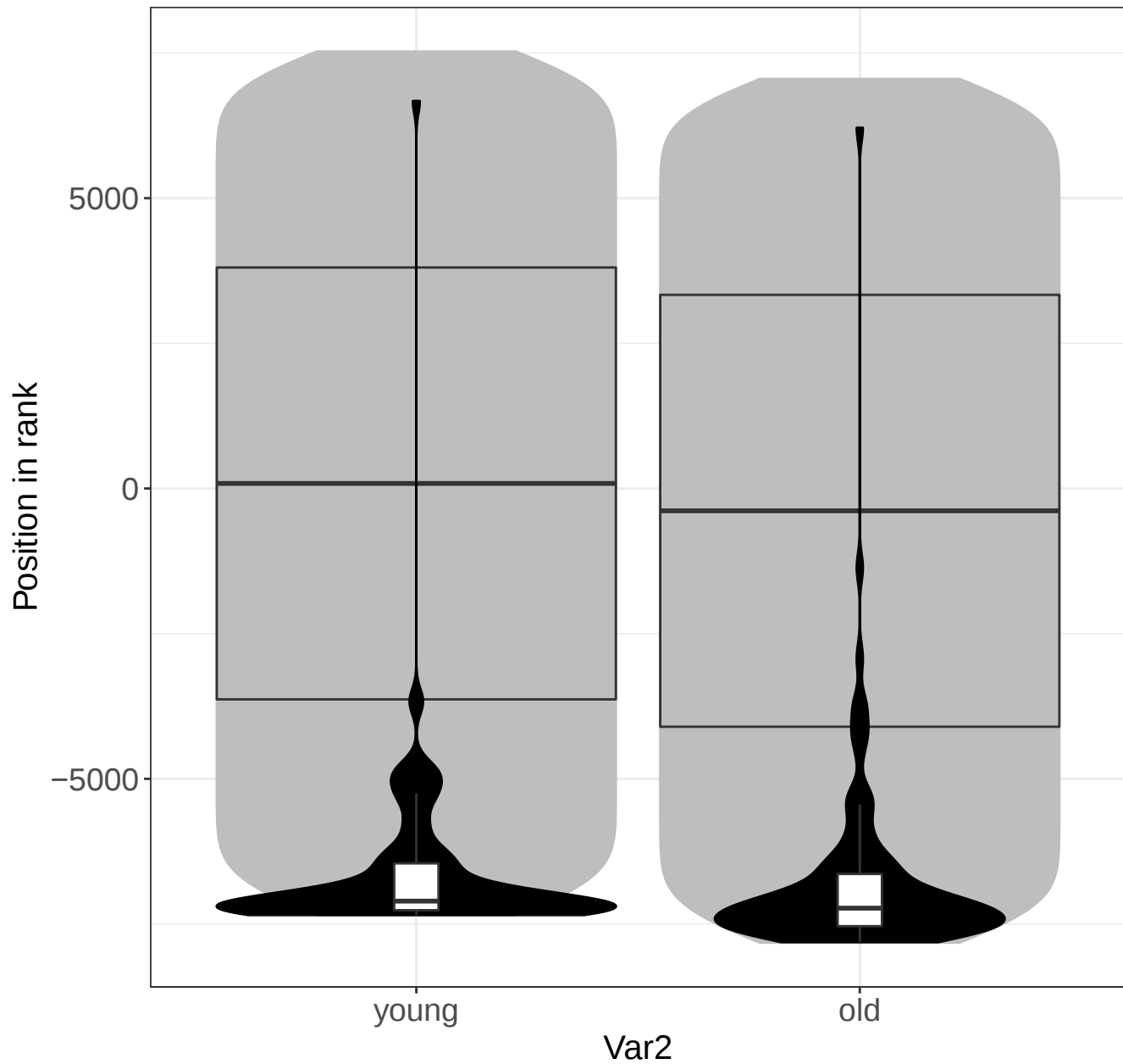
Respiratory-electron-transport



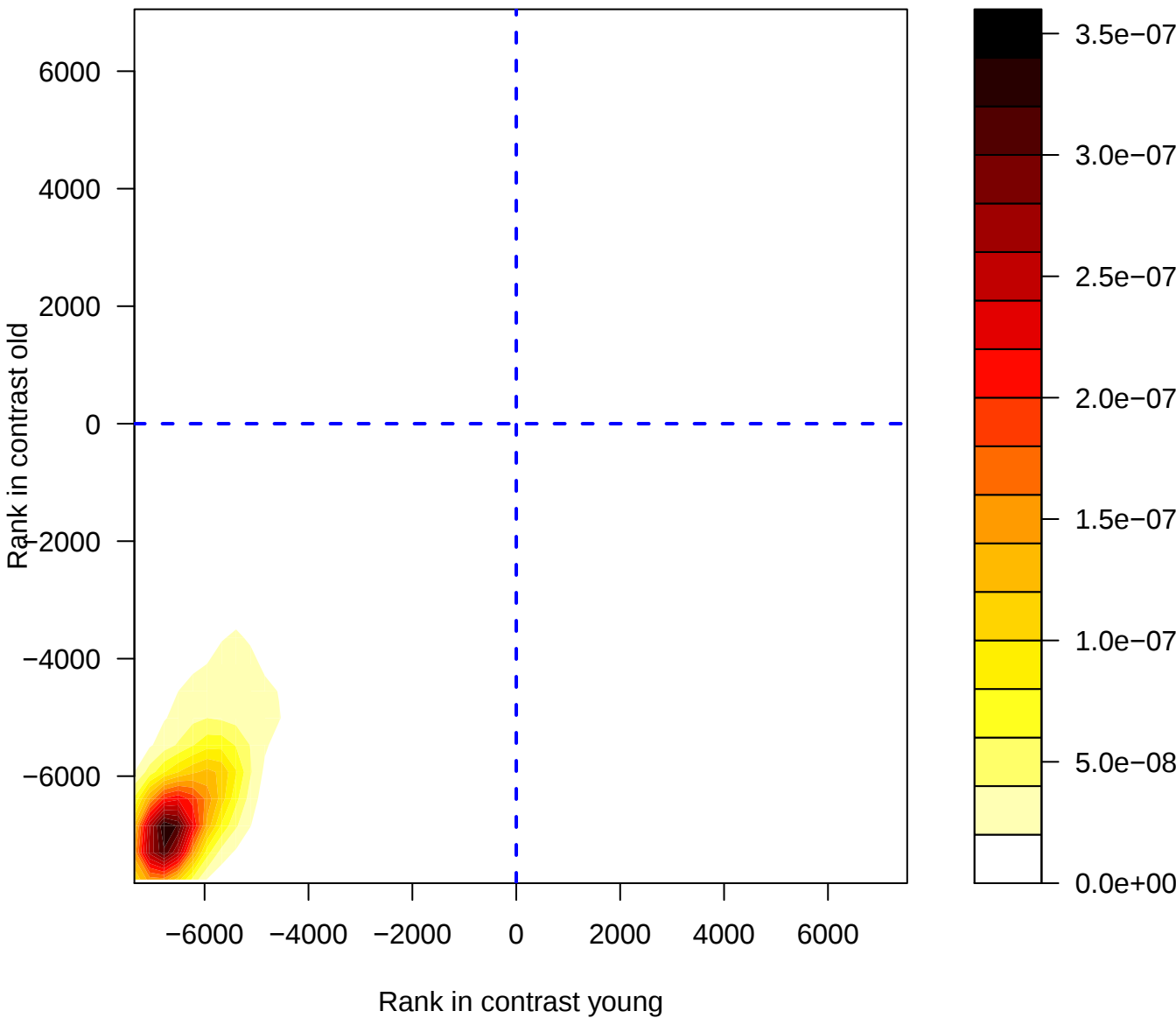
Respiratory-electron-transport



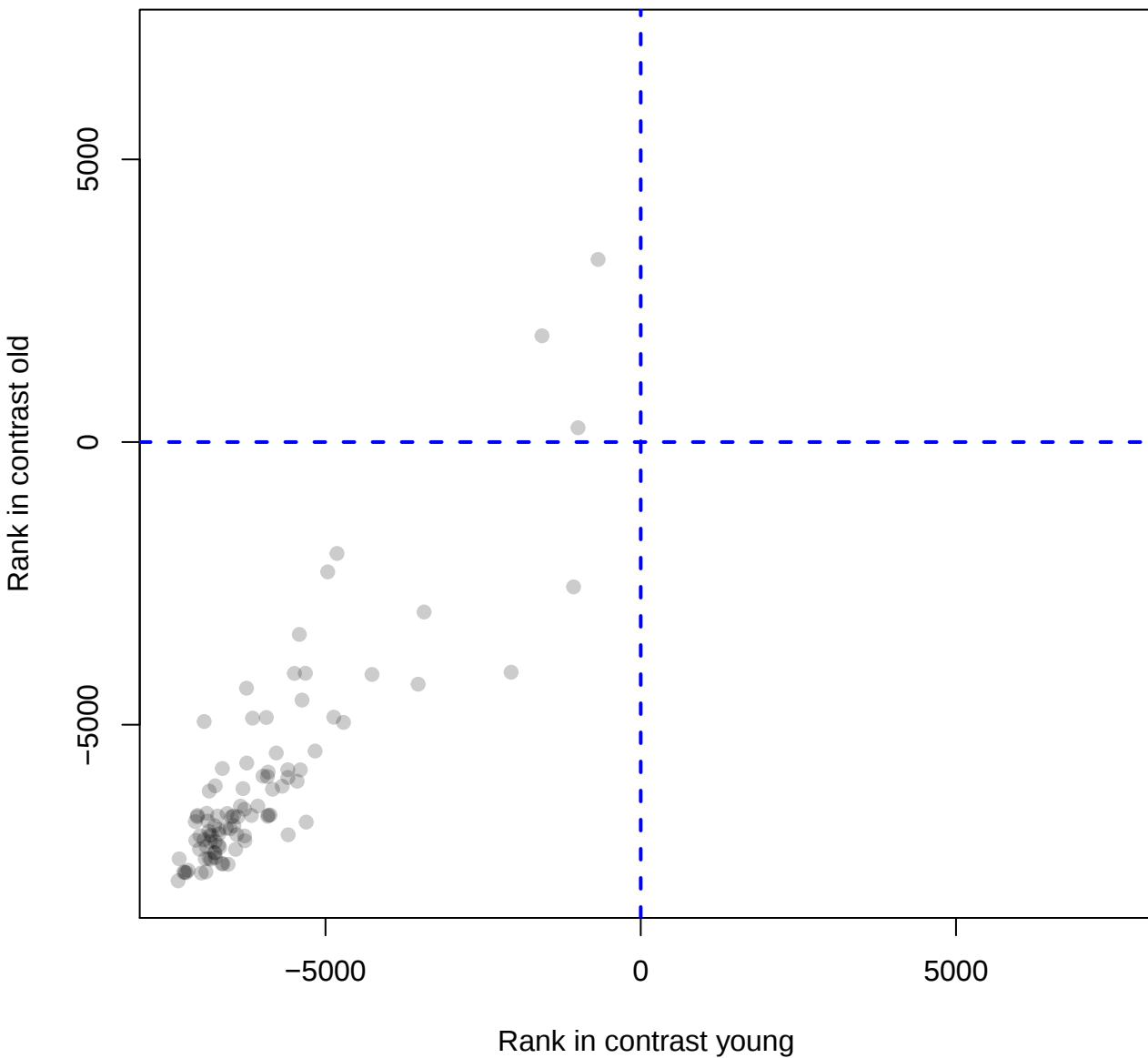
Respiratory-electron-transport



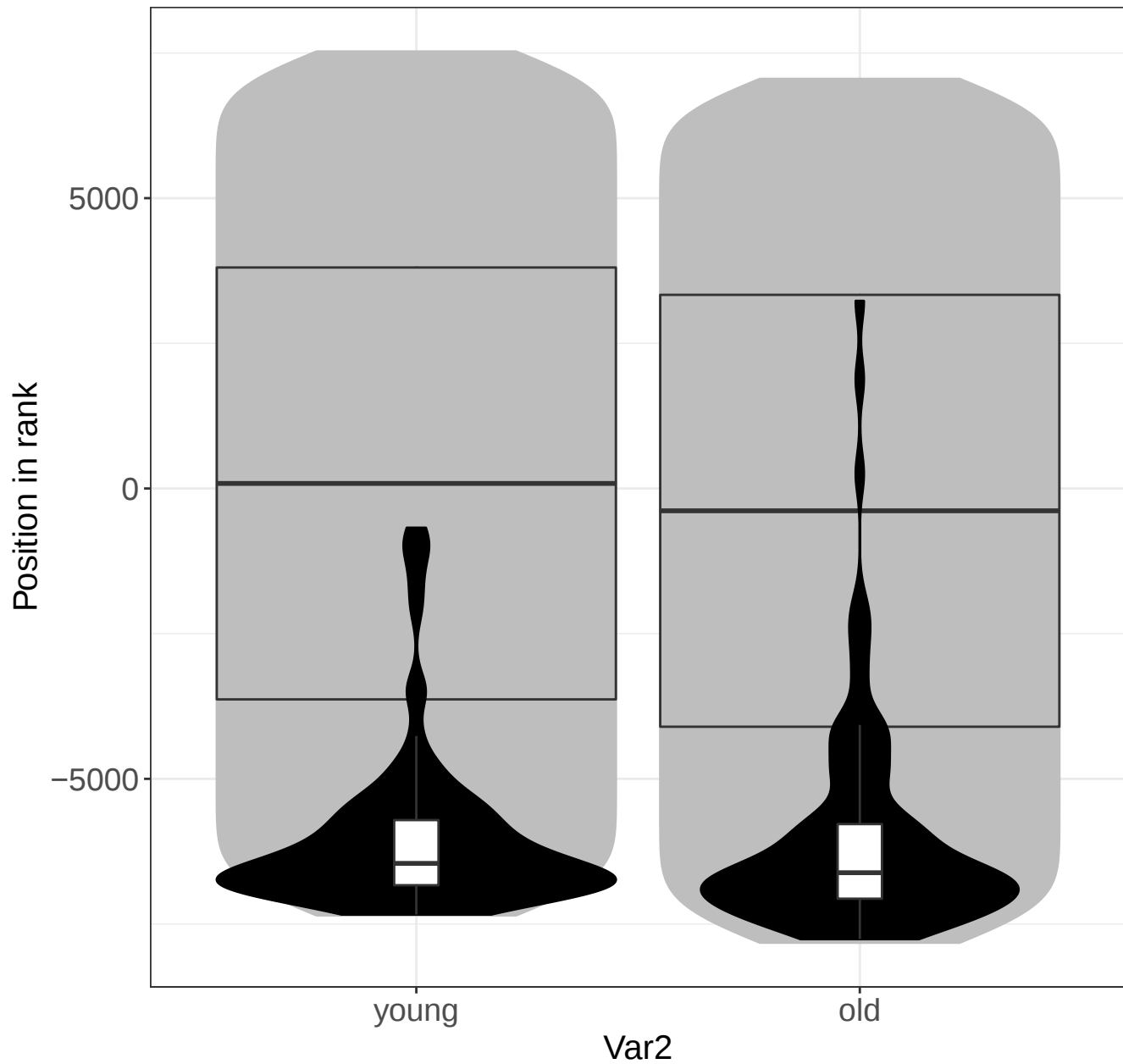
Mitochondrial-translation



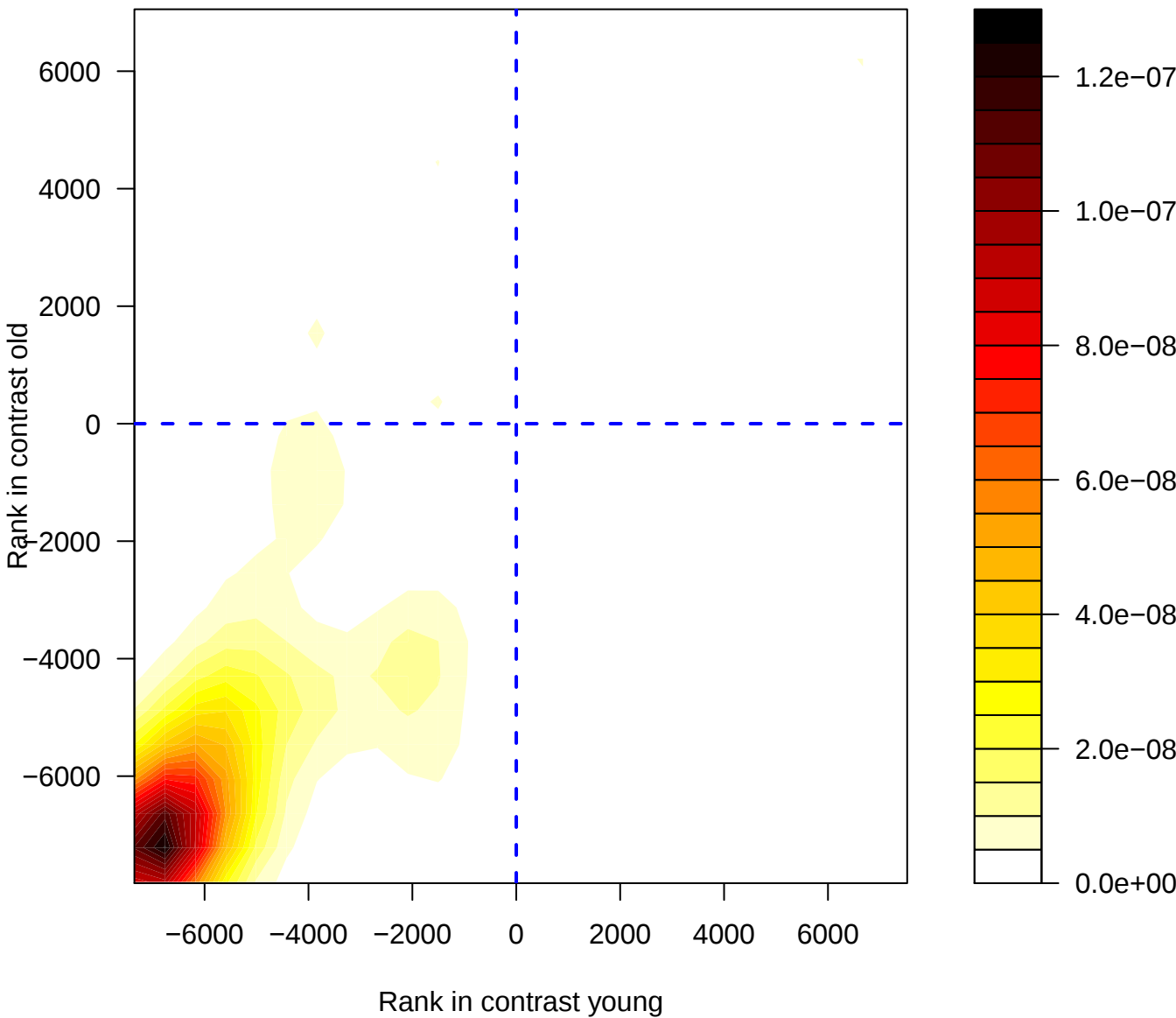
Mitochondrial-translation



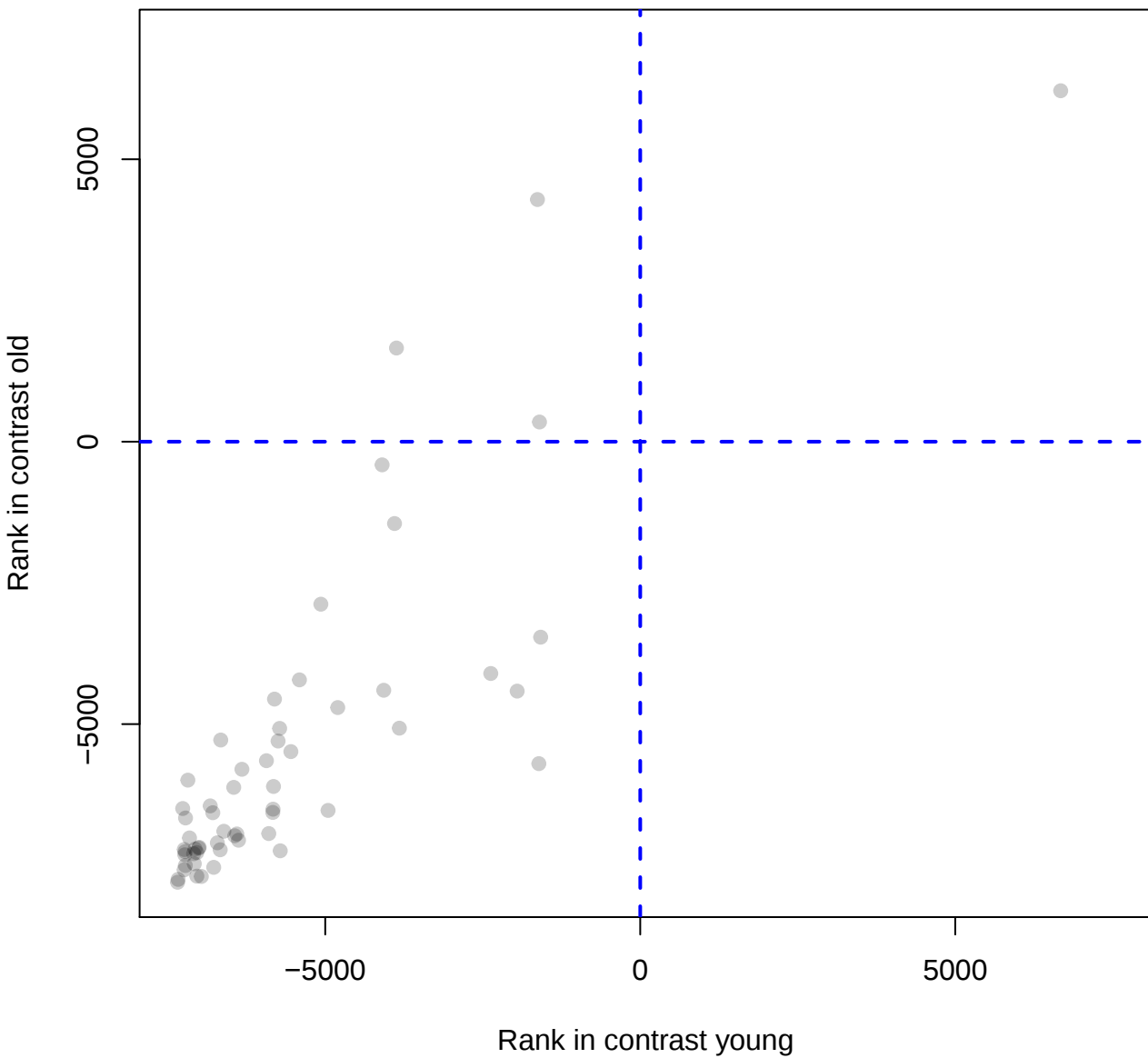
Mitochondrial-translation



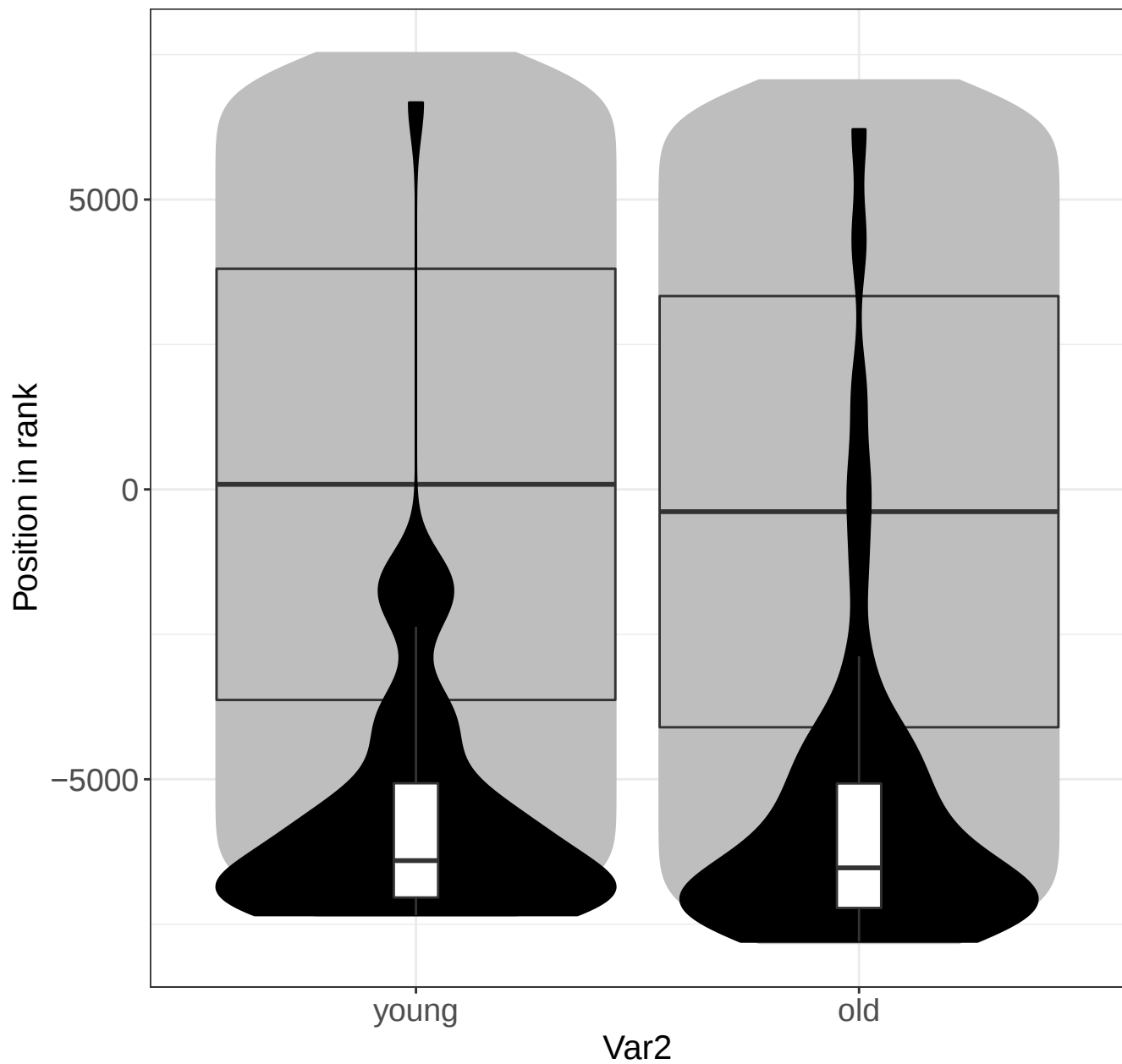
Mitochondrial-protein-import



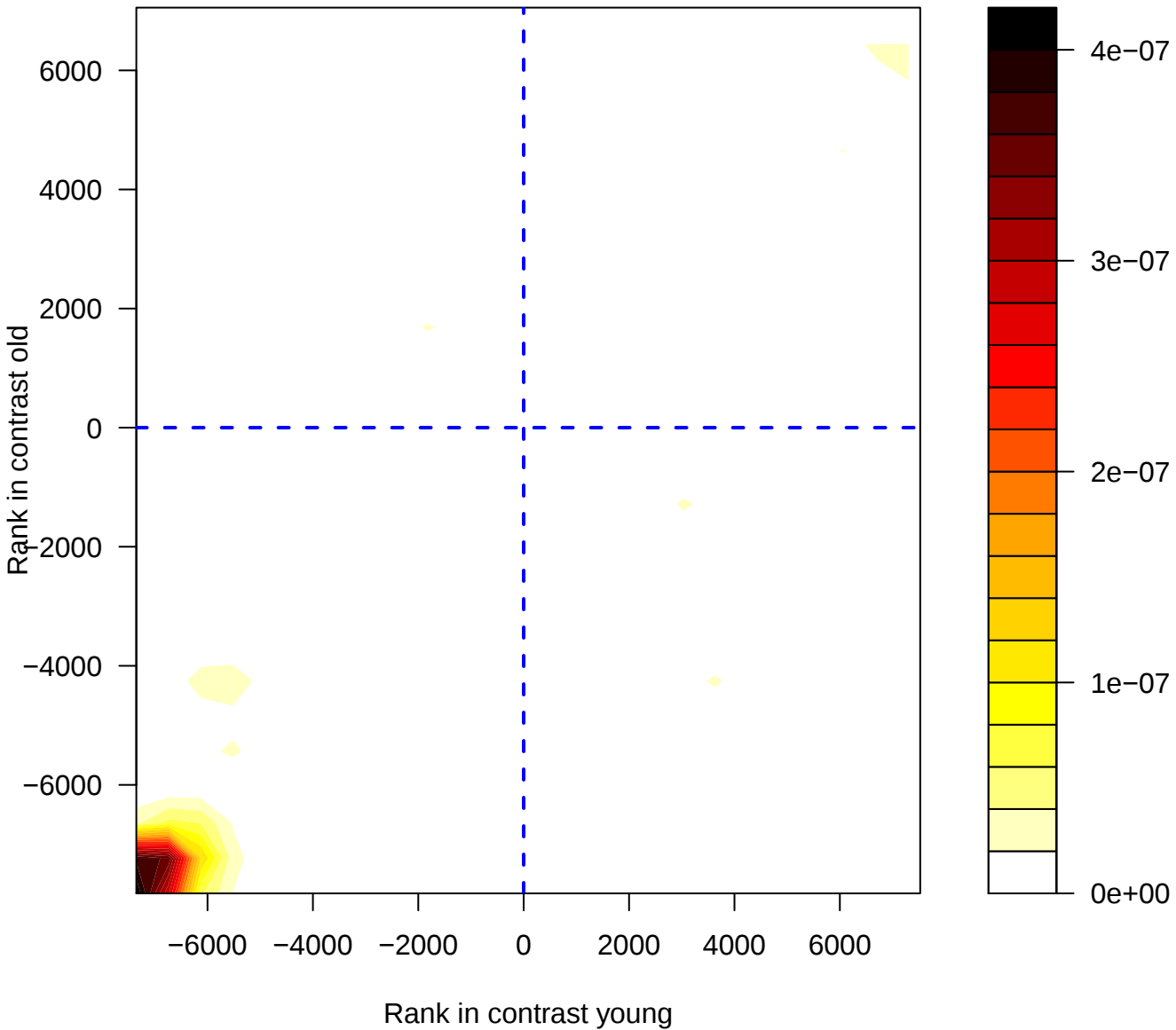
Mitochondrial-protein-import



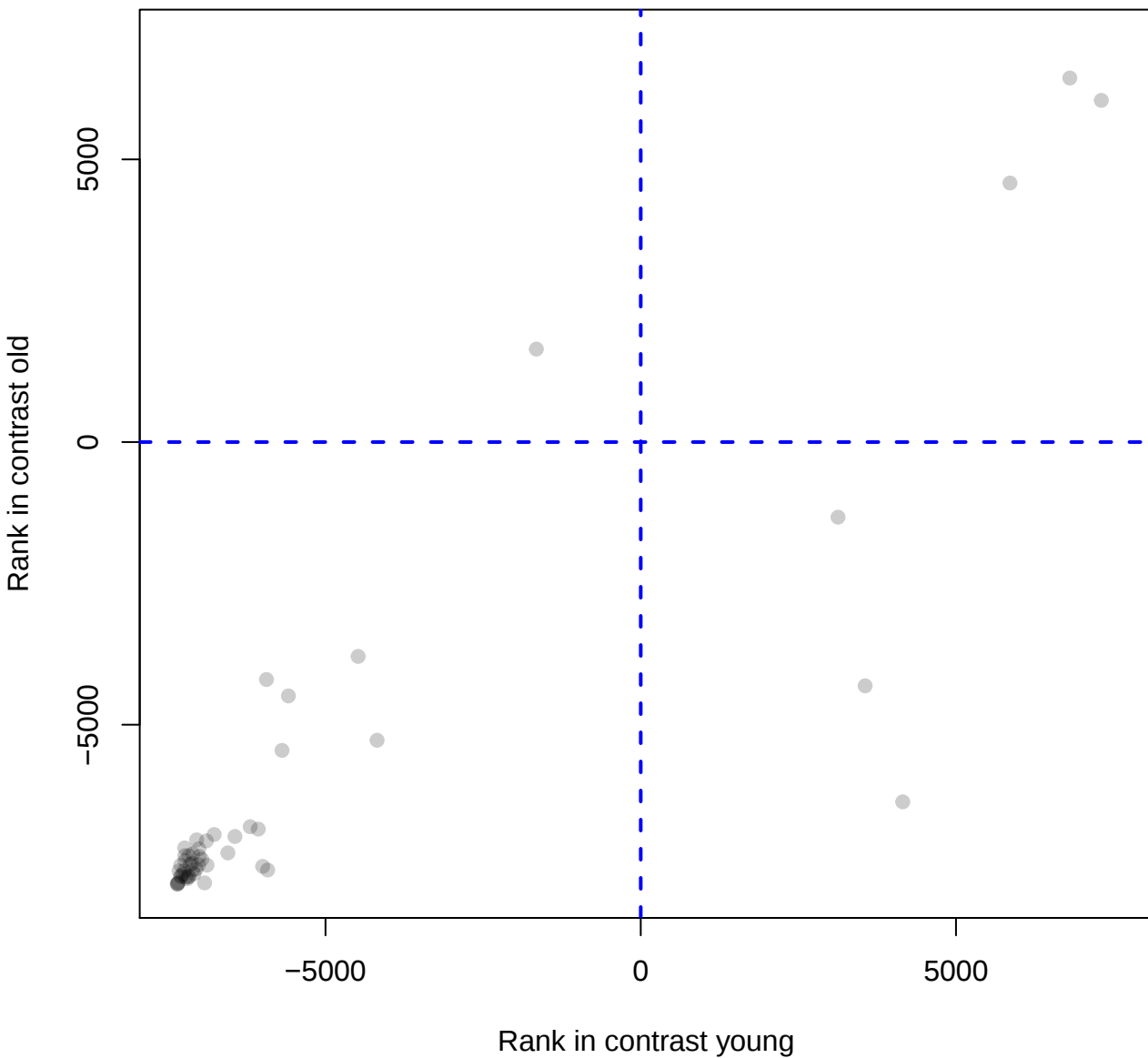
Mitochondrial-protein-import



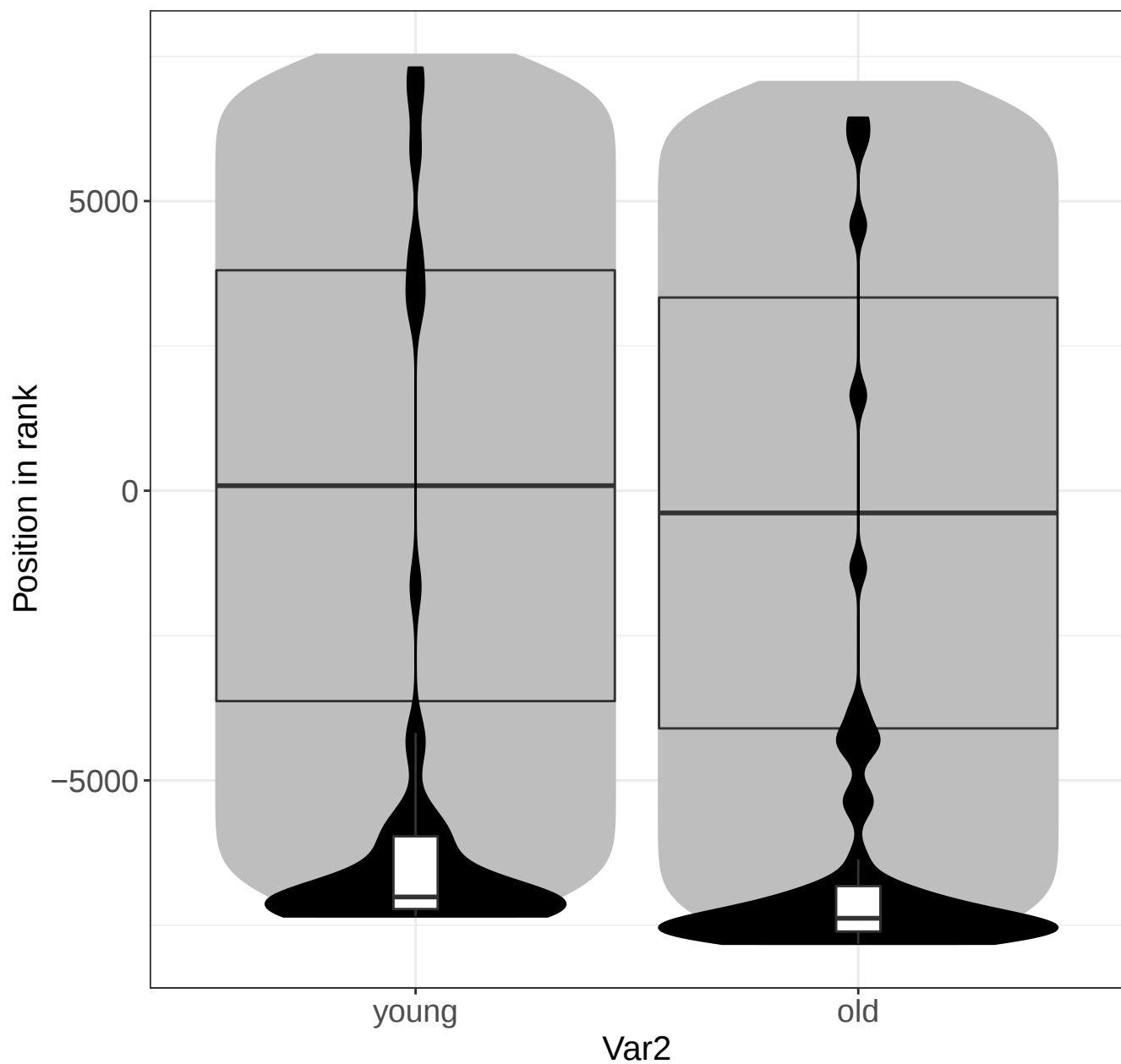
Pyruvate-metabolism-and-Citric-Acid-(TCA)-cycle



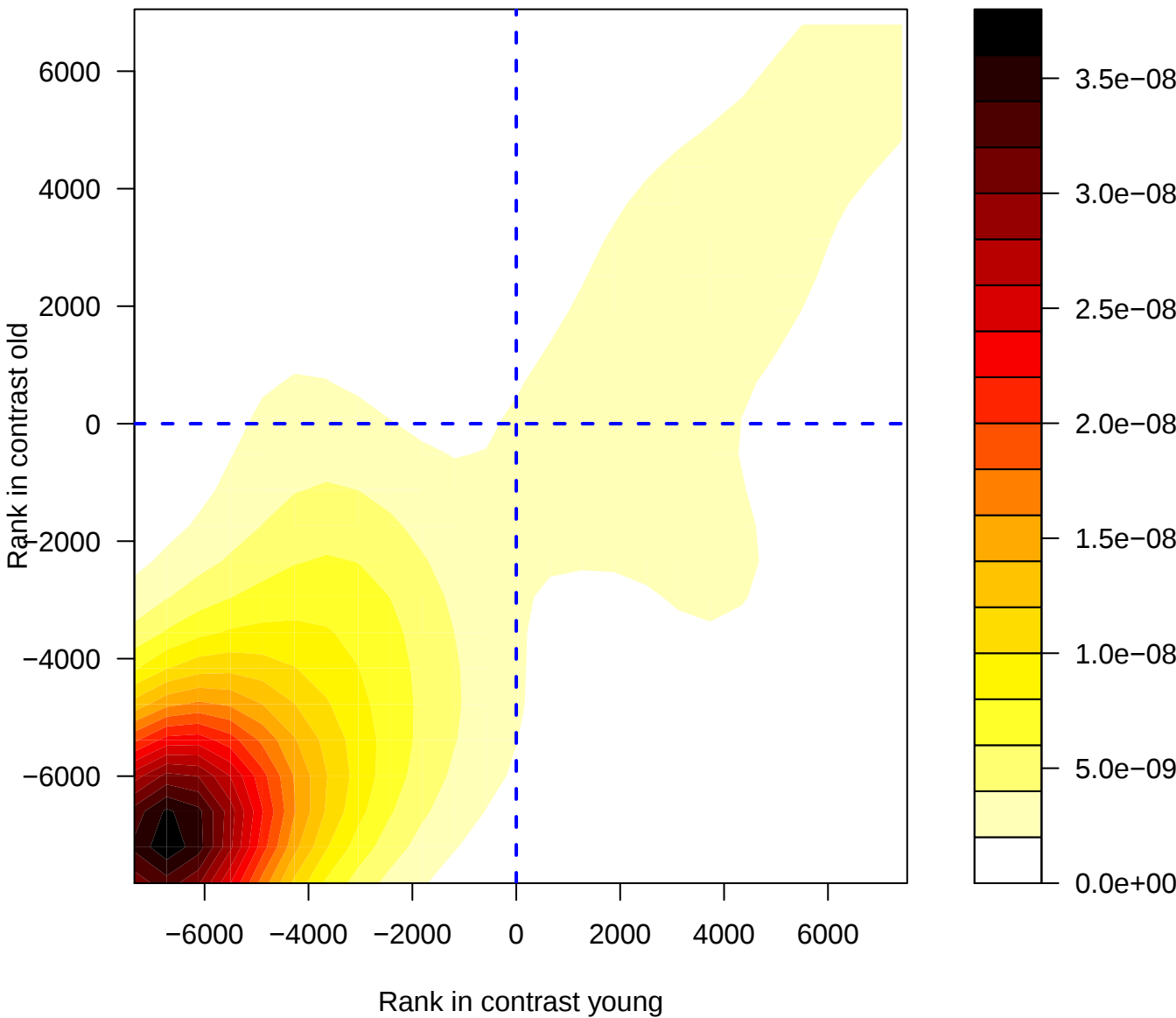
Pyruvate-metabolism-and-Citric-Acid-(TCA)-cycle



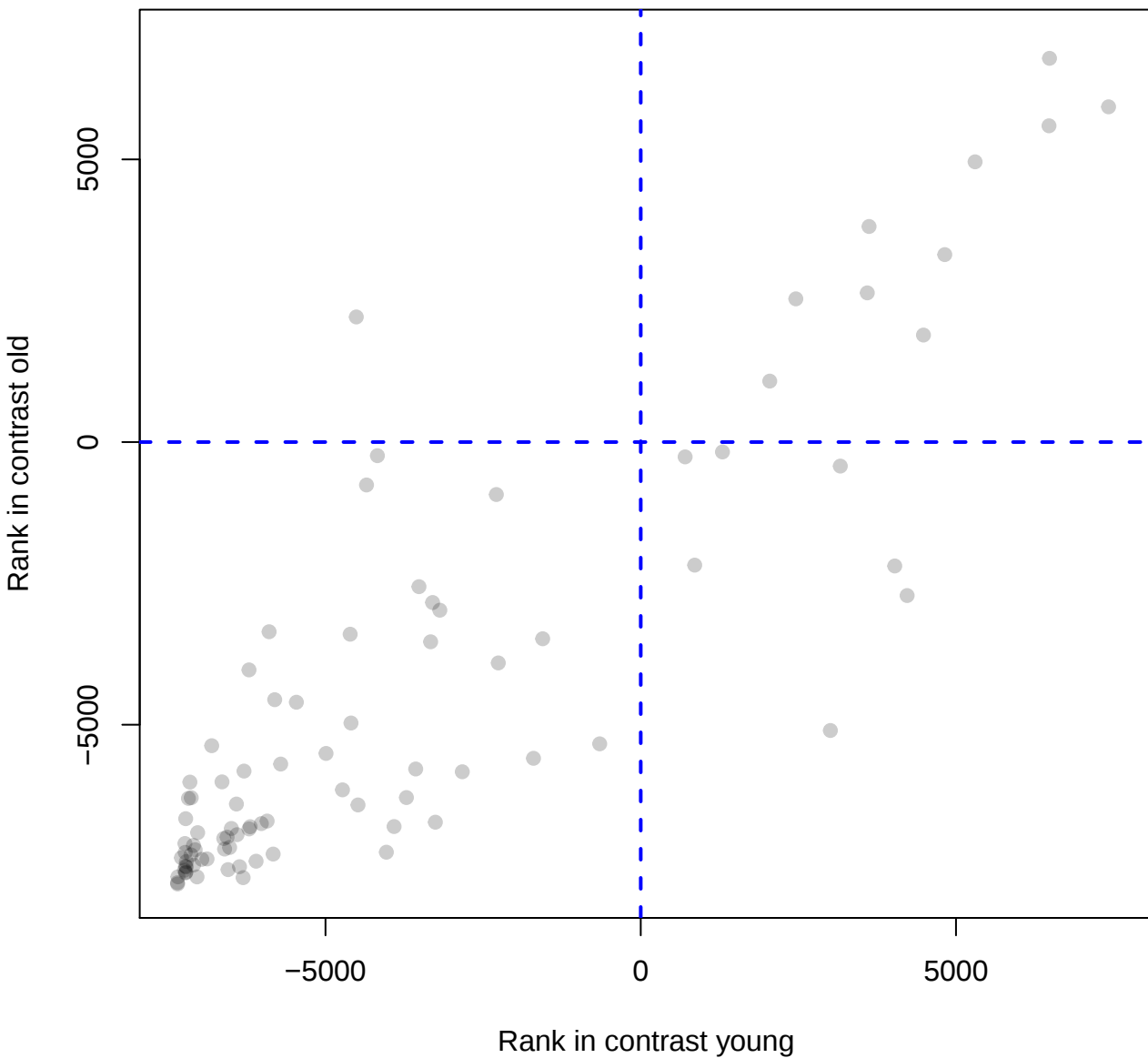
Pyruvate-metabolism-and-Citric-Acid-(TCA)-cy



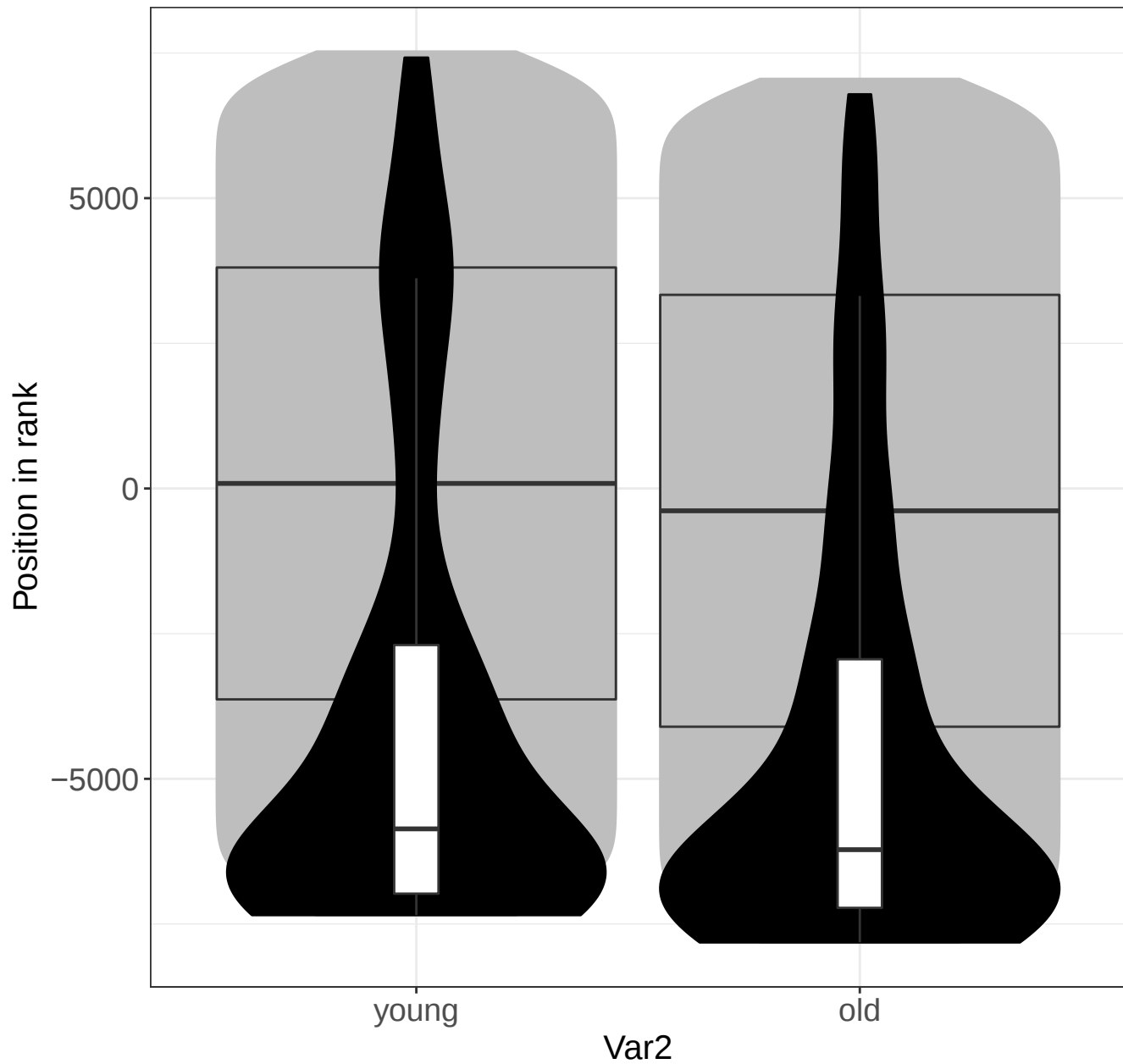
Mitochondrial-biogenesis



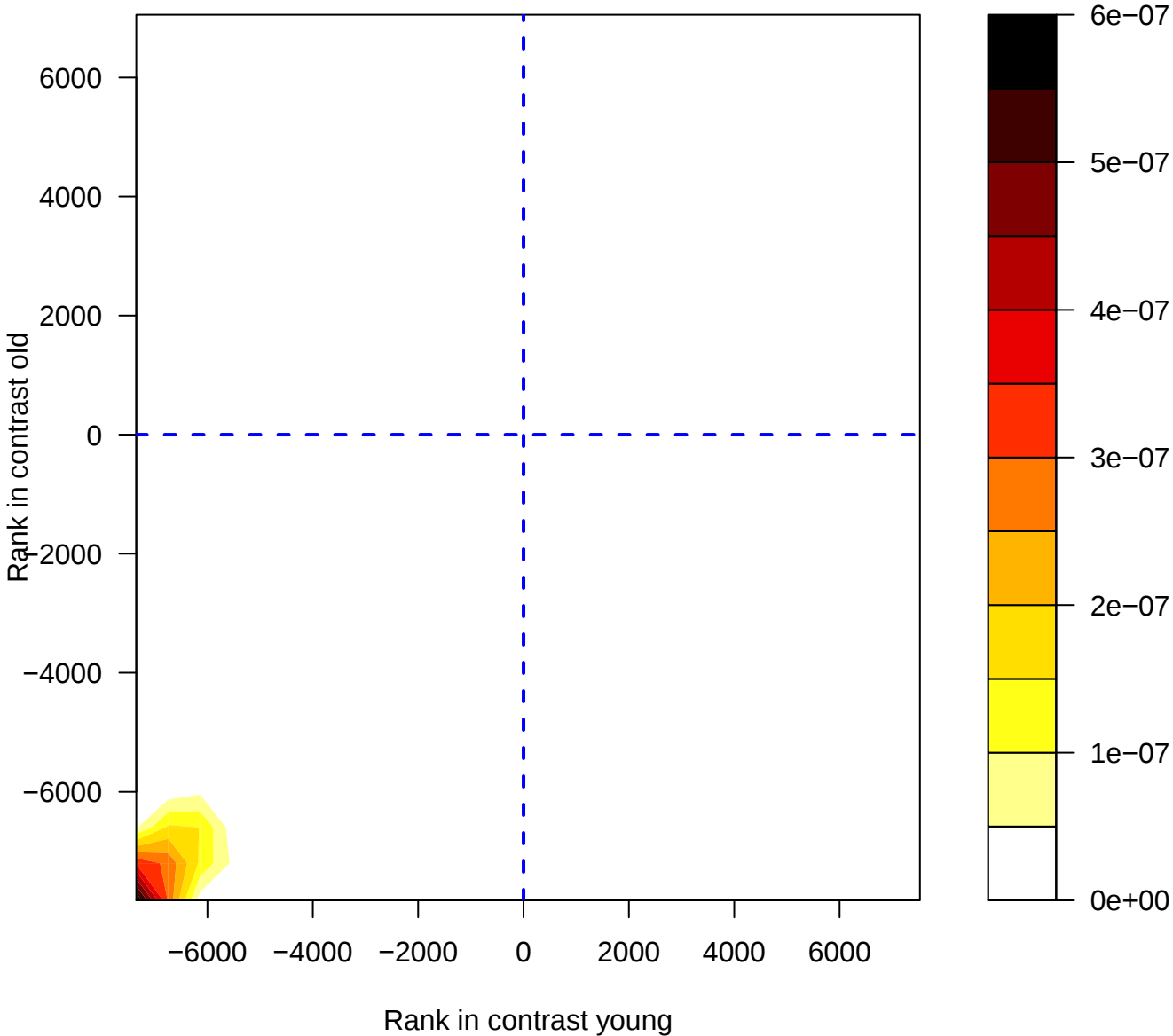
Mitochondrial-biogenesis



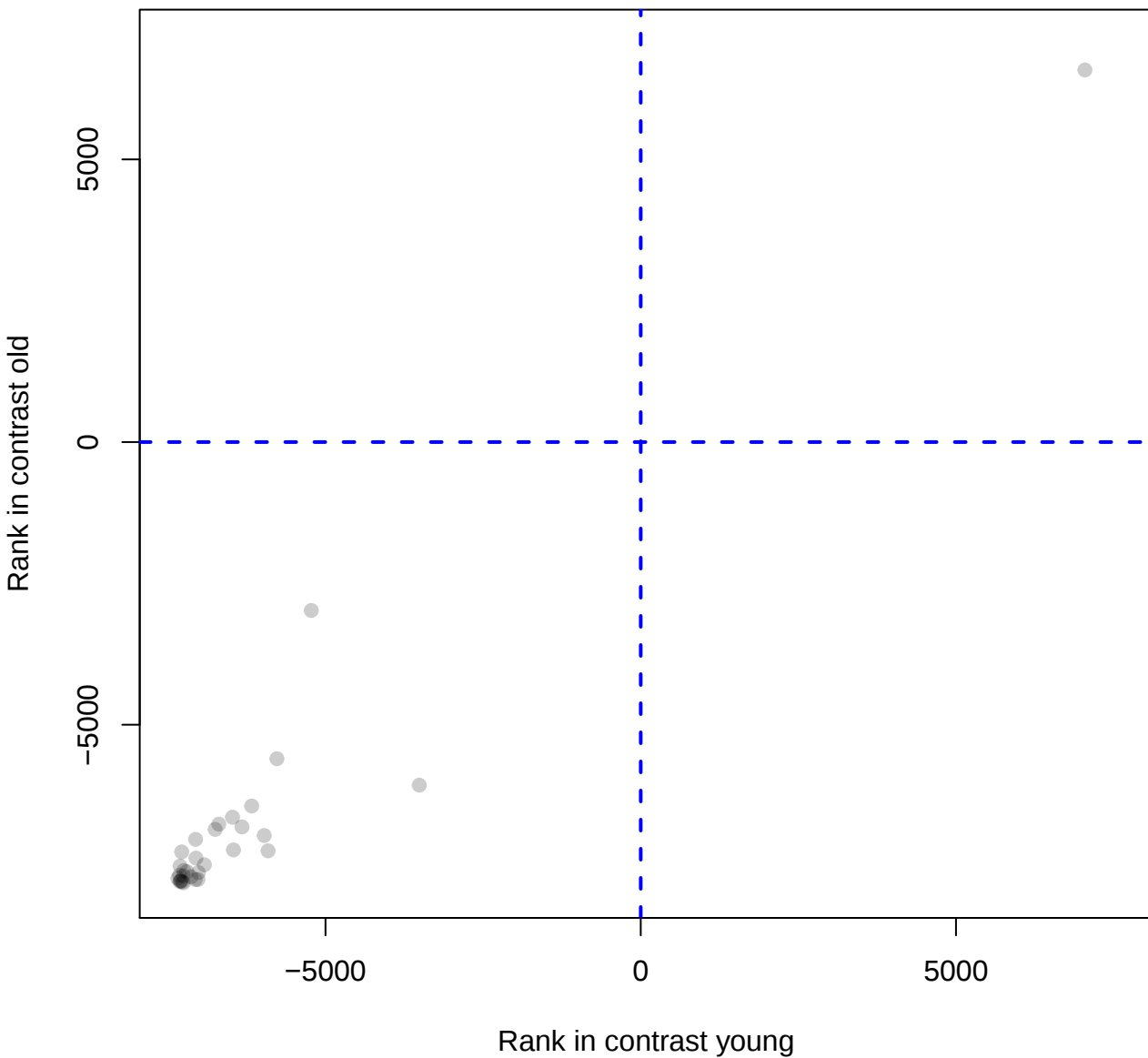
Mitochondrial-biogenesis



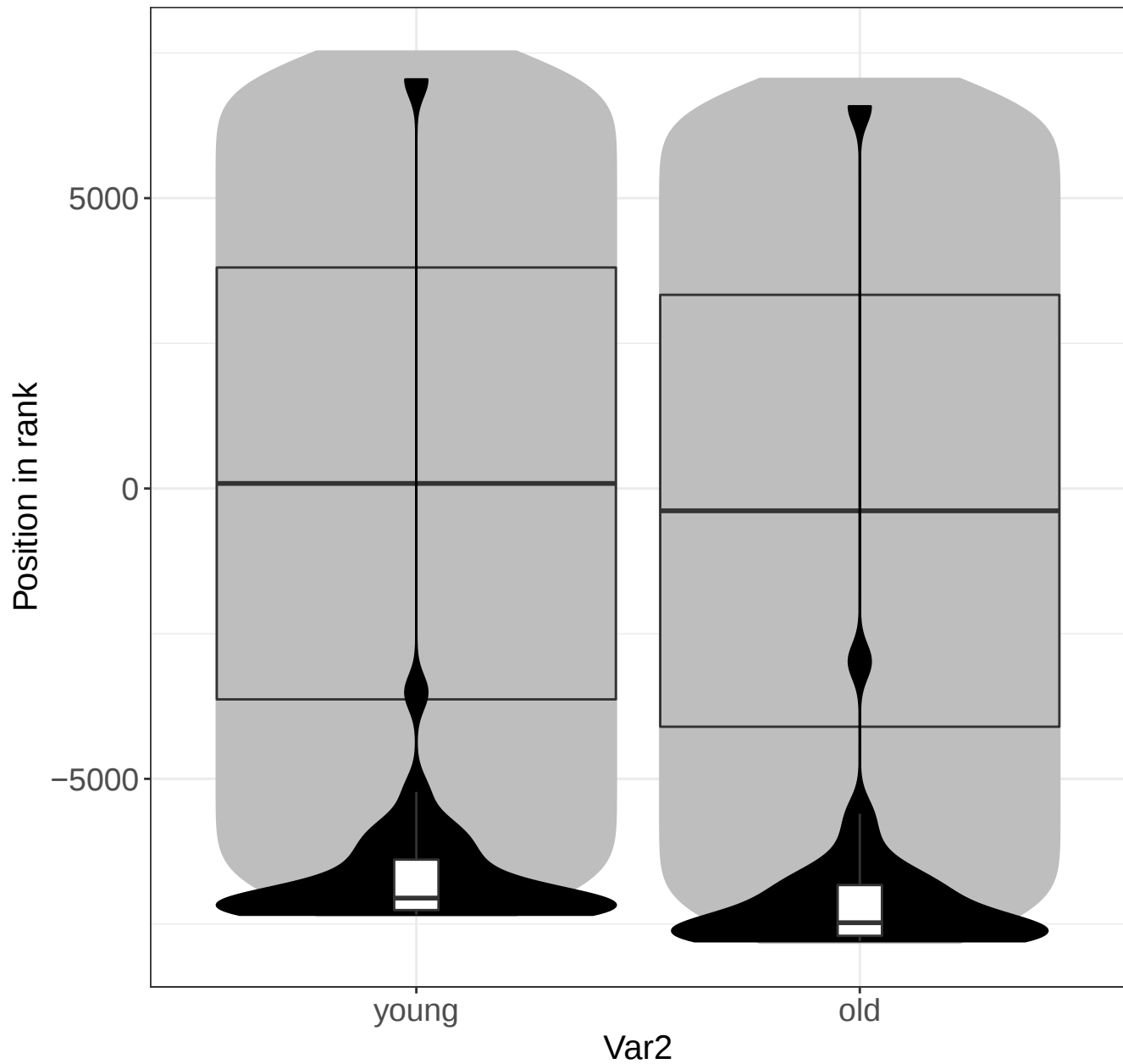
Mitochondrial-Fatty-Acid-Beta-Oxidation



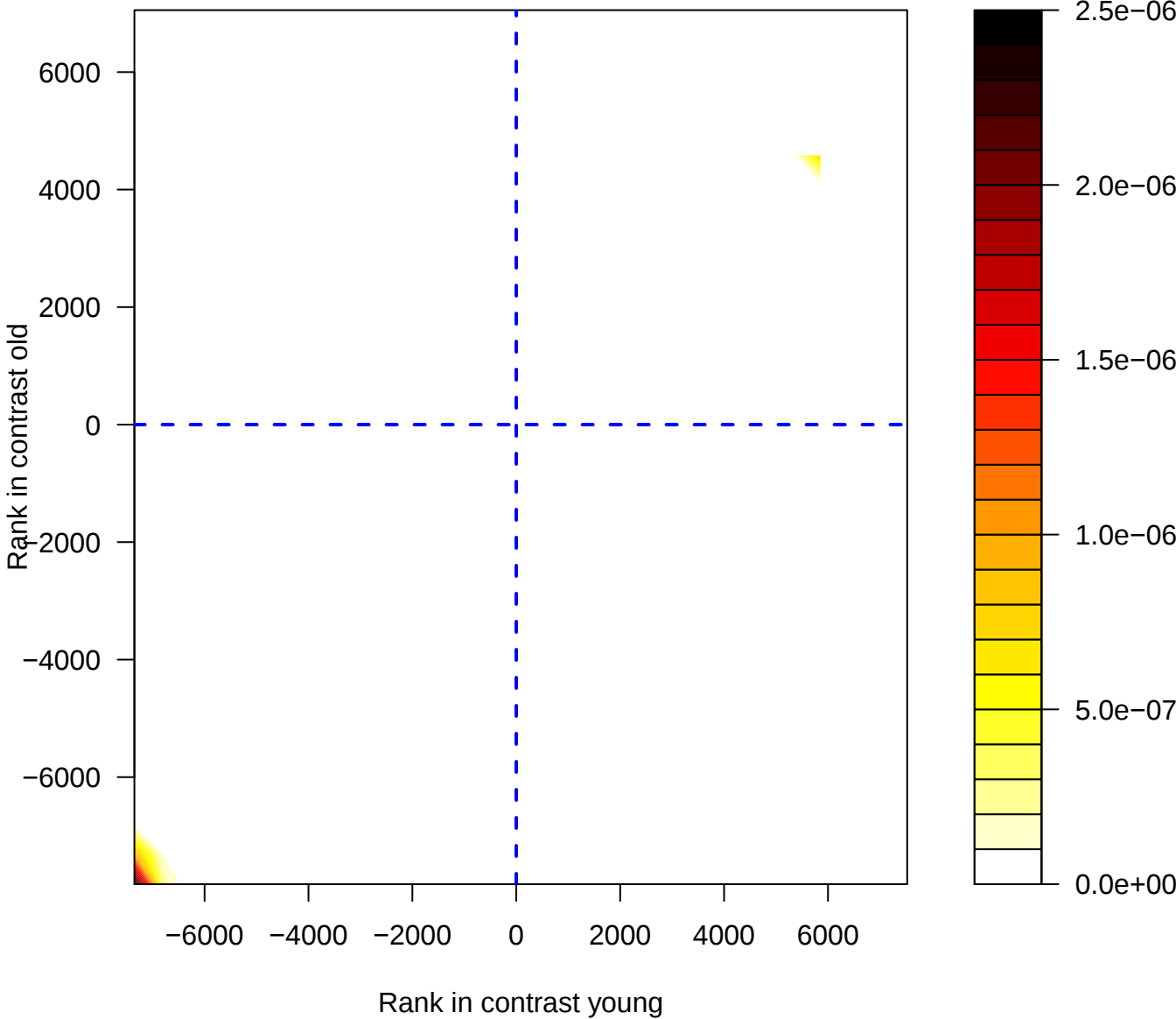
Mitochondrial-Fatty-Acid-Beta-Oxidation



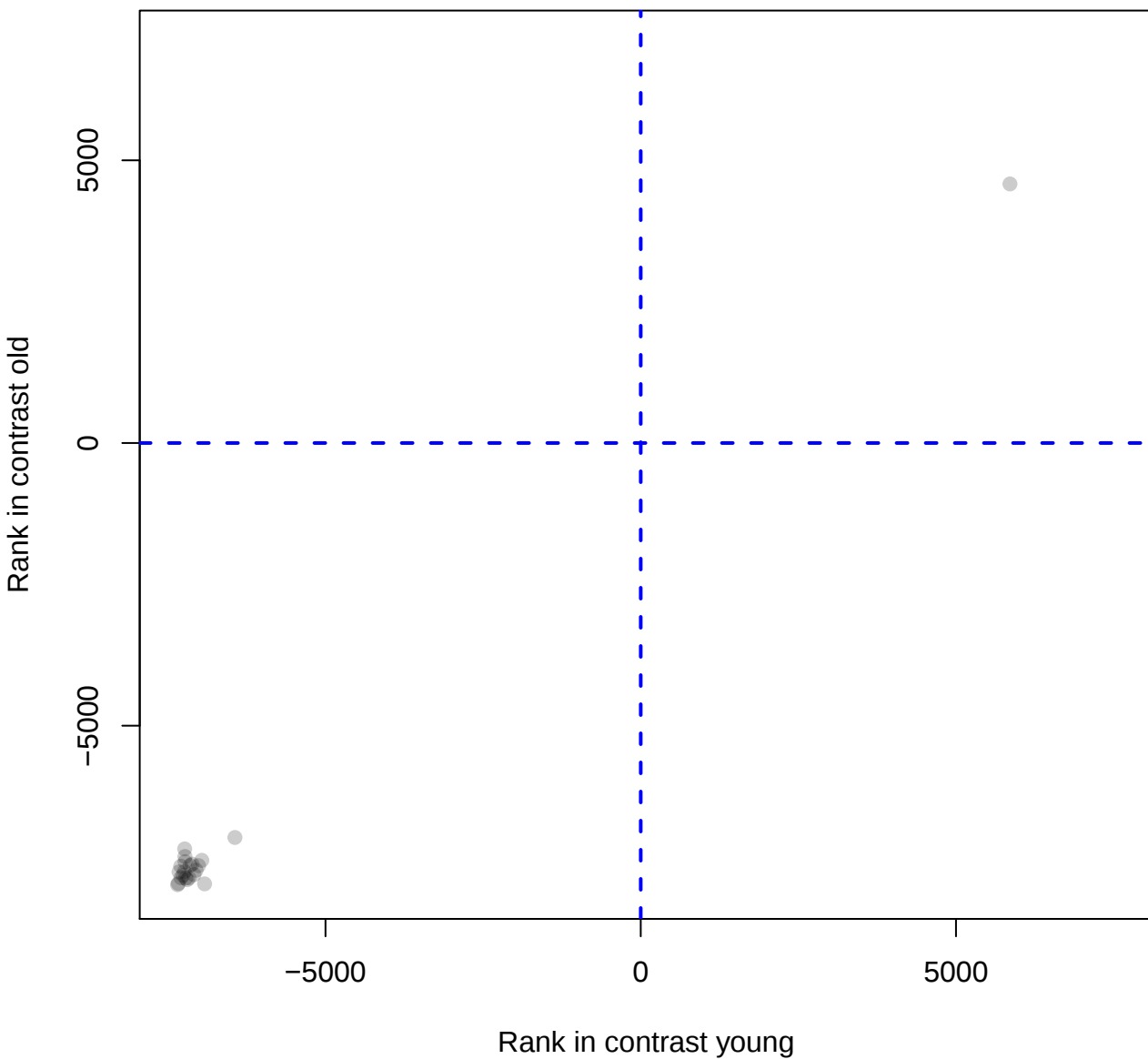
Mitochondrial-Fatty-Acid-Beta-Oxidation



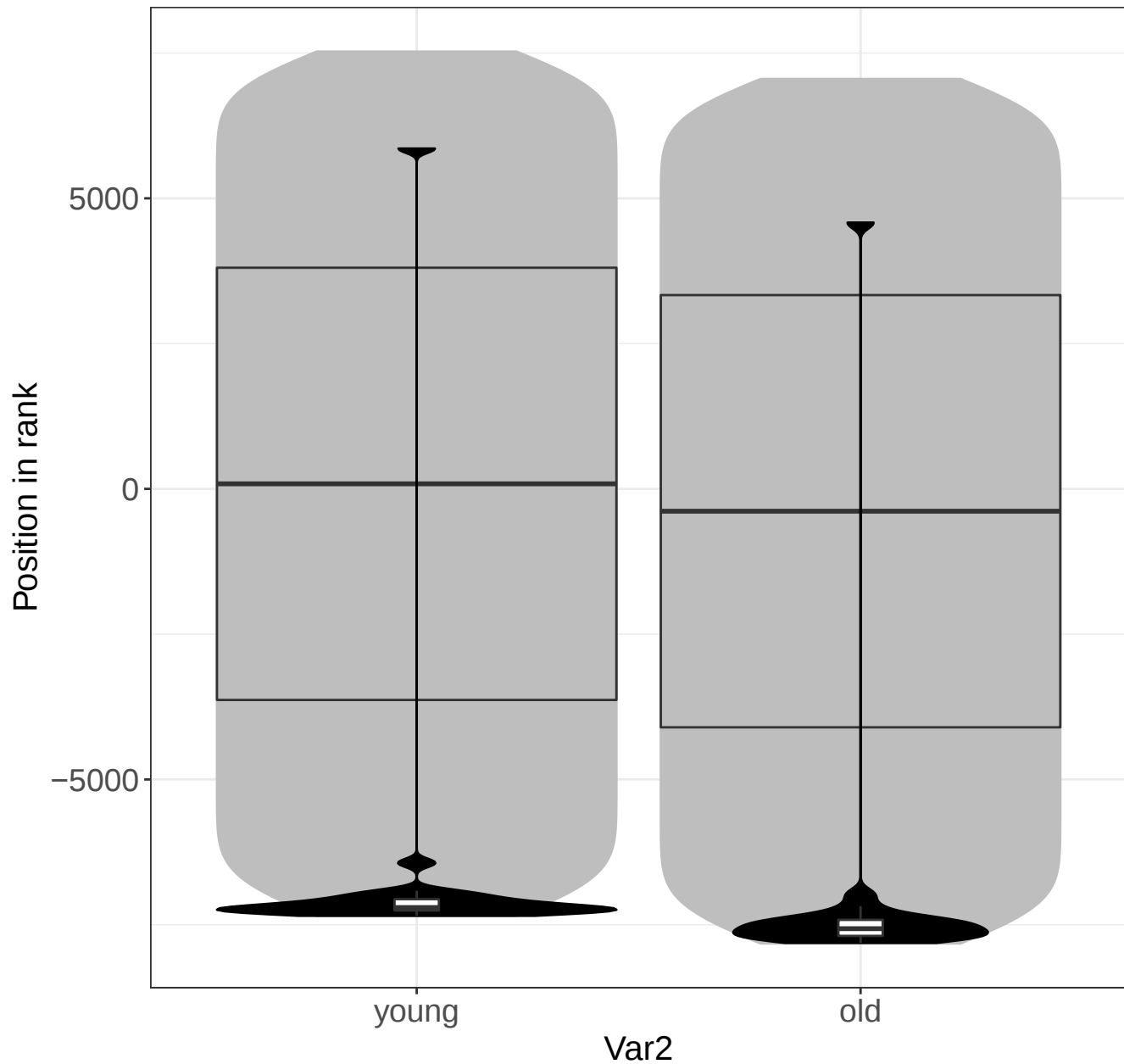
Citric-acid-cycle-(TCA-cycle)



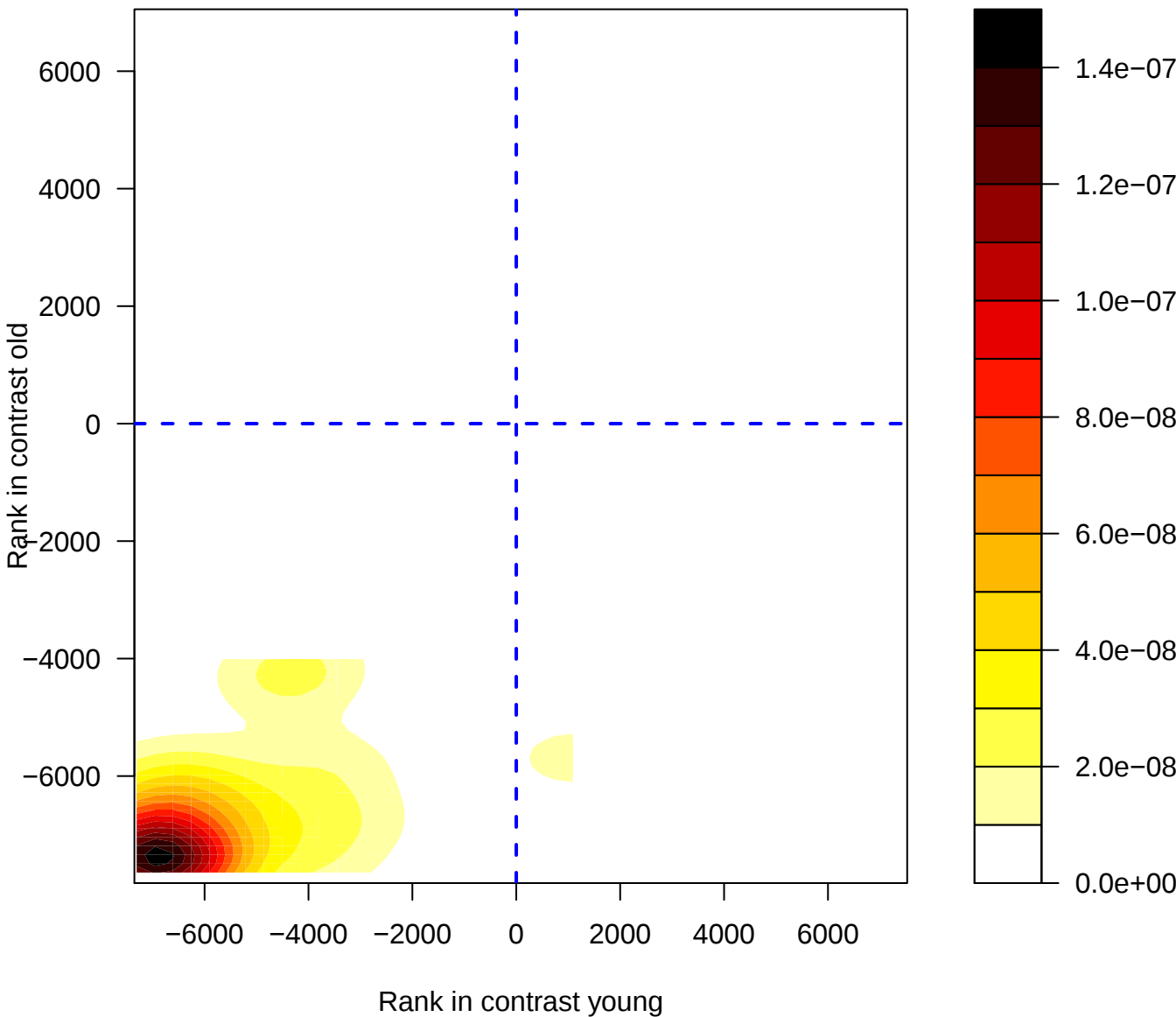
Citric-acid-cycle-(TCA-cycle)



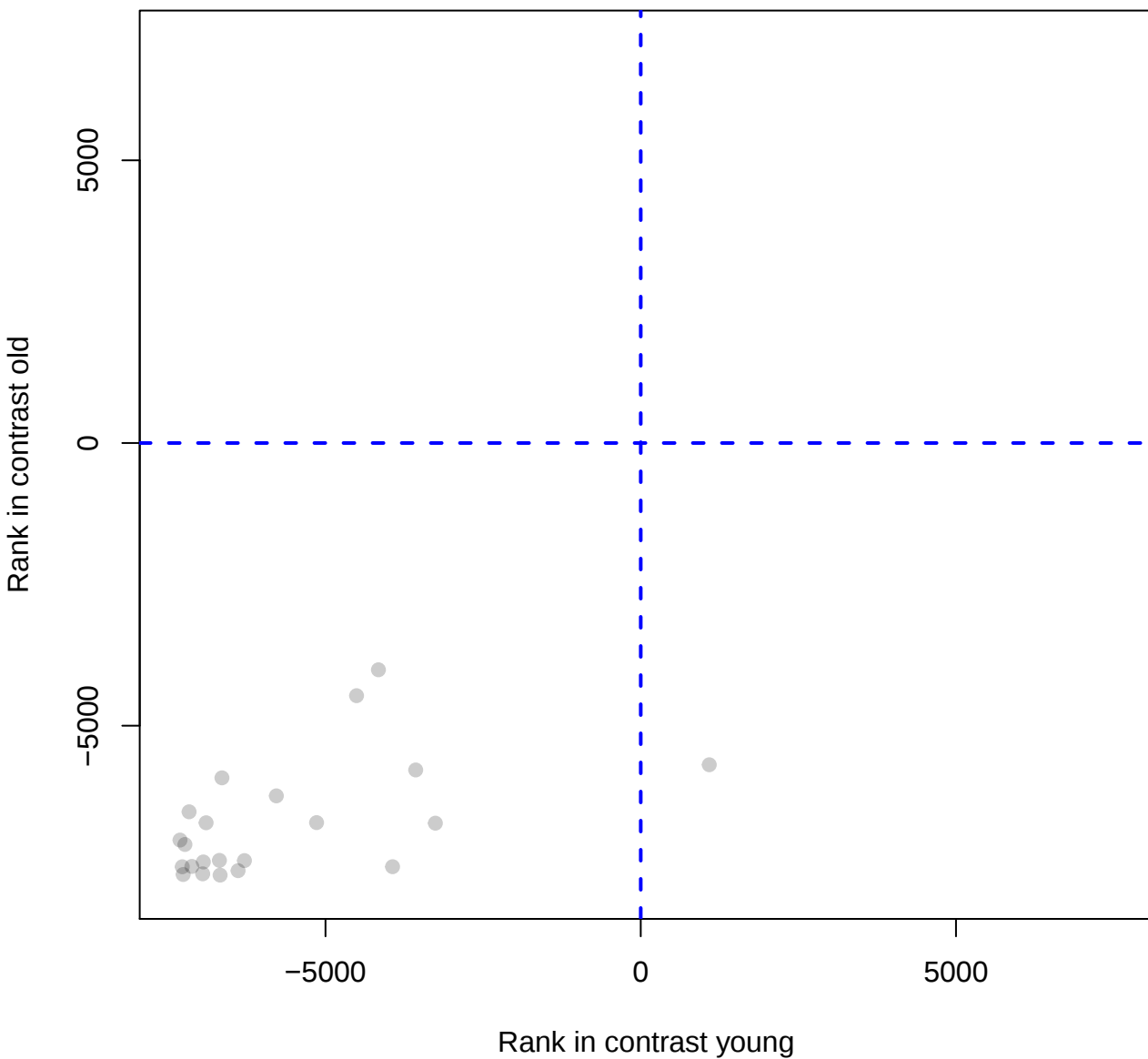
Citric-acid-cycle-(TCA-cycle)



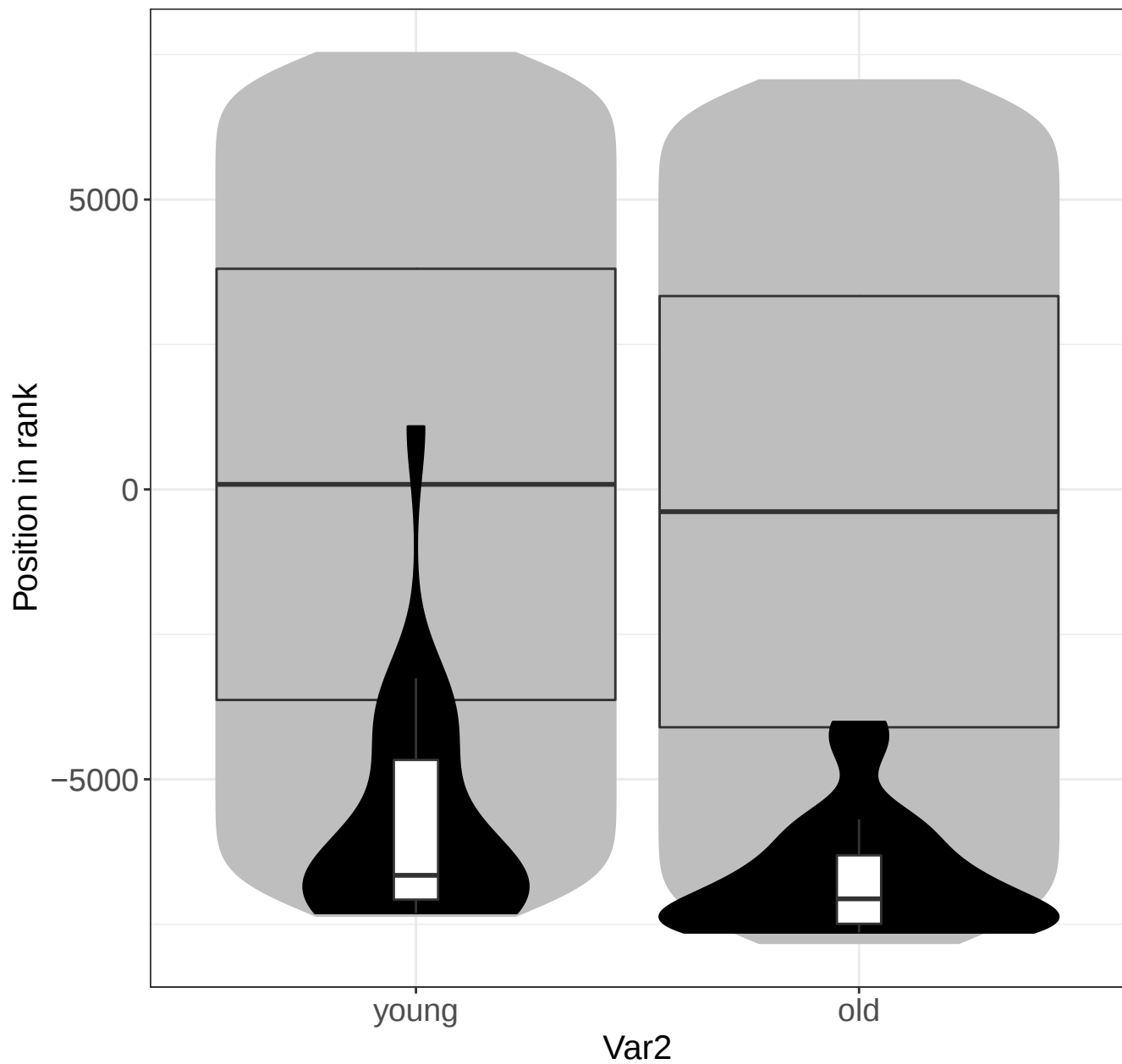
rRNA-processing-in-the-mitochondrion



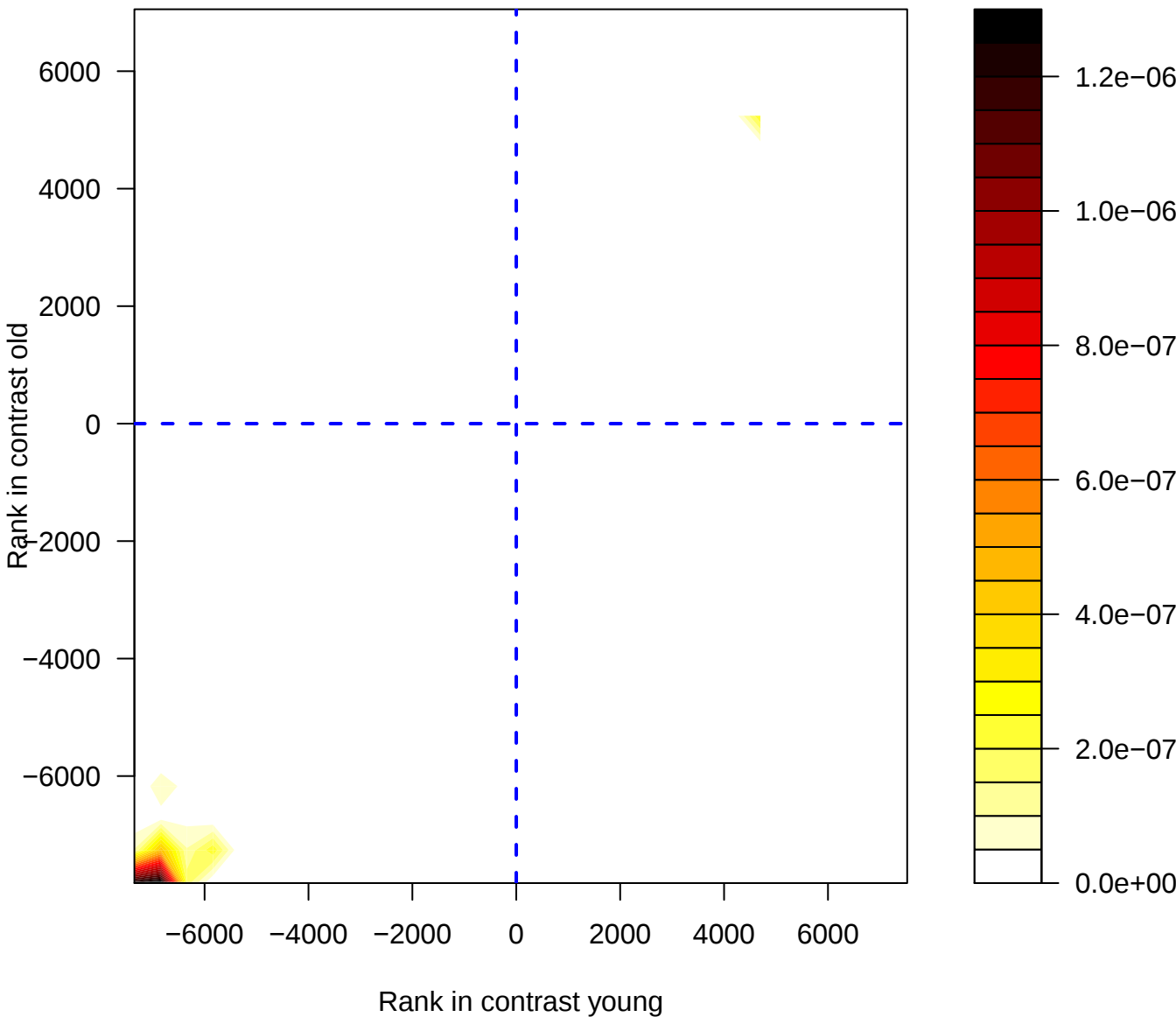
rRNA-processing-in-the-mitochondrion



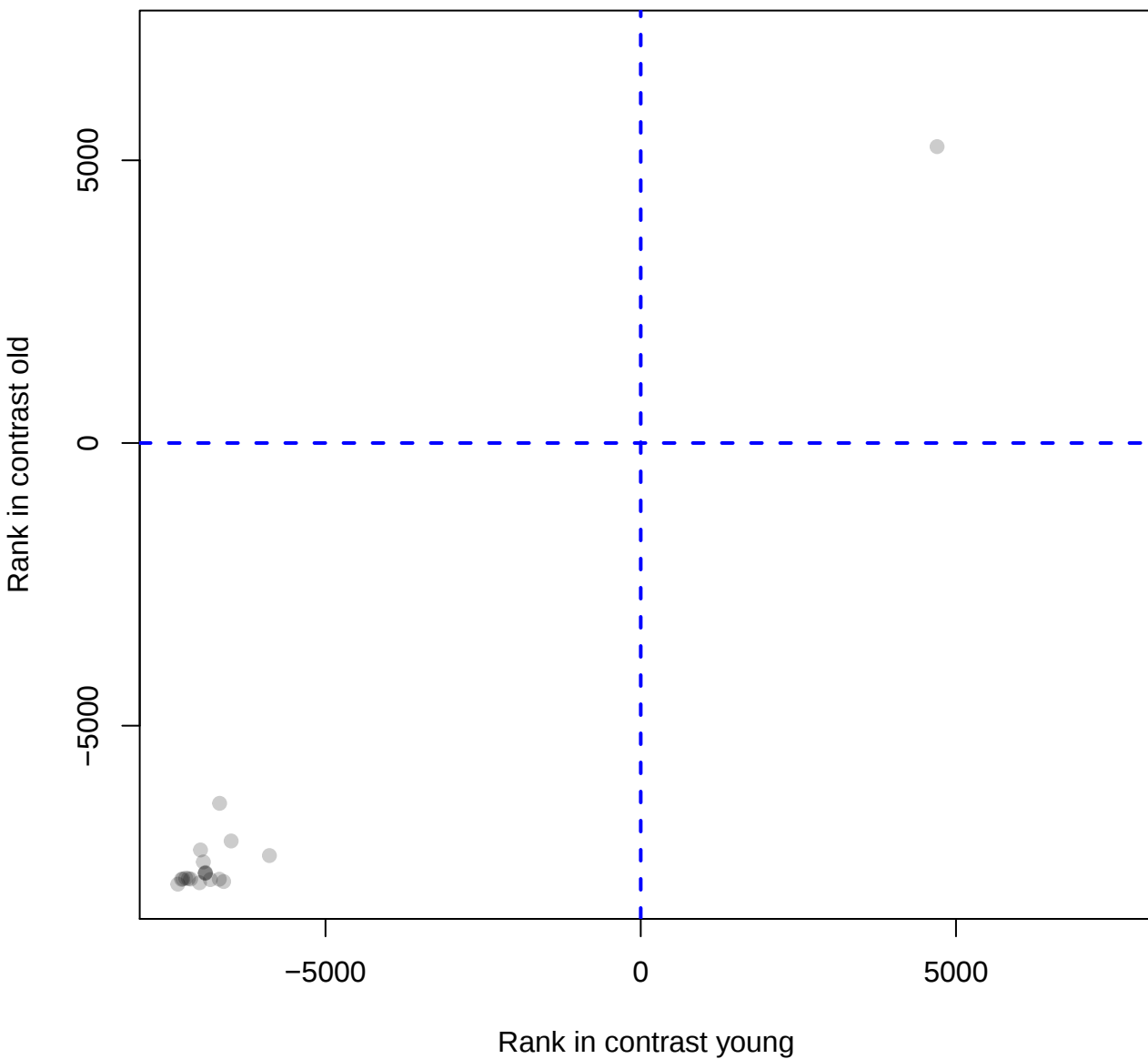
rRNA-processing-in-the-mitochondrion



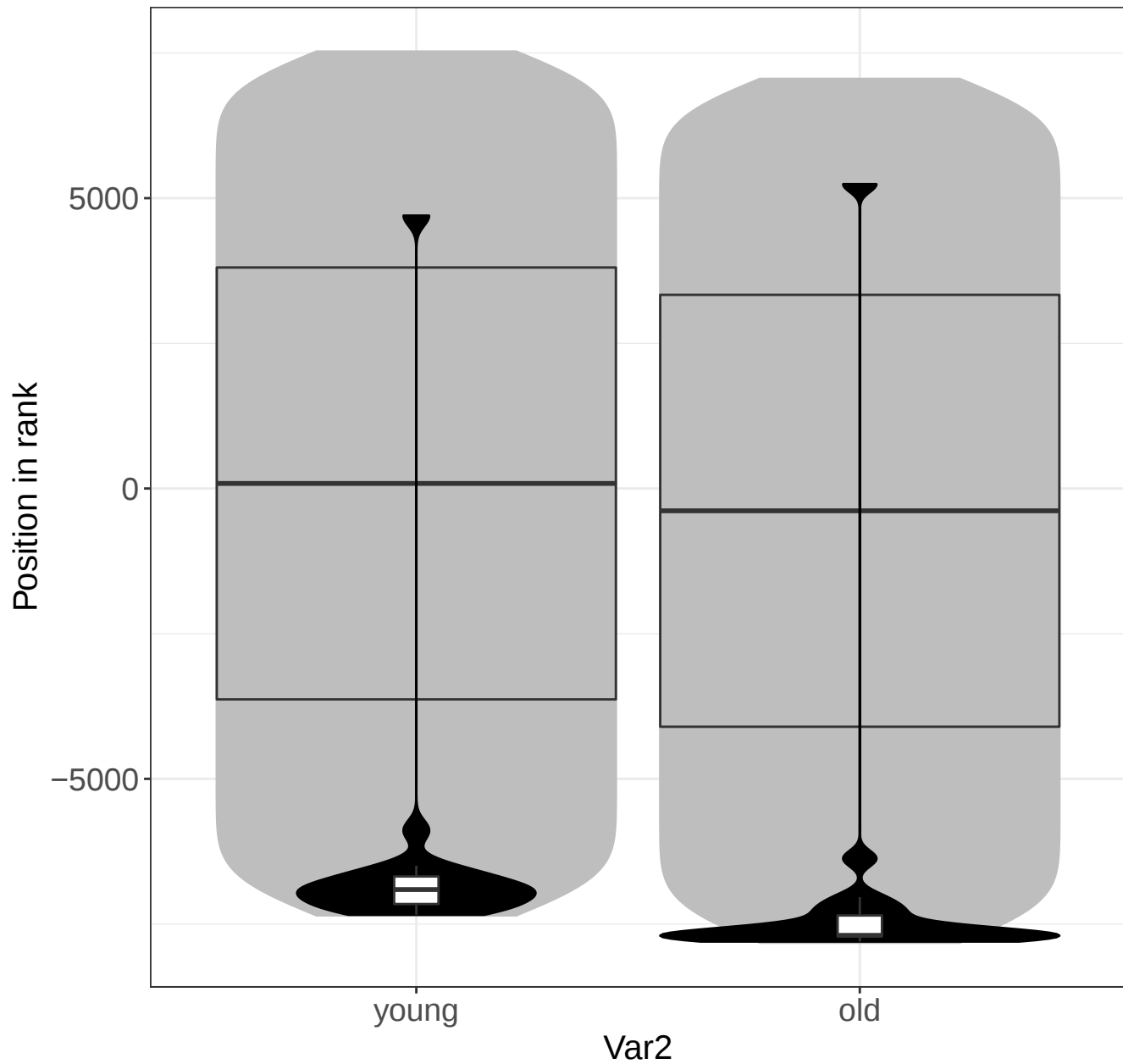
Branched-chain-amino-acid-catabolism



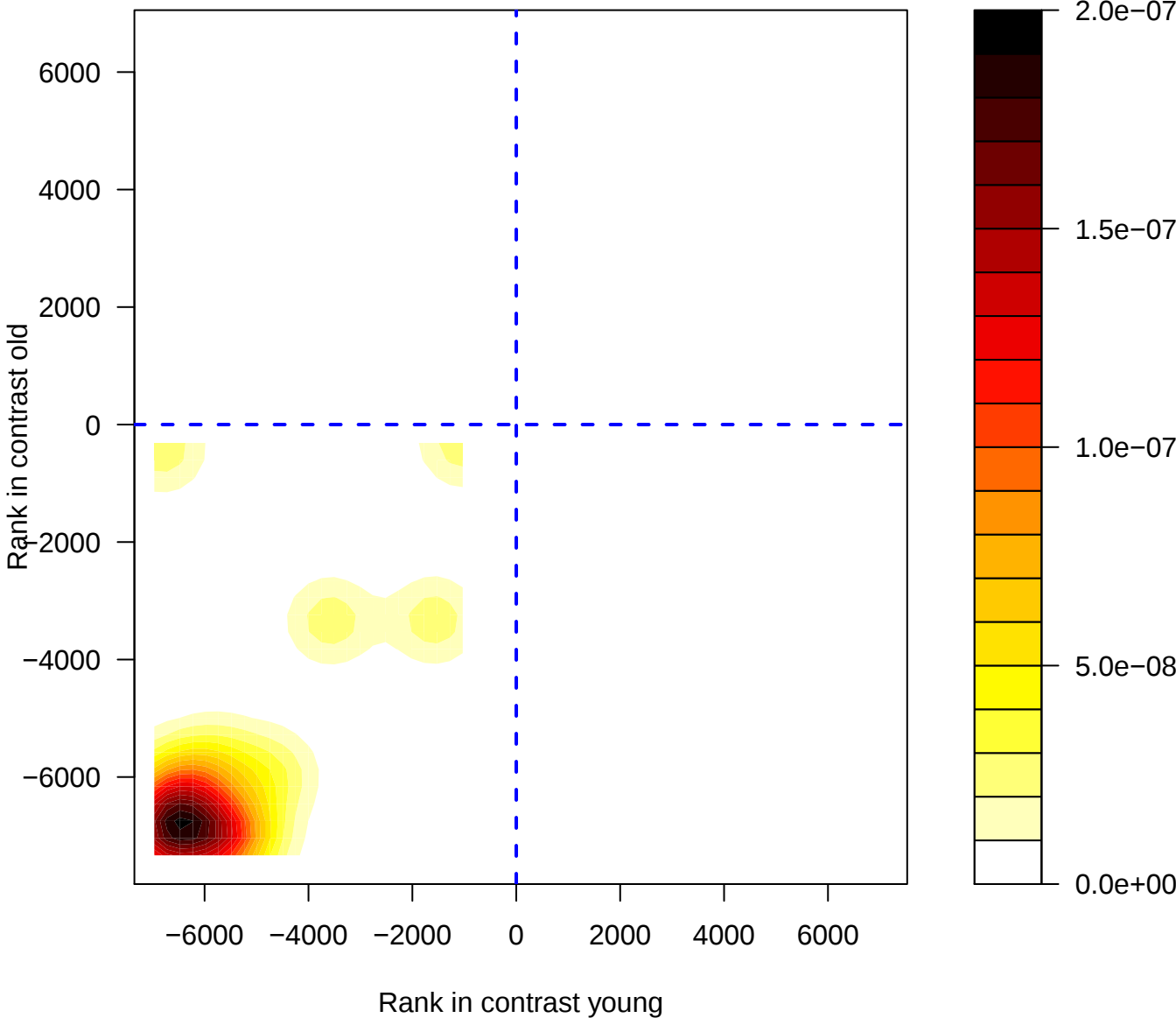
Branched-chain-amino-acid-catabolism



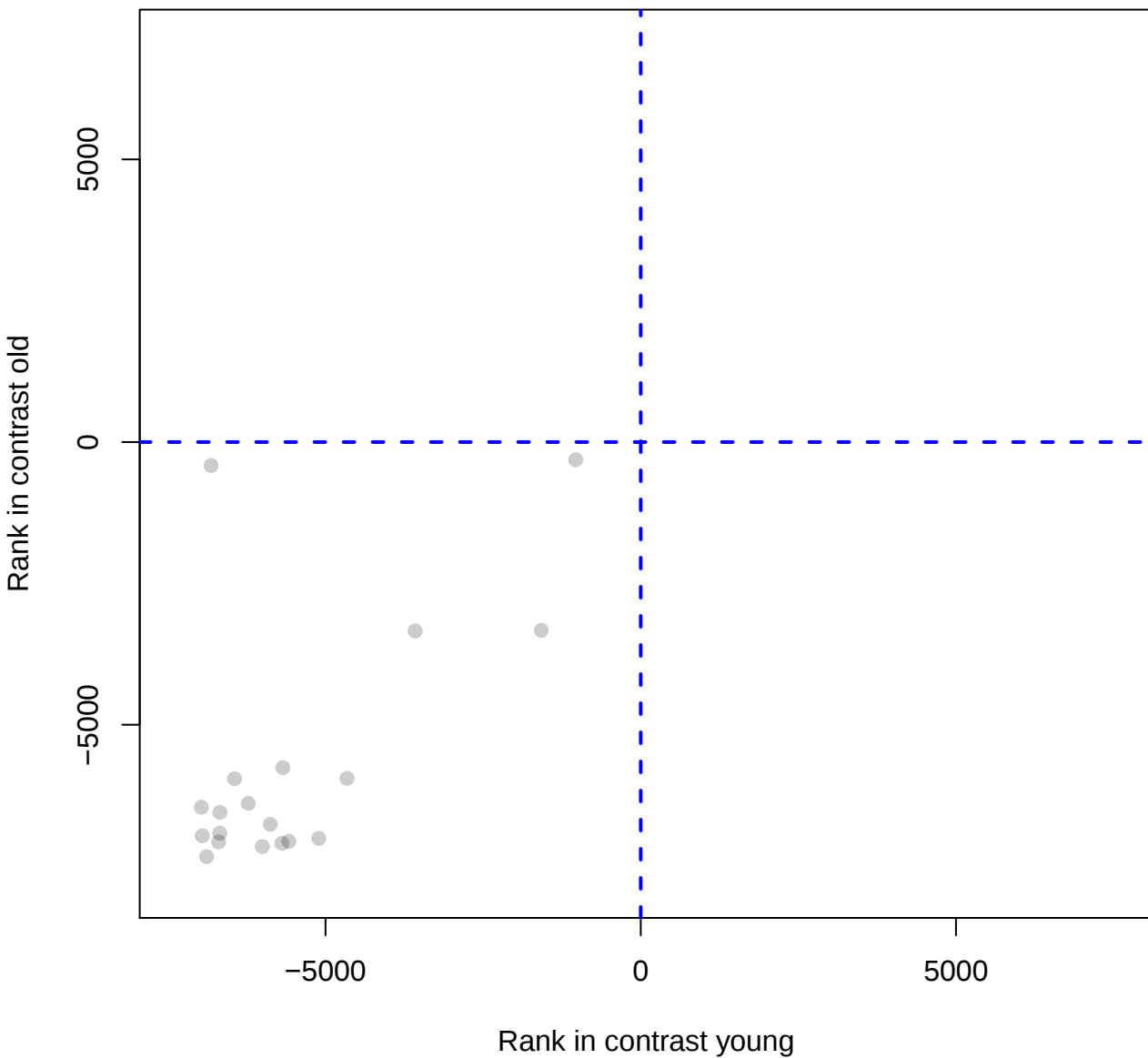
Branched-chain-amino-acid-catabolism



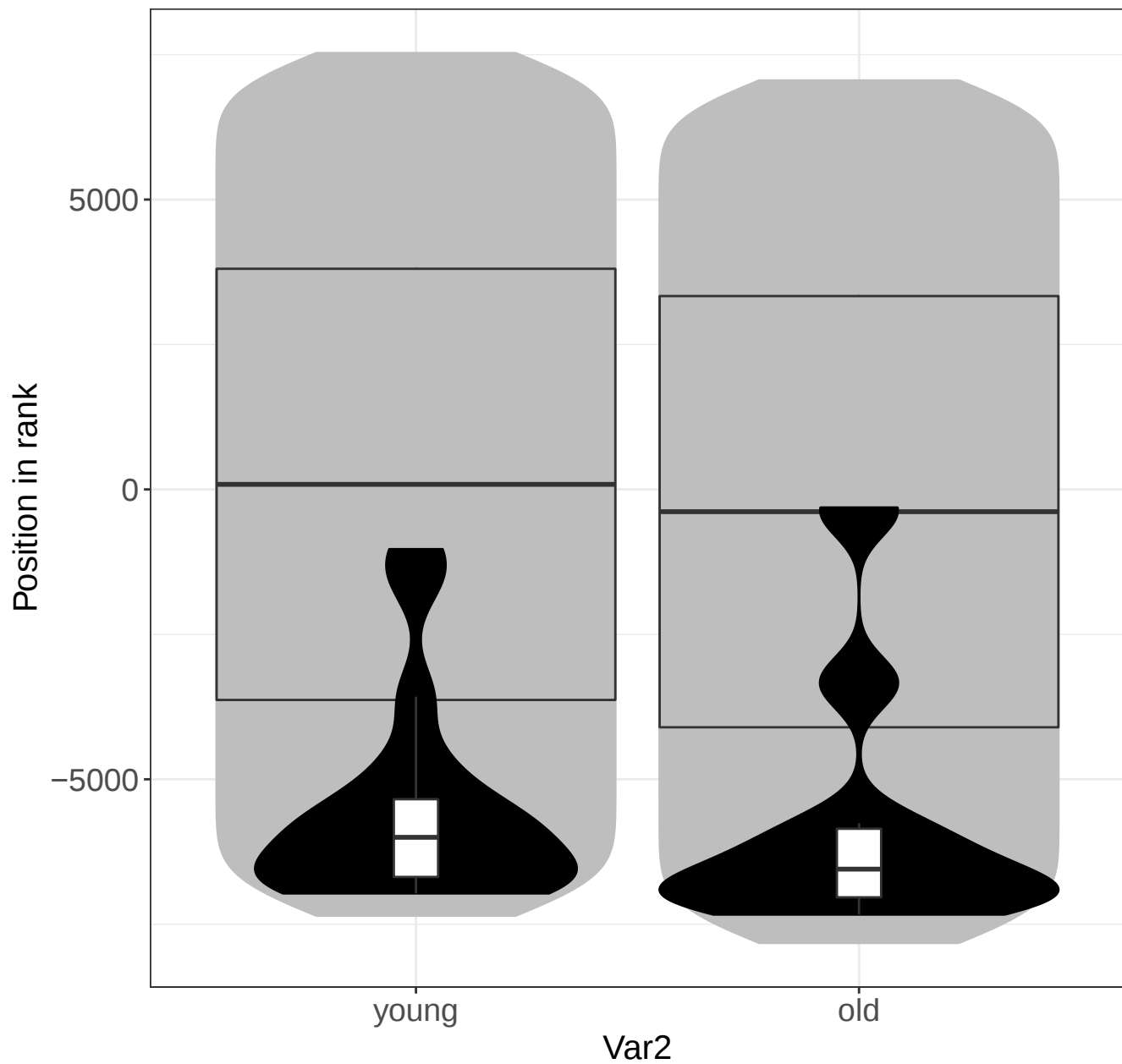
Mitochondrial-tRNA-aminoacylation



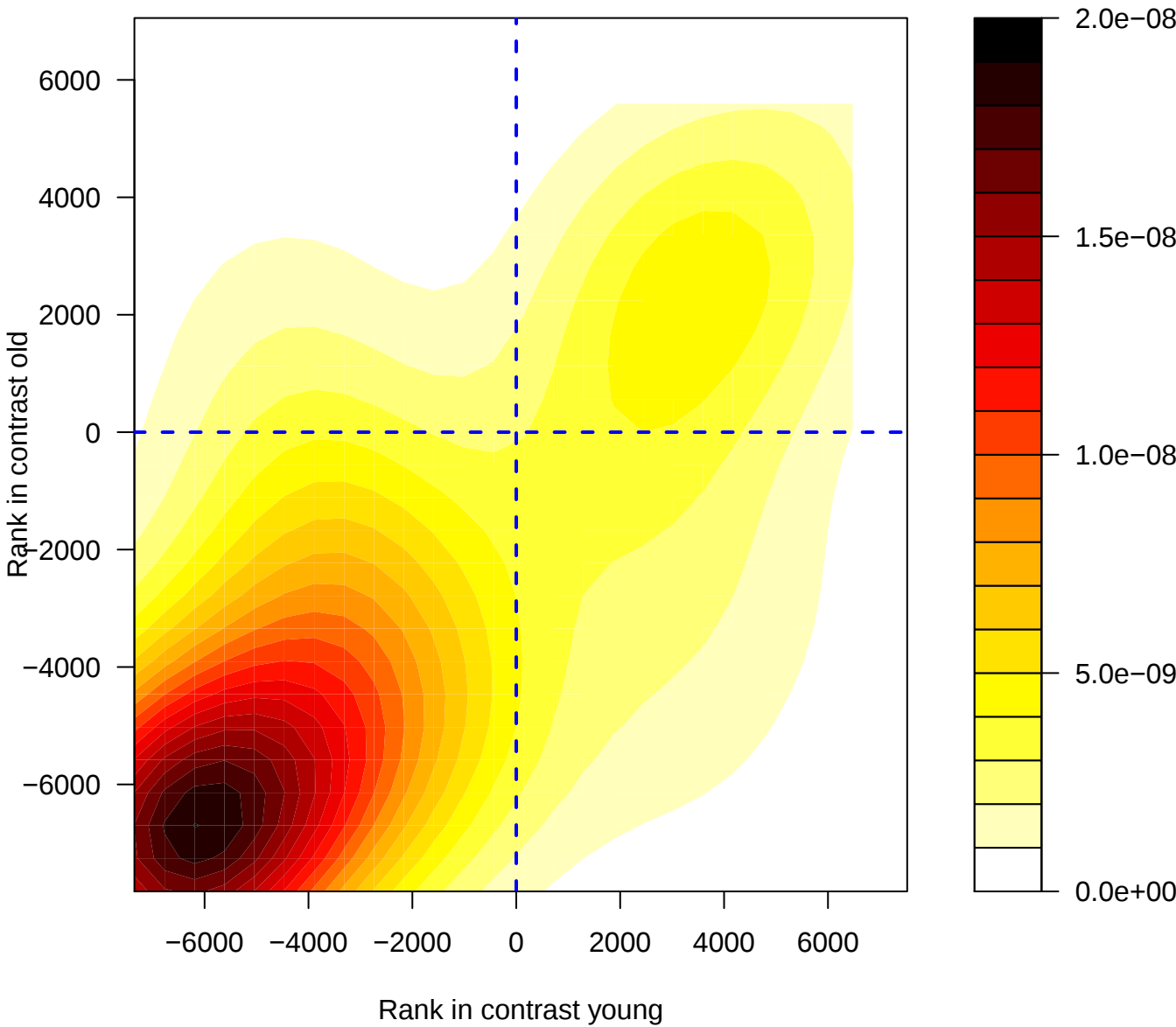
Mitochondrial-tRNA-aminoacylation



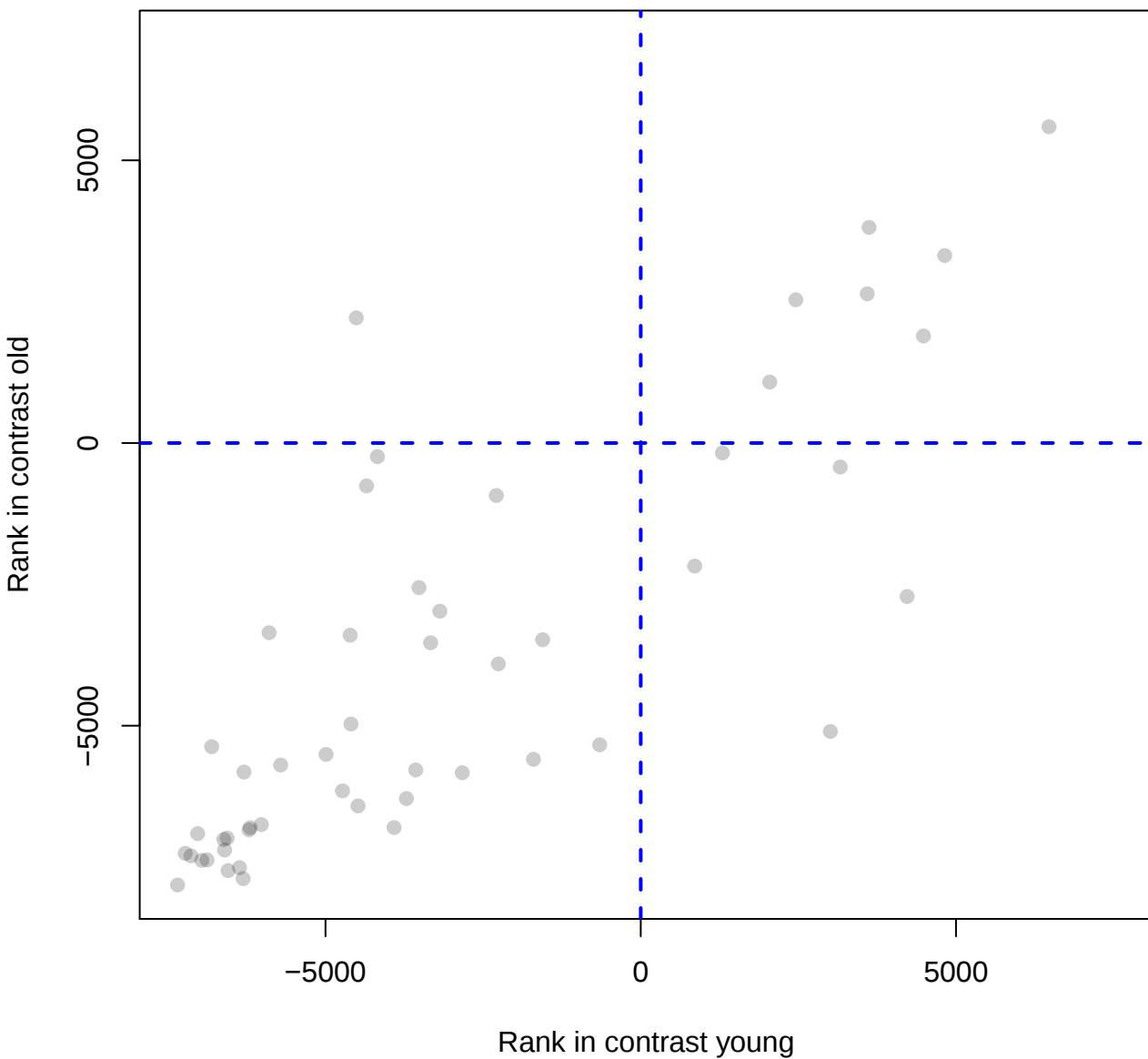
Mitochondrial-tRNA-aminoacylation



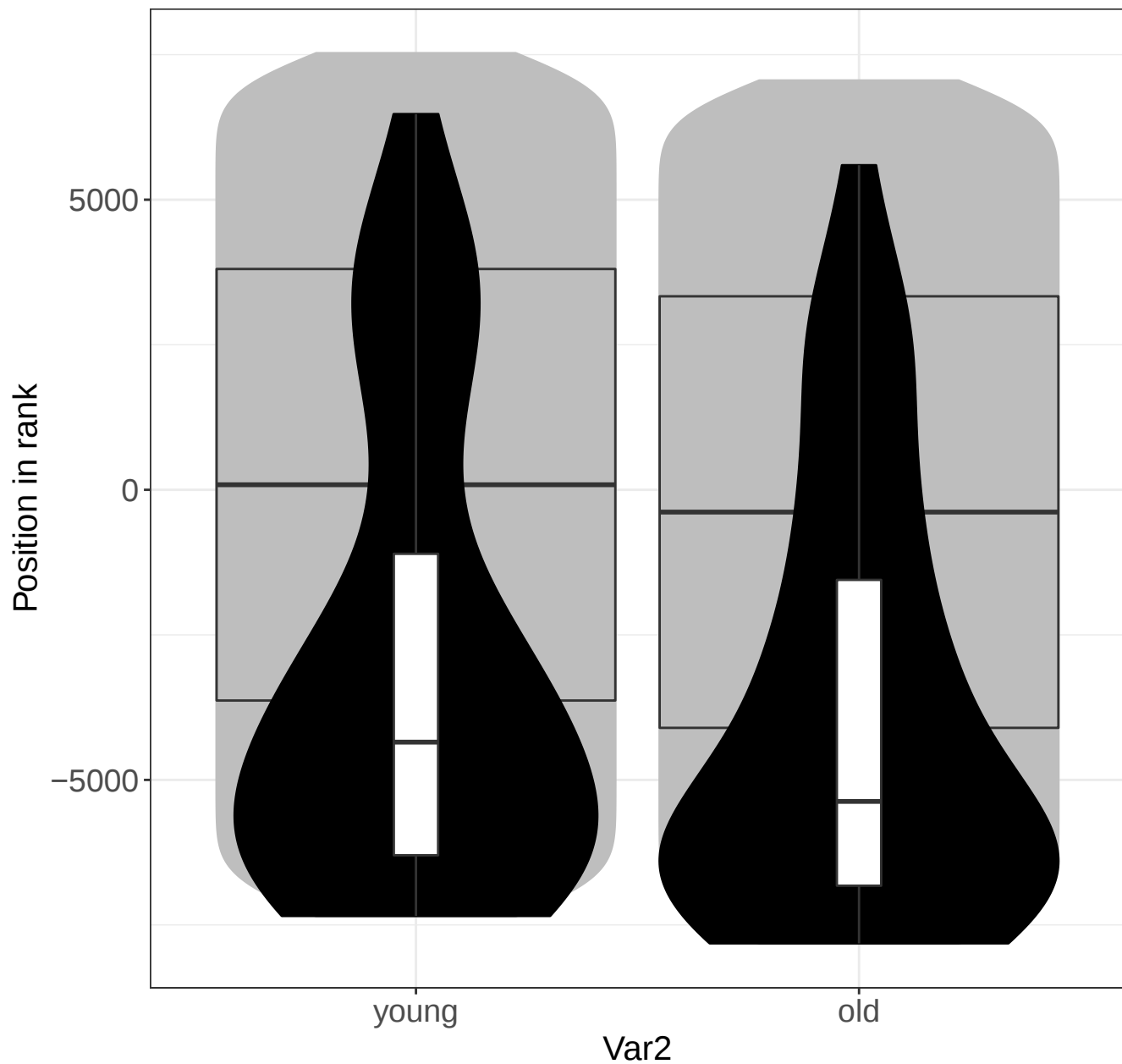
Transcriptional-activation-of-mitochondrial-biogenesis



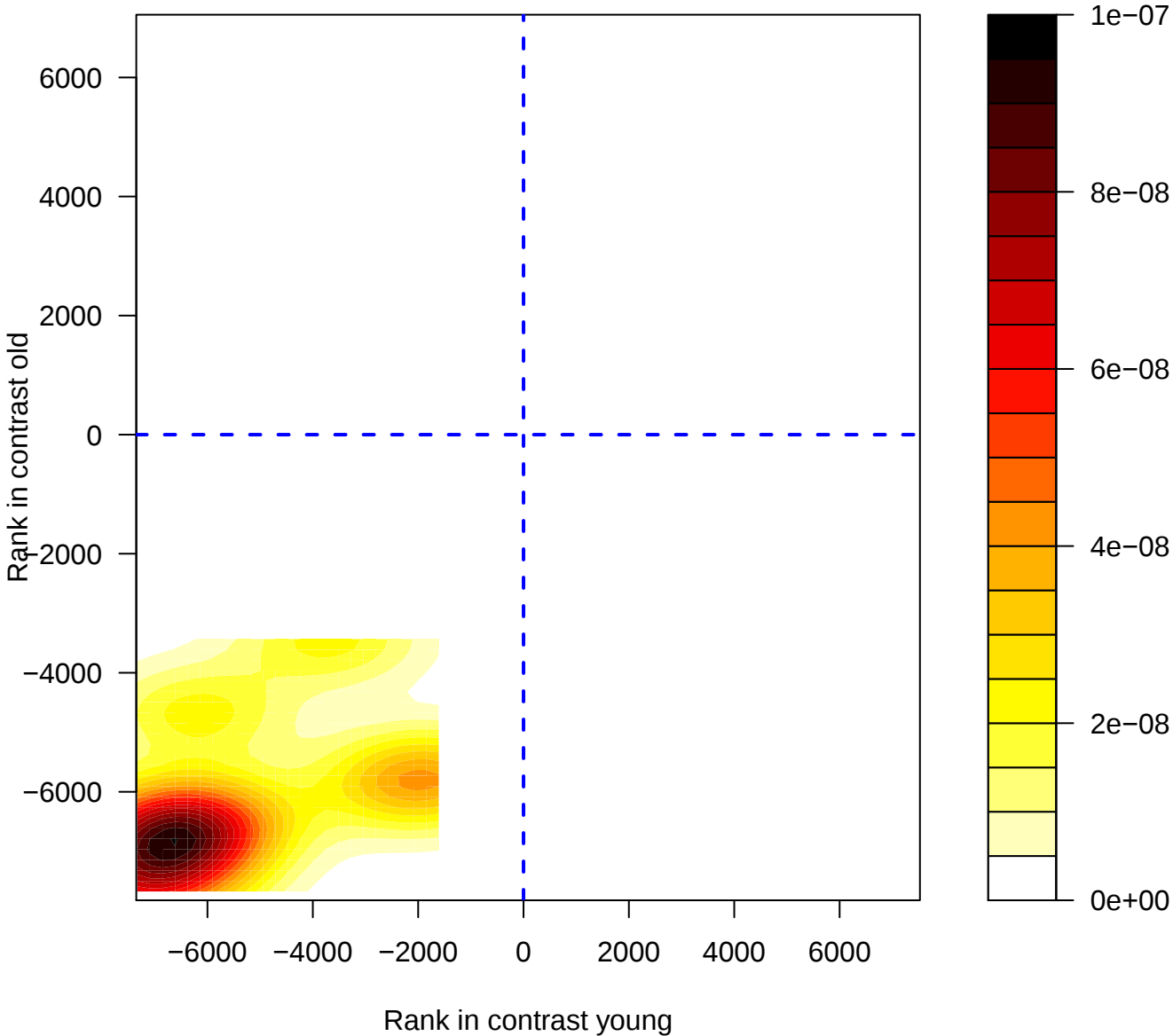
Transcriptional-activation-of-mitochondrial-biogenesis



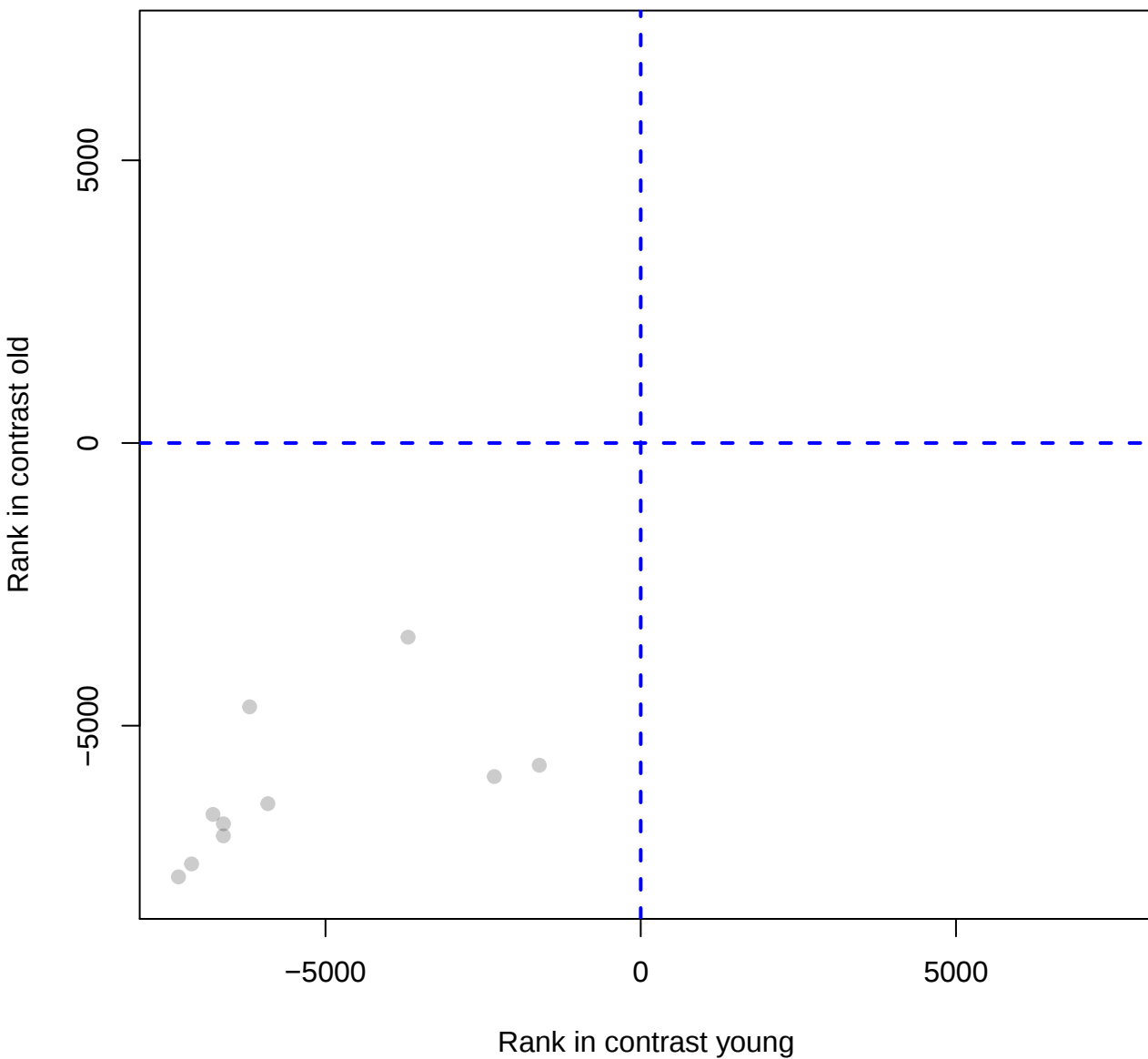
Transcriptional-activation-of-mitochondrial-bioge



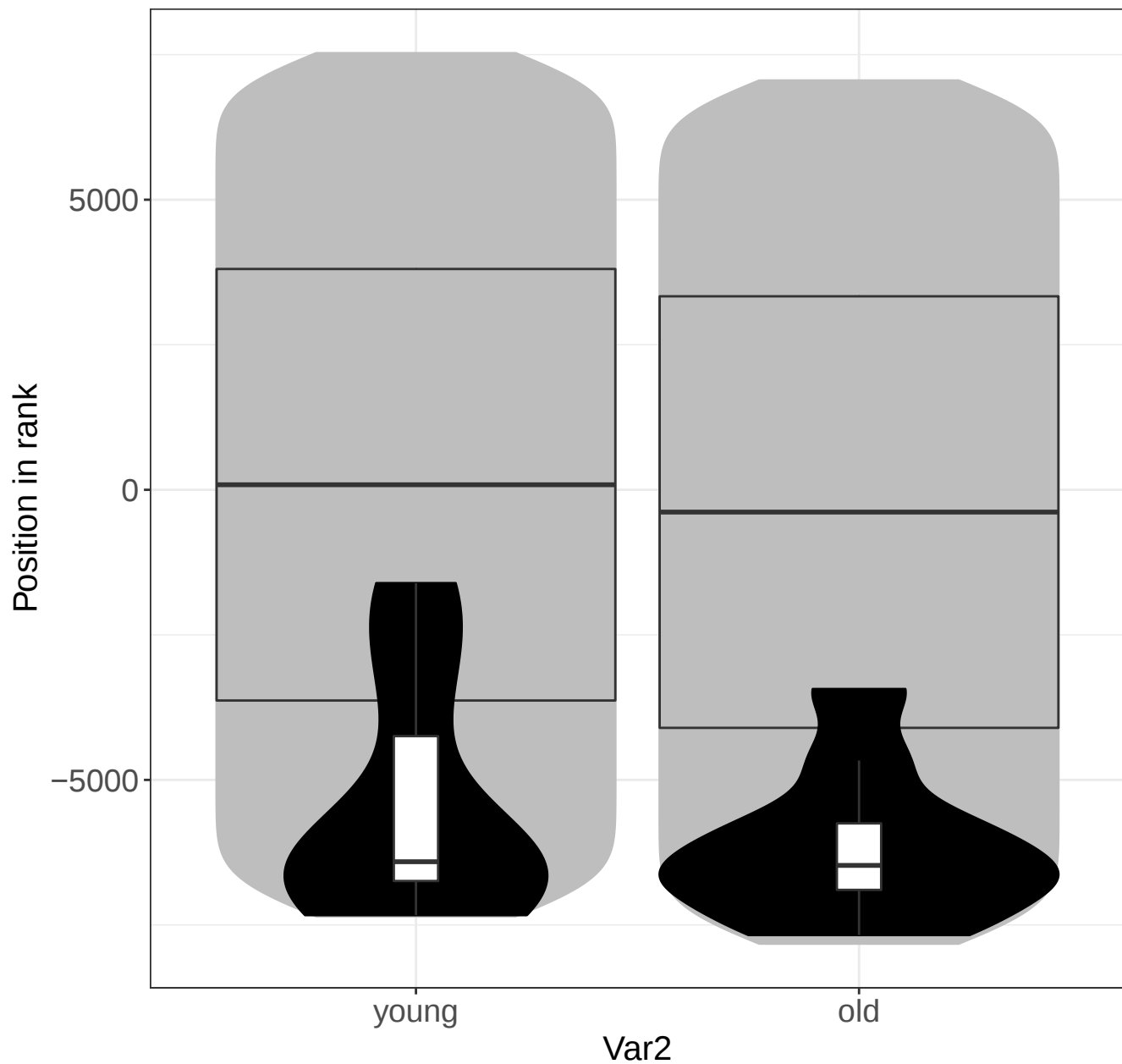
Mitochondrial-iron-sulfur-cluster-biogenesis



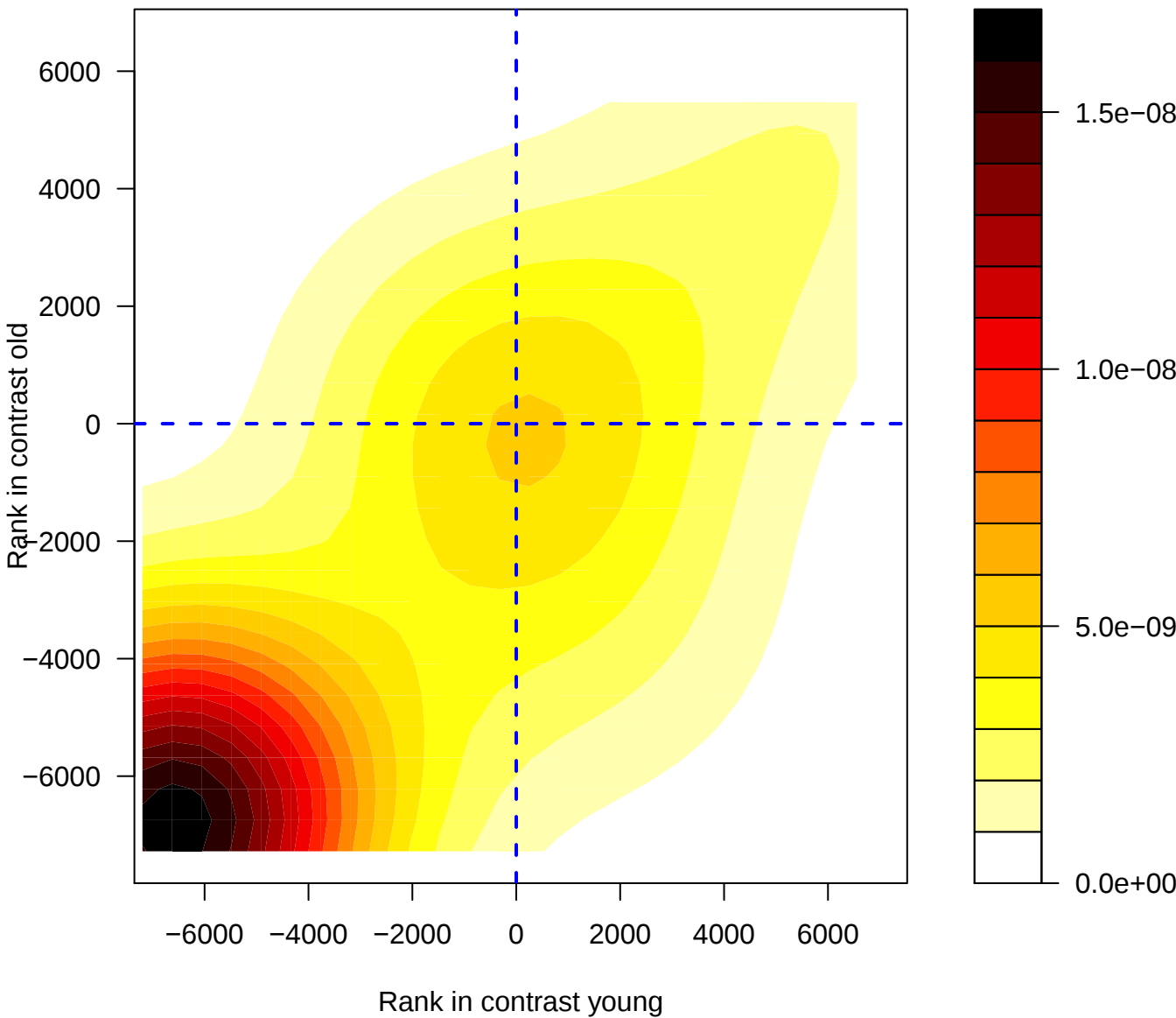
Mitochondrial-iron-sulfur-cluster-biogenesis



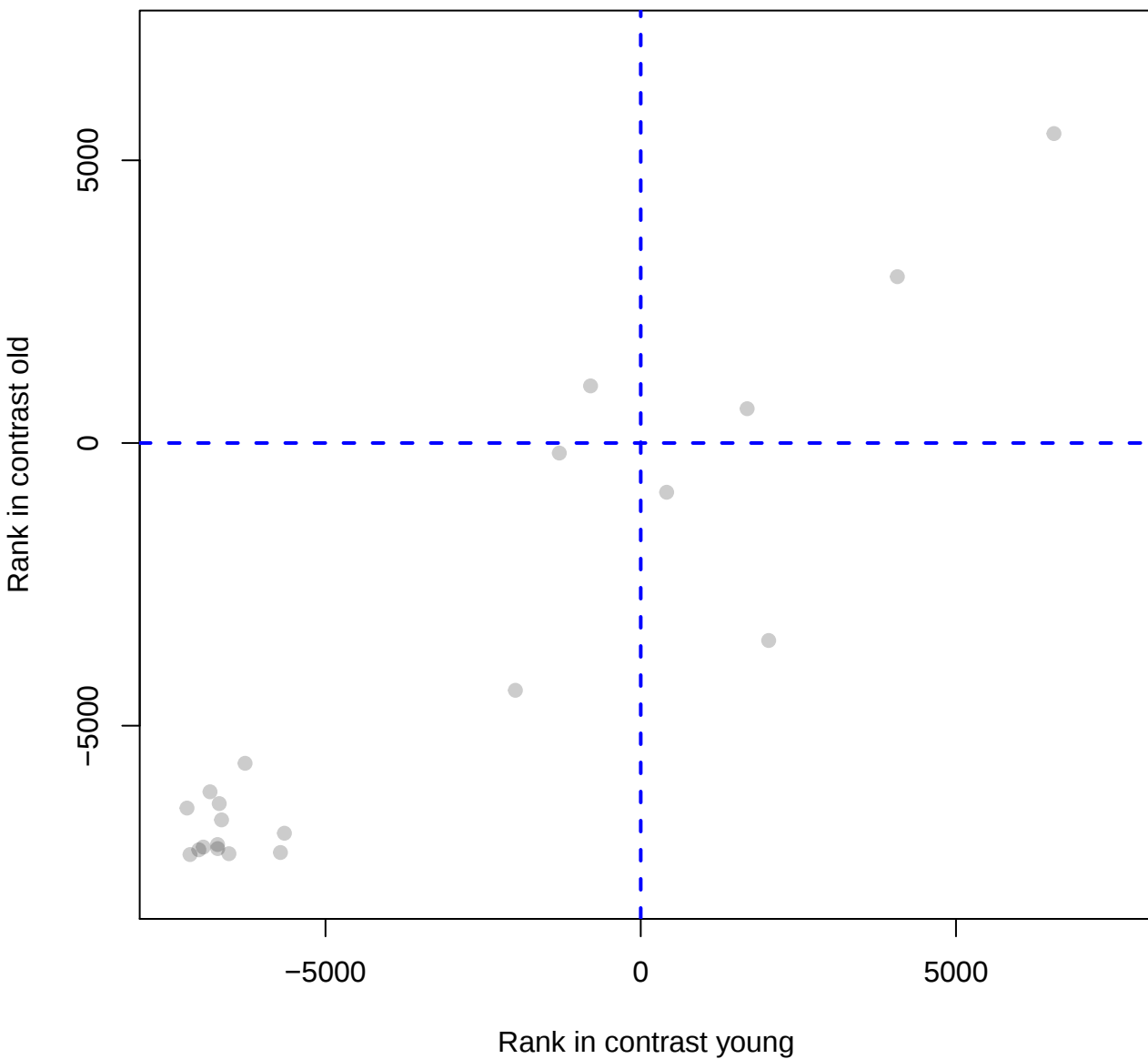
Mitochondrial-iron-sulfur-cluster-biogenesis



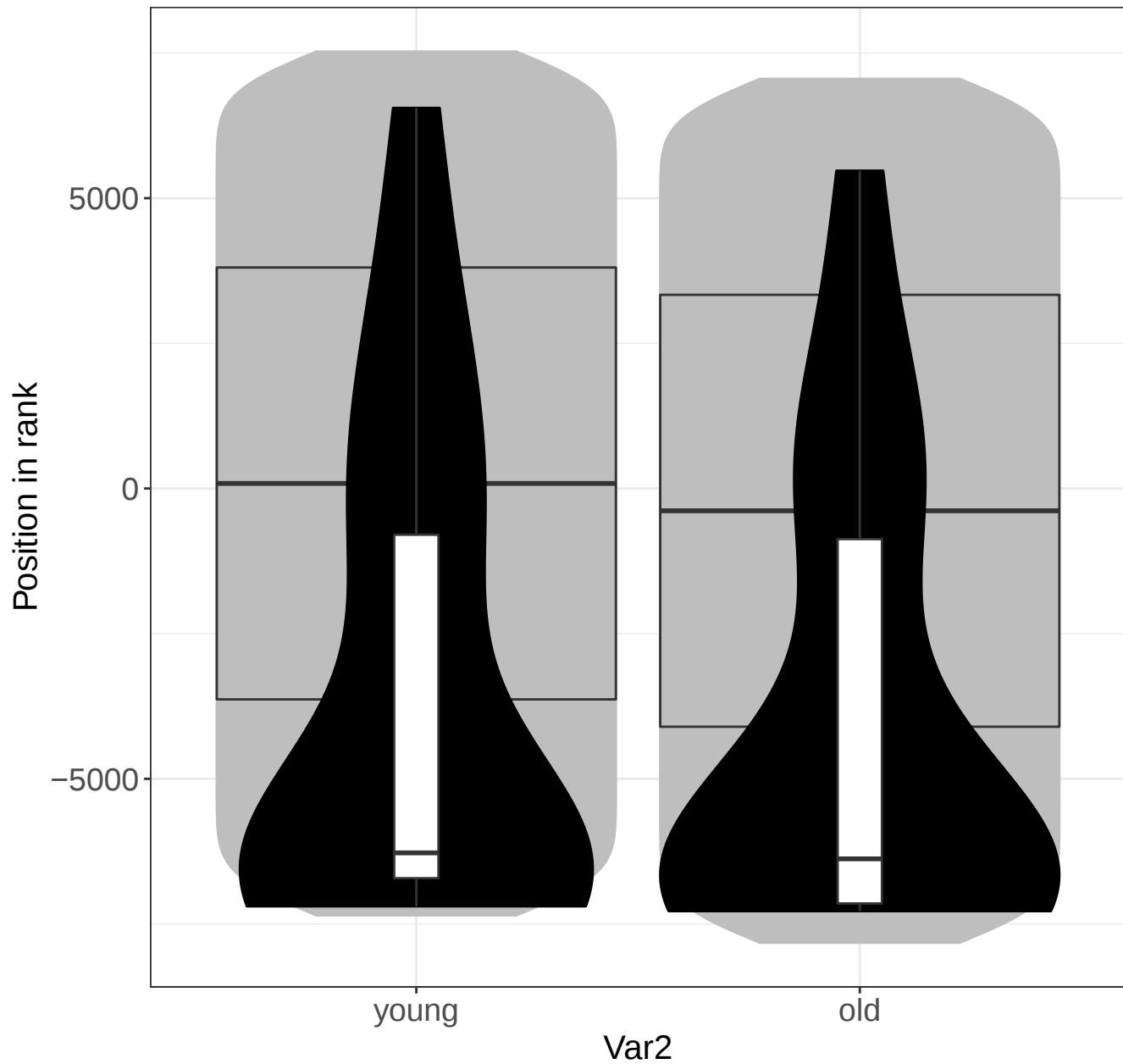
Mitochondrial-calcium-ion-transport



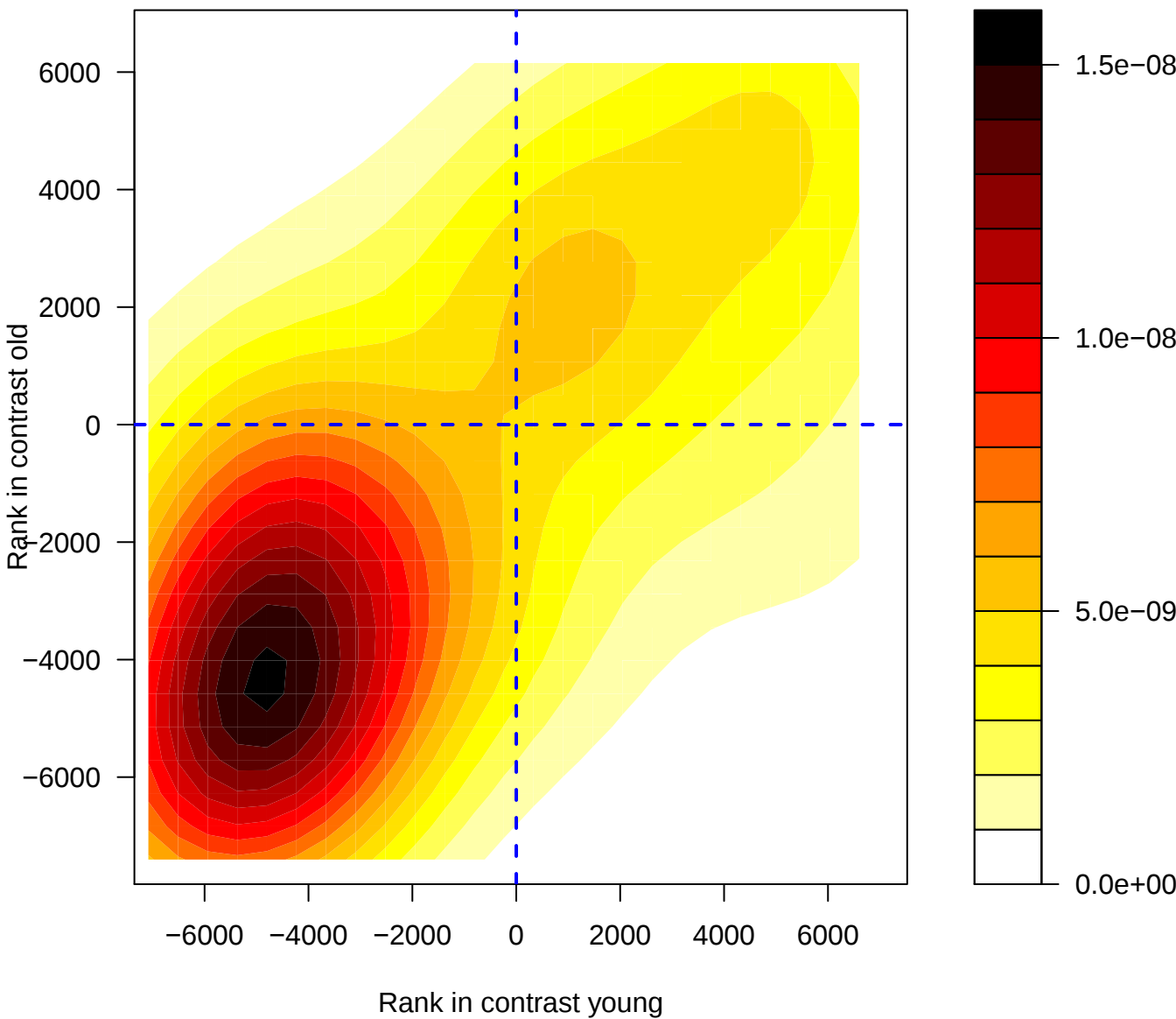
Mitochondrial-calcium-ion-transport



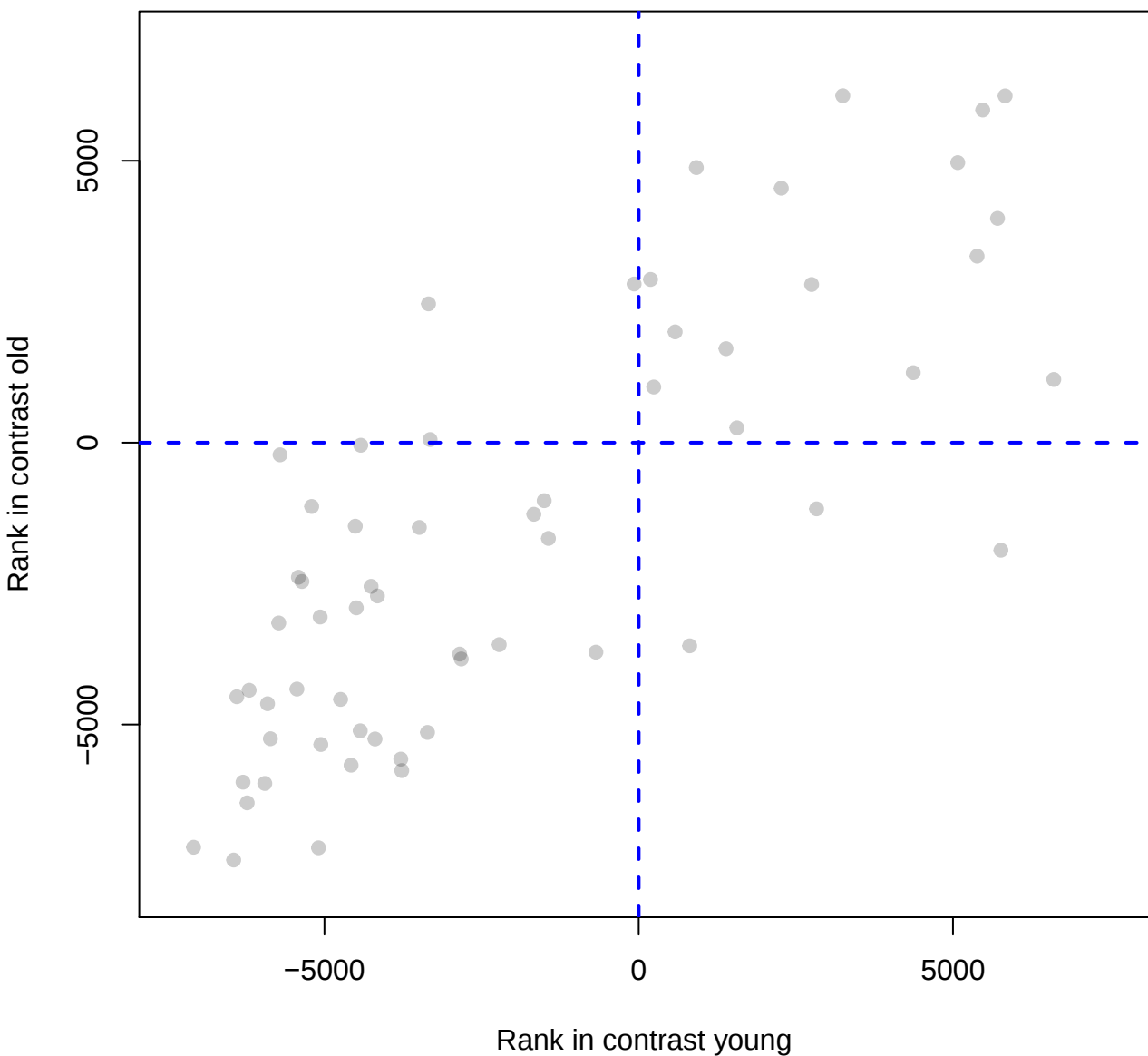
Mitochondrial-calcium-ion-transport



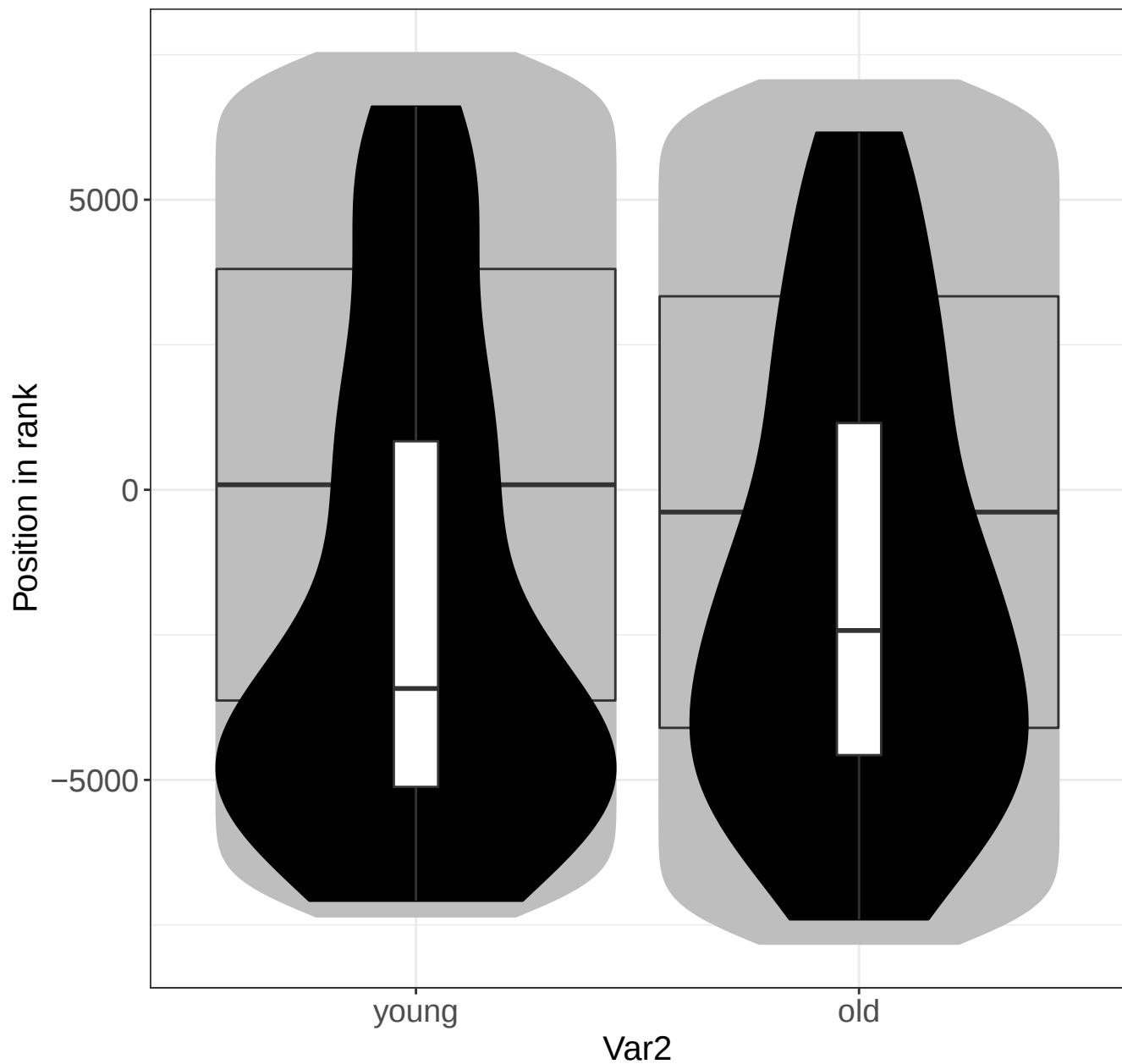
n-dependent-proline-hydroxylation-of-Hypoxia-inducible-I



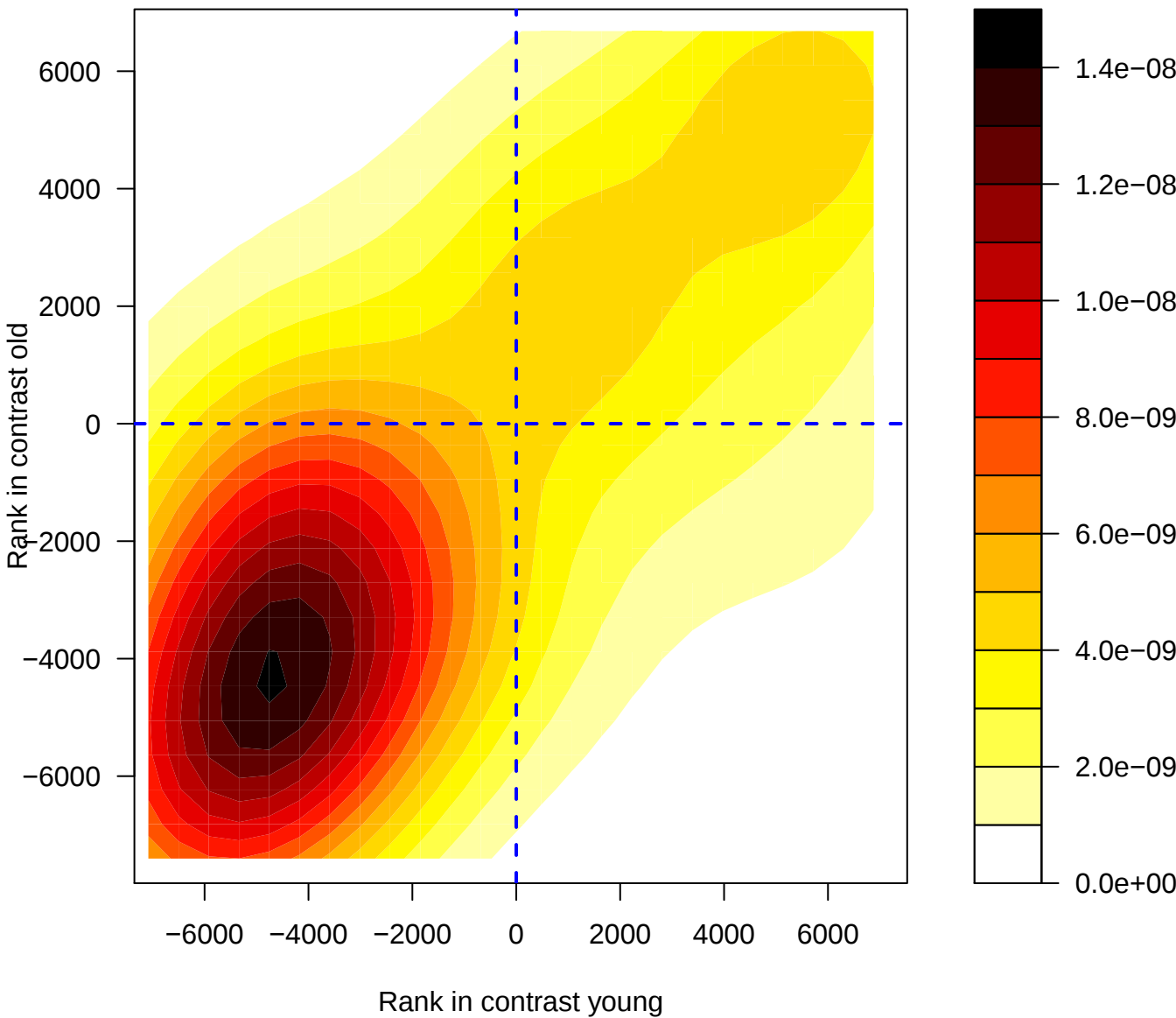
Oxygen-dependent-proline-hydroxylation-of-Hypoxia-inducible-Factor-A



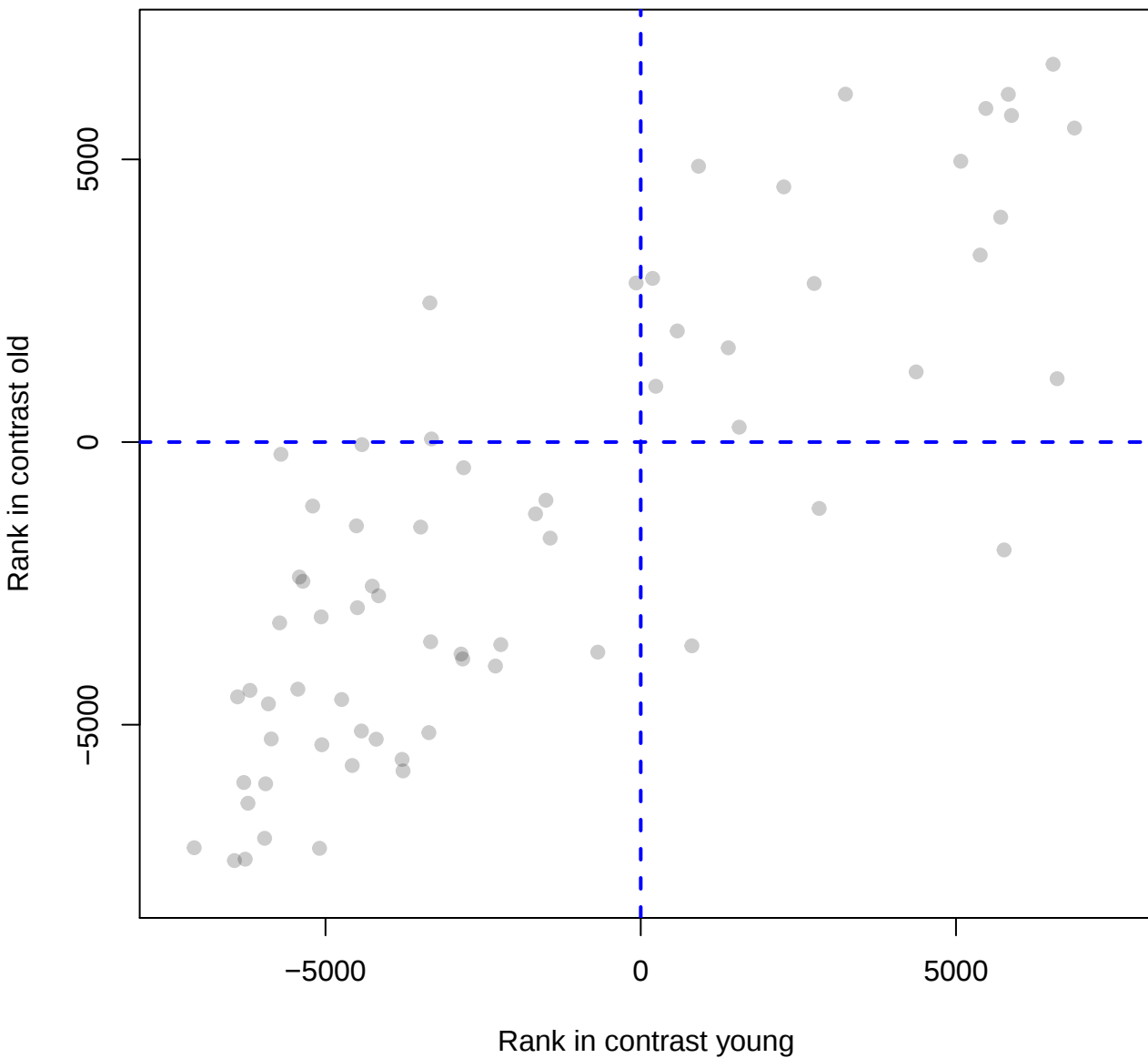
Oxygen-dependent-proline-hydroxylation-of-Hy



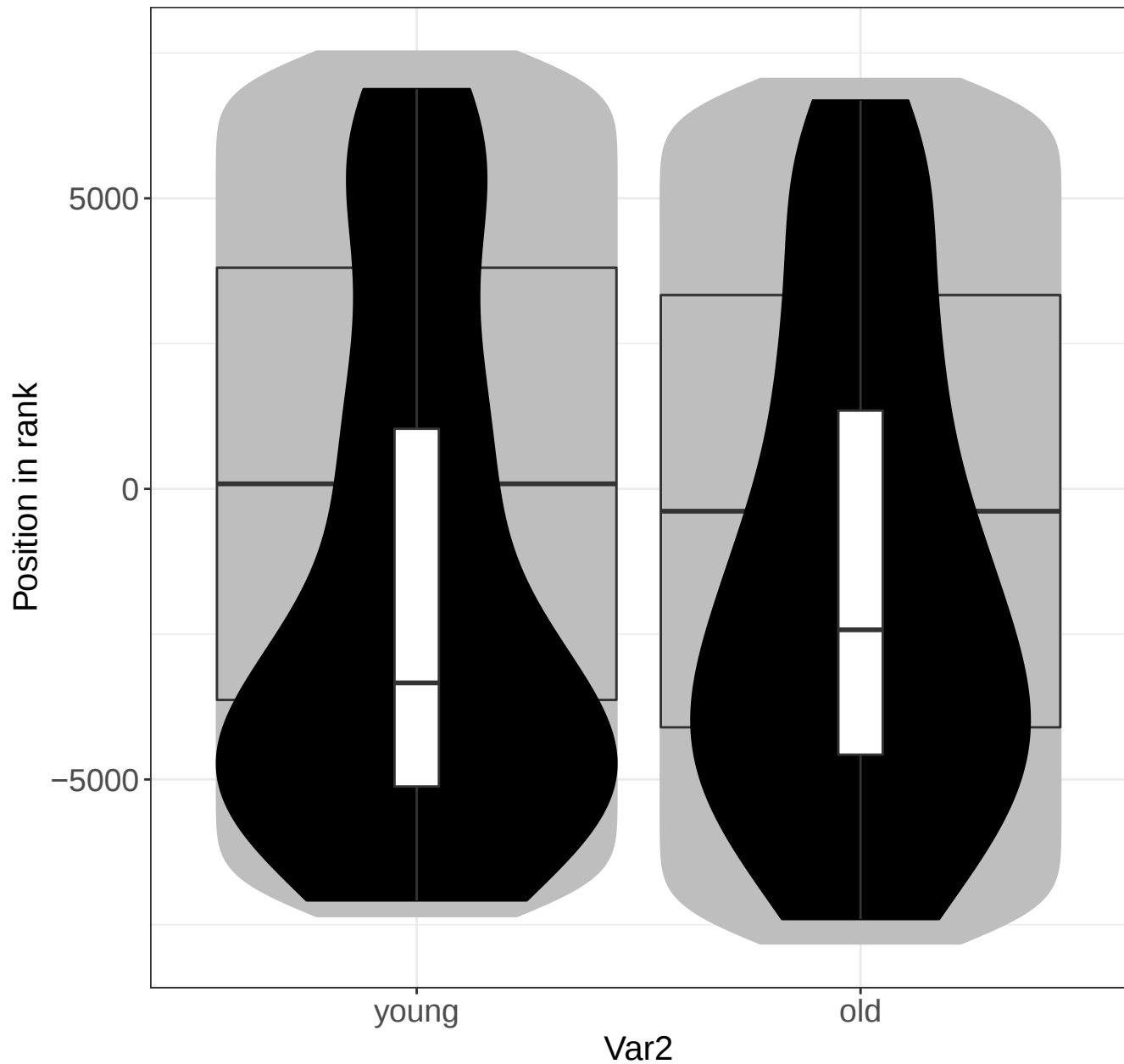
Cellular-response-to-hypoxia



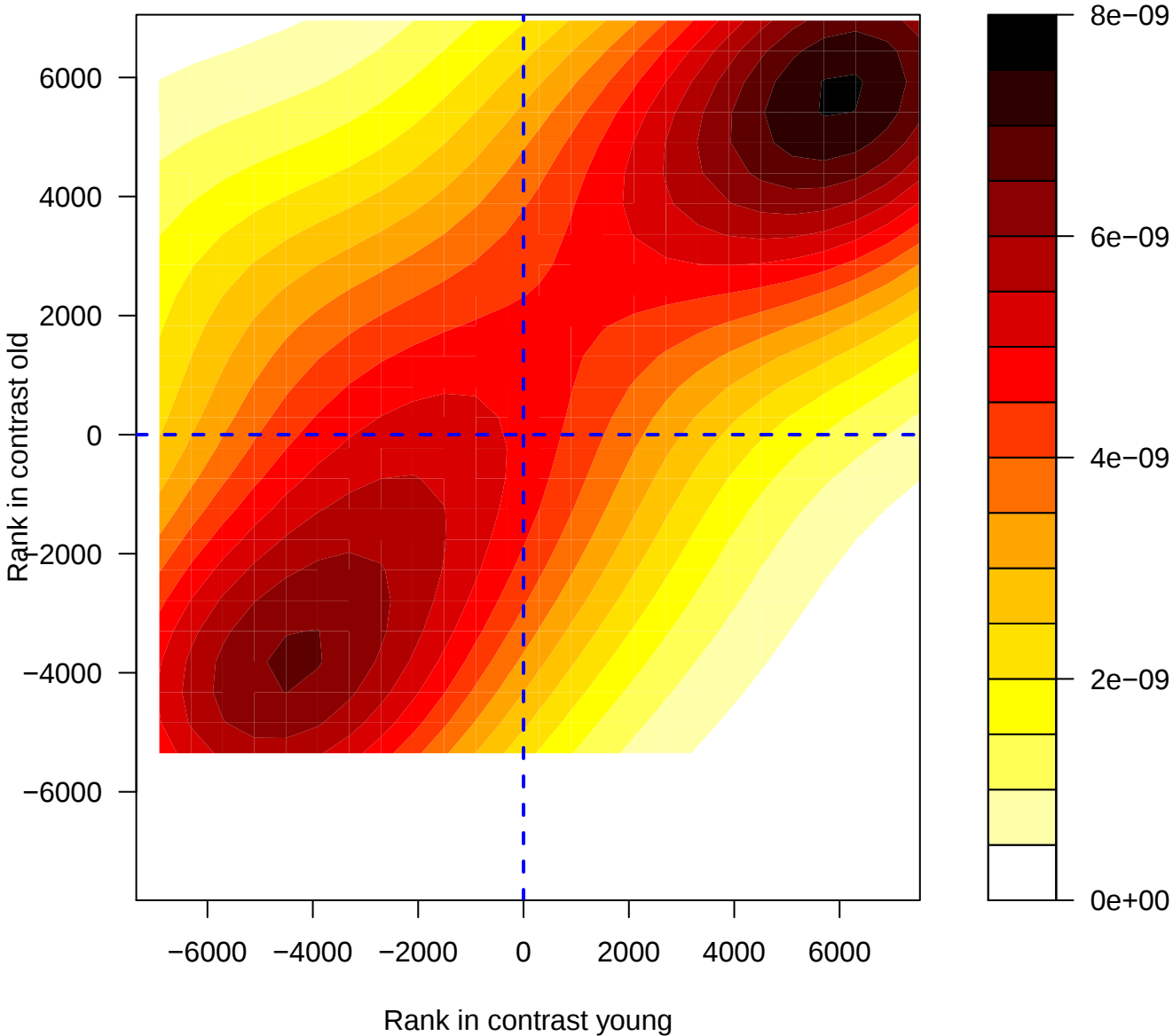
Cellular-response-to-hypoxia



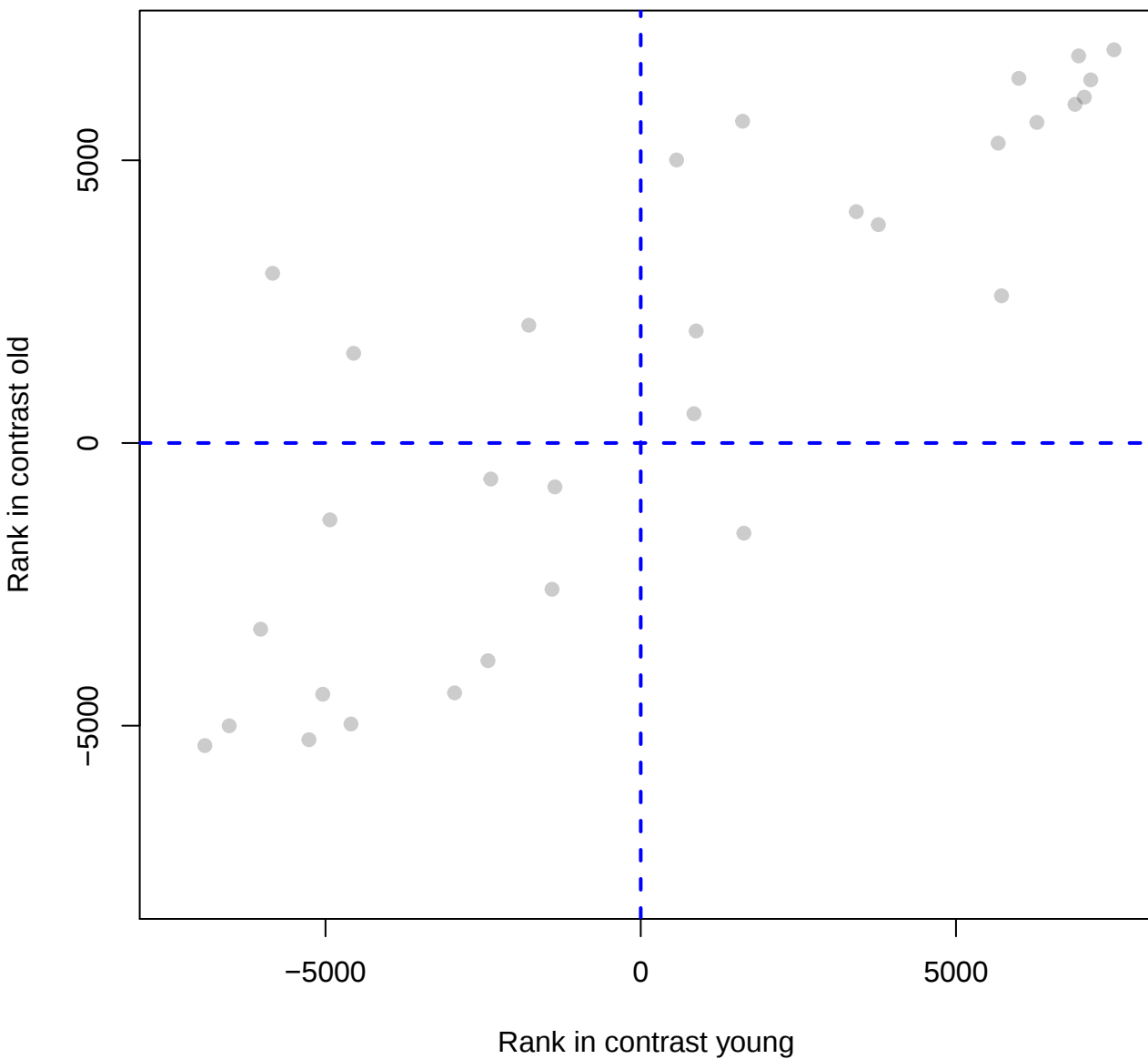
Cellular-response-to-hypoxia



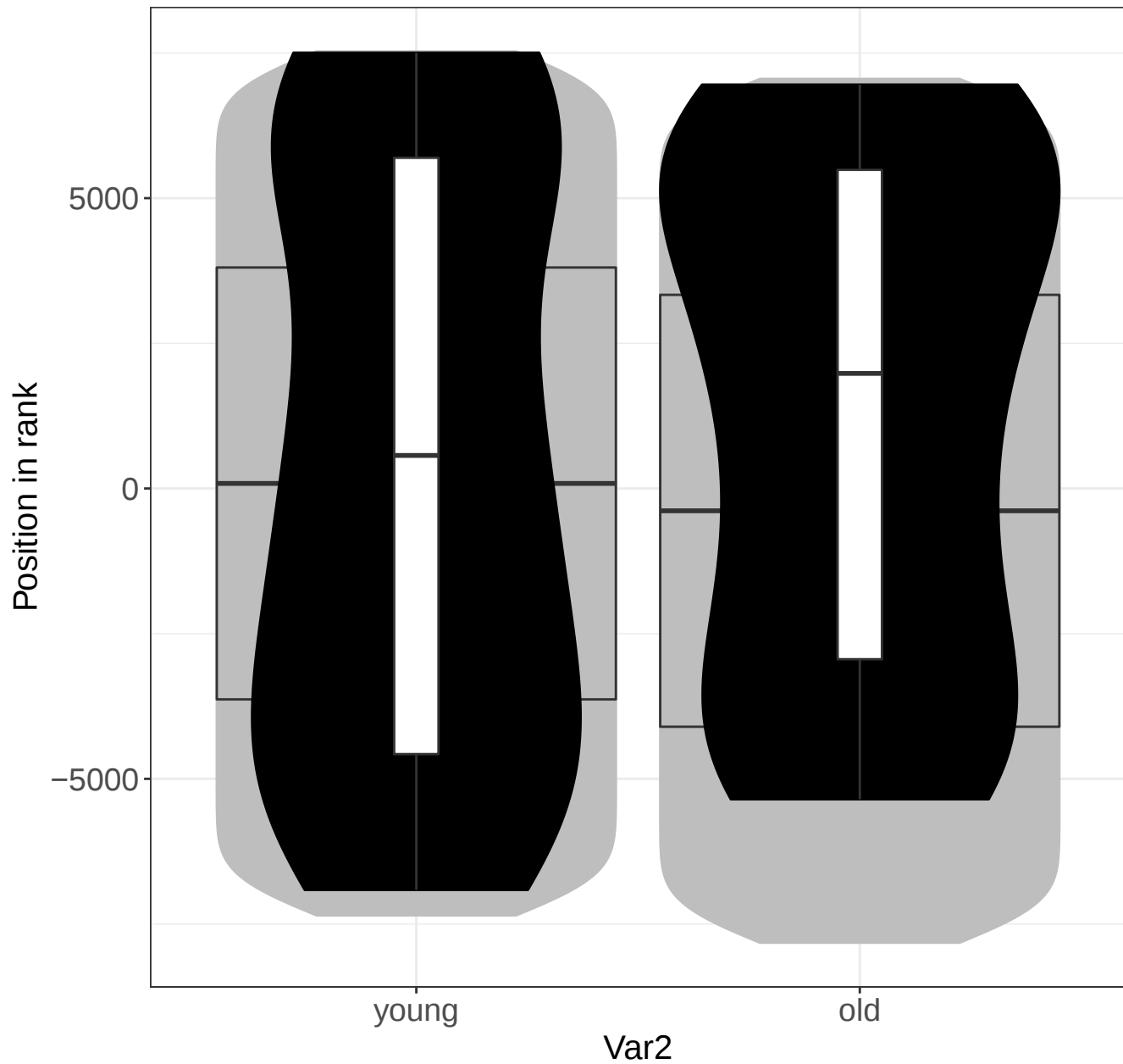
Smooth-Muscle-Contraction



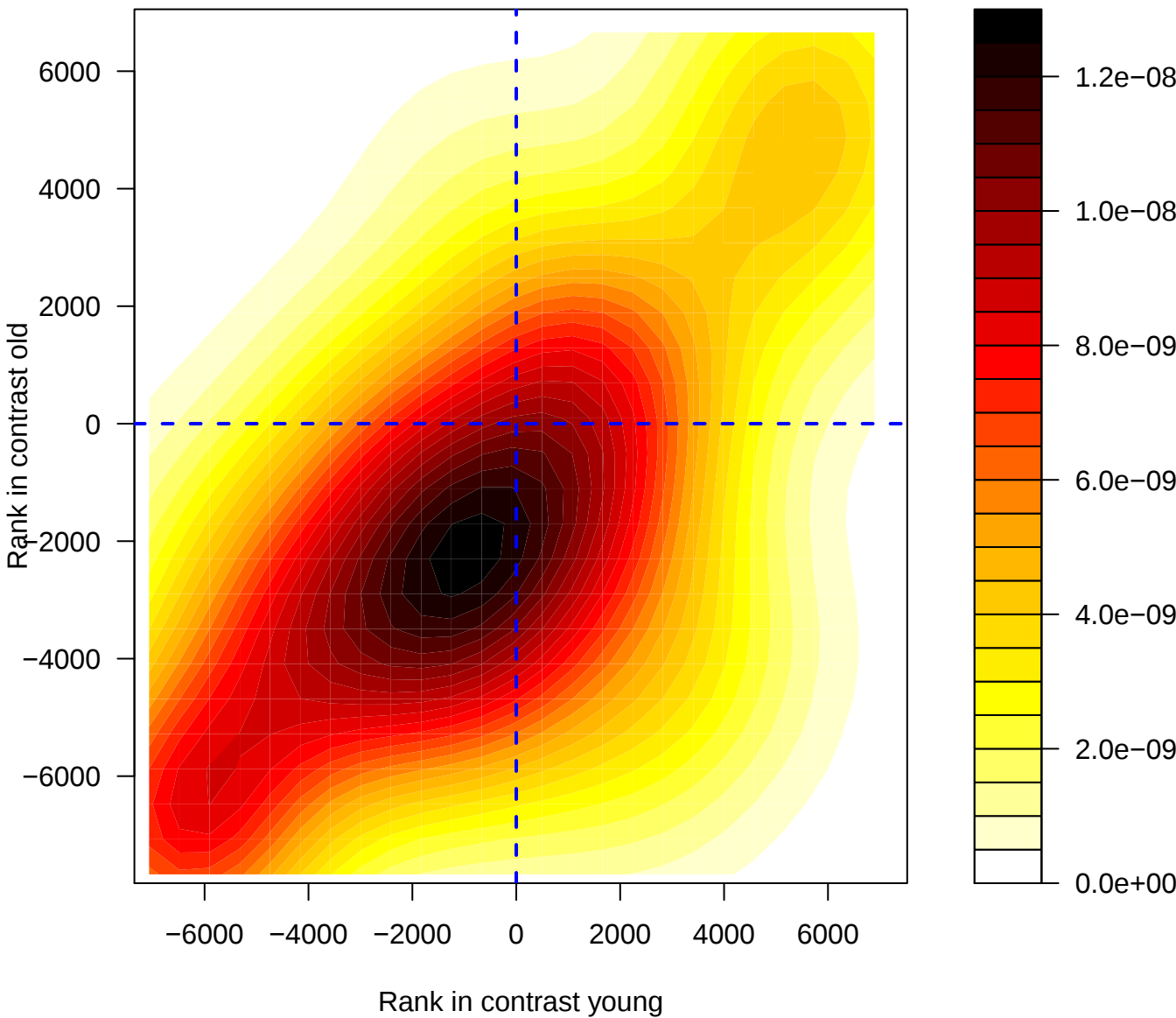
Smooth-Muscle-Contraction



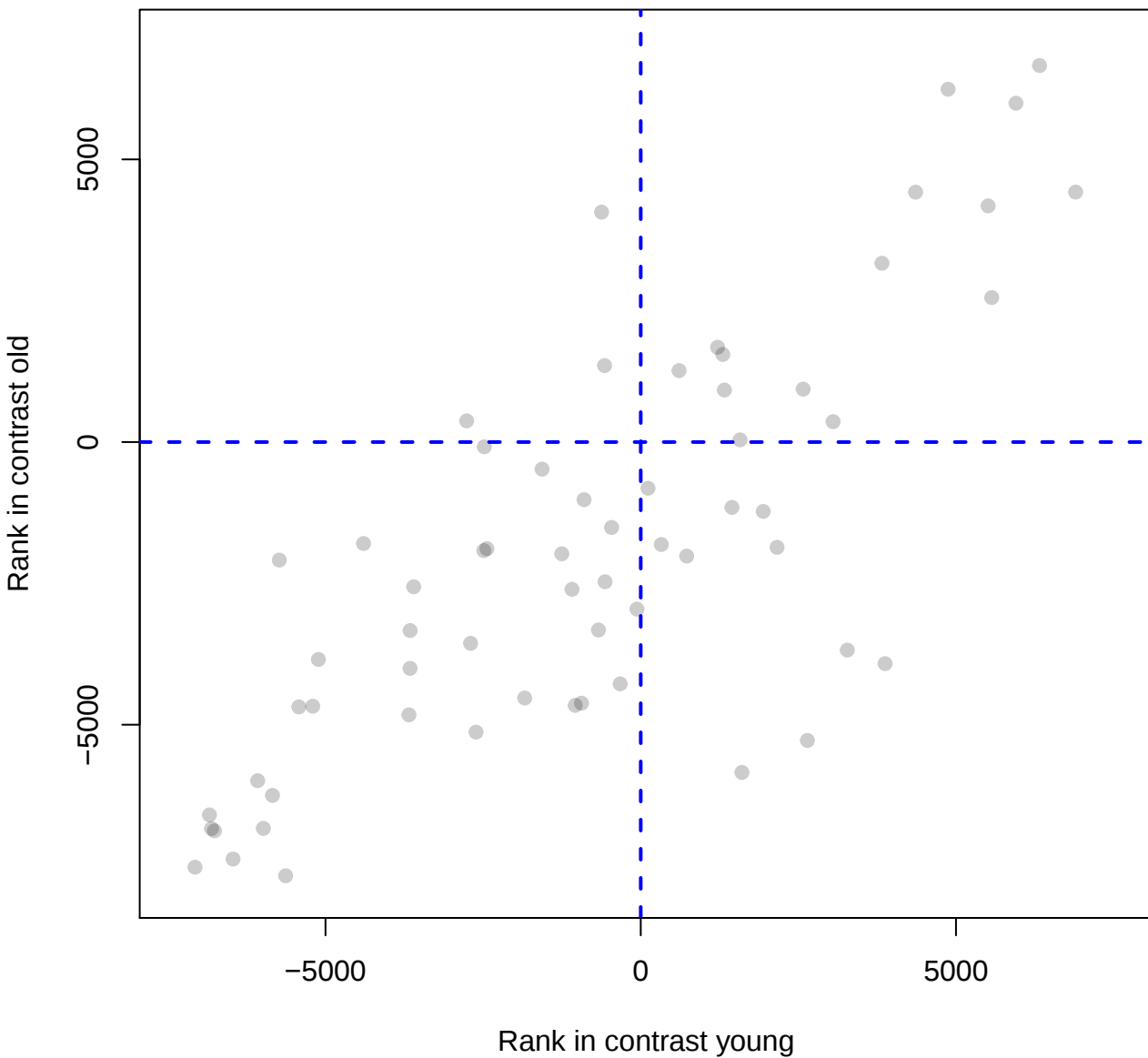
Smooth-Muscle-Contraction



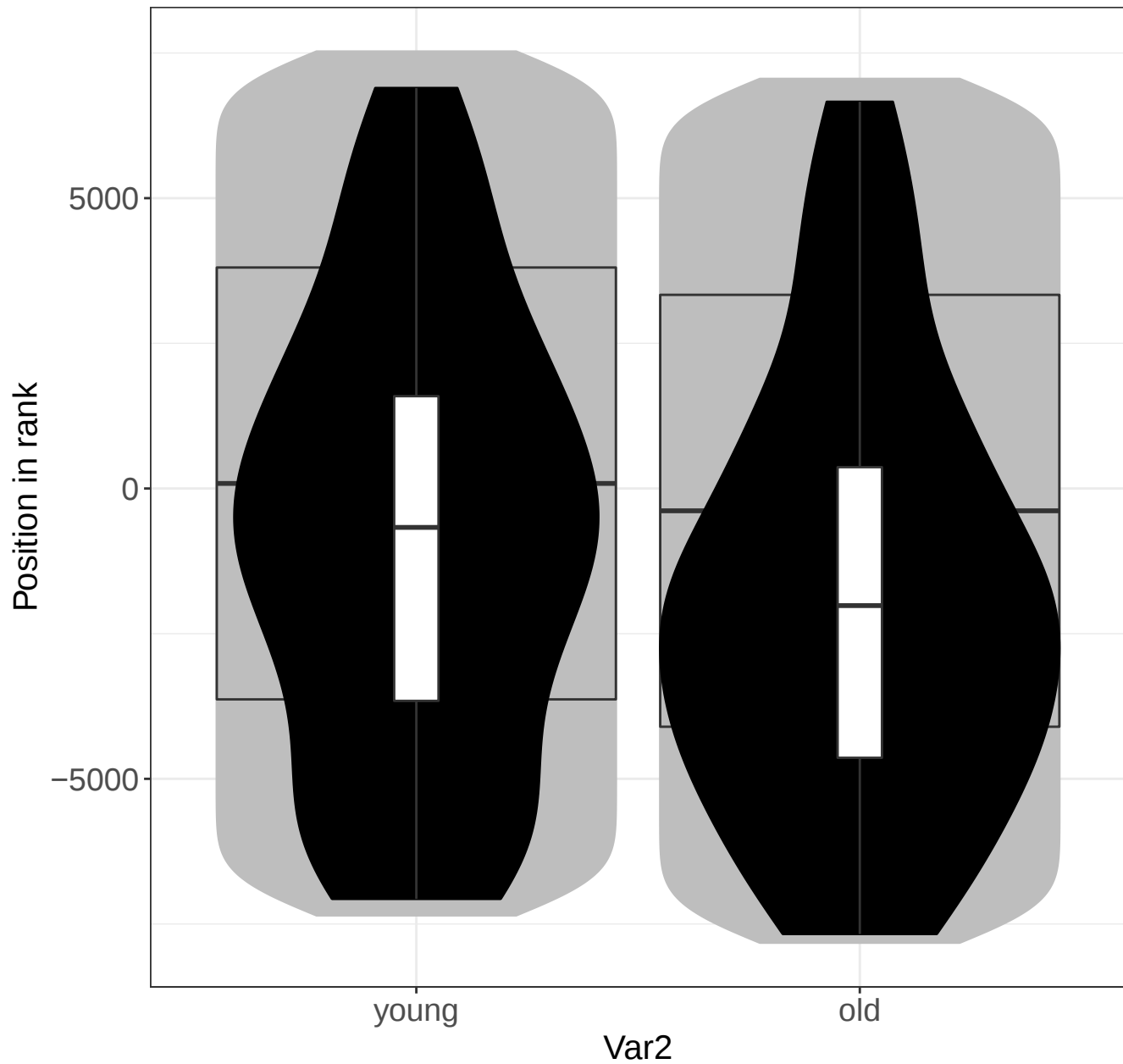
Glycolysis



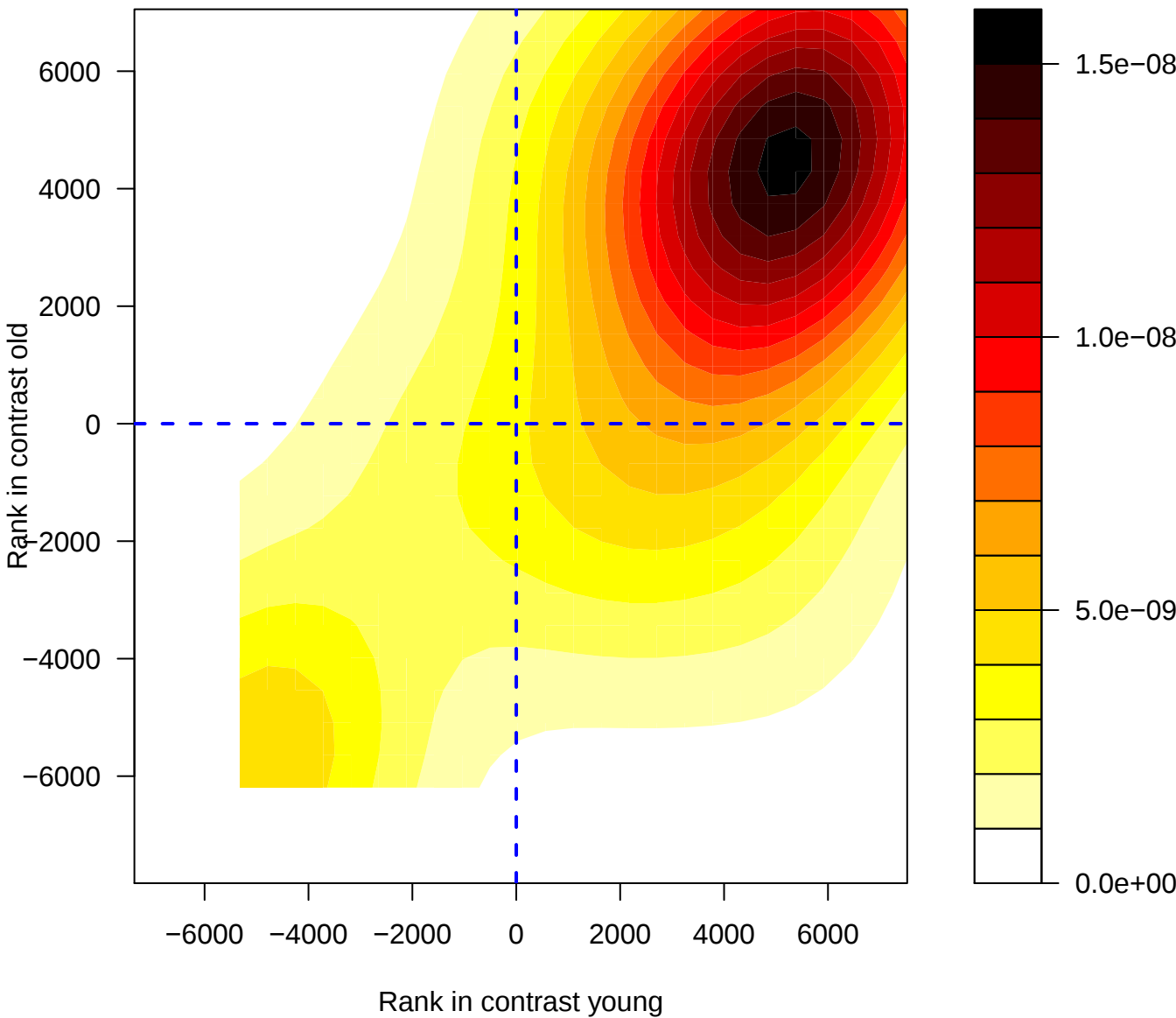
Glycolysis



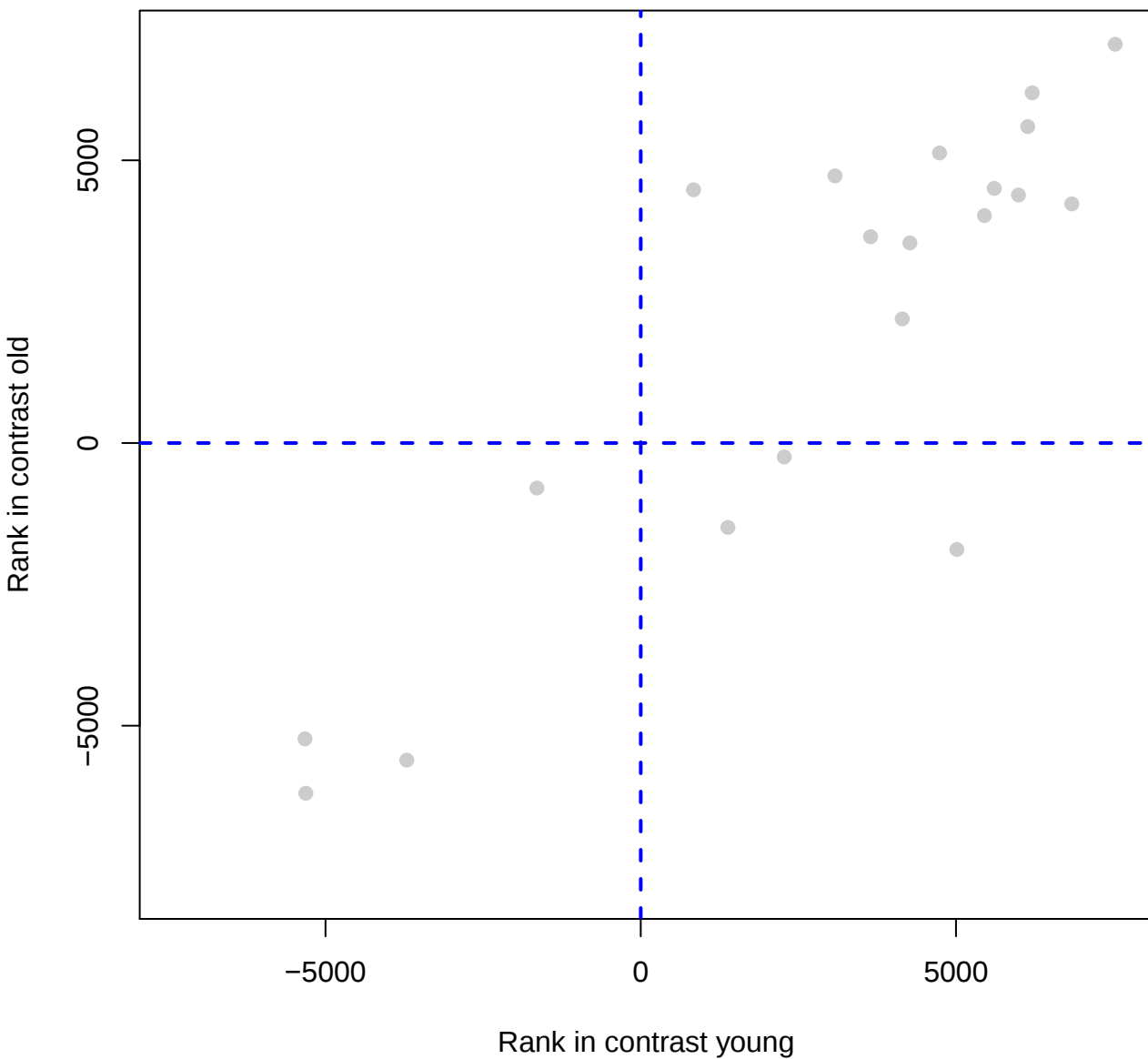
Glycolysis



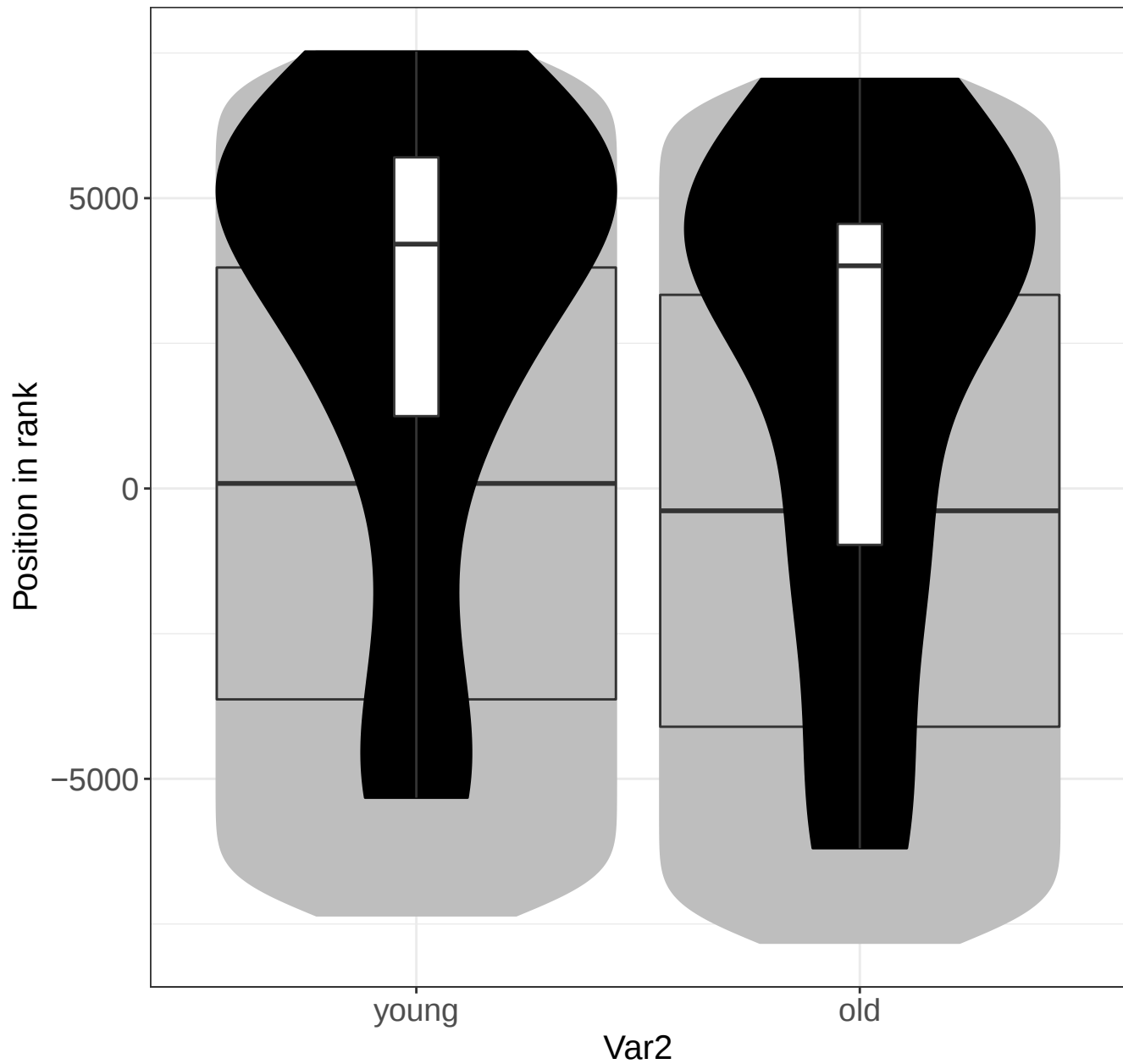
Signaling-by-Hippo



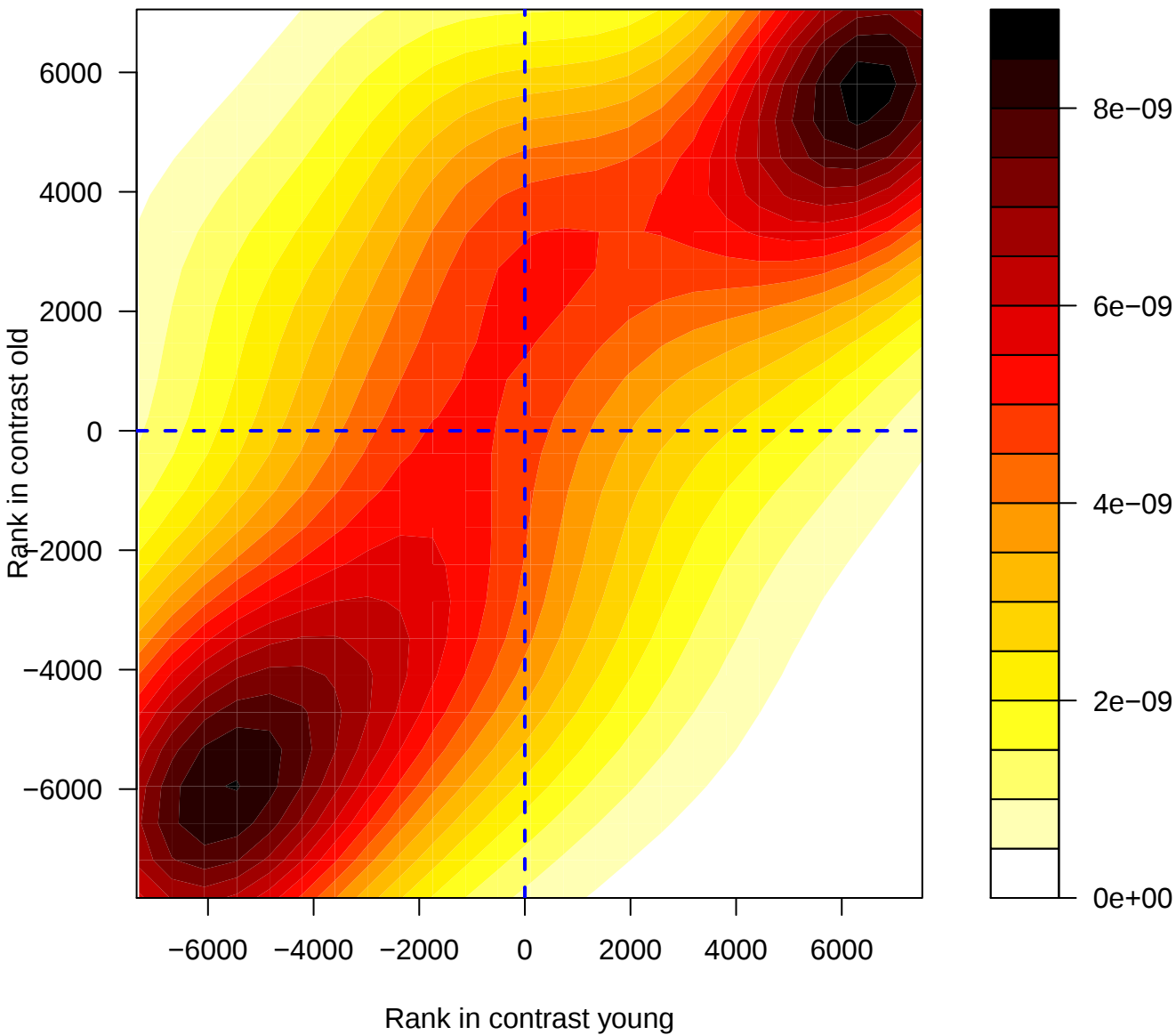
Signaling-by-Hippo



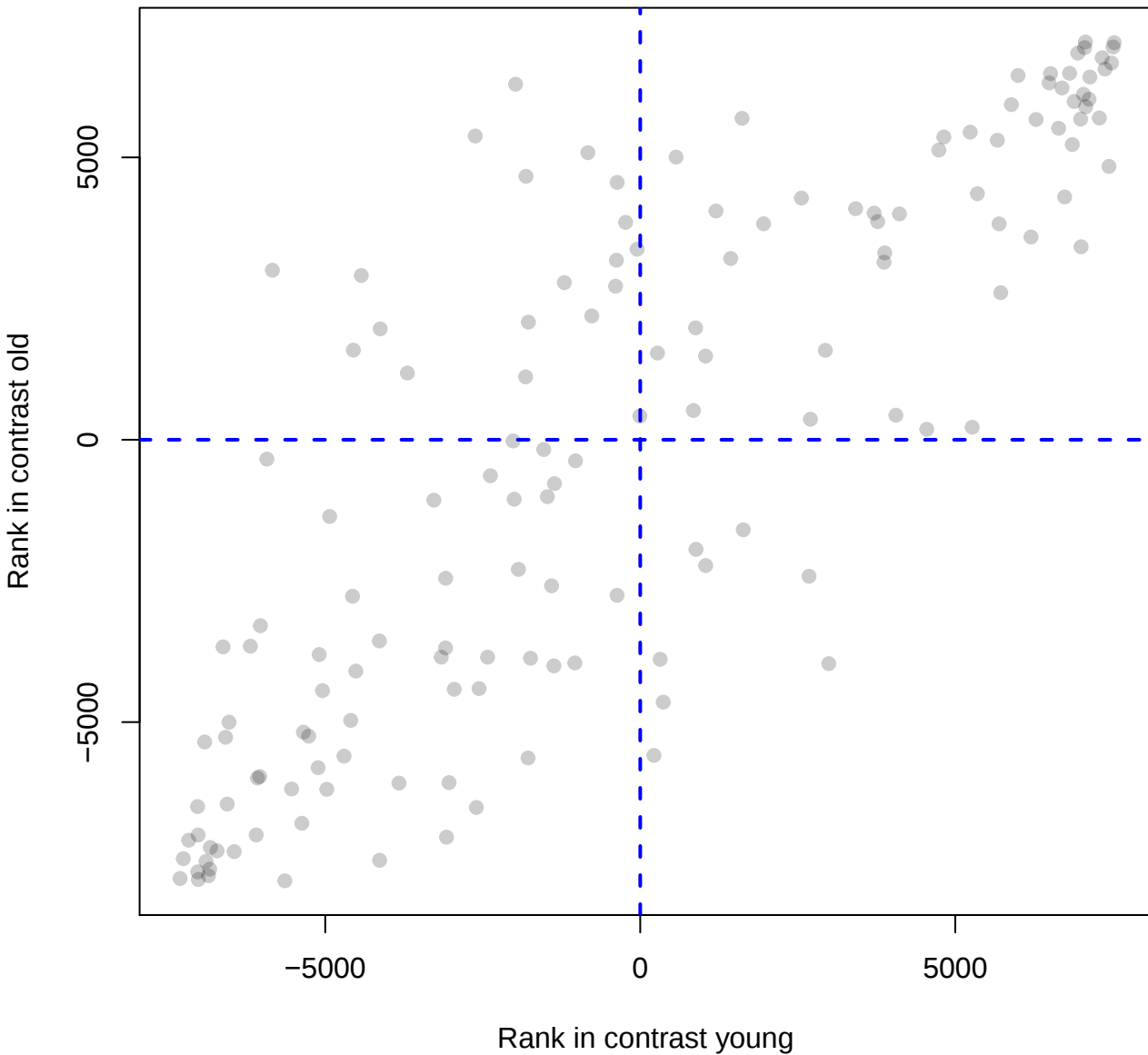
Signaling-by-Hippo



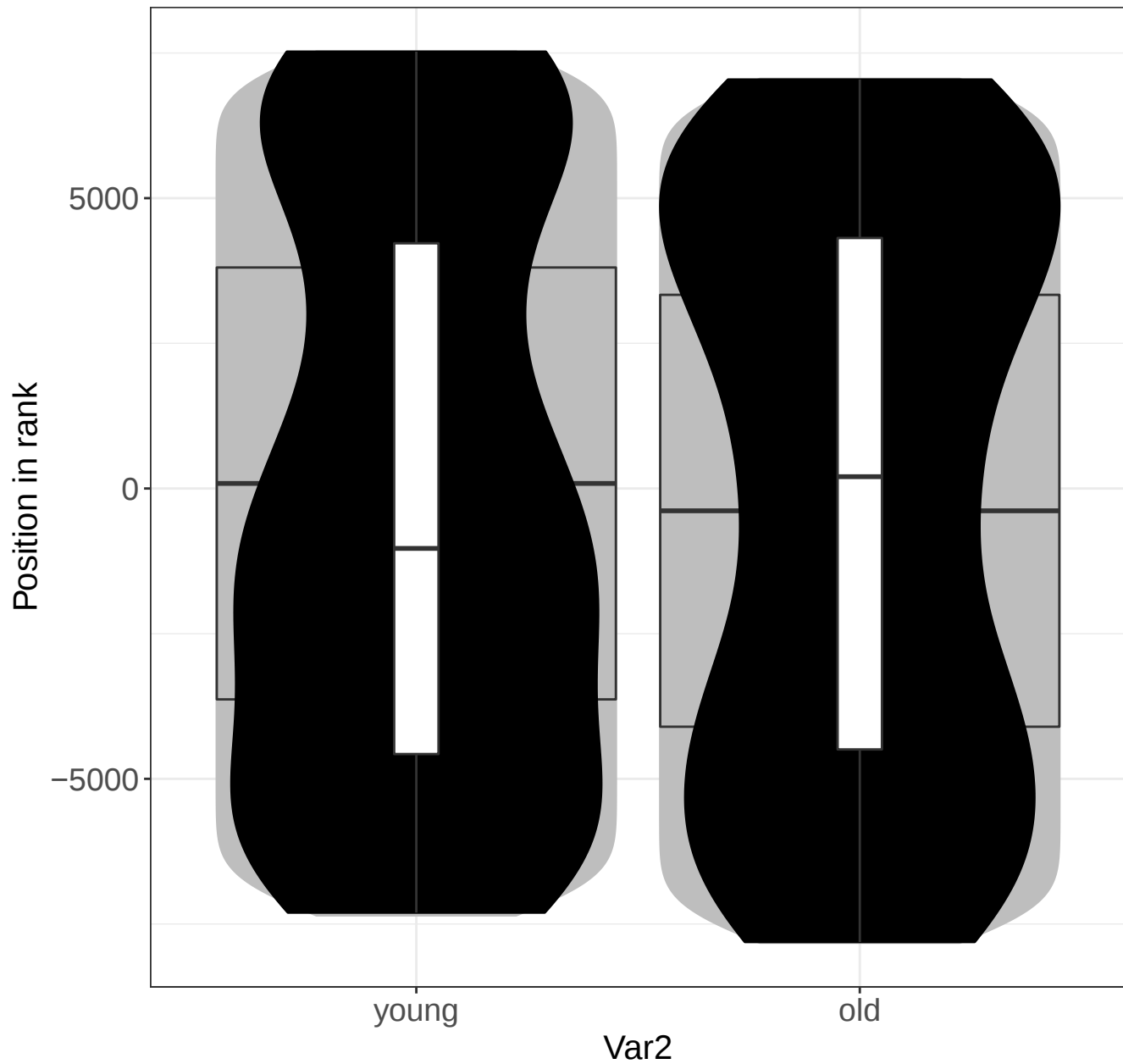
Muscle-contraction



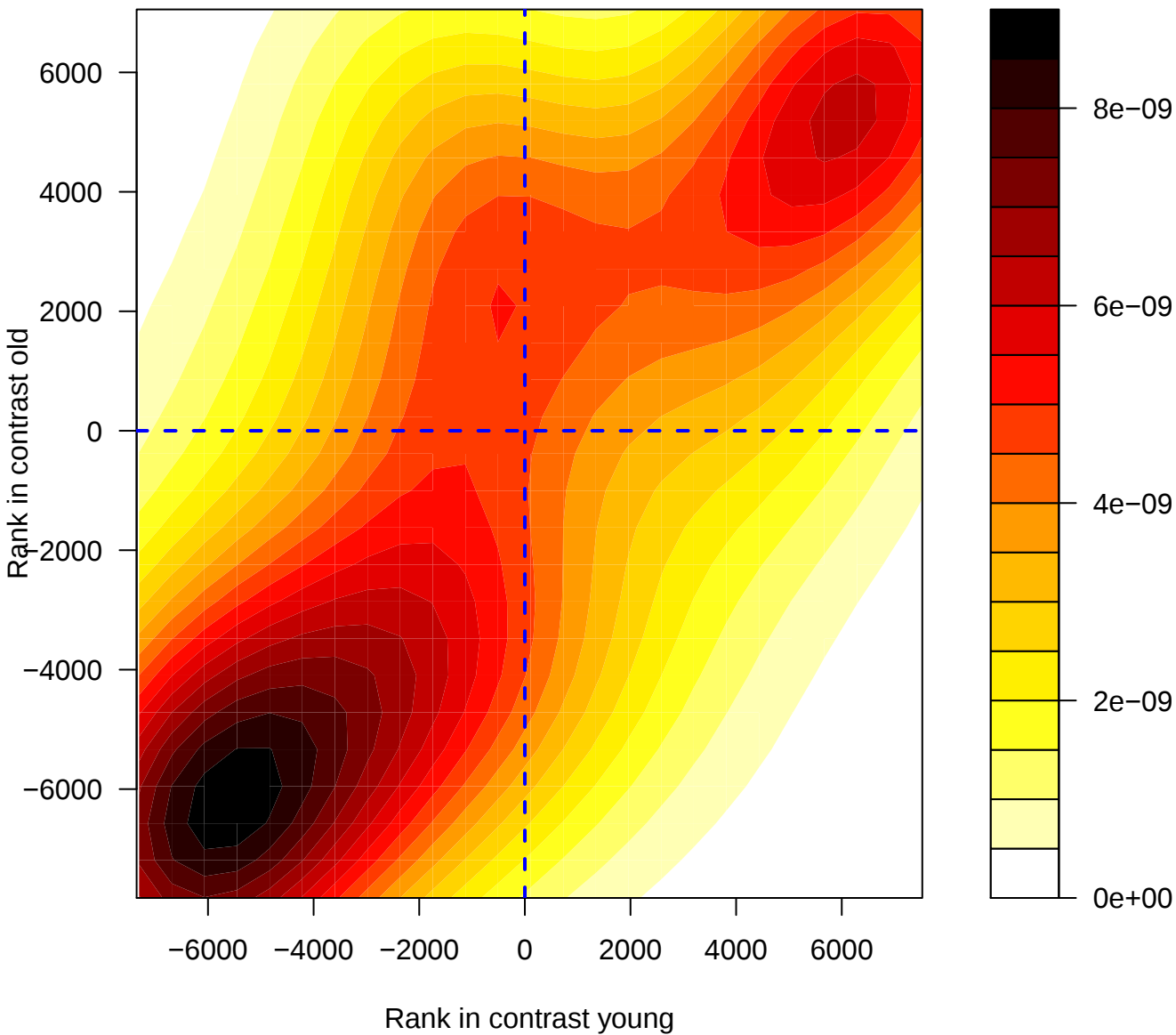
Muscle-contraction



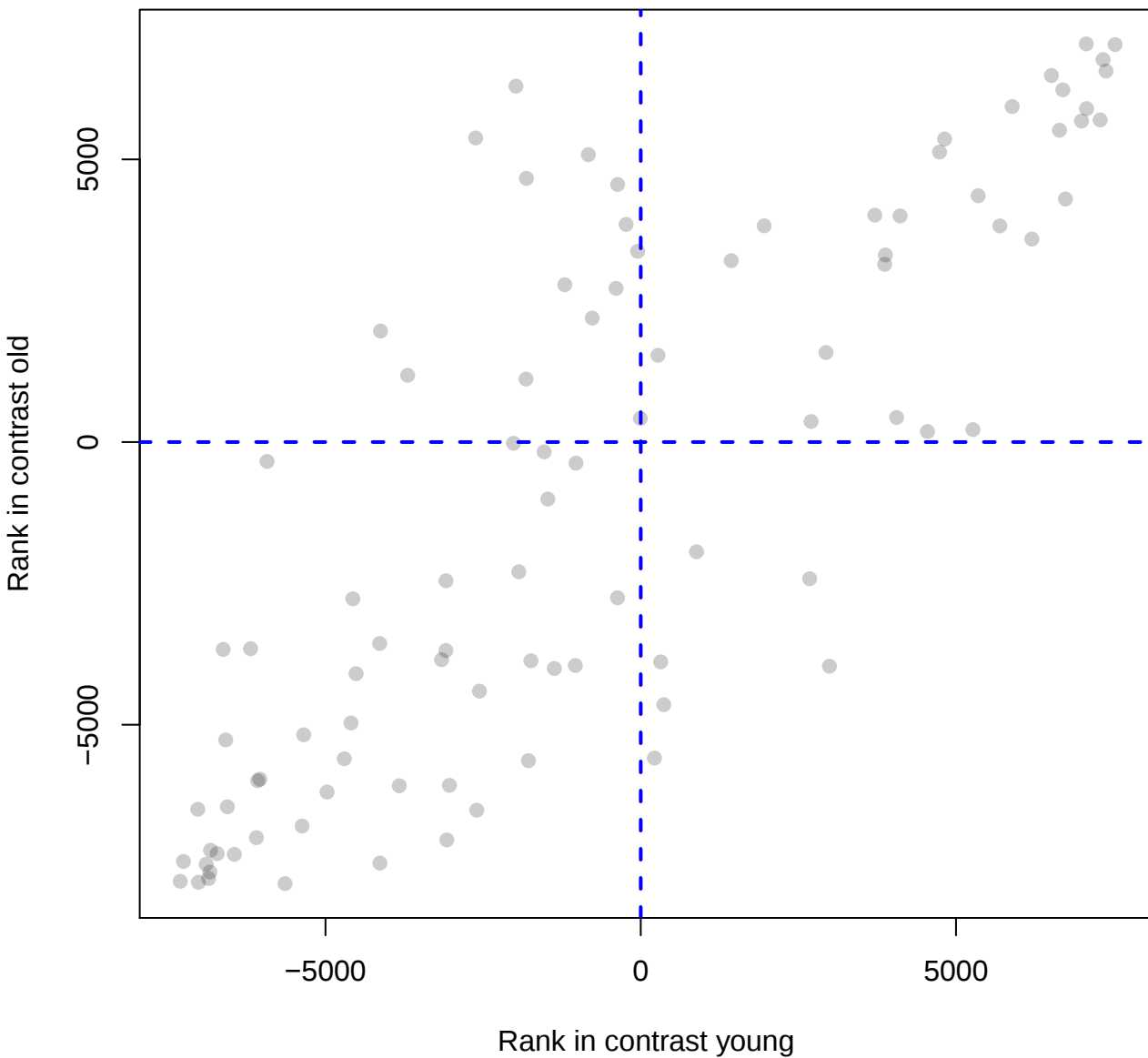
Muscle-contraction



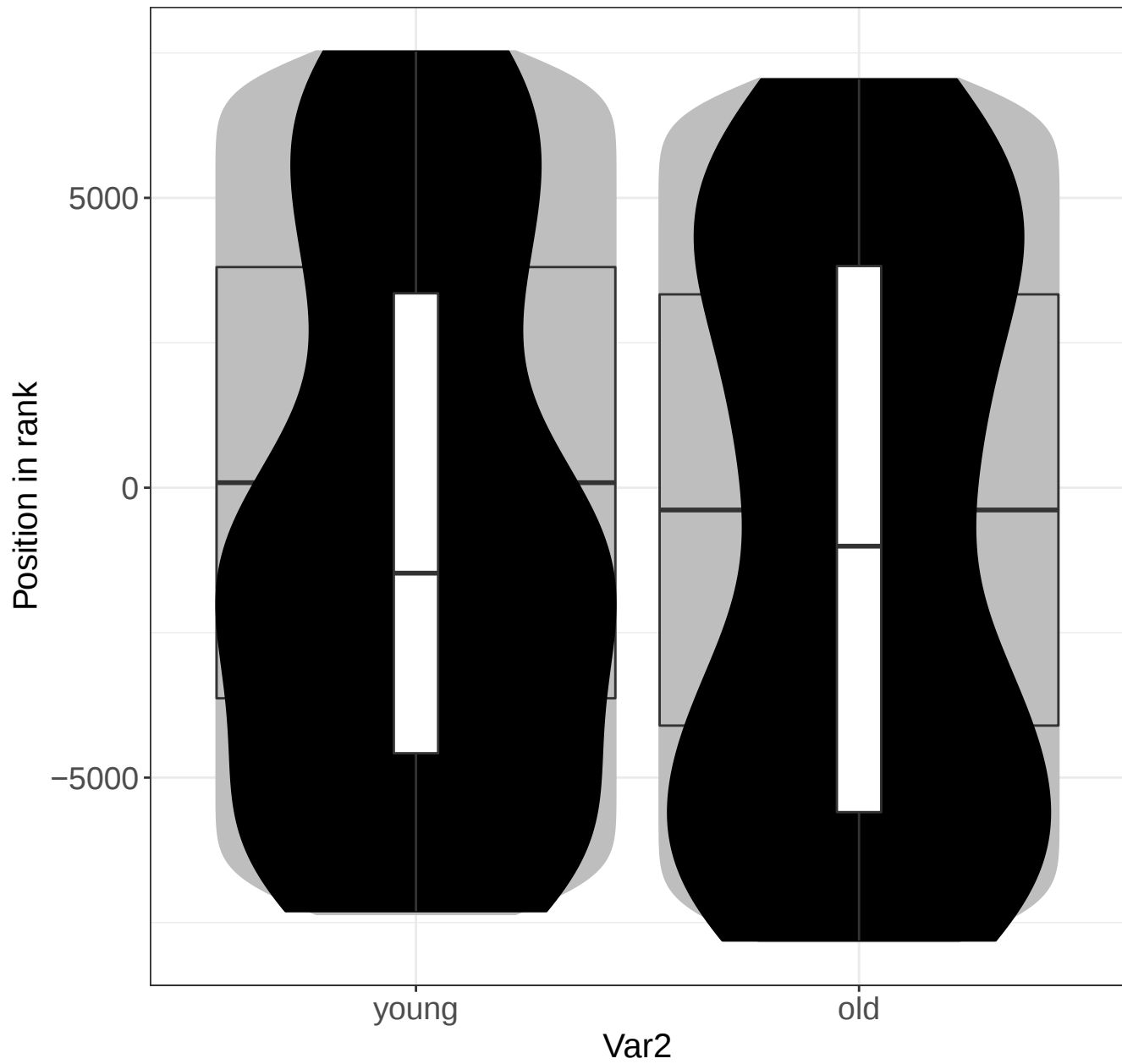
Cardiac-conduction



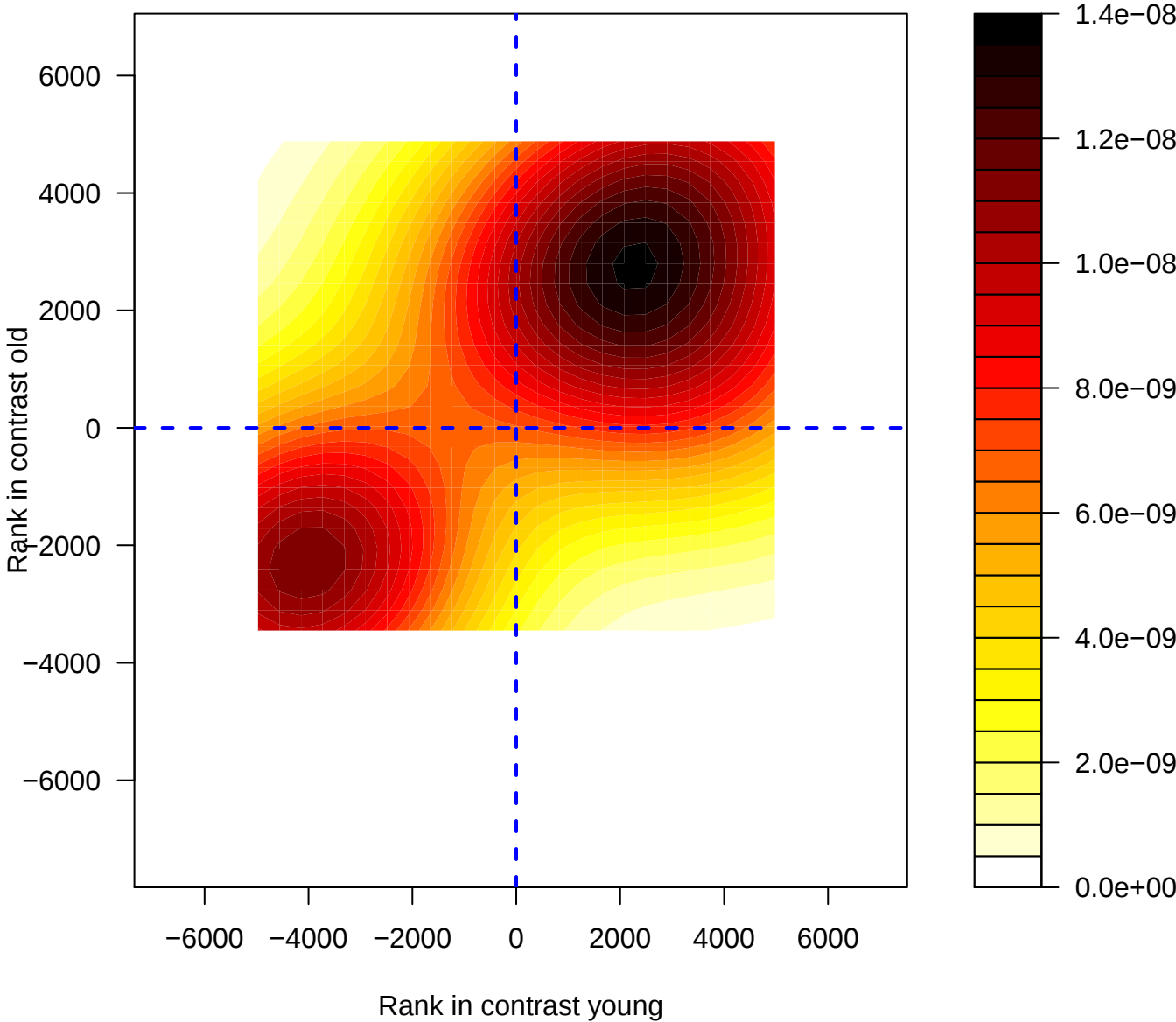
Cardiac-conduction



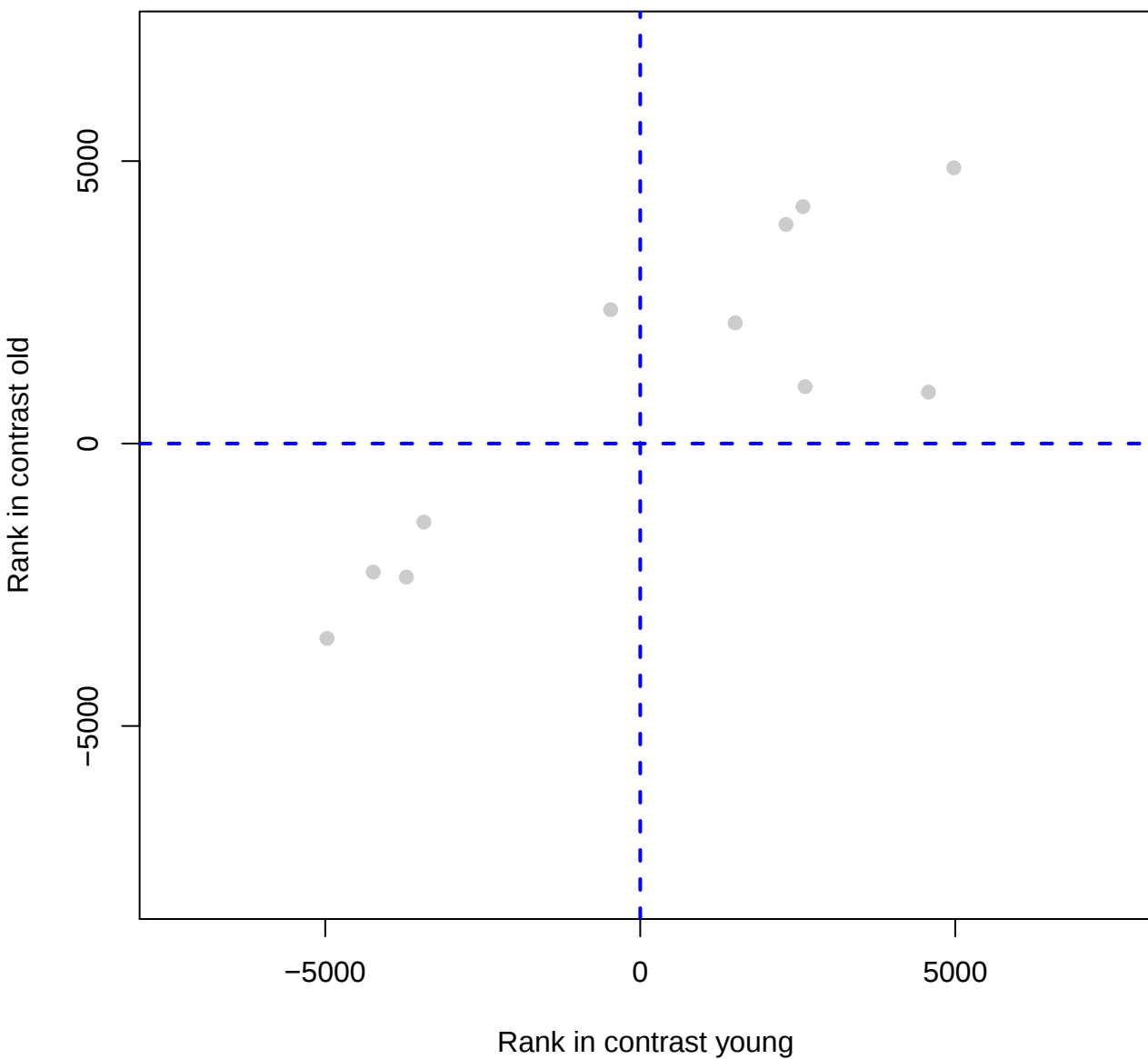
Cardiac-conduction



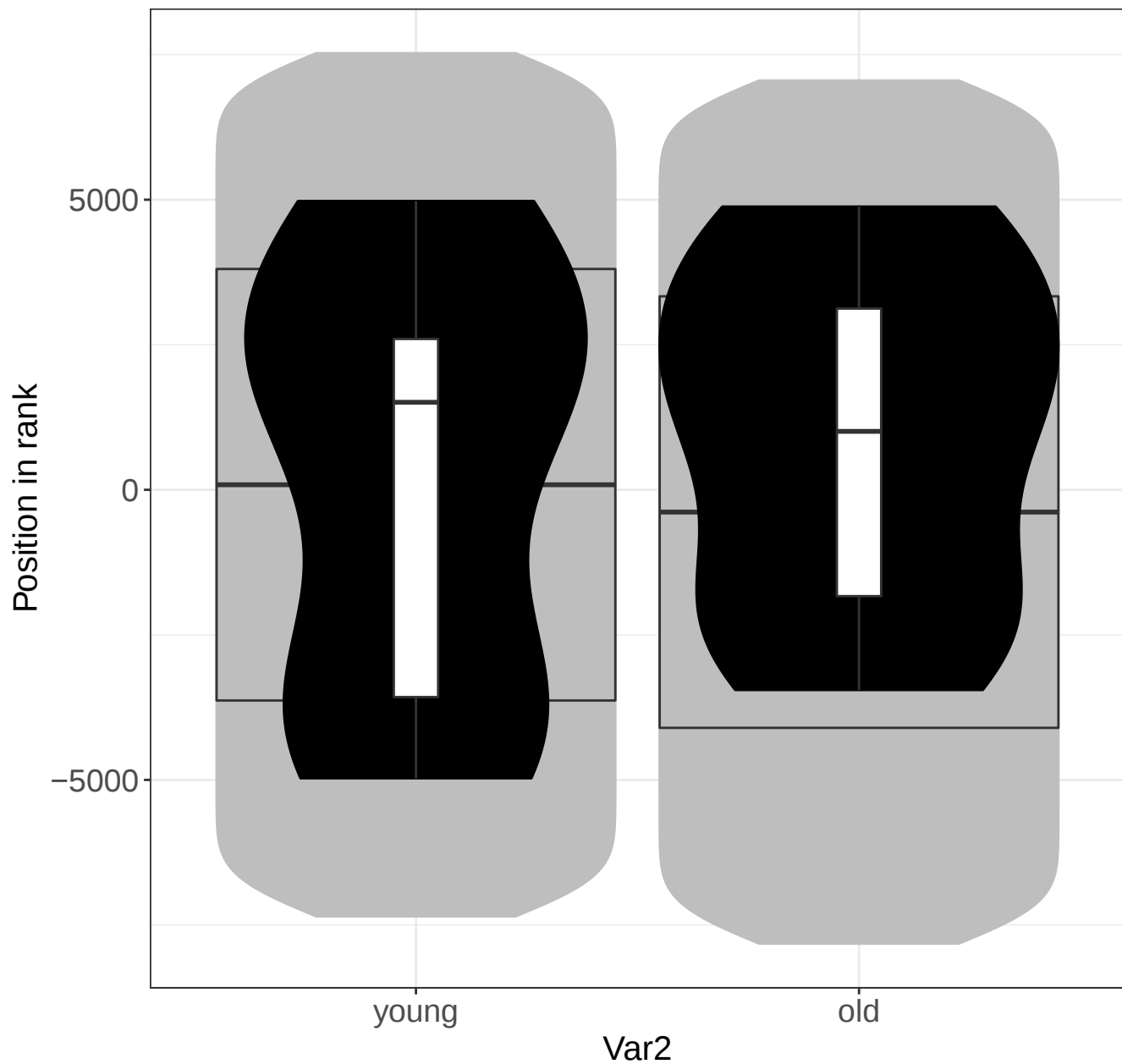
ral-additional-cell-death-genes-whose-specific-roles-in-l



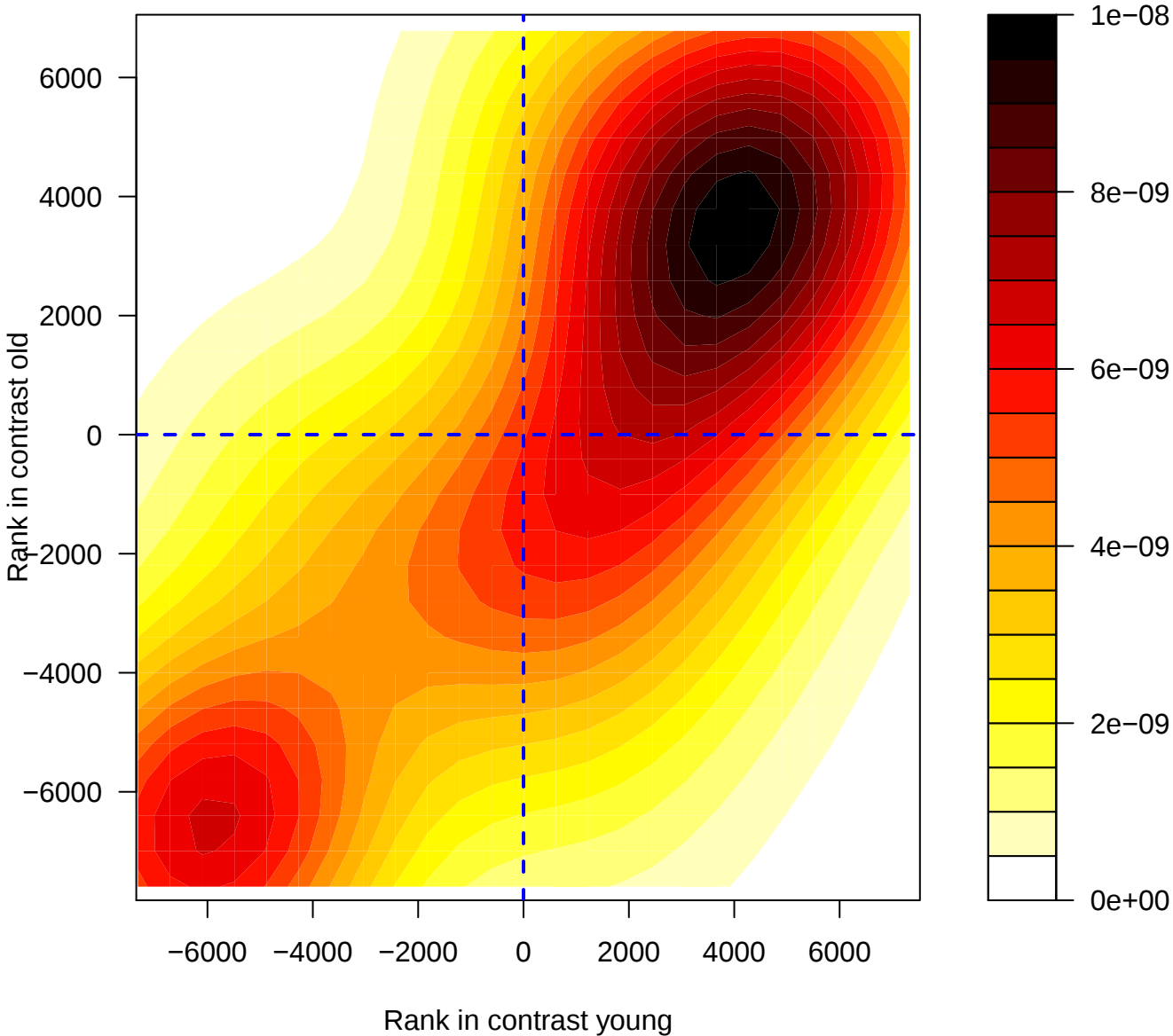
f-several-additional-cell-death-genes-whose-specific-roles-in-p53-depe



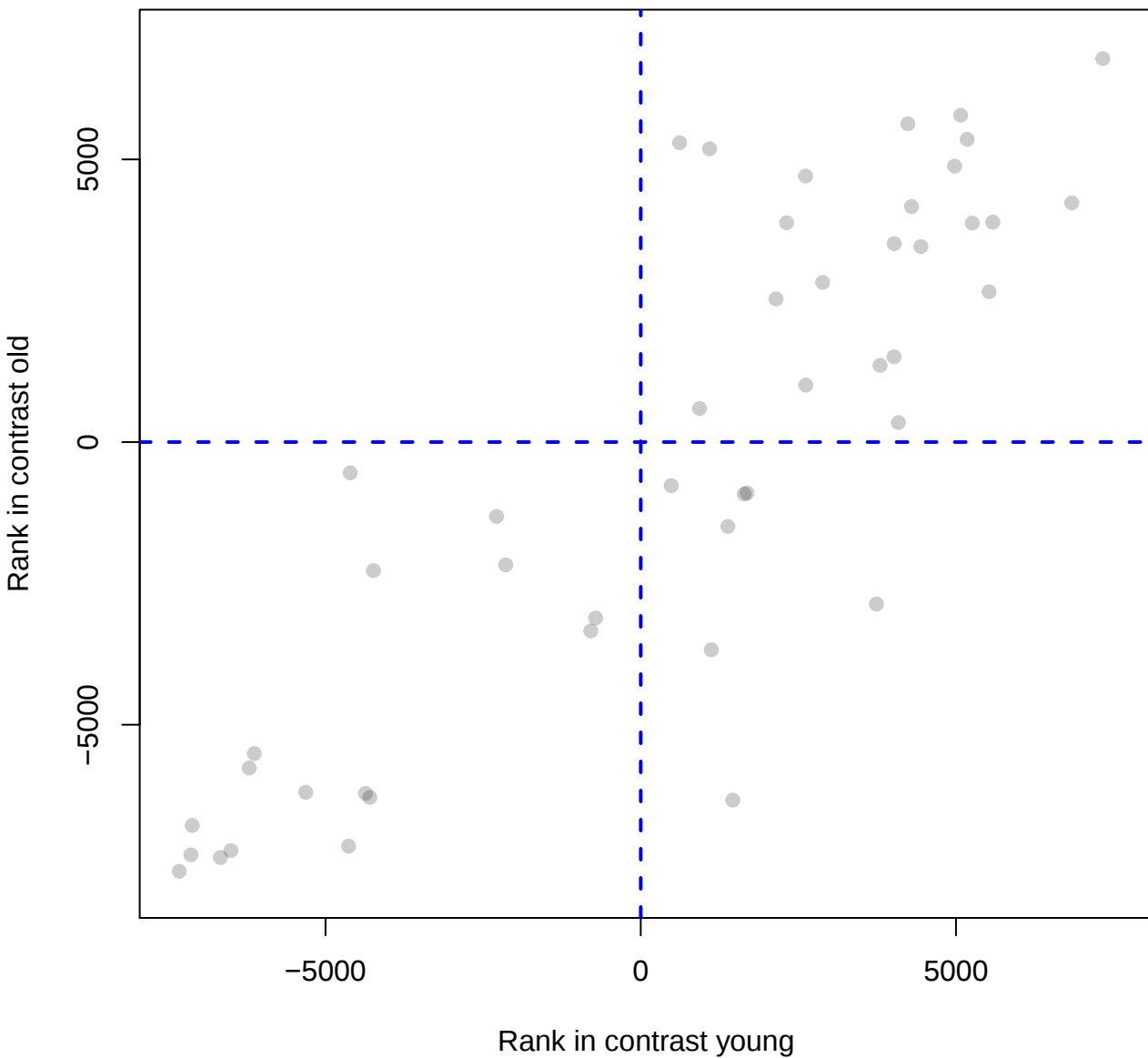
TP53-regulates-transcription-of-several-additio



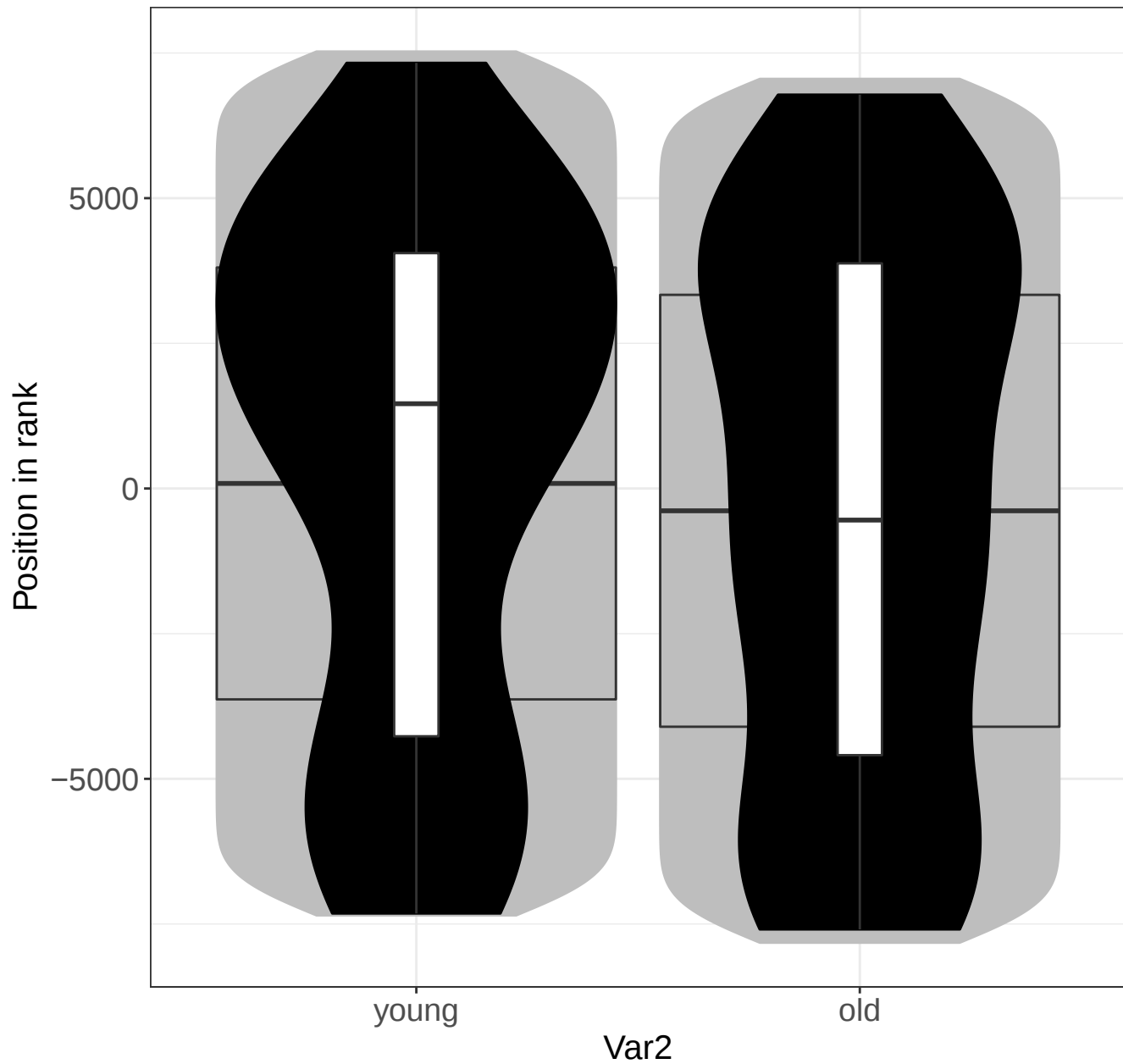
Intrinsic-Pathway-for-Apoptosis



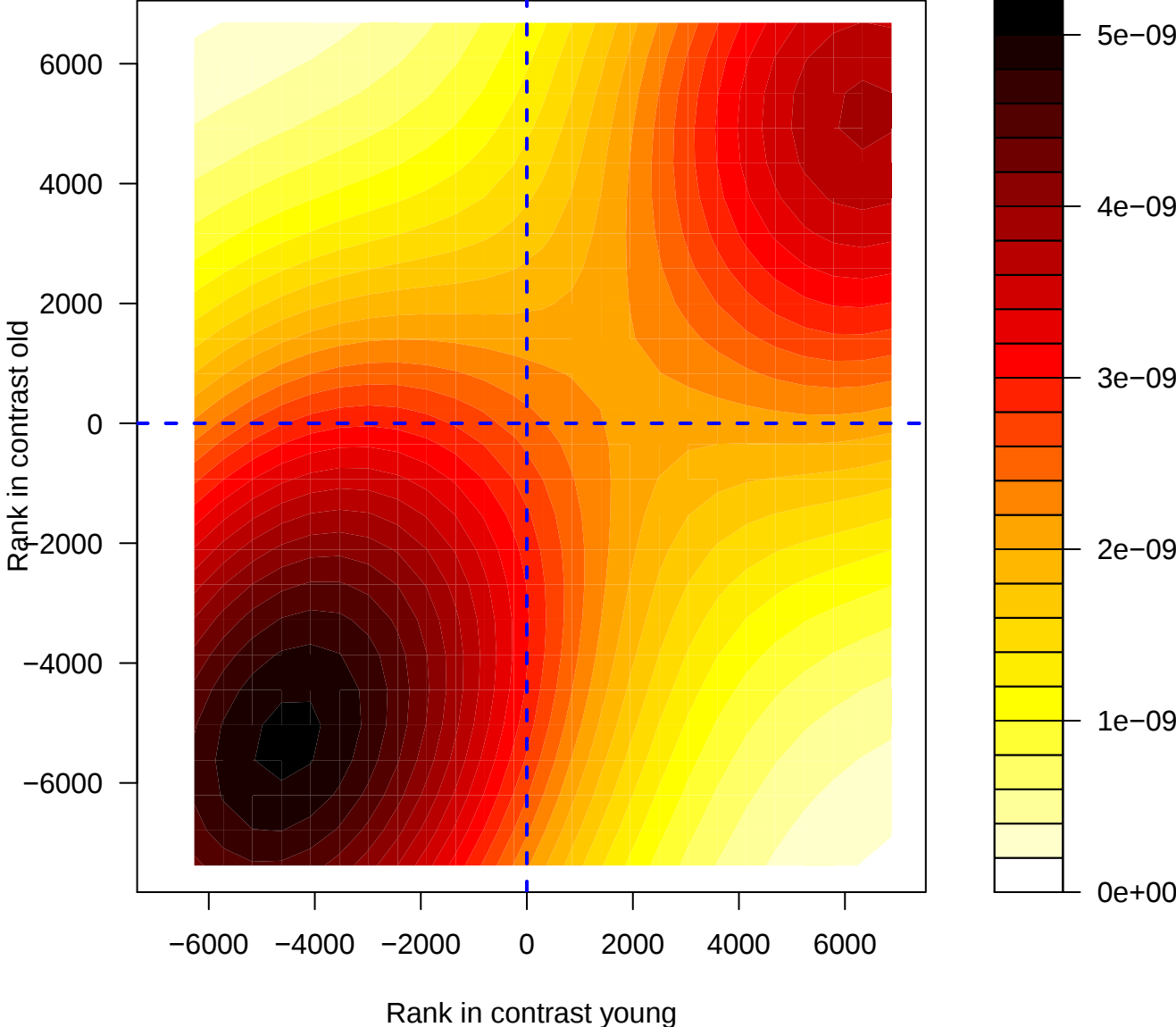
Intrinsic-Pathway-for-Apoptosis



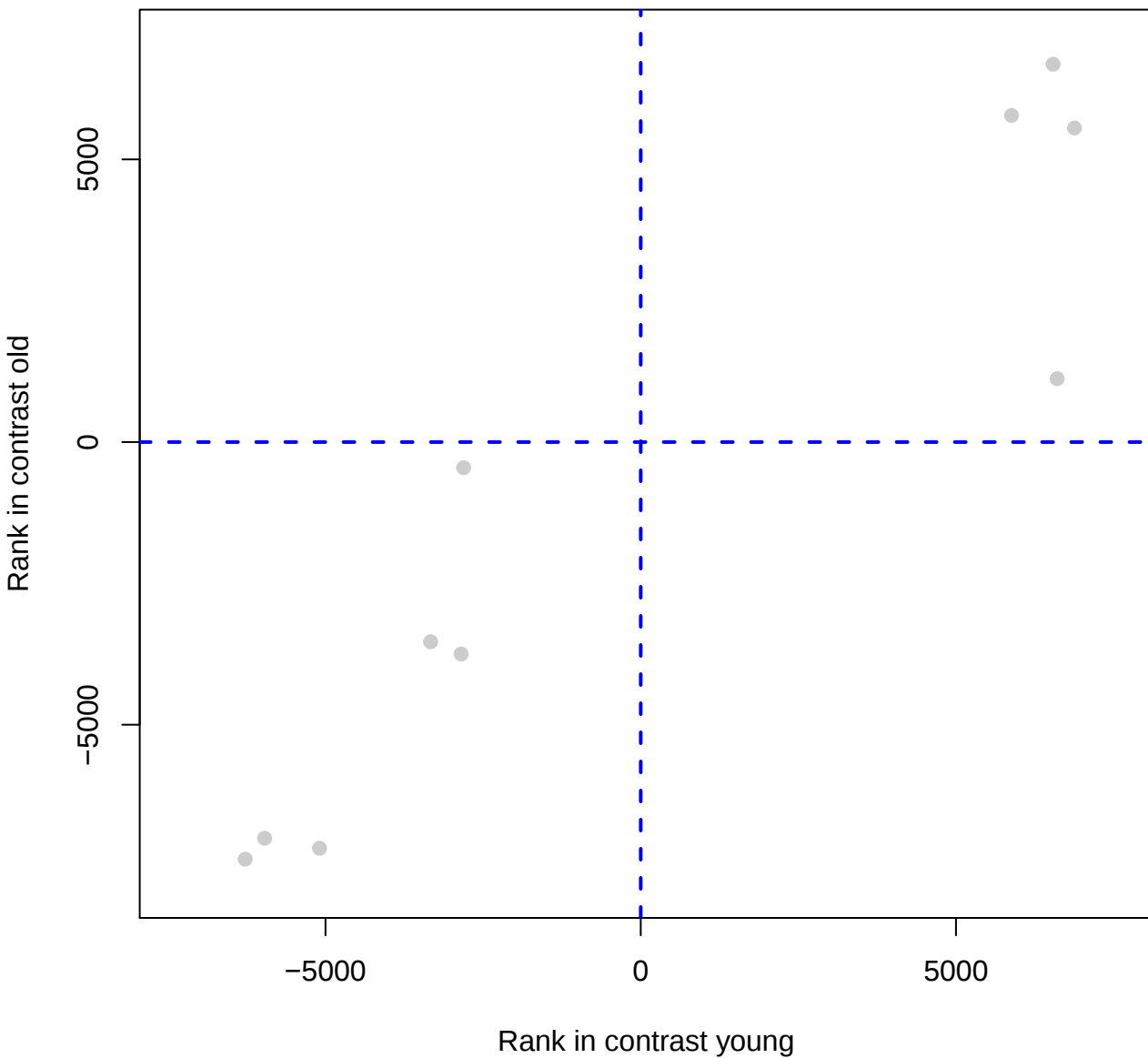
Intrinsic-Pathway-for-Apoptosis



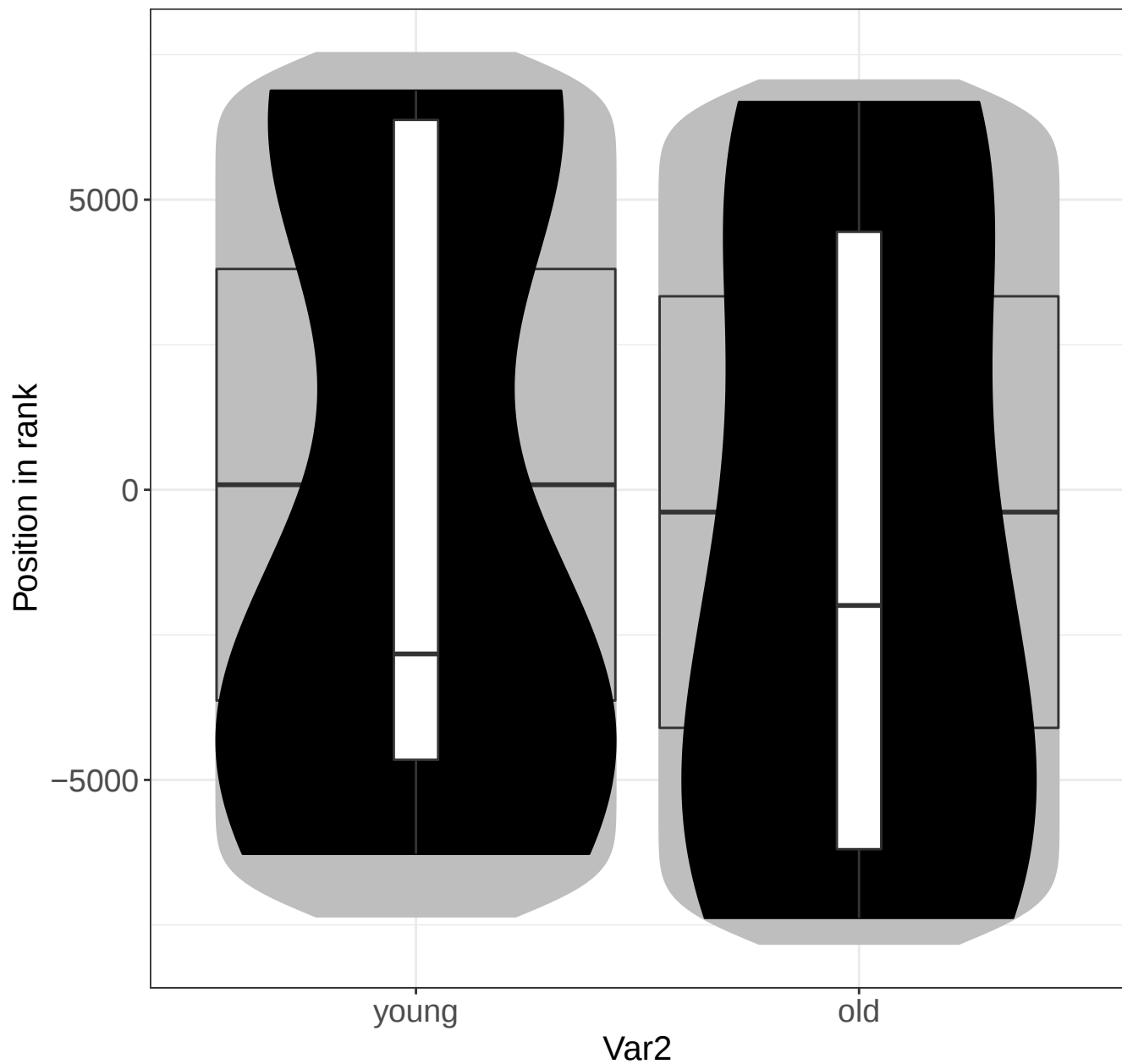
Regulation-of-gene-expression-by-Hypoxia-inducible-Fa



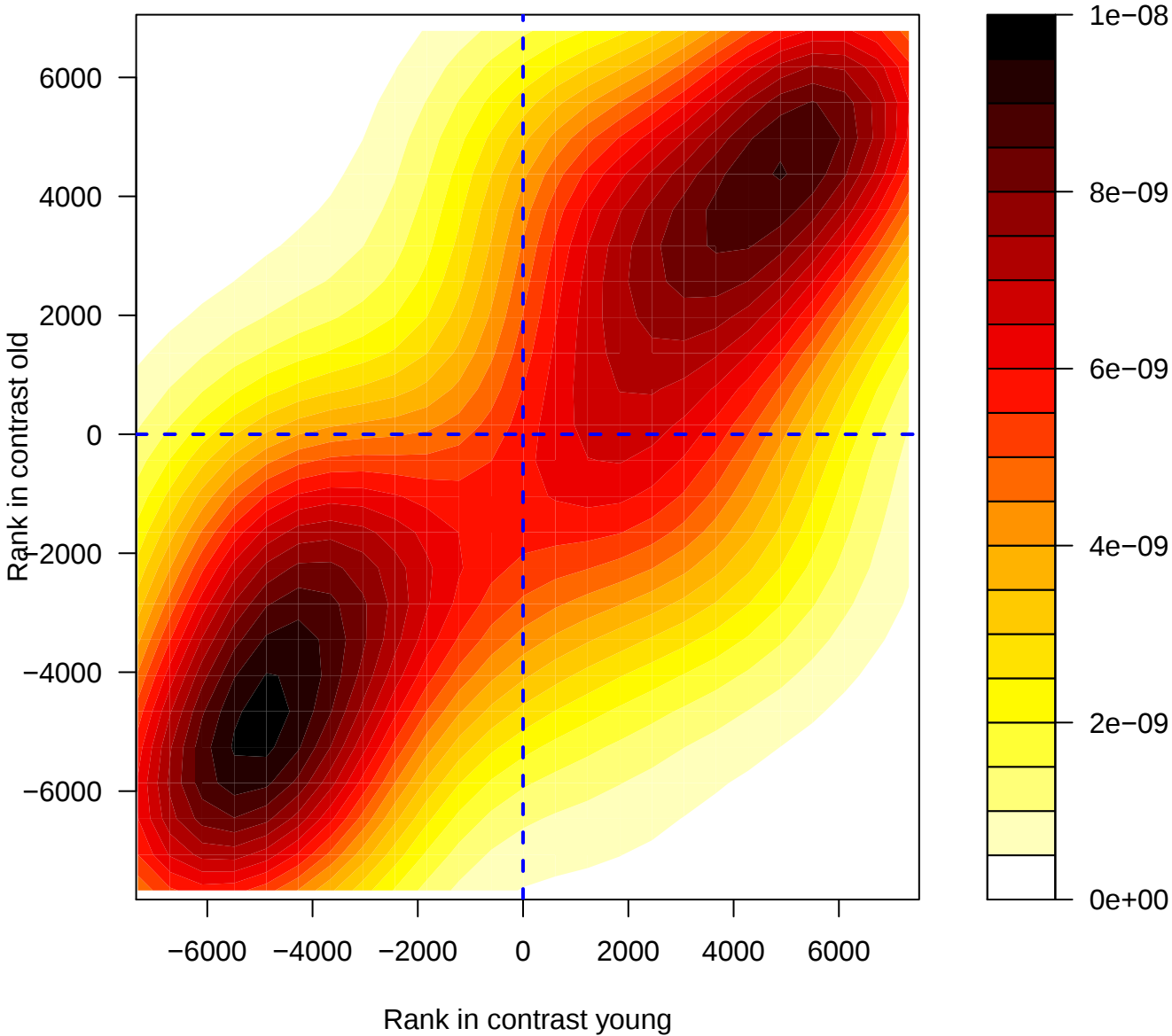
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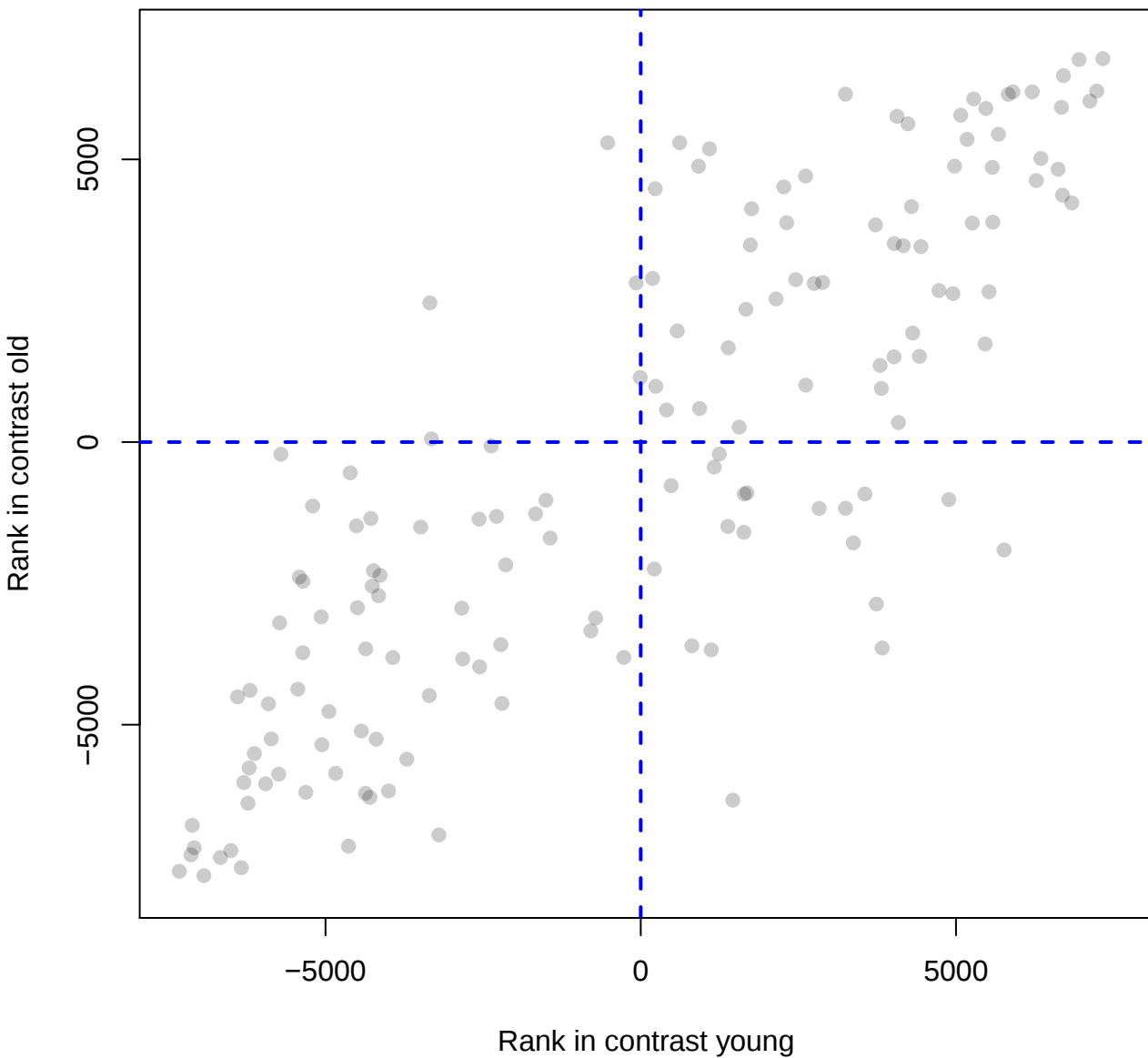
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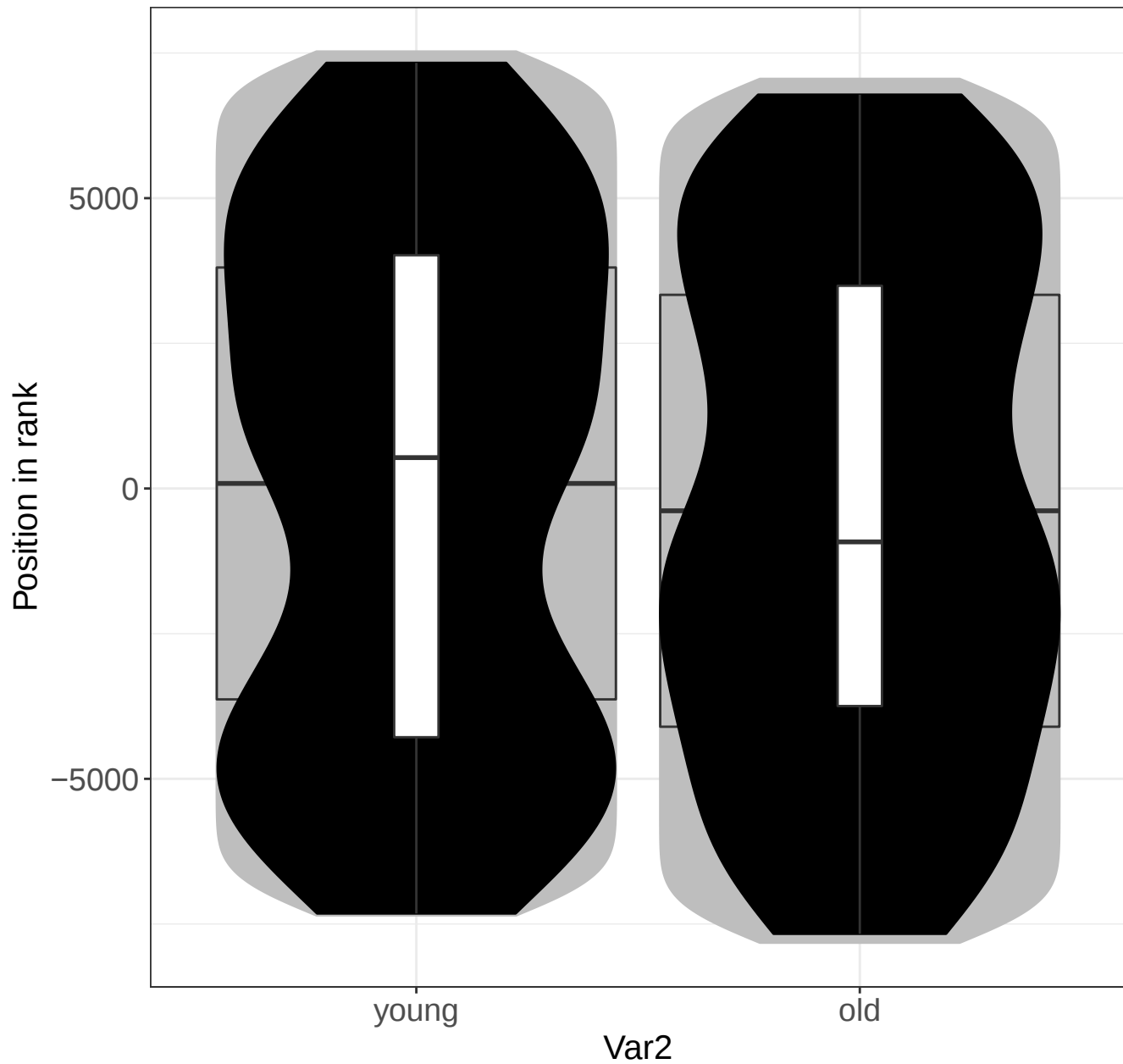
Apoptosis



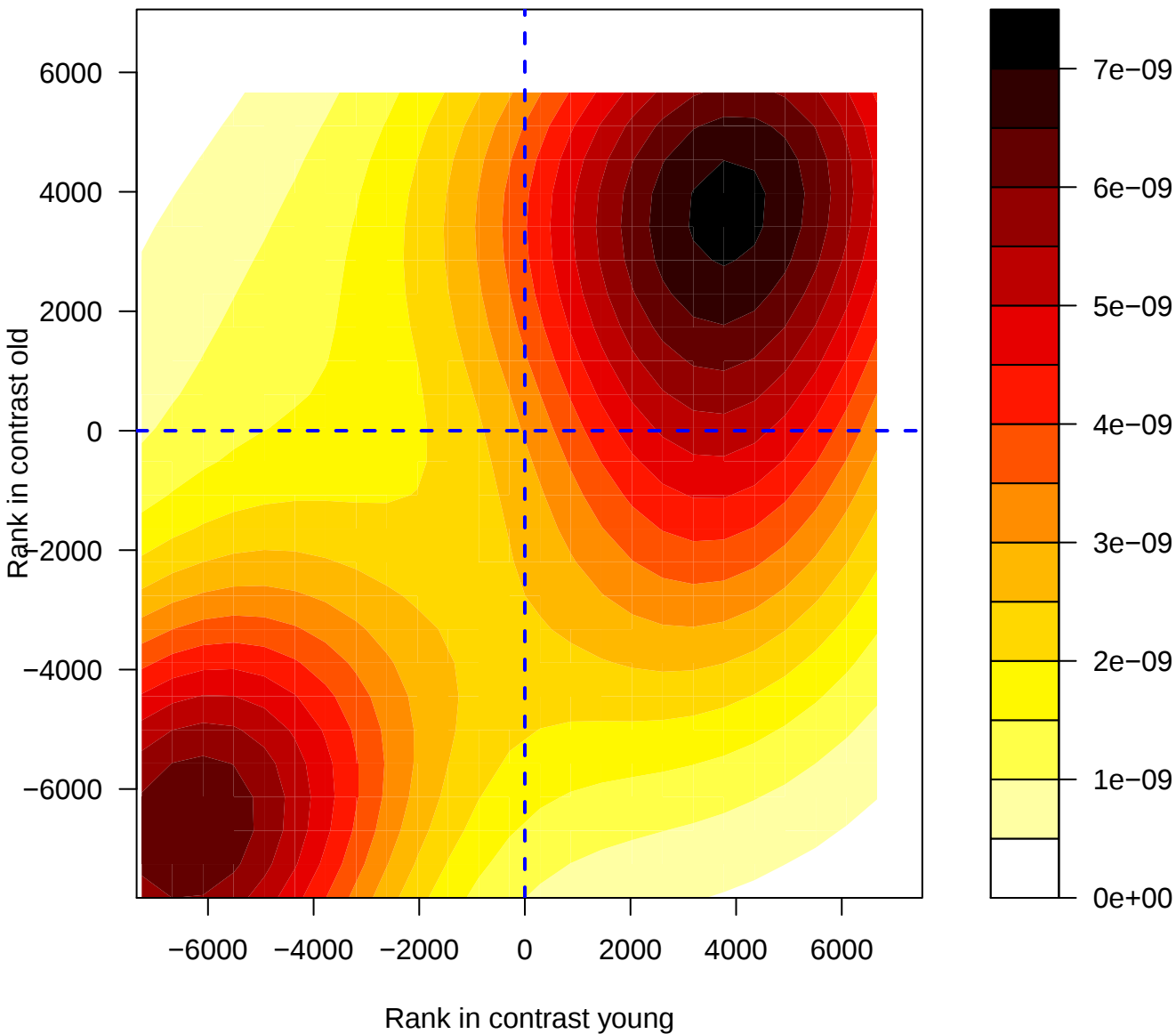
Apoptosis



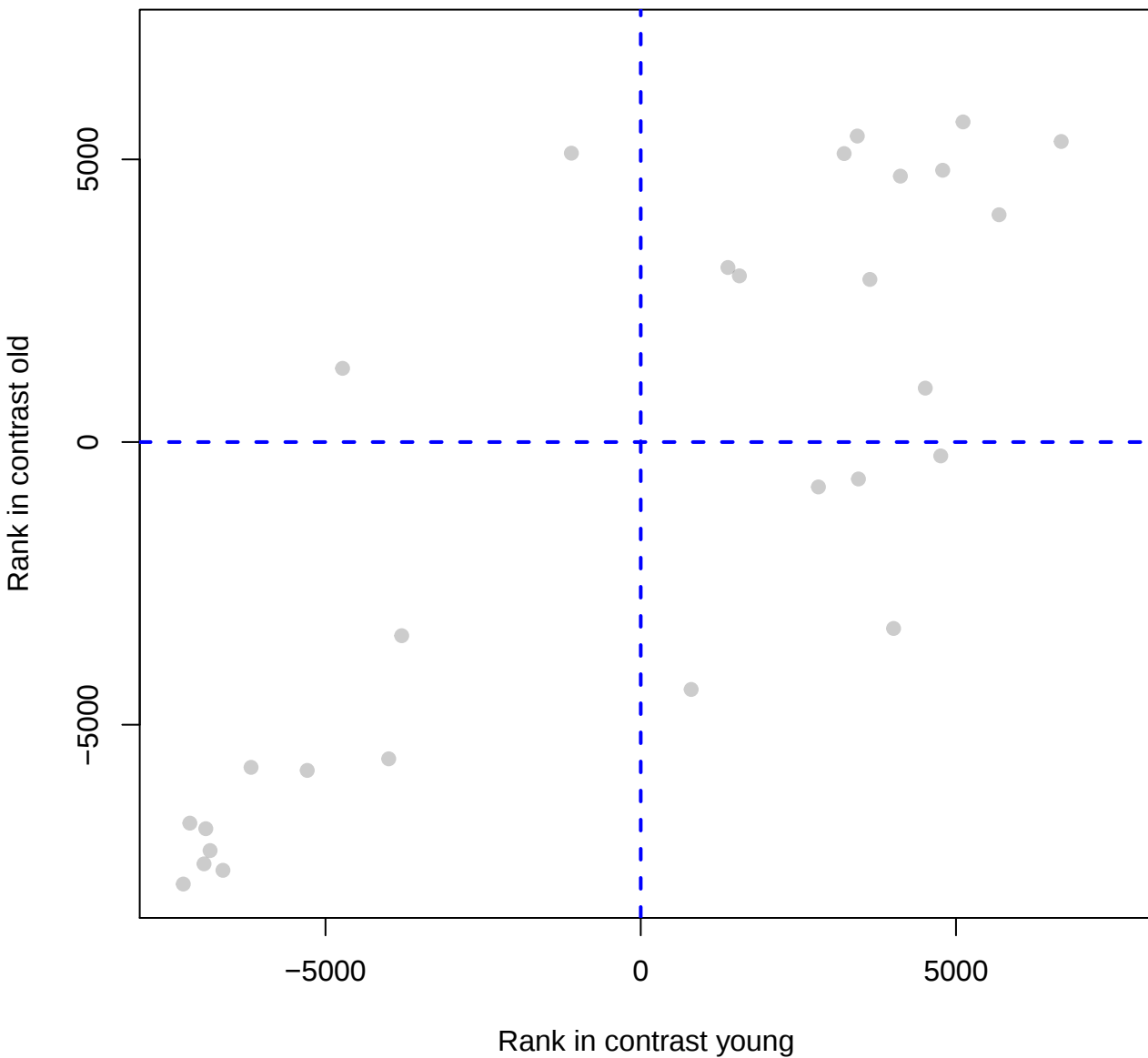
Apoptosis



Fatty-acyl-CoA-biosynthesis



Fatty-acyl-CoA-biosynthesis



Fatty-acyl-CoA-biosynthesis

