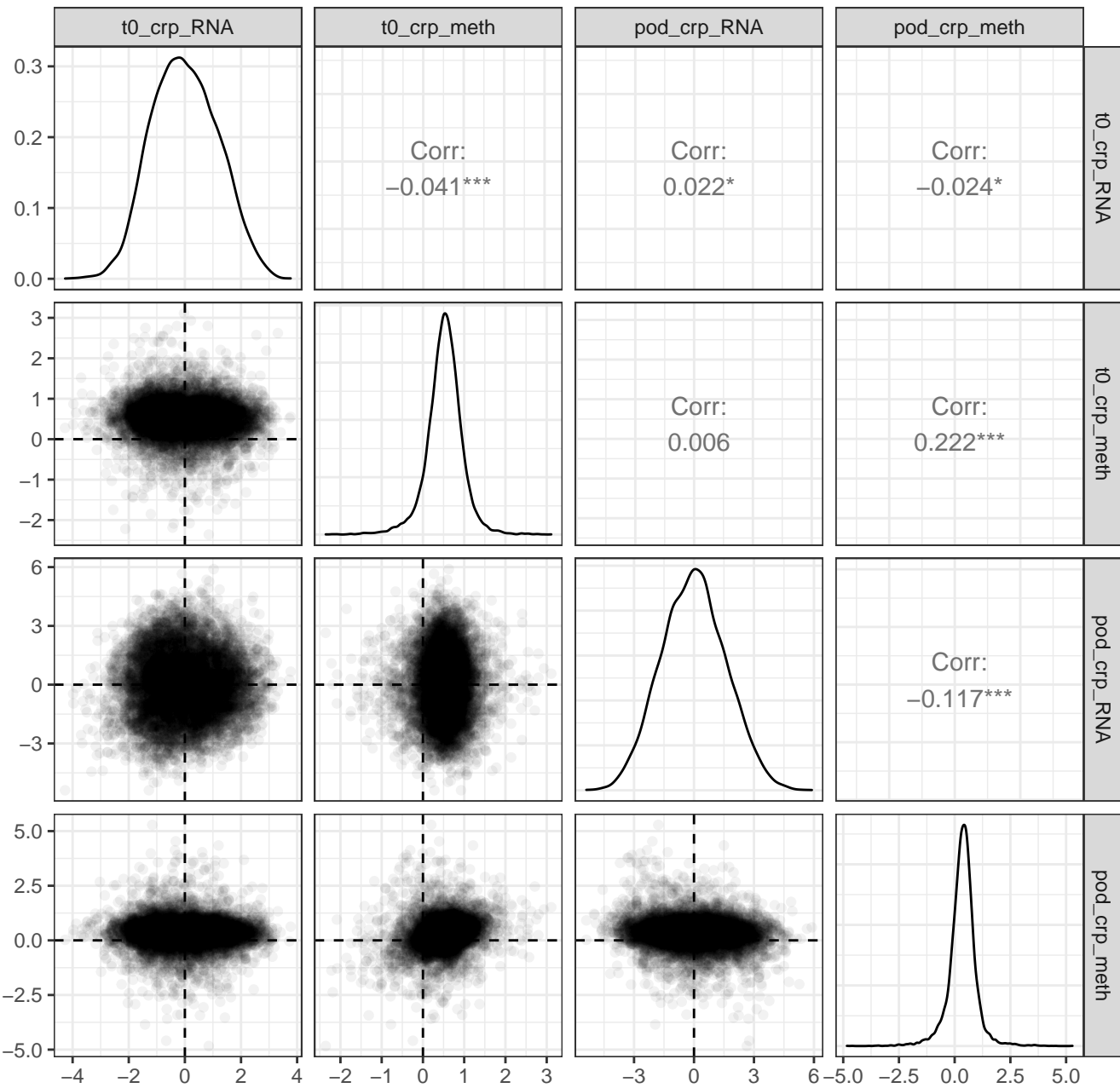
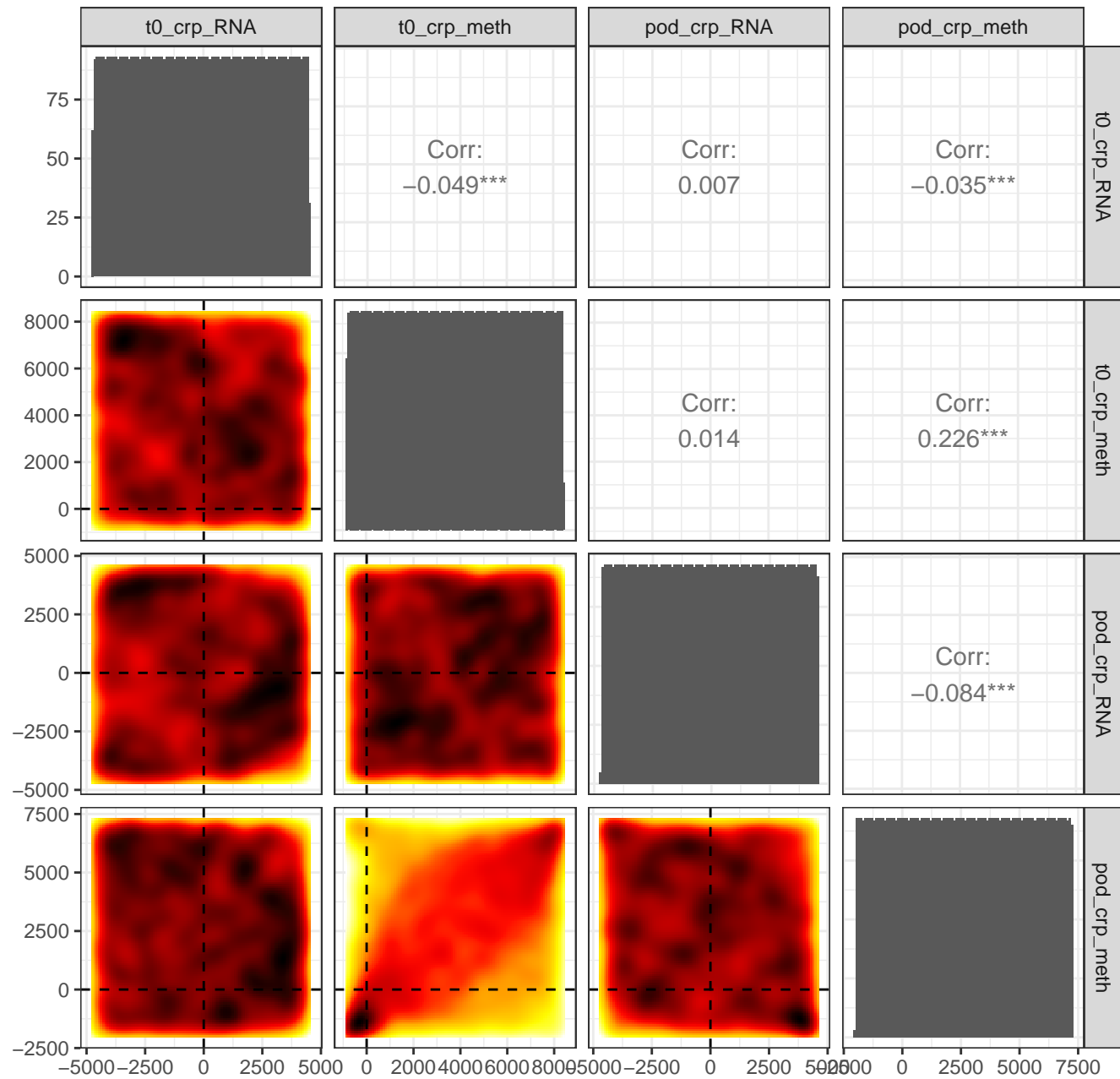


Scatterplot of all genes

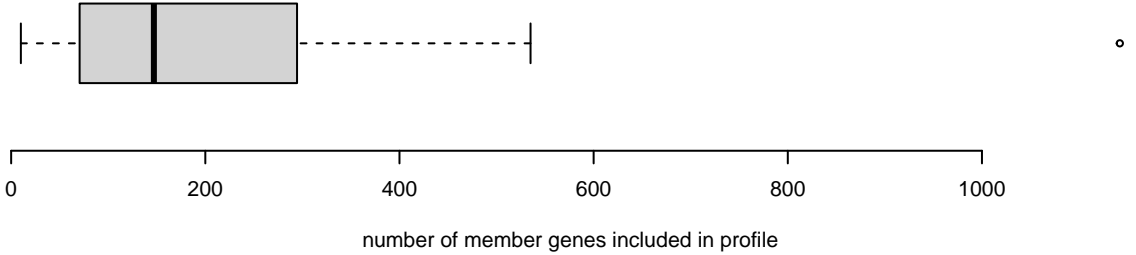


Contour plot of all genes after ranking

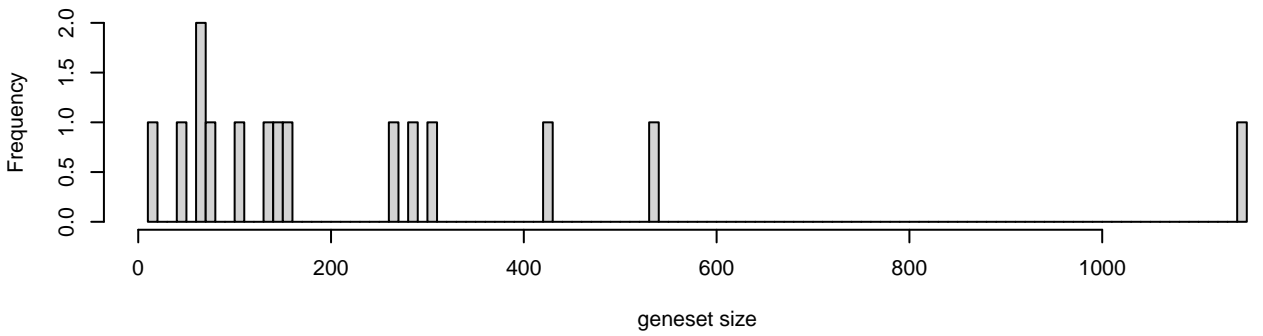


	t0_crp_RNA	t0_crp_meth	pod_crp_RNA	pod_crp_meth	Number of genes in each sector
1	-1	-1	-1	-1	101
2	1	-1	-1	-1	92
3	-1	1	-1	-1	375
4	1	1	-1	-1	344
5	-1	-1	1	-1	101
6	1	-1	1	-1	120
7	-1	1	1	-1	450
8	1	1	1	-1	415
9	-1	-1	1	0	1
10	-1	-1	-1	1	122
11	1	-1	-1	1	122
12	-1	0	-1	1	1
13	-1	1	-1	1	1763
14	0	1	-1	1	1
15	1	1	-1	1	1770
16	-1	1	0	1	1
17	-1	-1	1	1	101
18	1	-1	1	1	101
19	-1	1	1	1	1732
20	1	1	1	1	1568

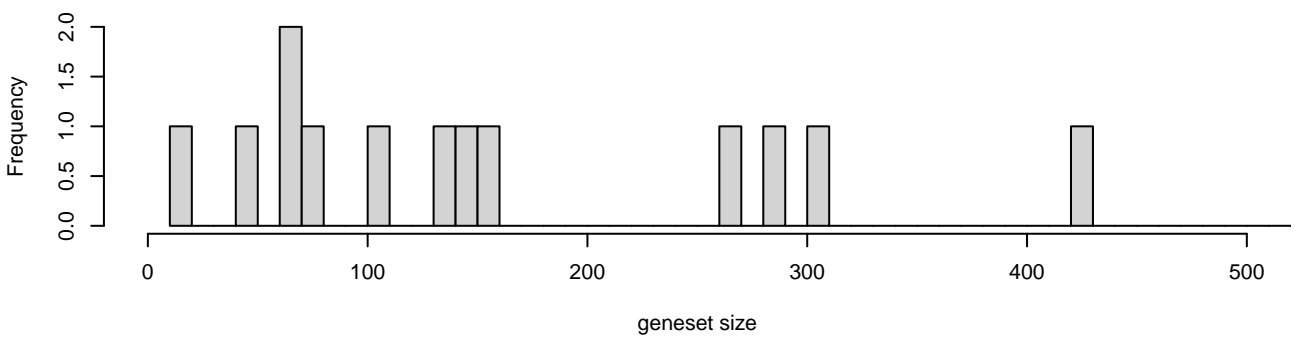
Gene set size



Histogram of geneset size

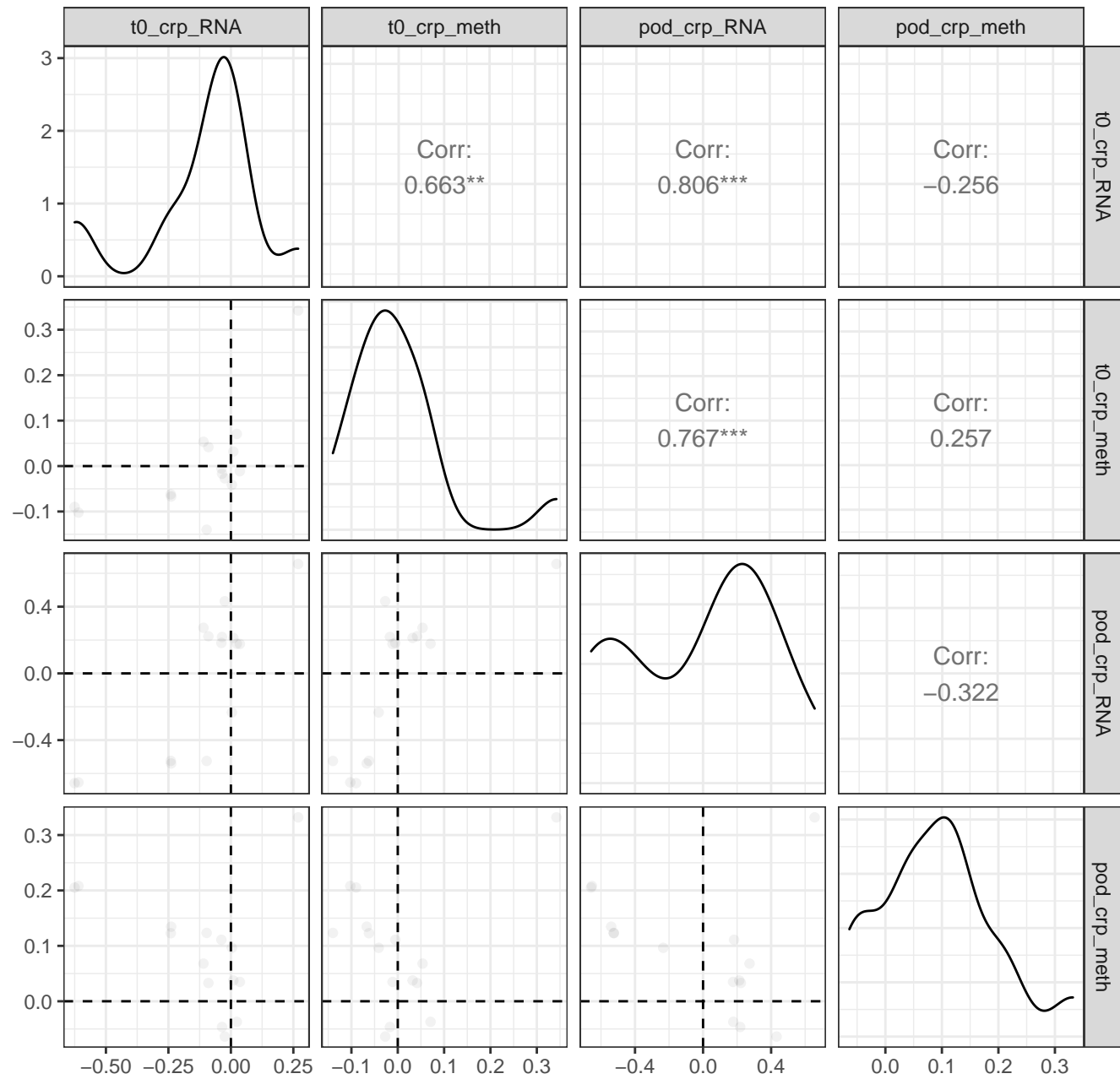


Trimmed histogram of geneset size

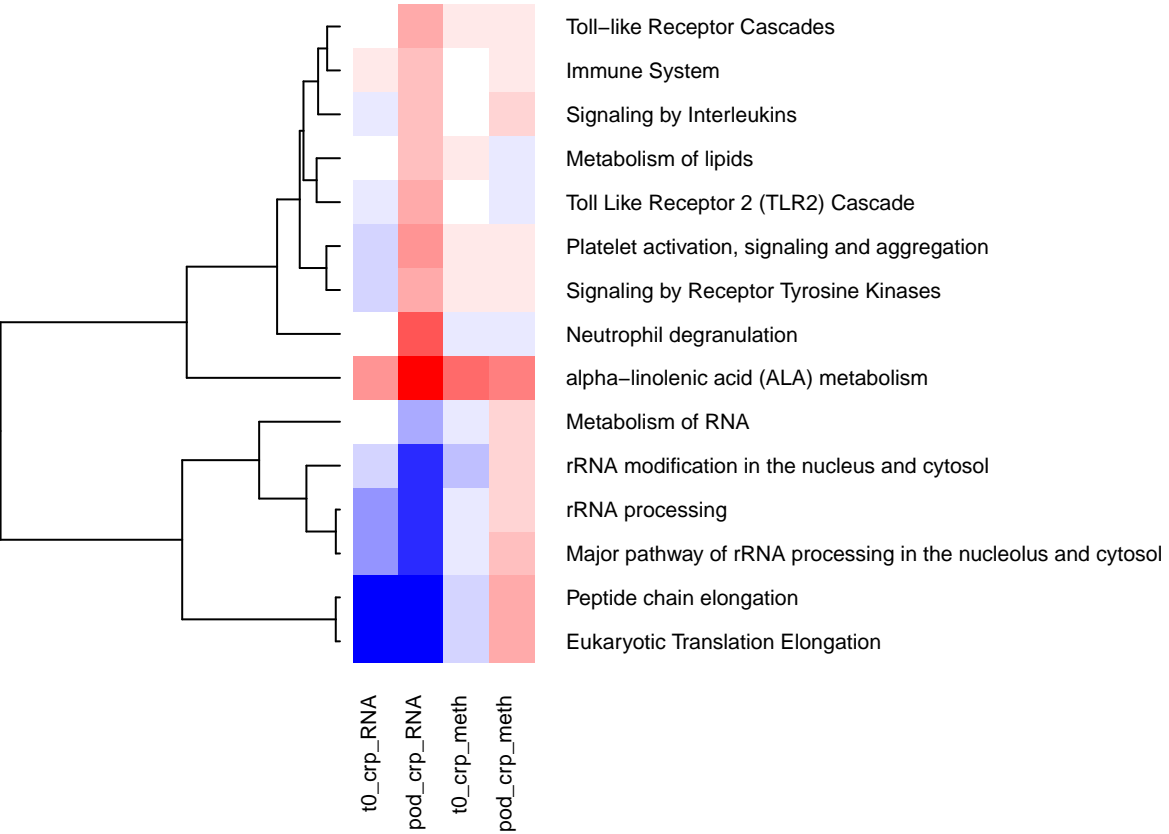
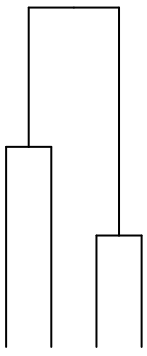
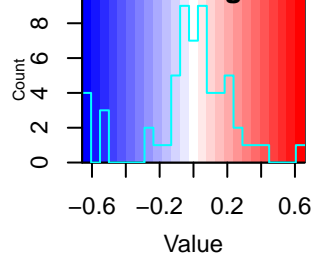


	s.t0_crp_RNA	s.t0_crp_meth	s.pod_crp_RNA	s.pod_crp_meth	Number of gene sets in each sector
1	-1	-1	1	-1	2
2	1	1	1	-1	1
3	-1	-1	-1	1	5
4	1	-1	-1	1	1
5	-1	-1	1	1	1
6	1	-1	1	1	1
7	-1	1	1	1	2
8	1	1	1	1	2

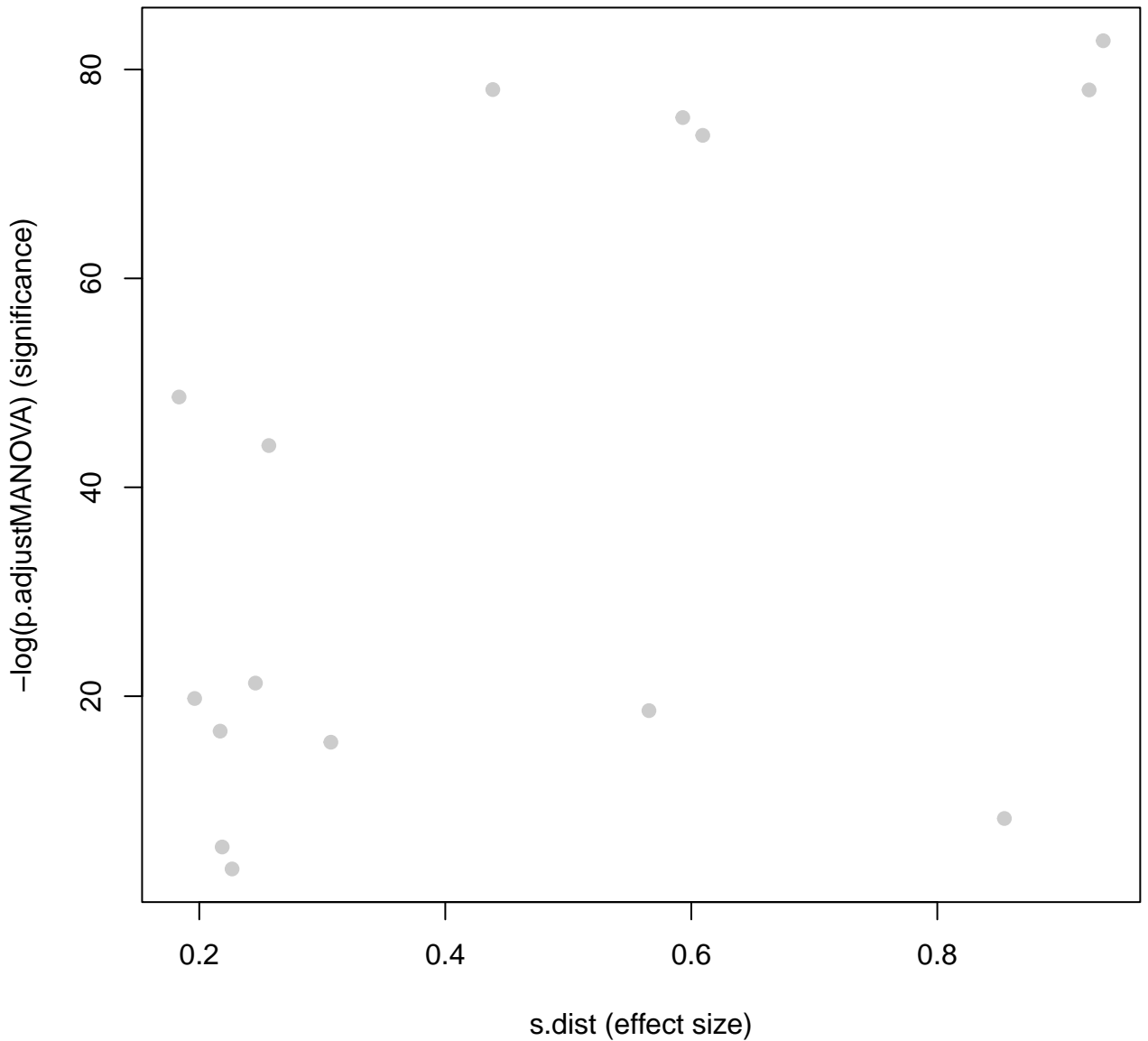
Scatterplot of all genesets; FDR<0.05 in red



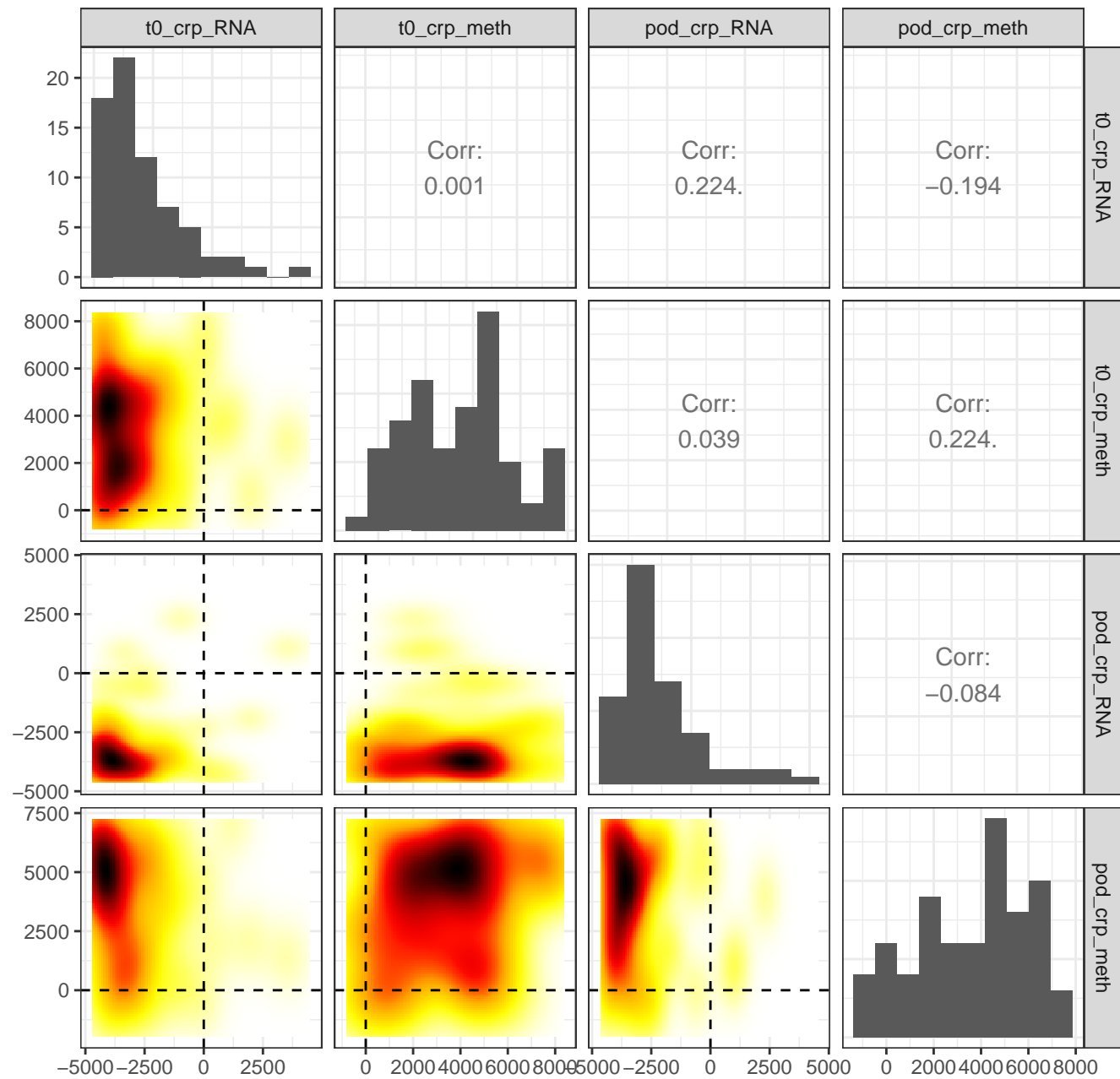
Color Key and Histogram



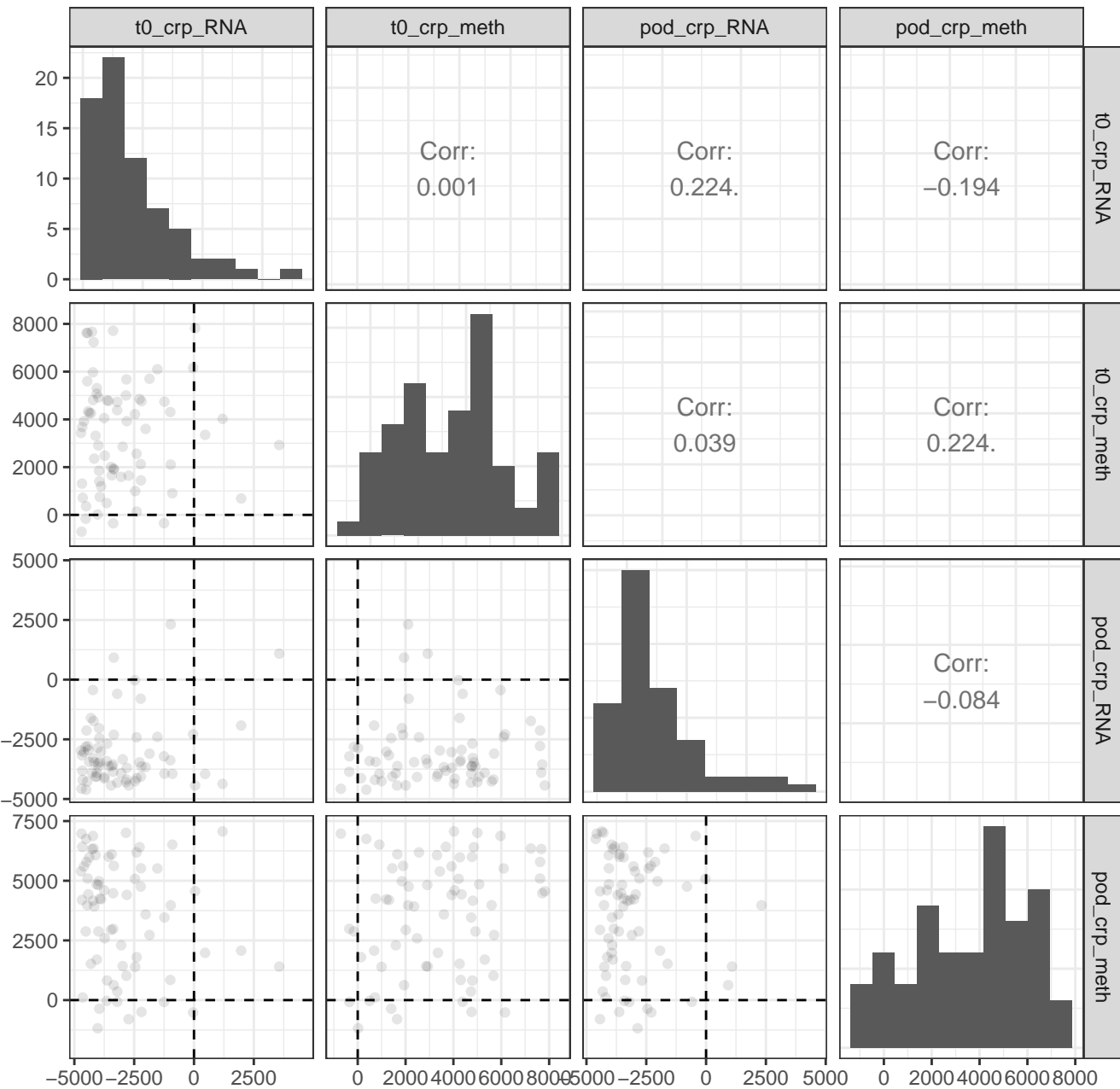
effect size versus statistical significance



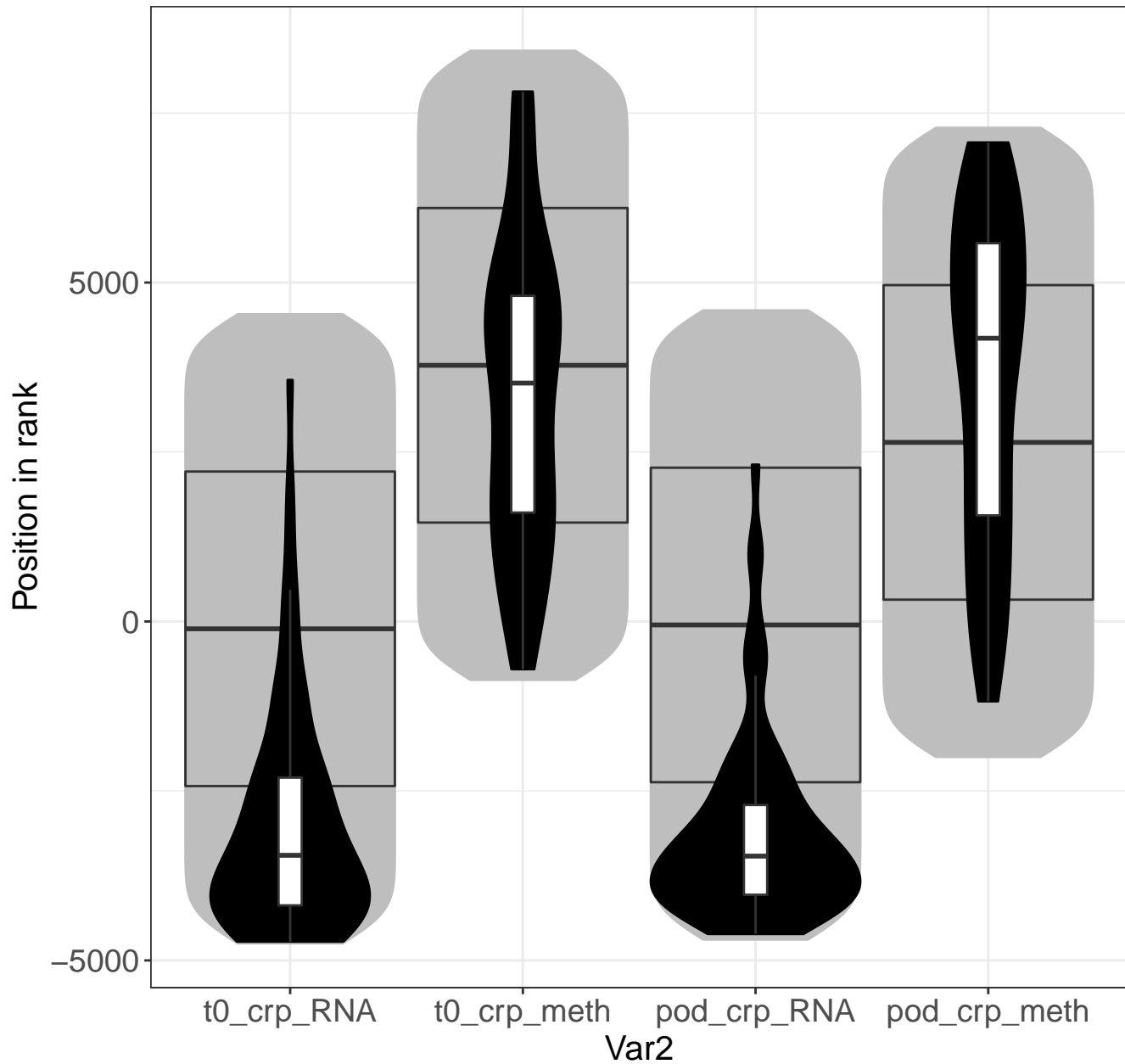
Eukaryotic Translation Elongation



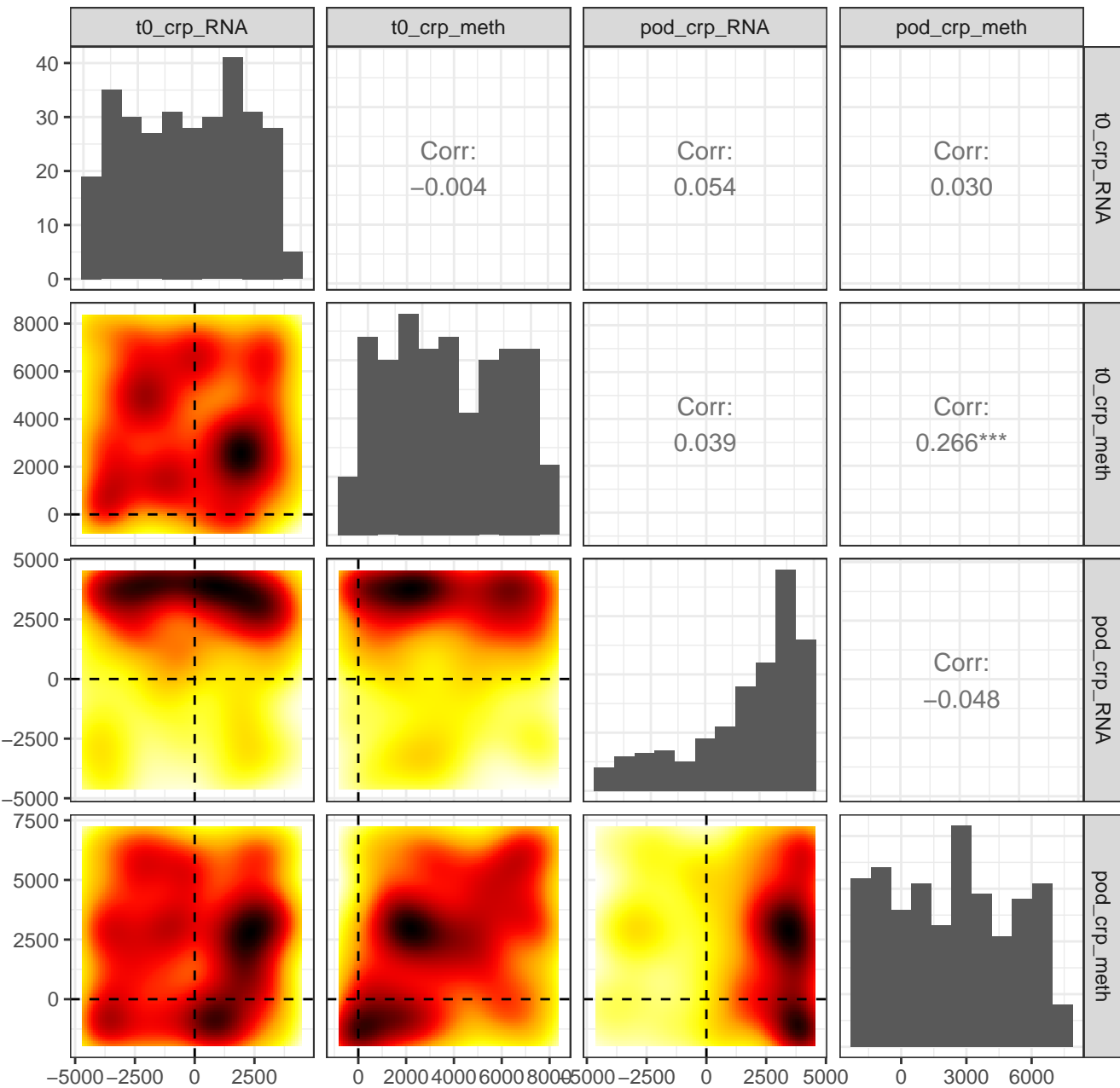
Eukaryotic Translation Elongation



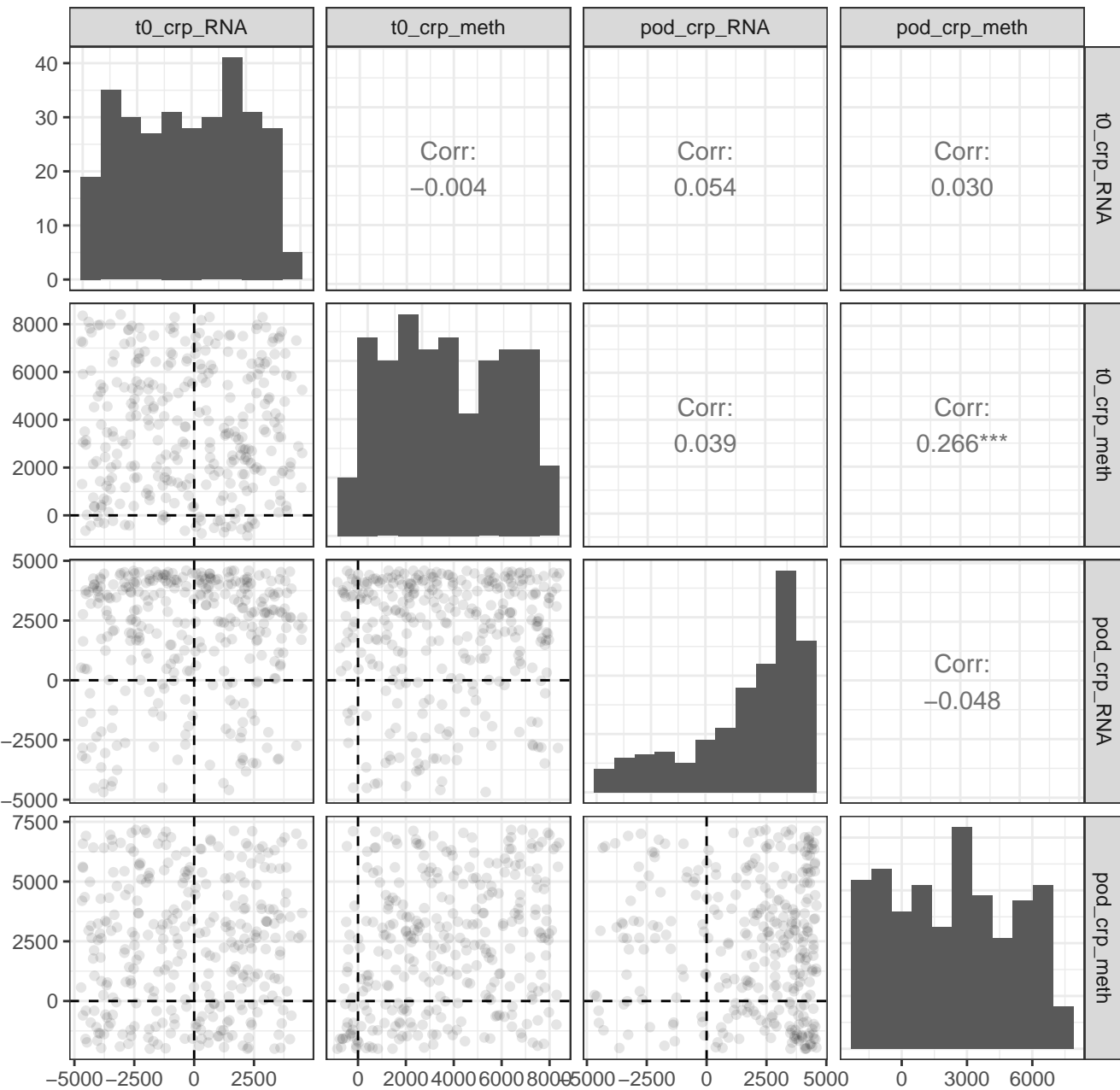
Eukaryotic Translation Elongation



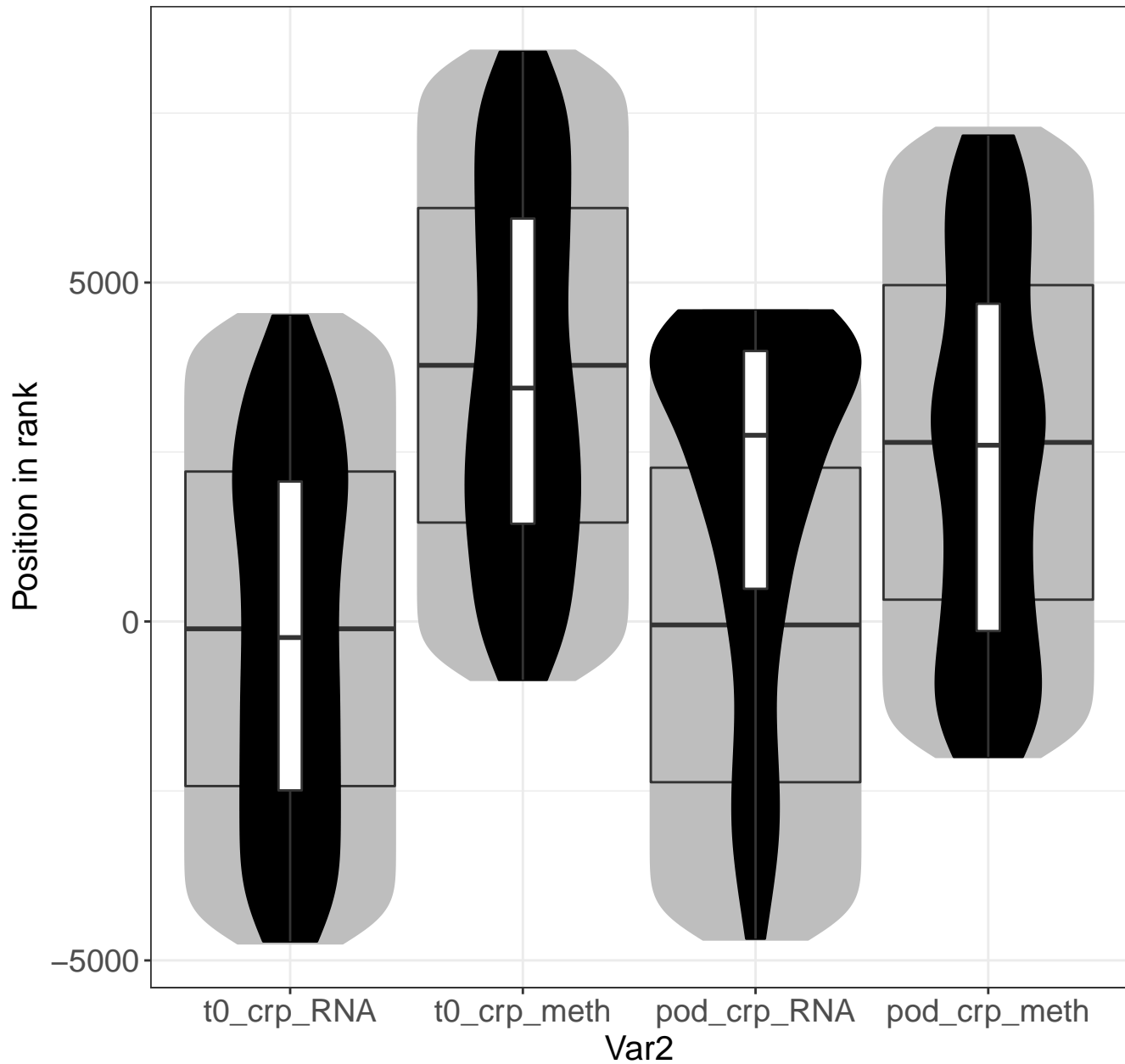
Neutrophil degranulation



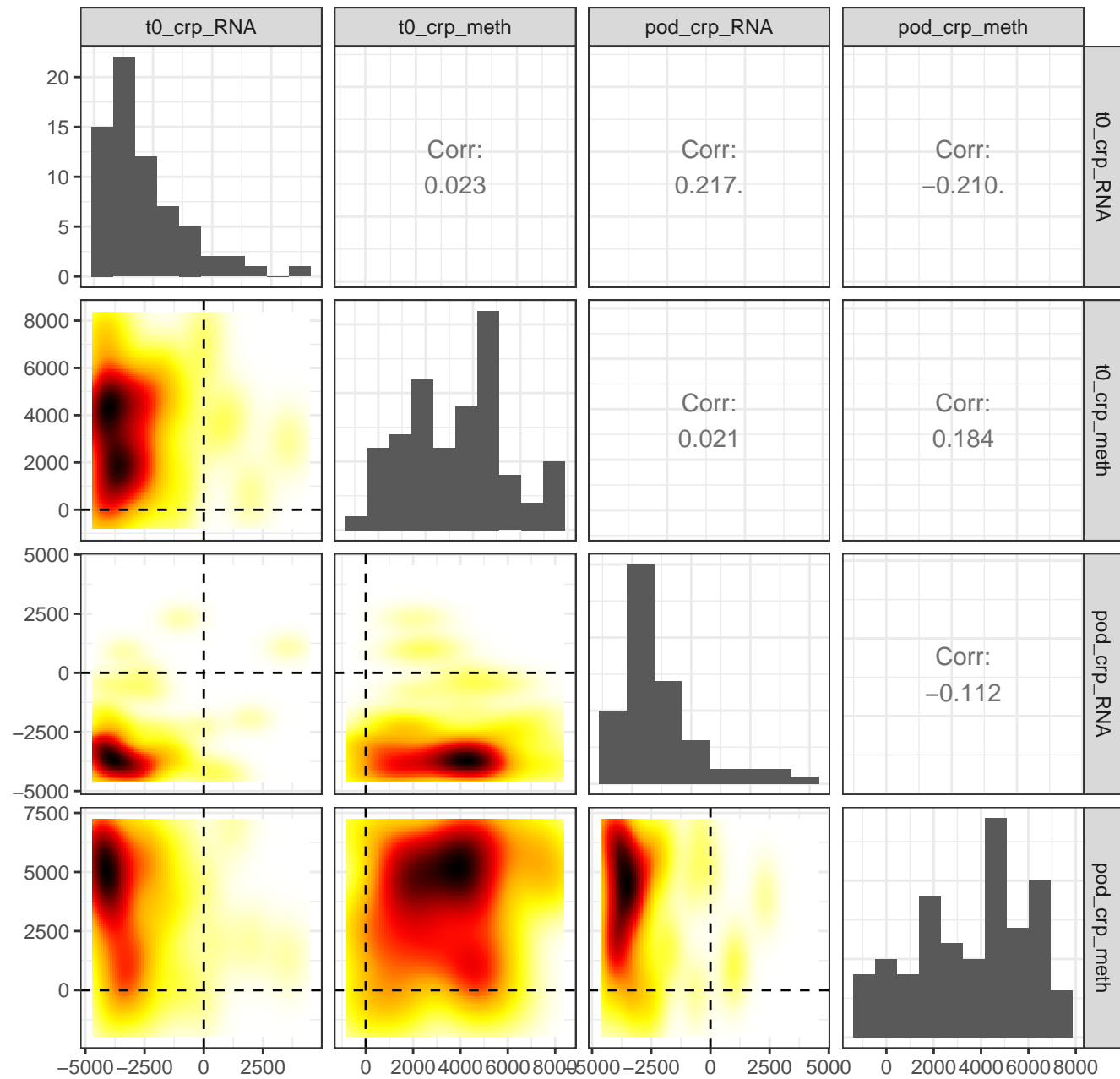
Neutrophil degranulation



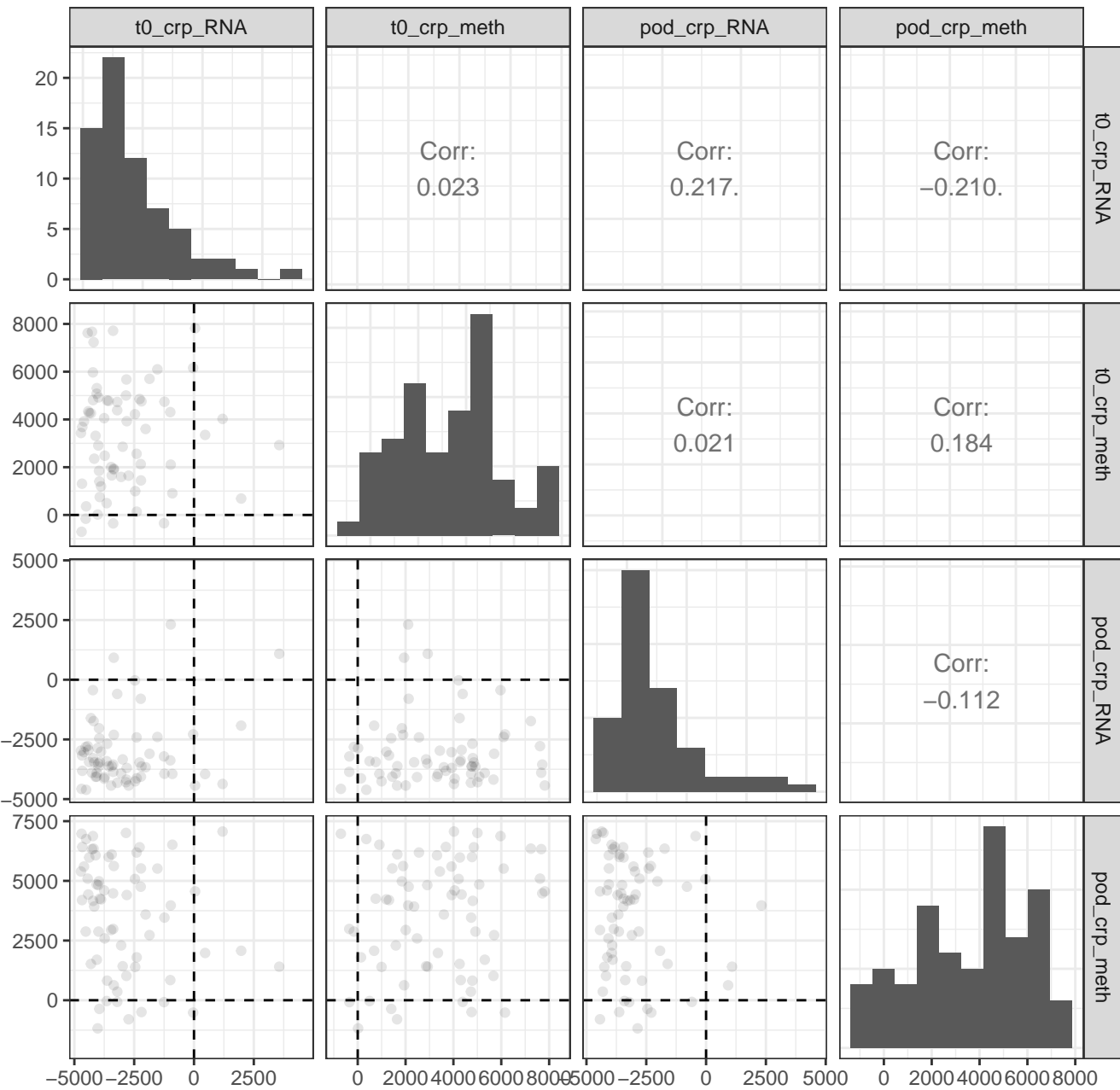
Neutrophil degranulation



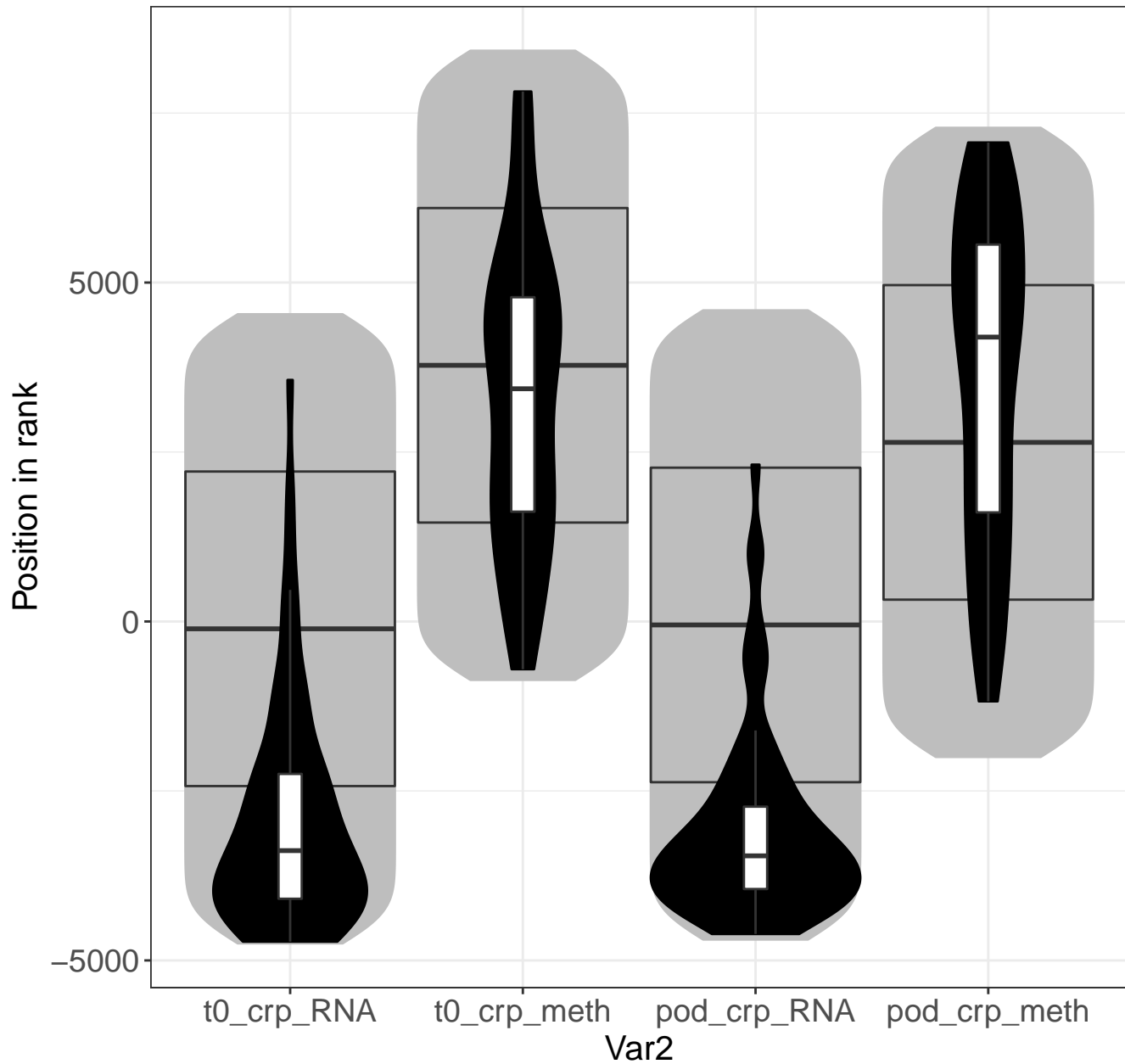
Peptide chain elongation



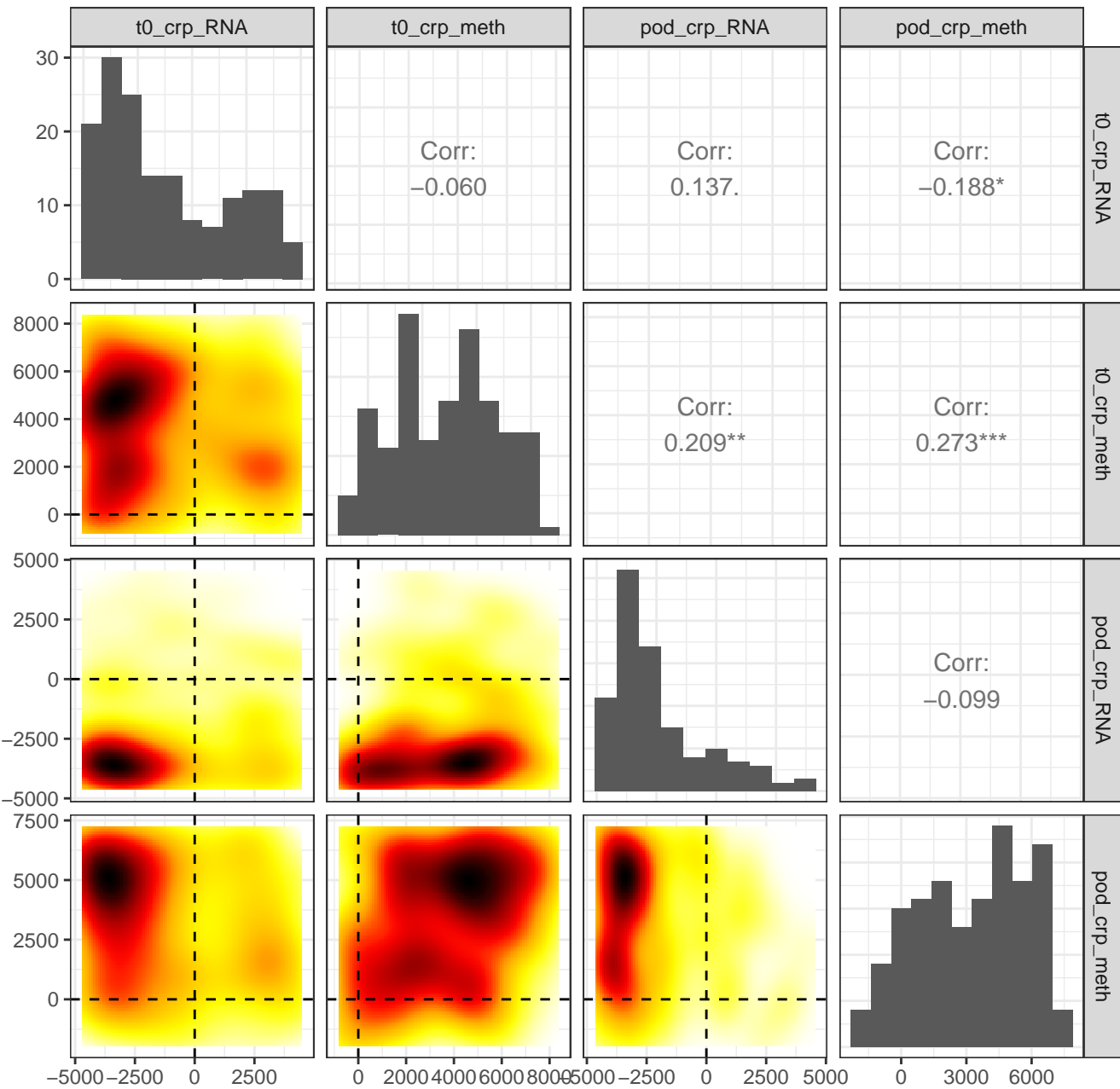
Peptide chain elongation



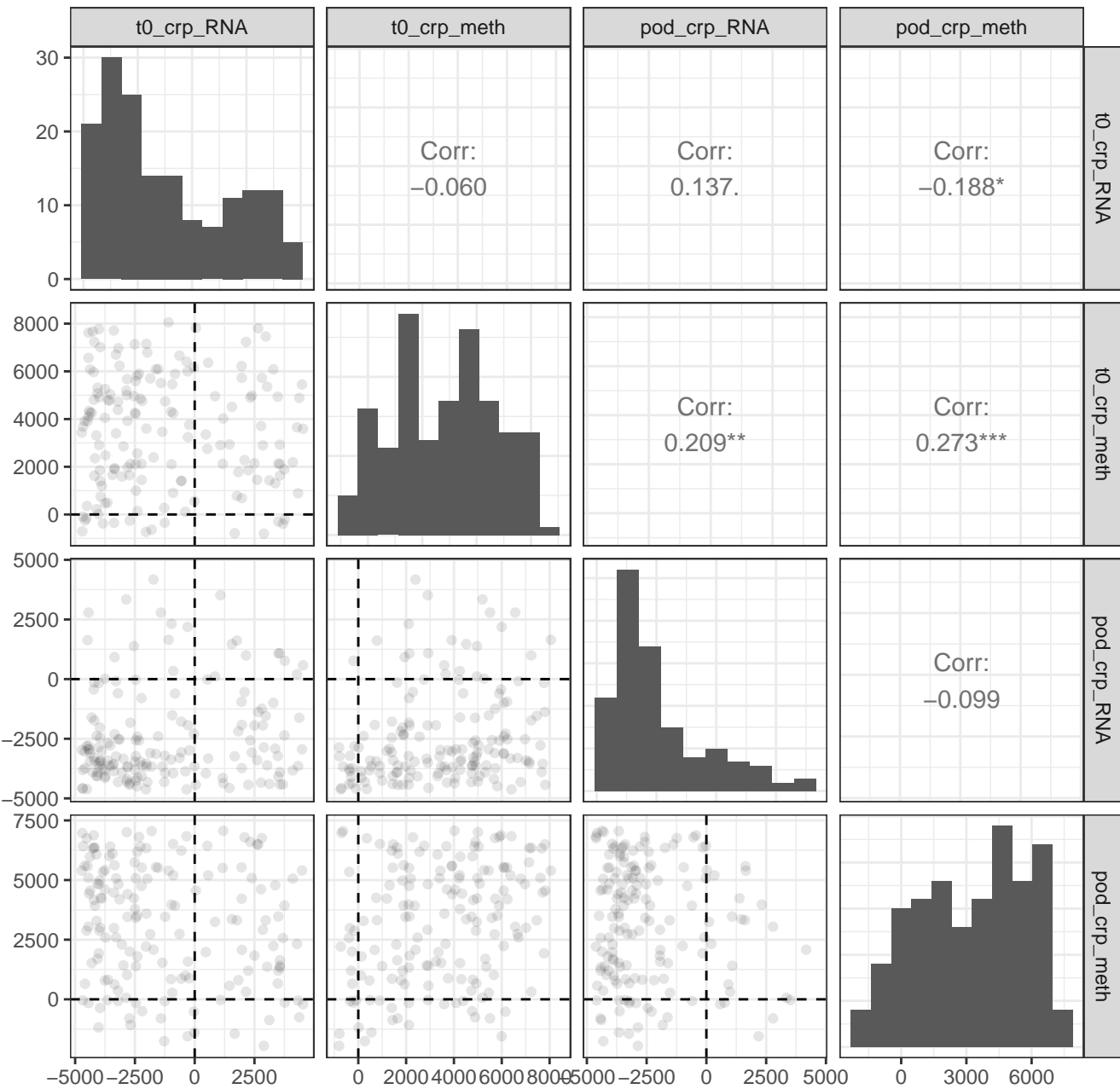
Peptide chain elongation



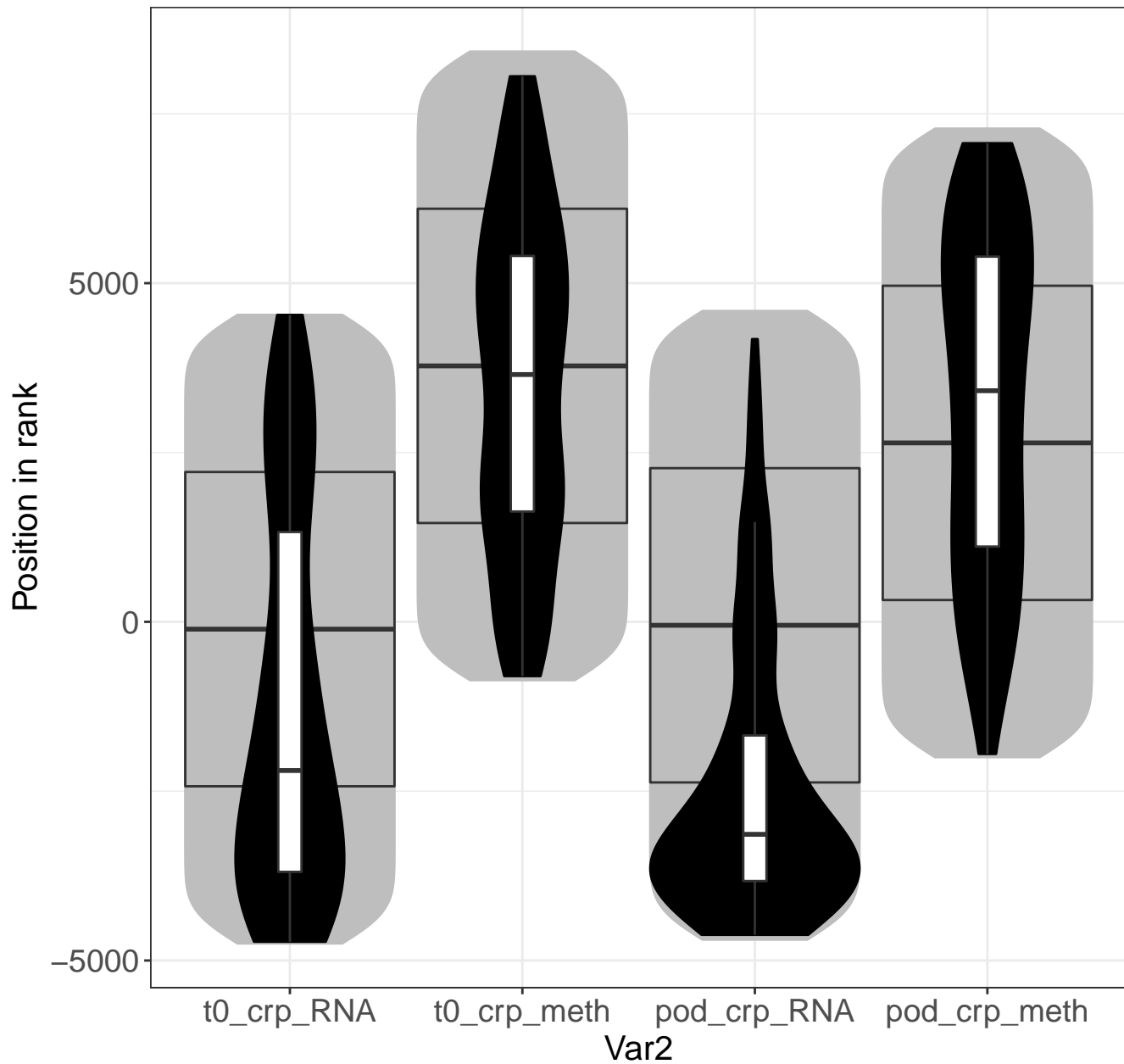
rRNA processing



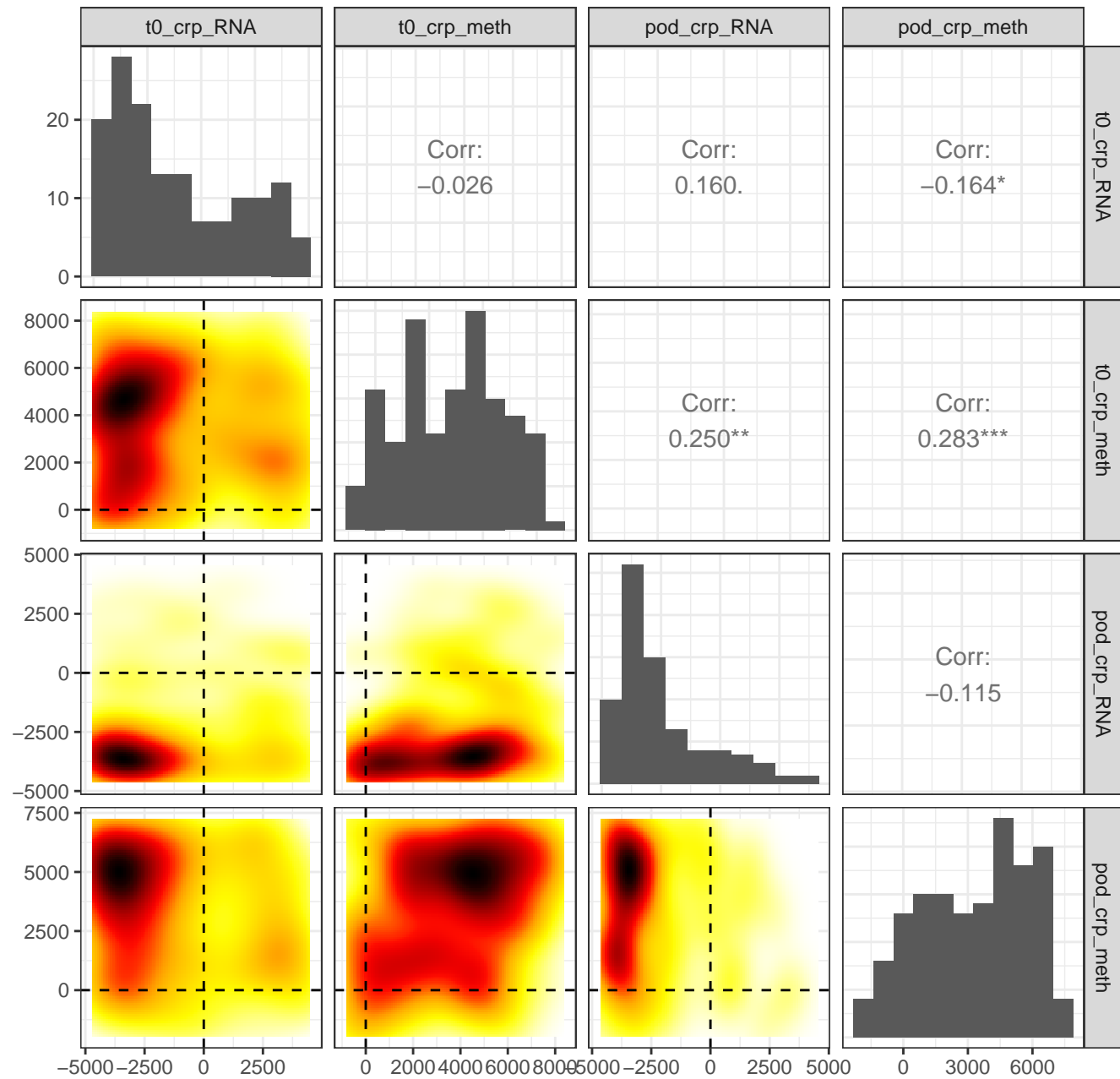
rRNA processing



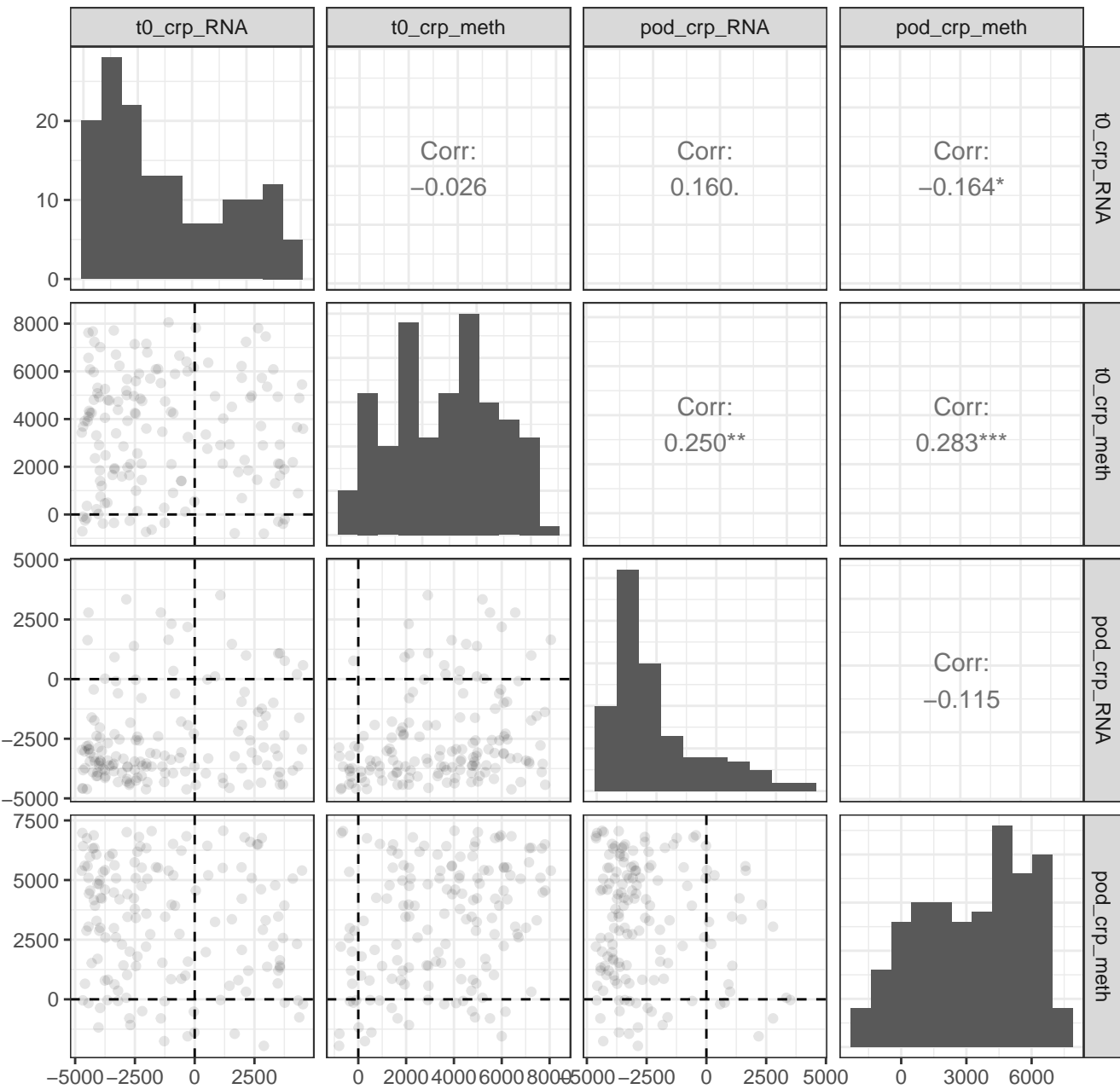
rRNA processing



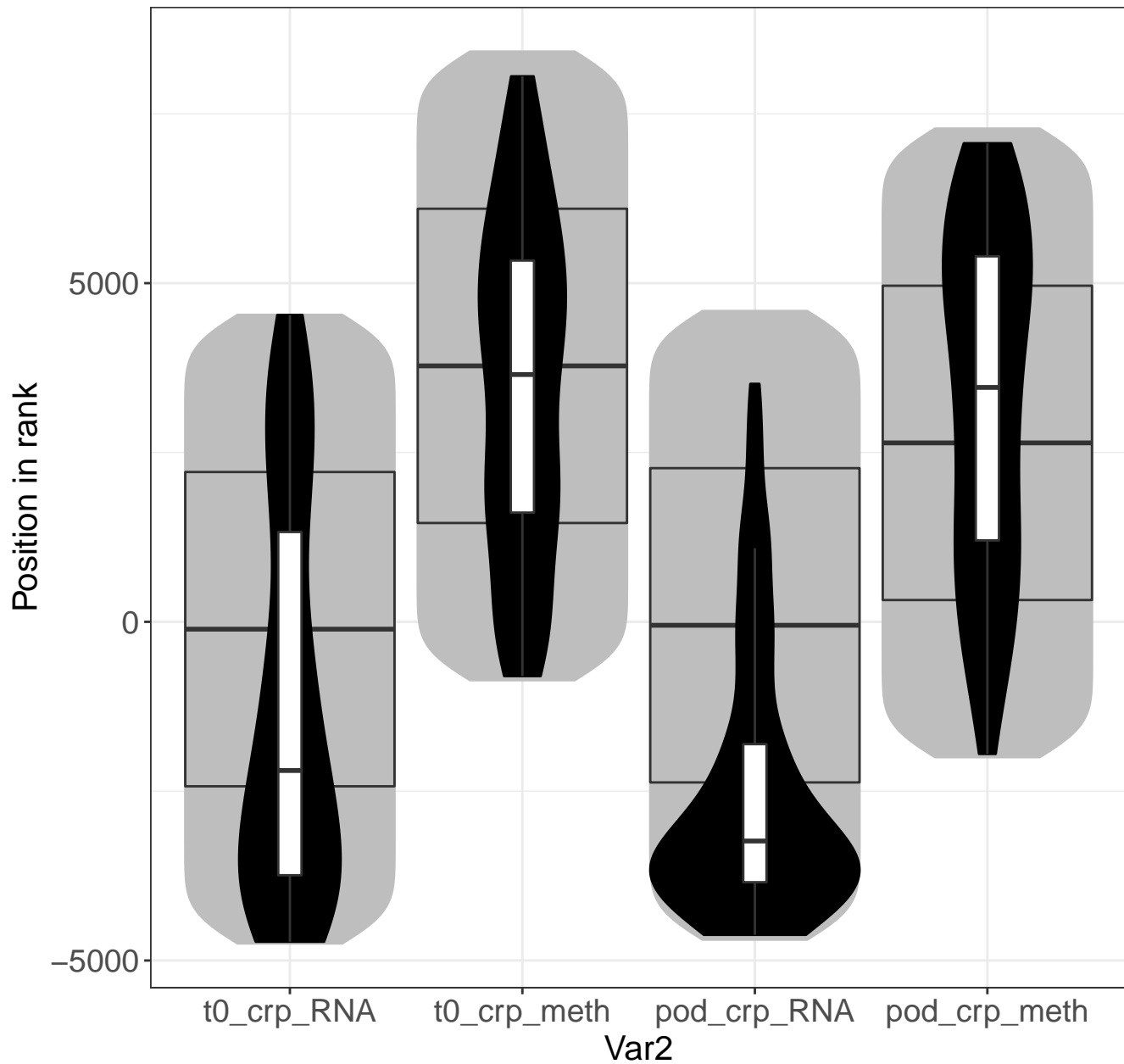
Major pathway of rRNA processing in the nucleolus and cytosol



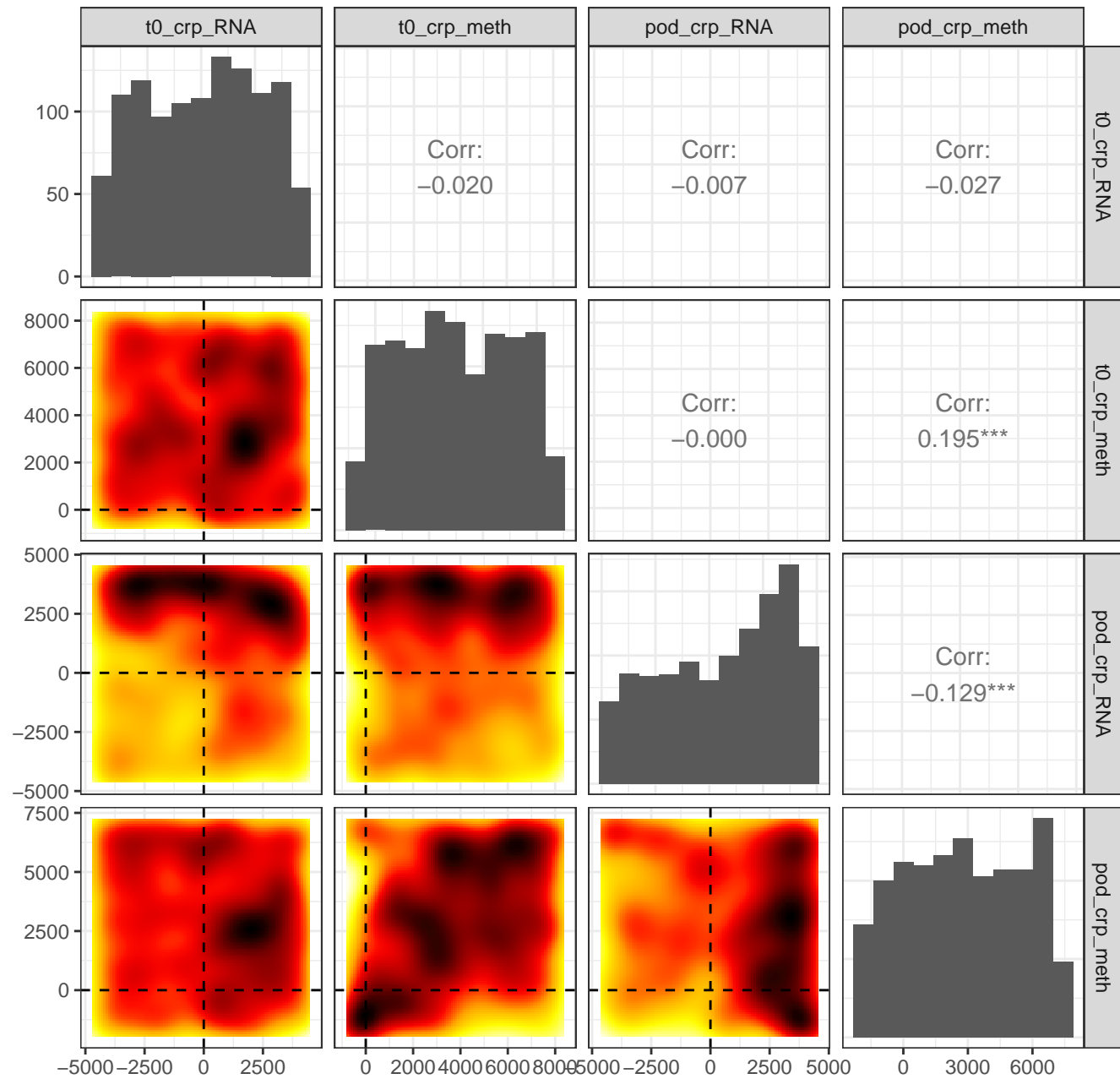
Major pathway of rRNA processing in the nucleolus and cytosol



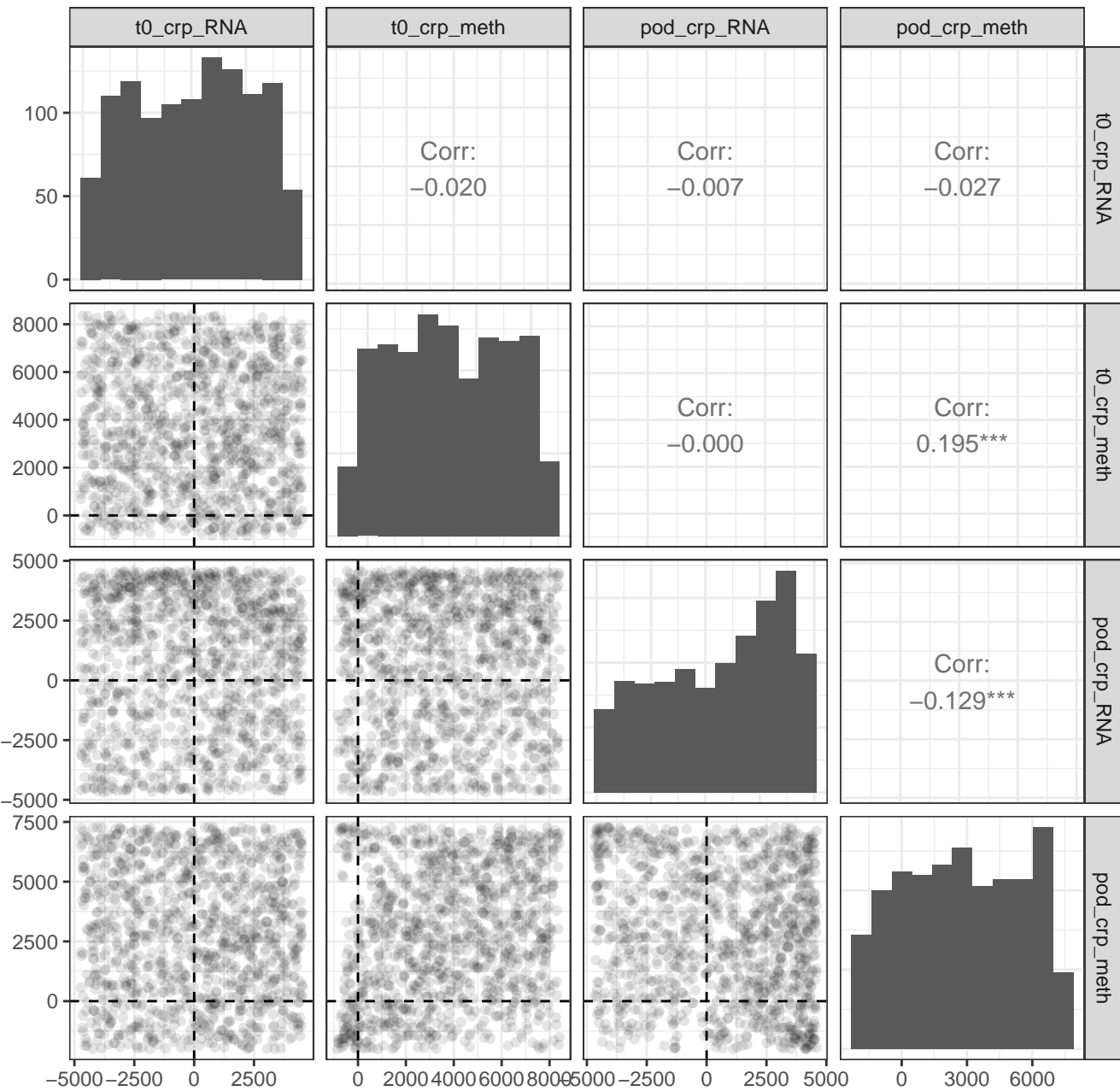
Major pathway of rRNA processing in the nucleolus



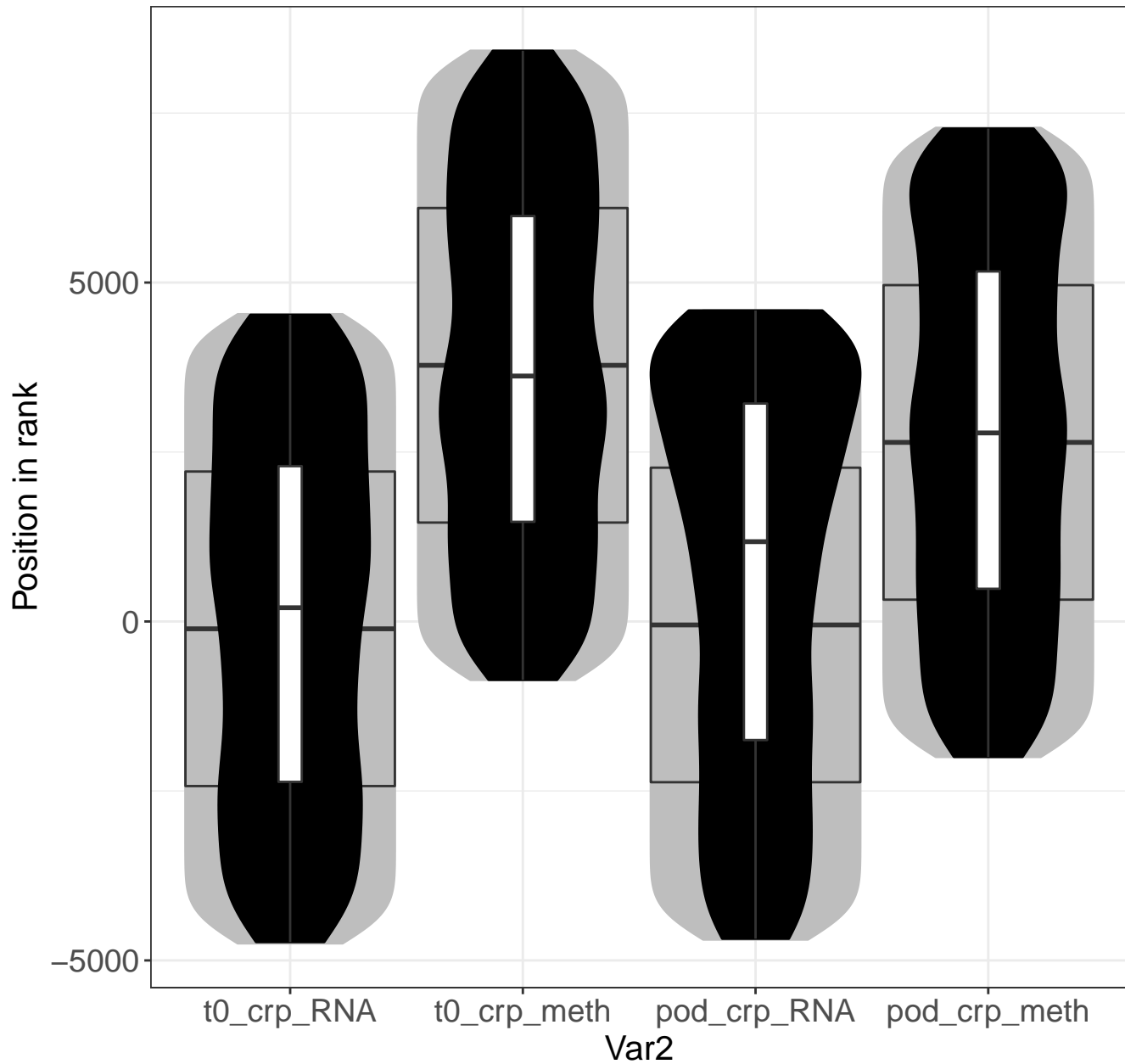
Immune System



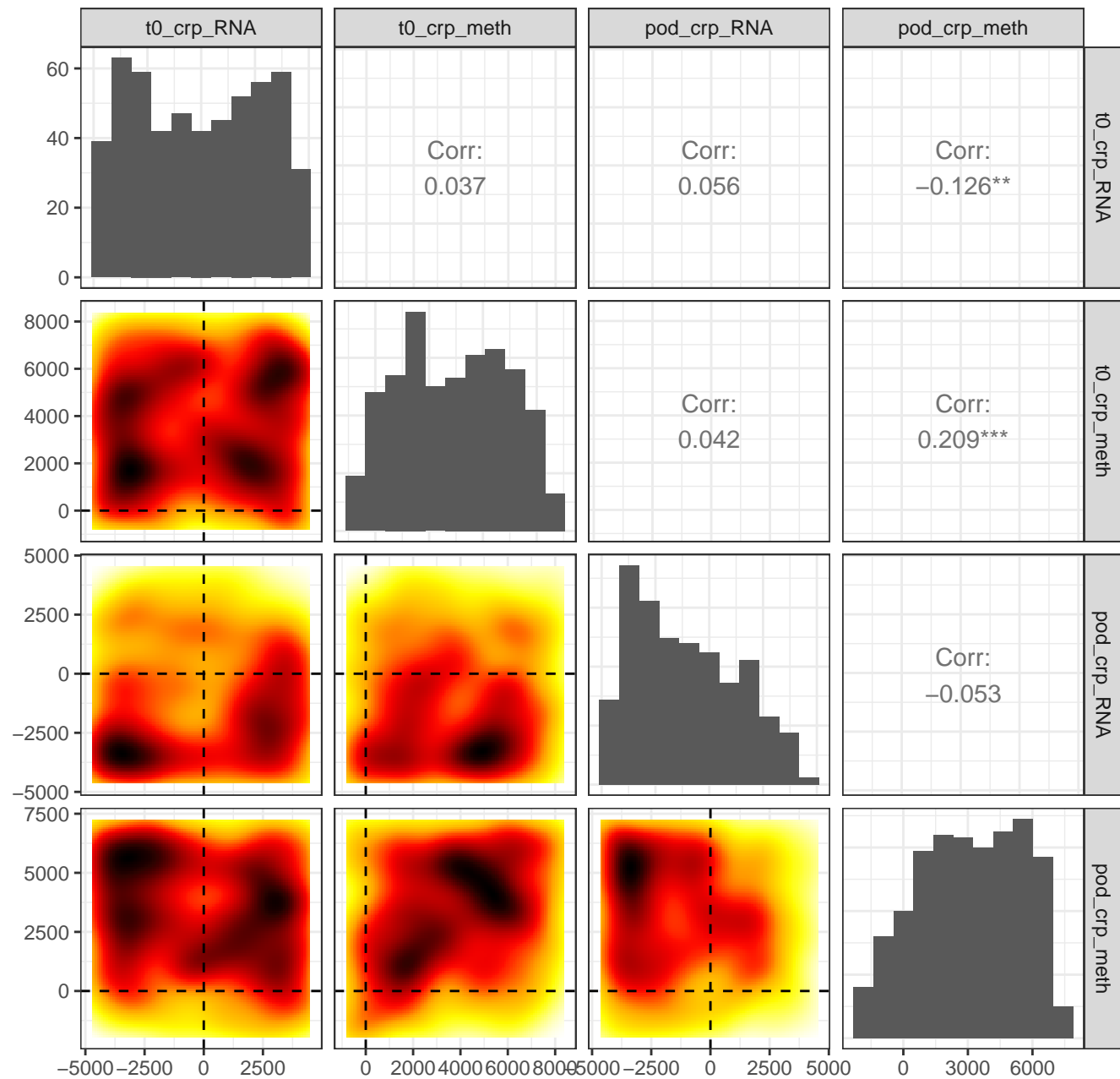
Immune System



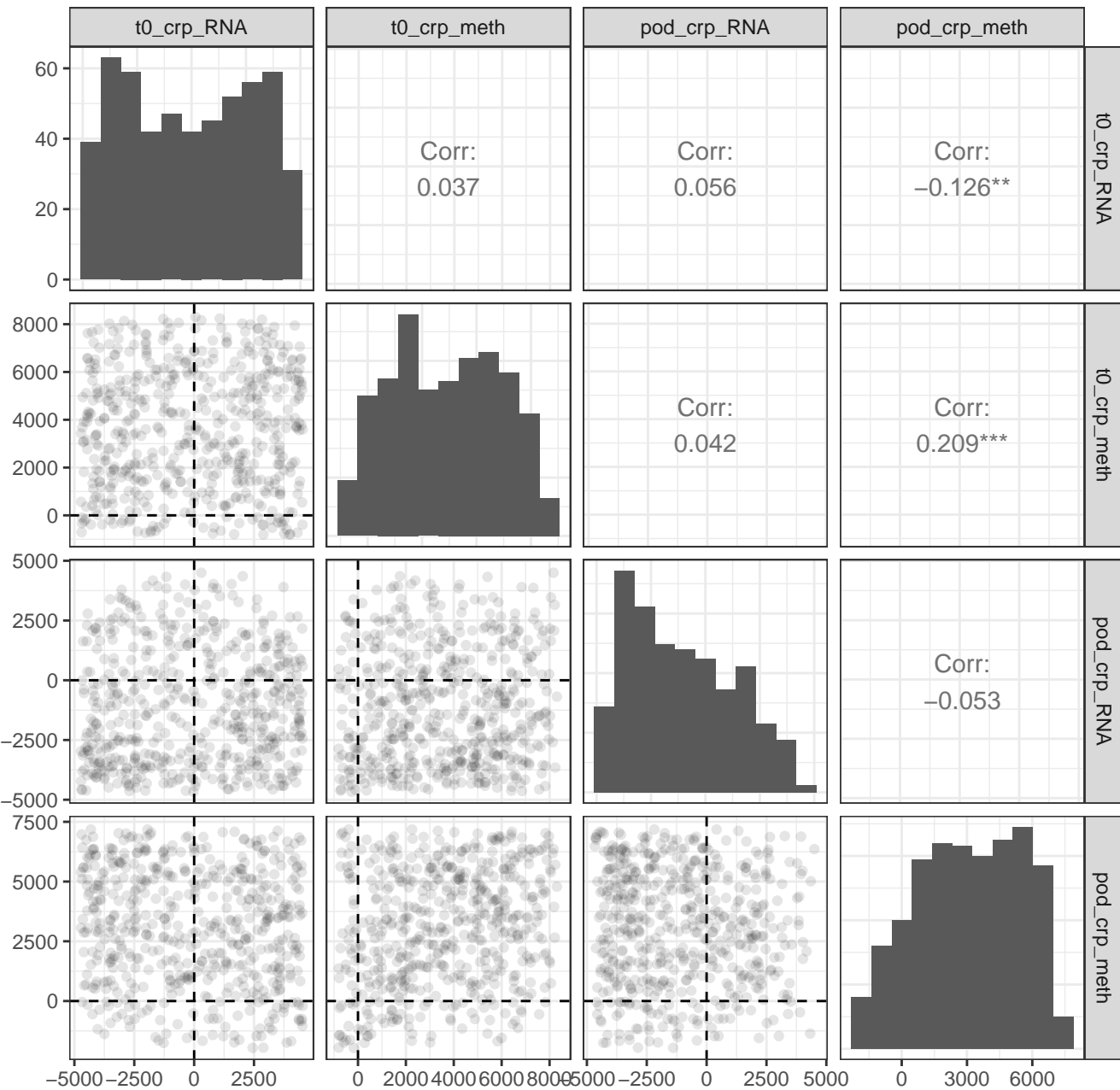
Immune System



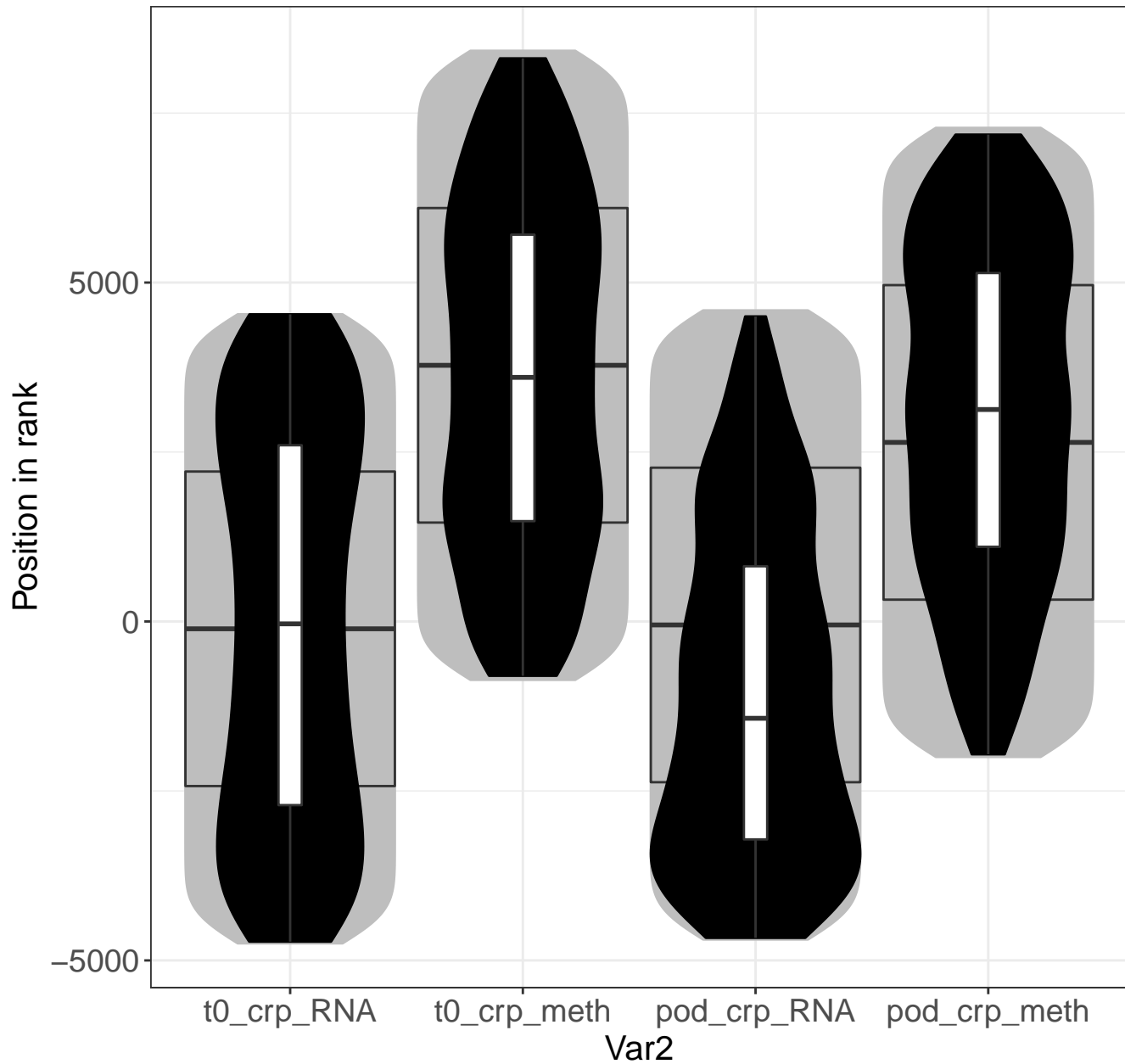
Metabolism of RNA



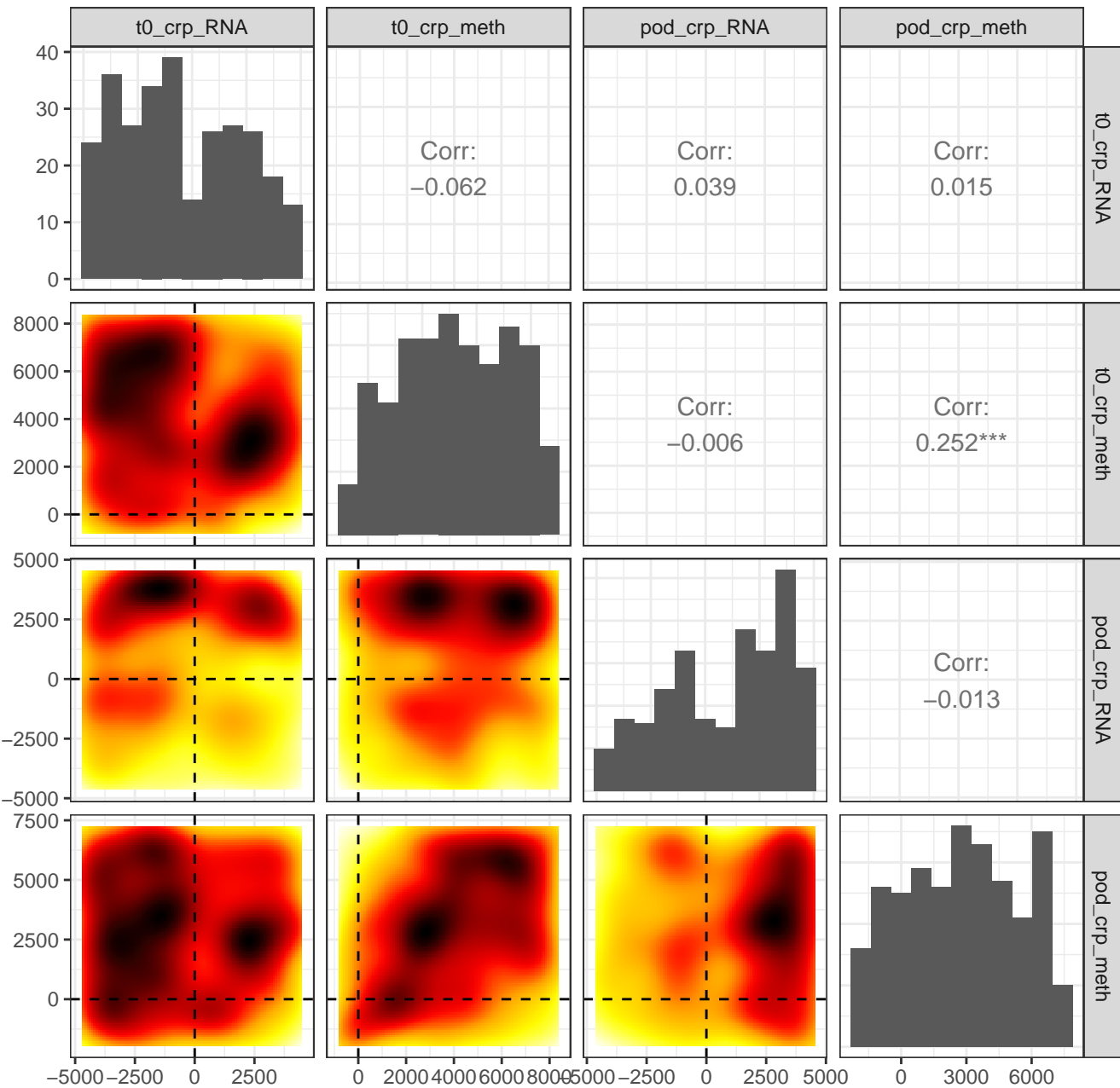
Metabolism of RNA



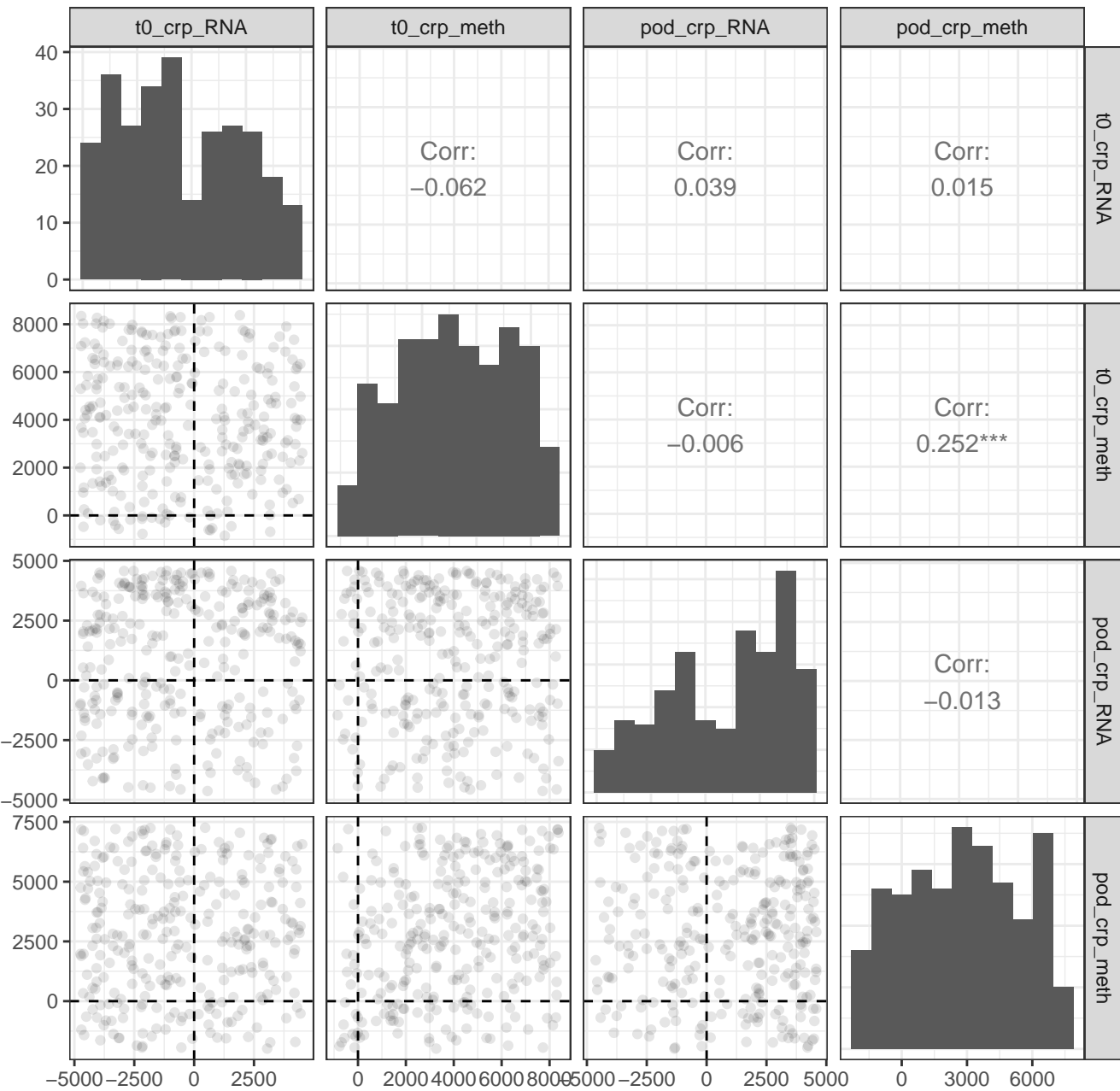
Metabolism of RNA



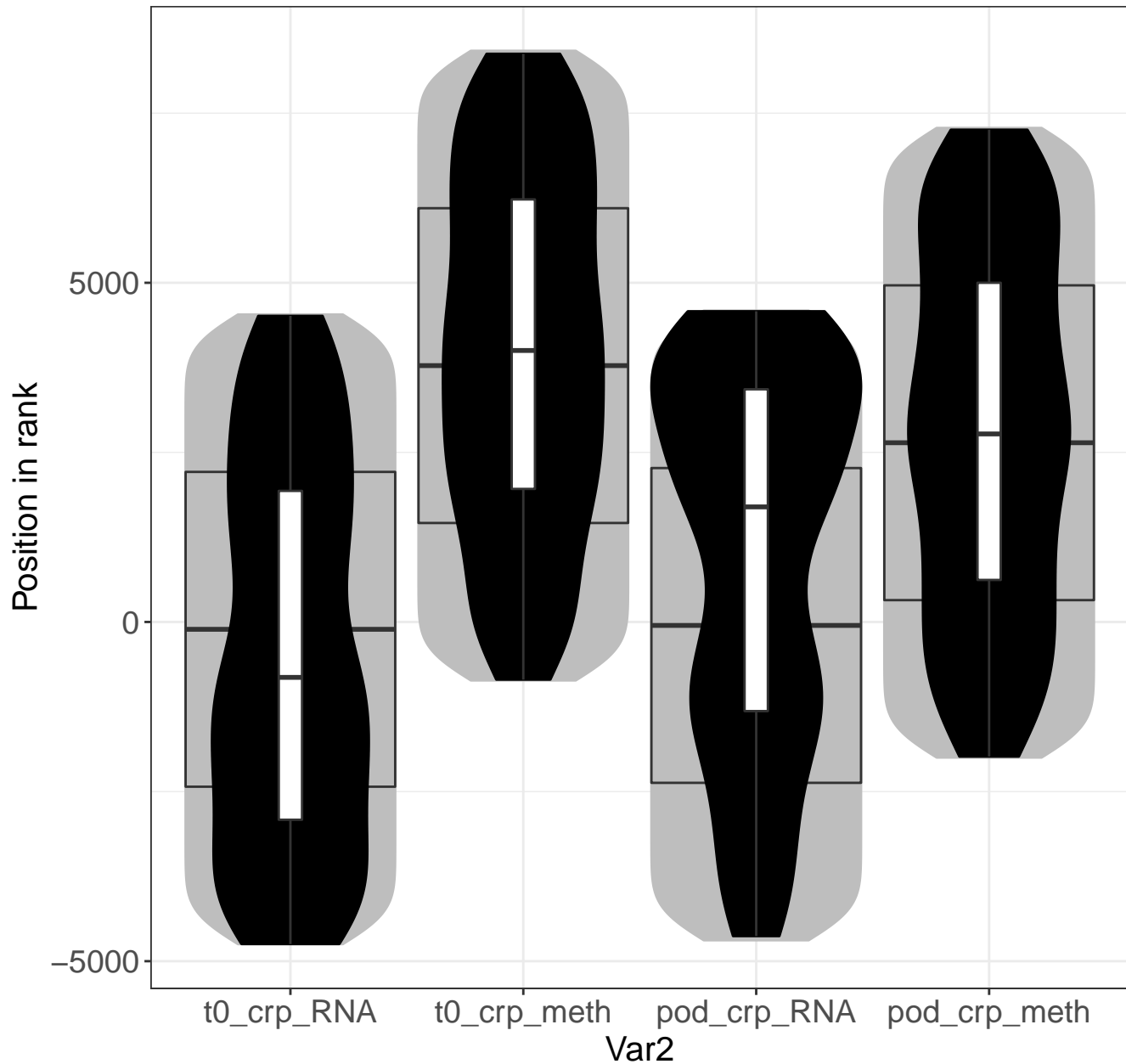
Signaling by Receptor Tyrosine Kinases



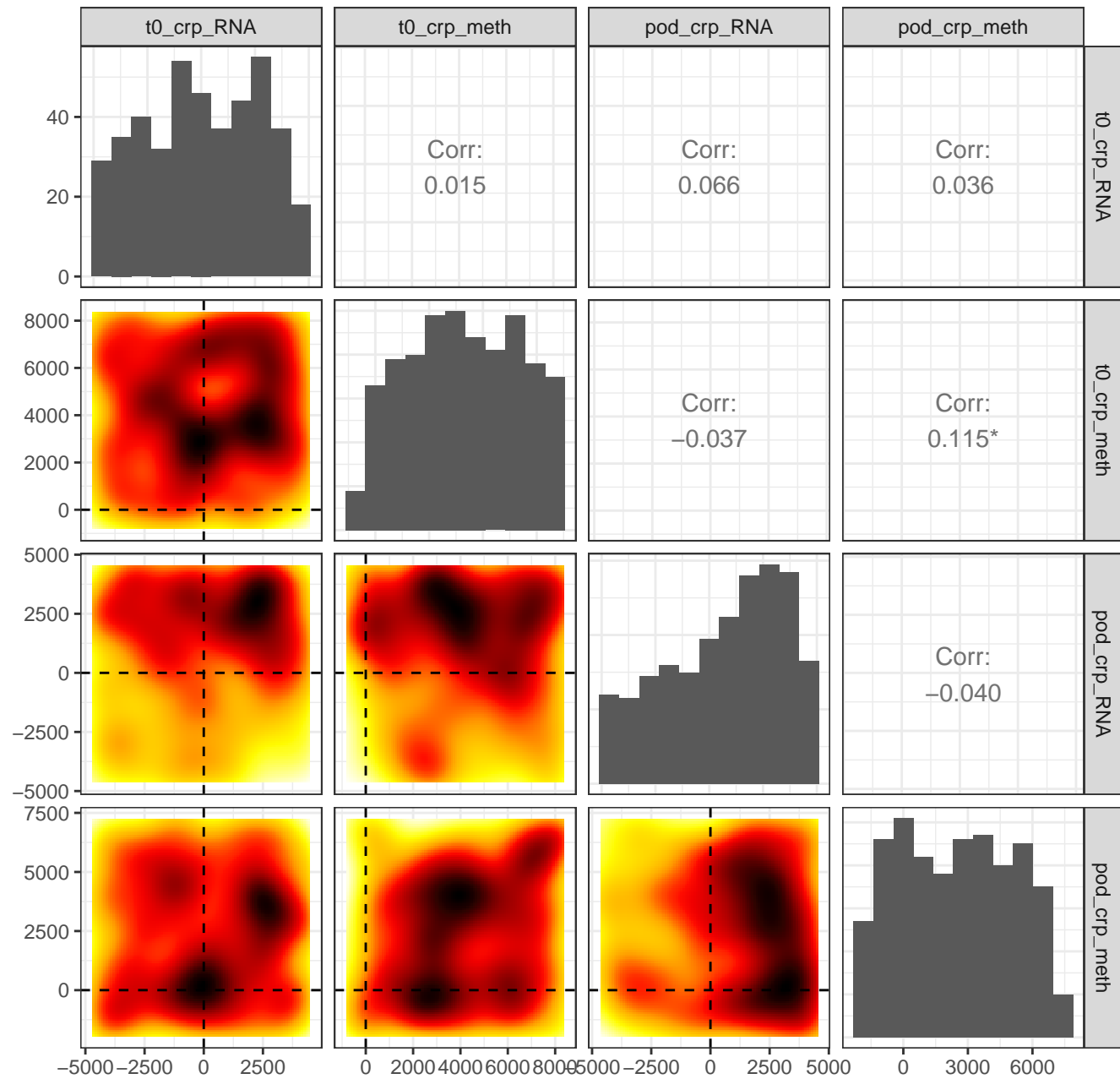
Signaling by Receptor Tyrosine Kinases



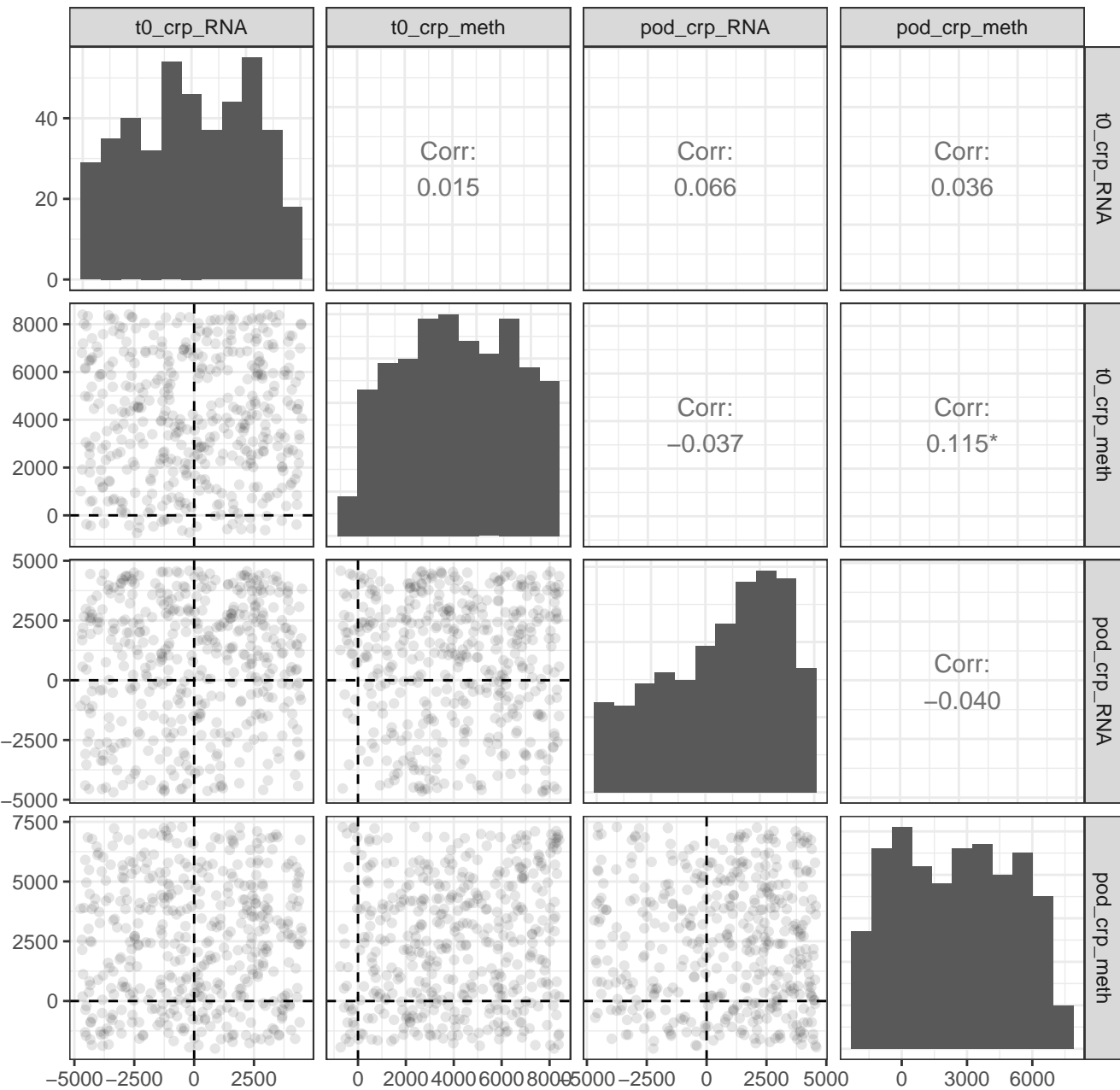
Signaling by Receptor Tyrosine Kinases



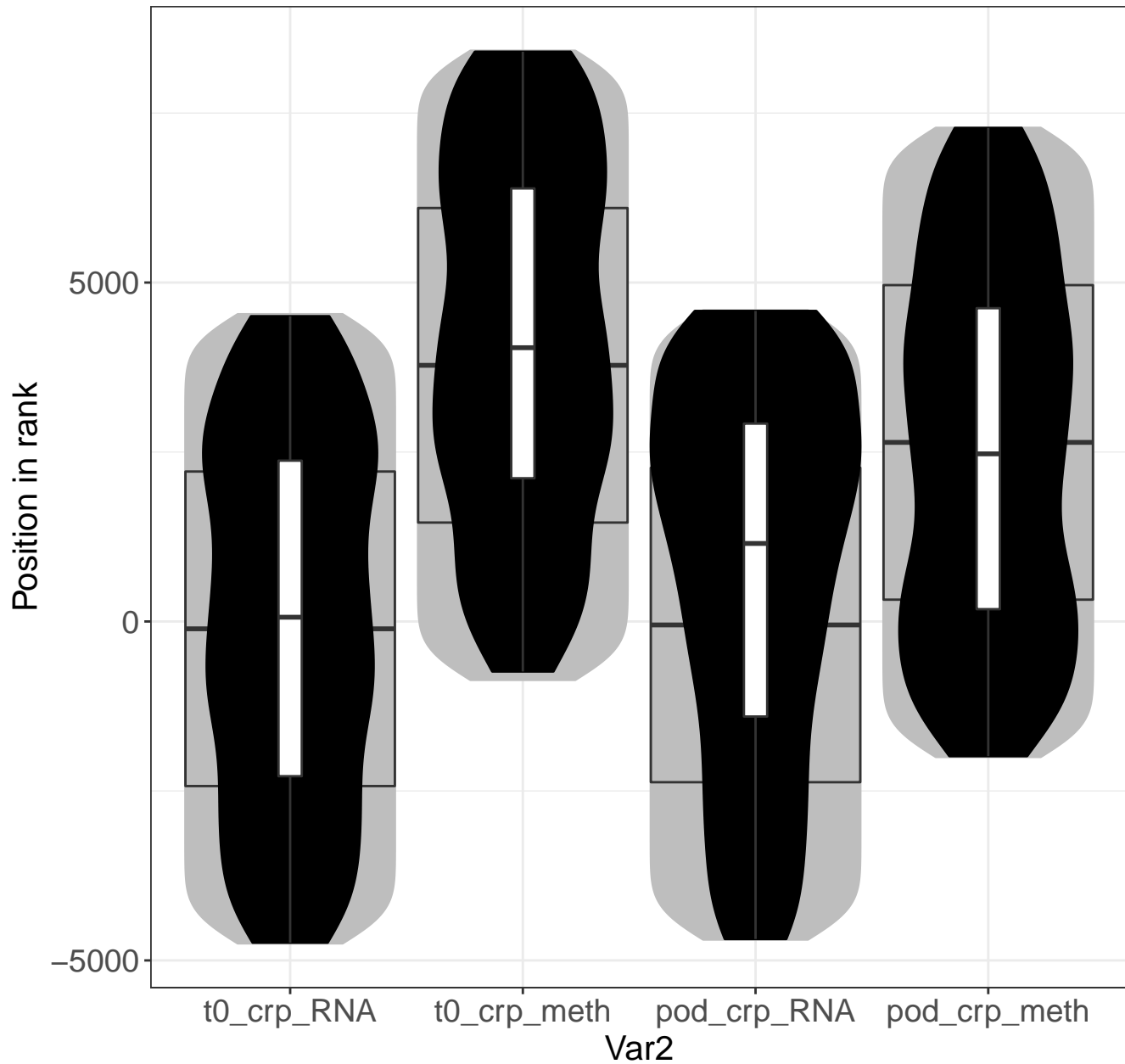
Metabolism of lipids



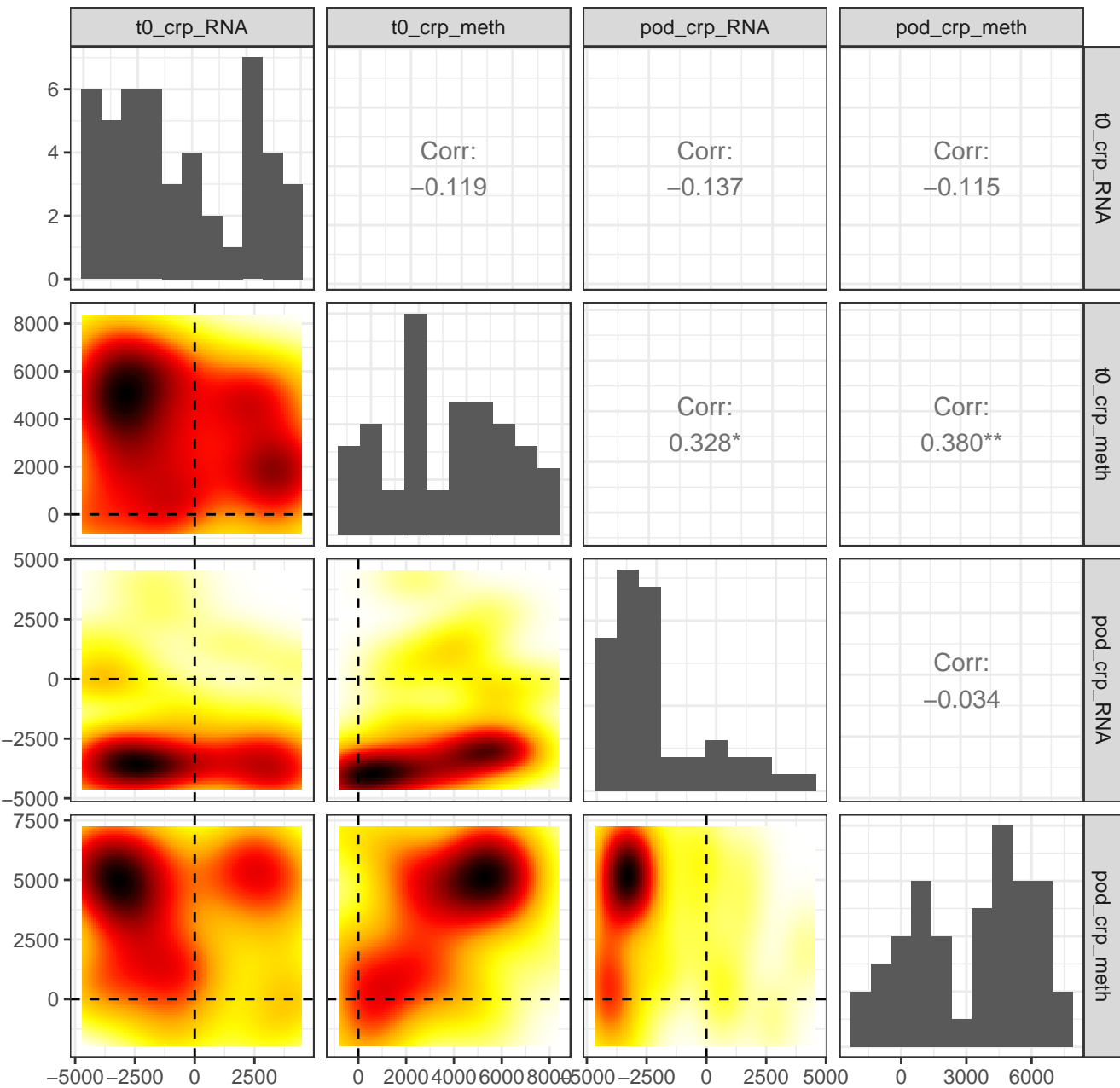
Metabolism of lipids



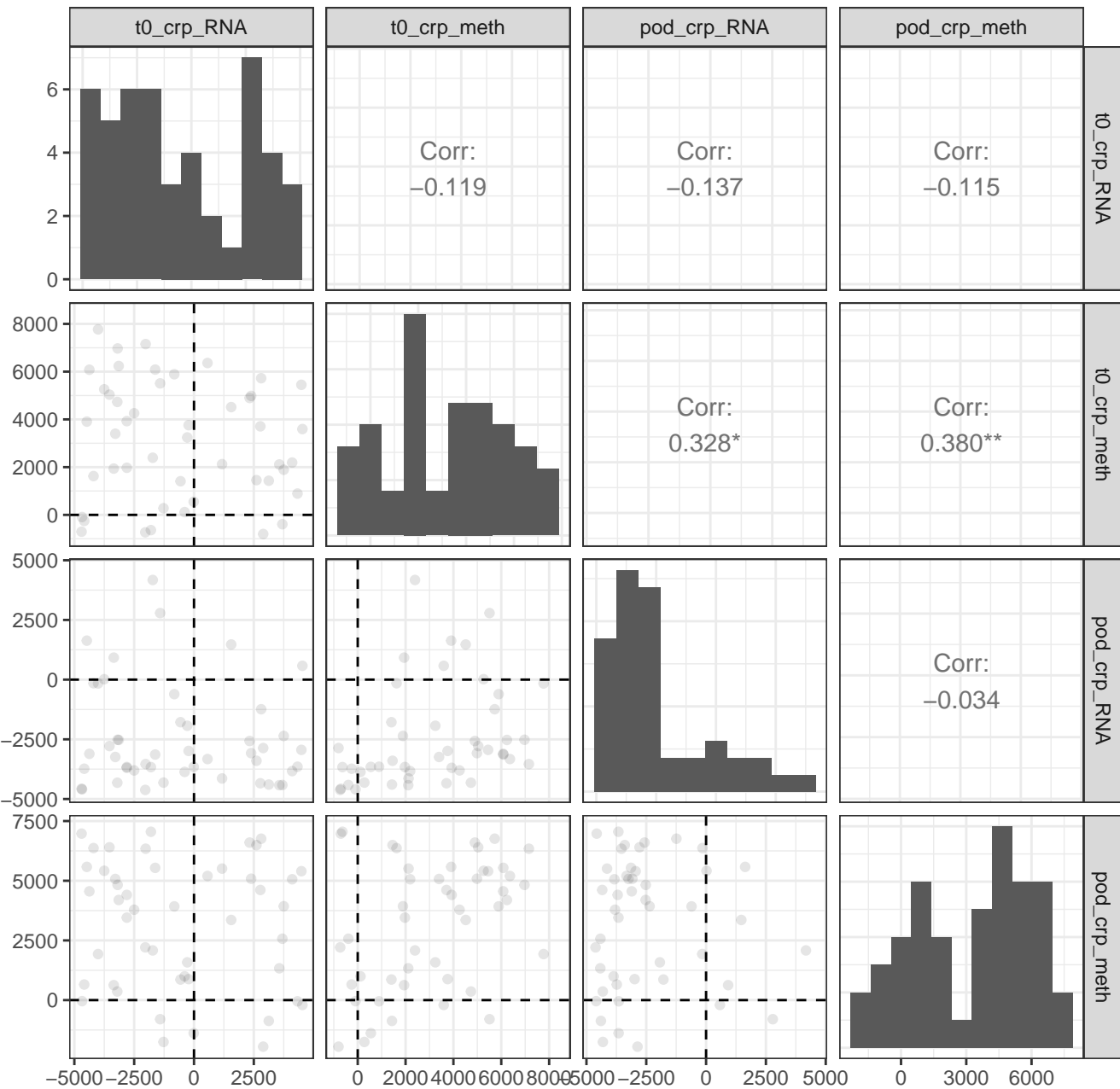
Metabolism of lipids



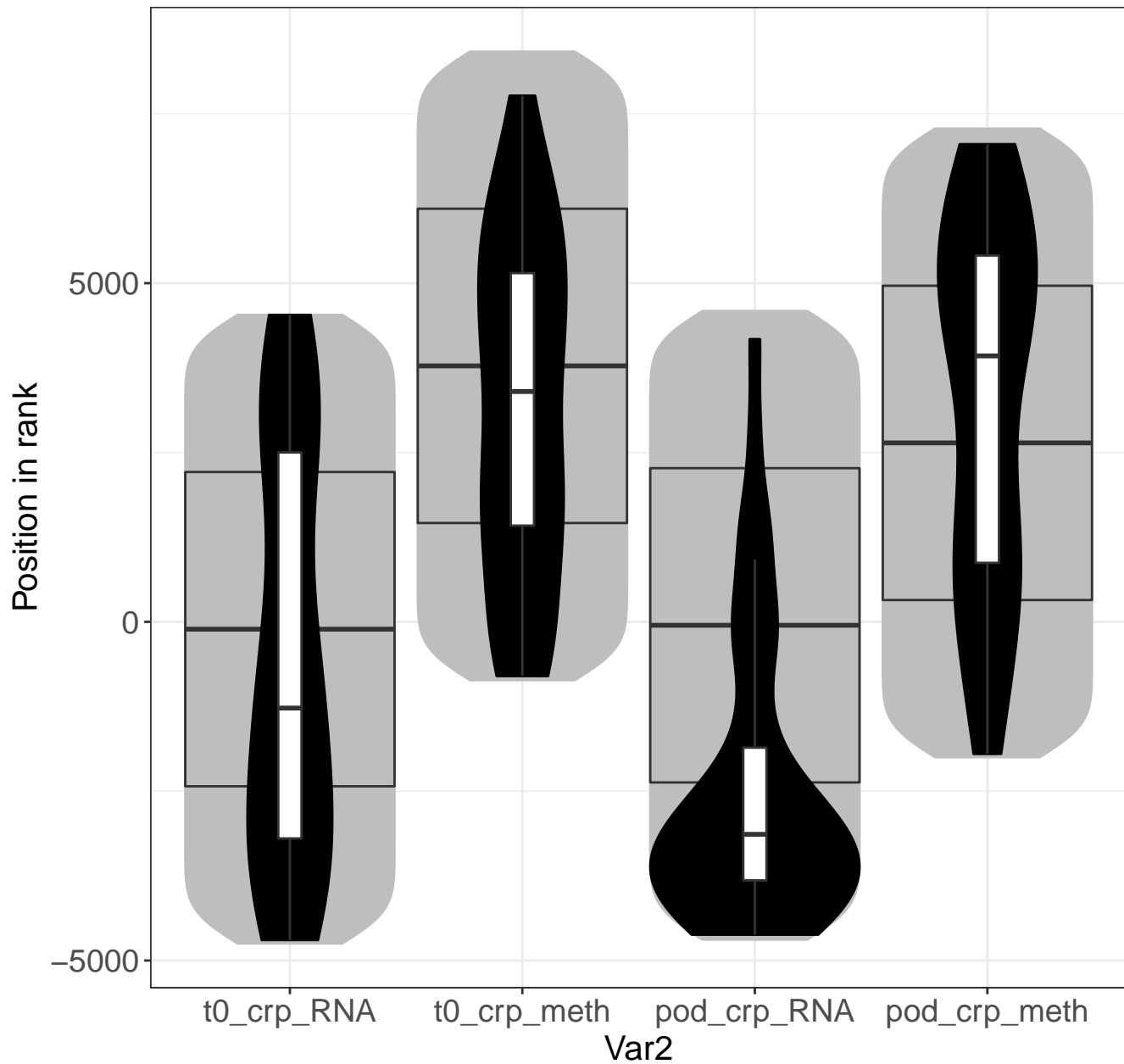
rRNA modification in the nucleus and cytosol



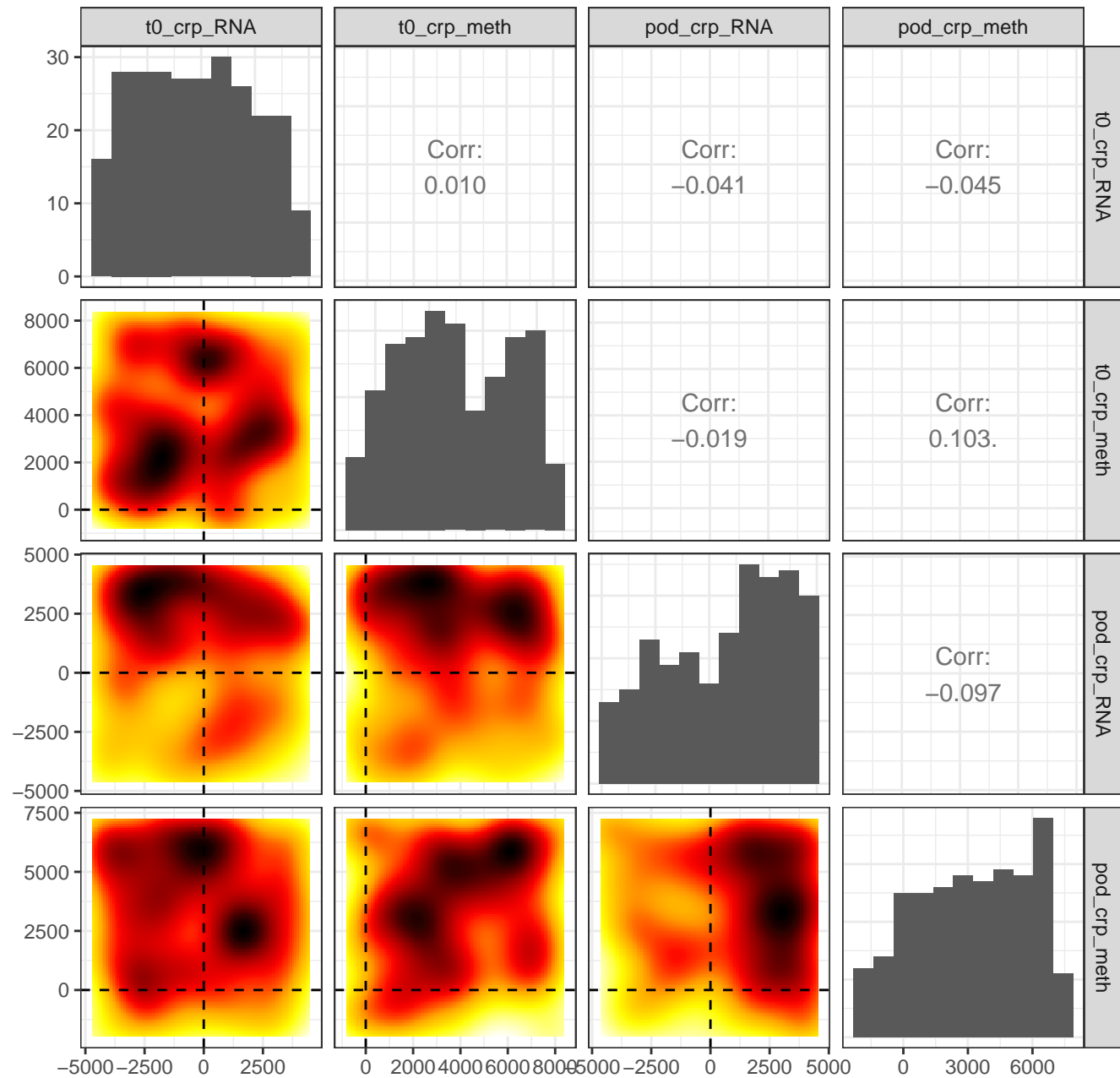
rRNA modification in the nucleus and cytosol



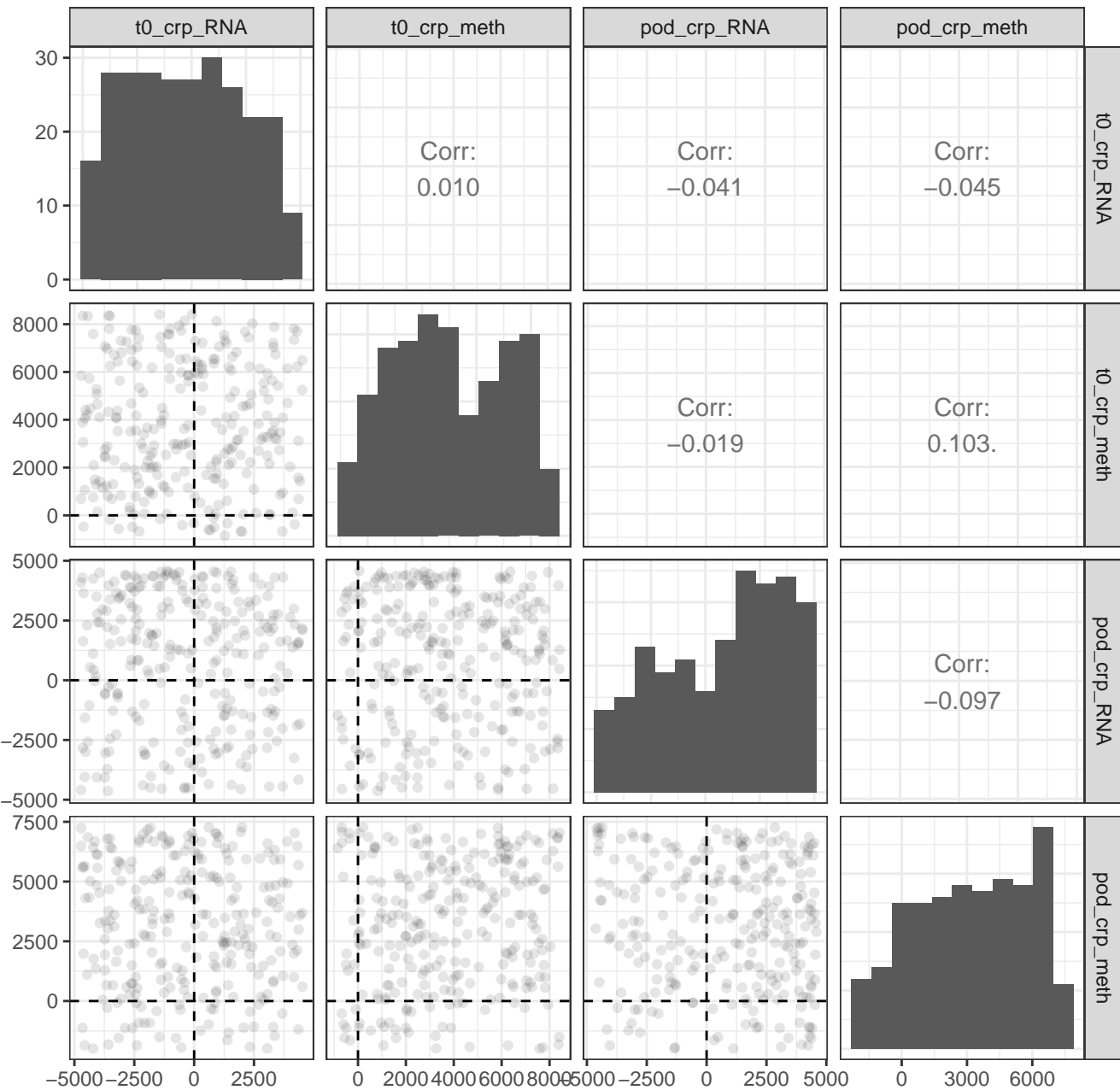
rRNA modification in the nucleus and cytosol



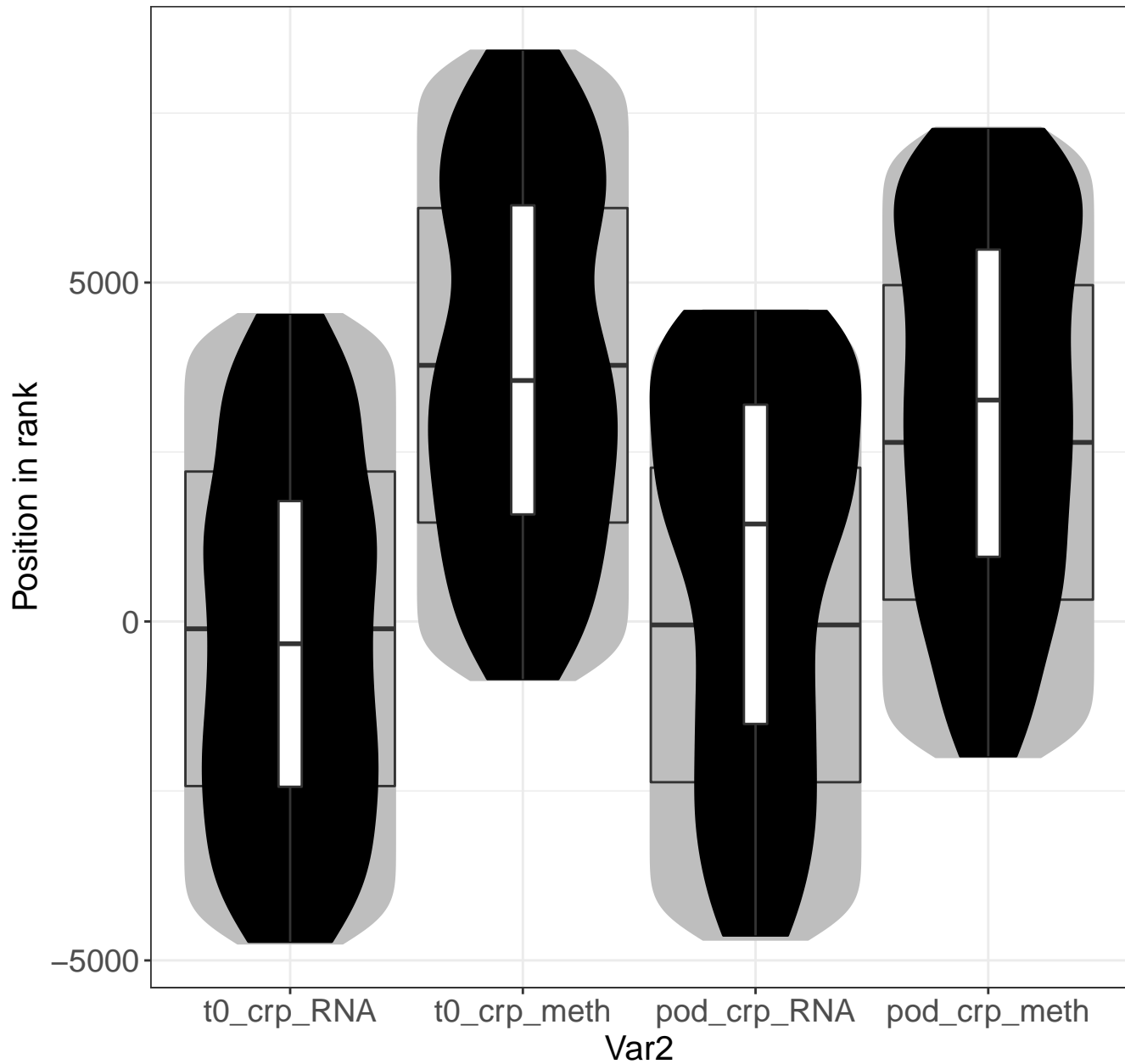
Signaling by Interleukins



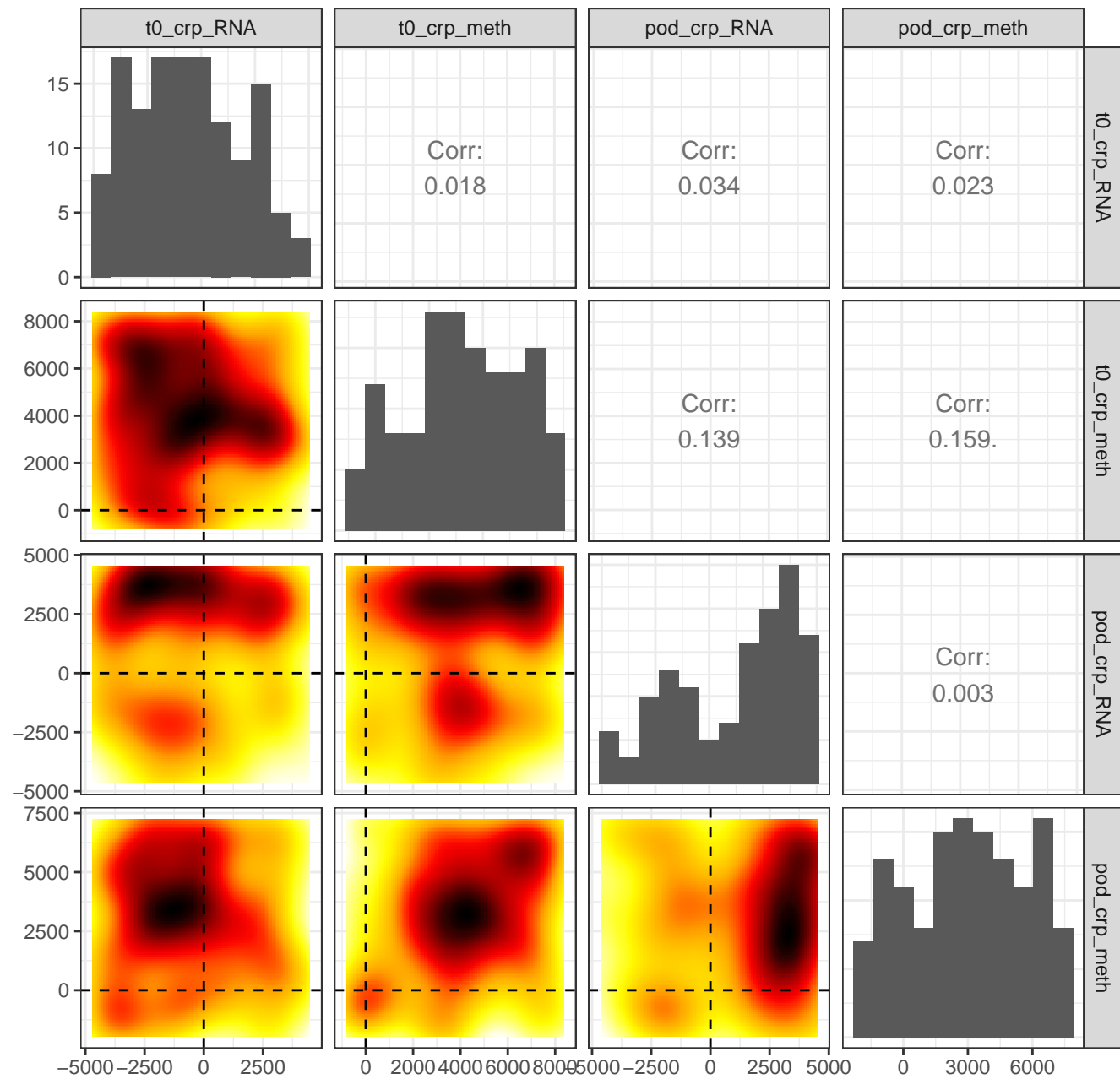
Signaling by Interleukins



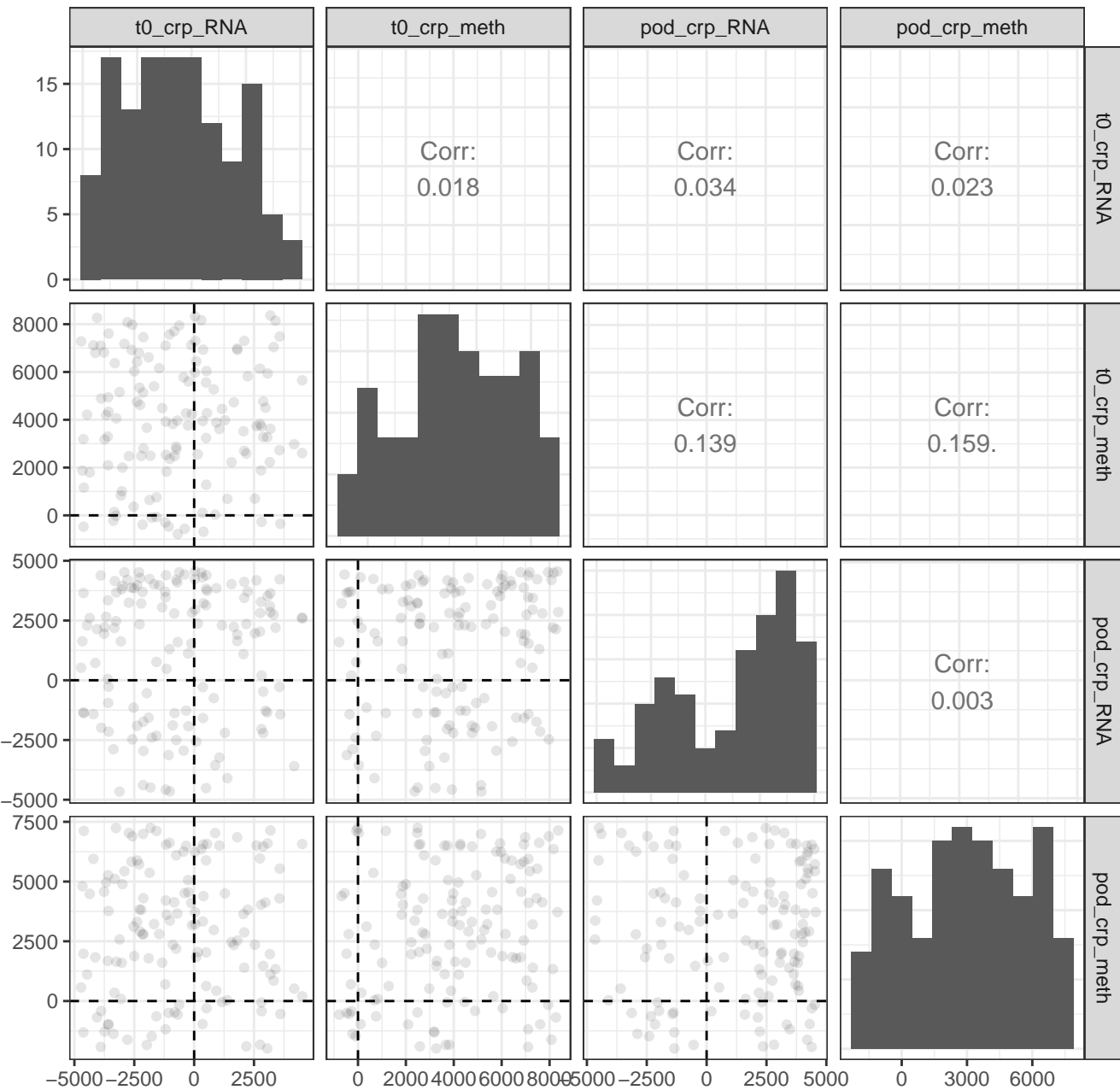
Signaling by Interleukins



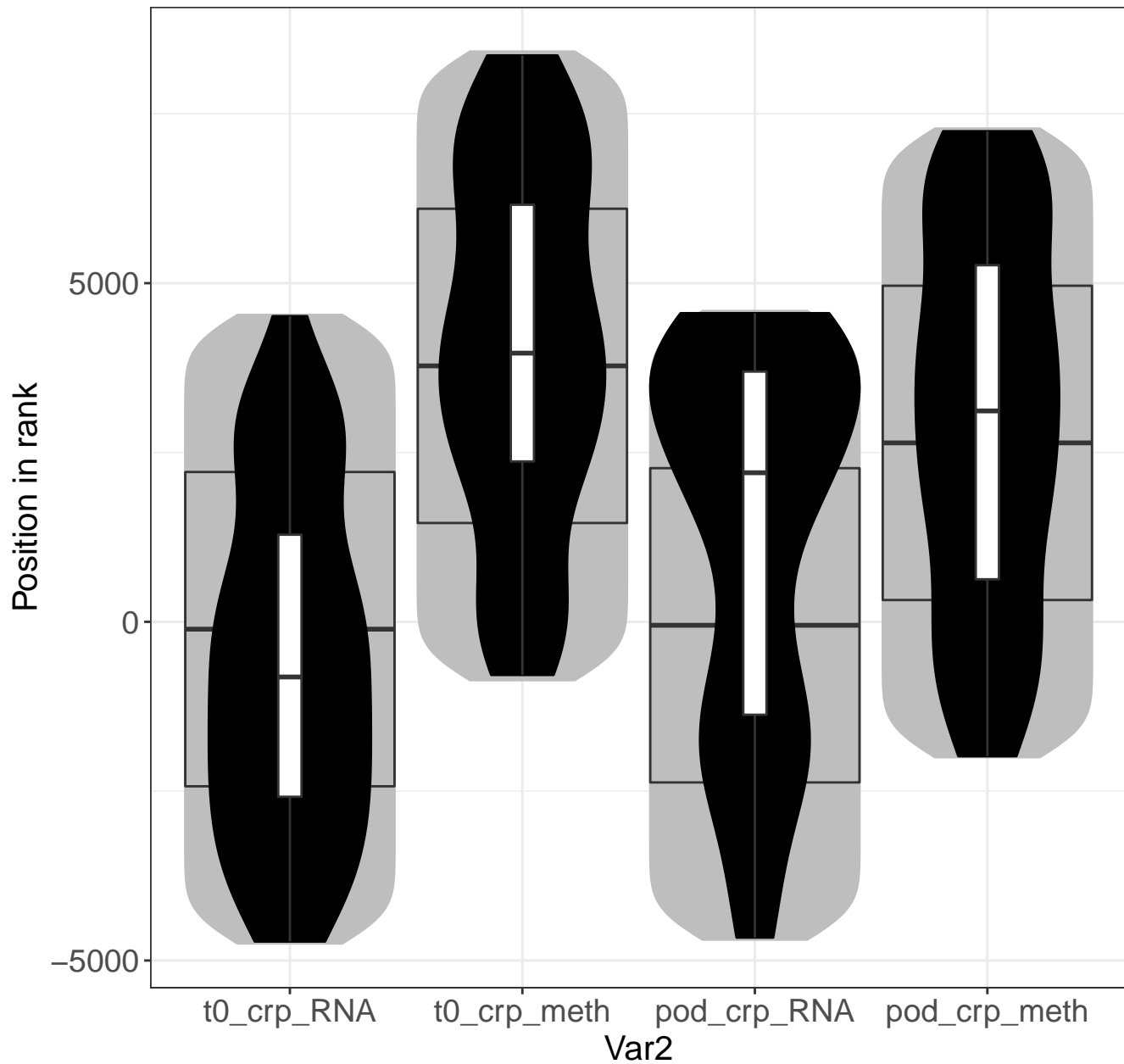
Platelet activation, signaling and aggregation



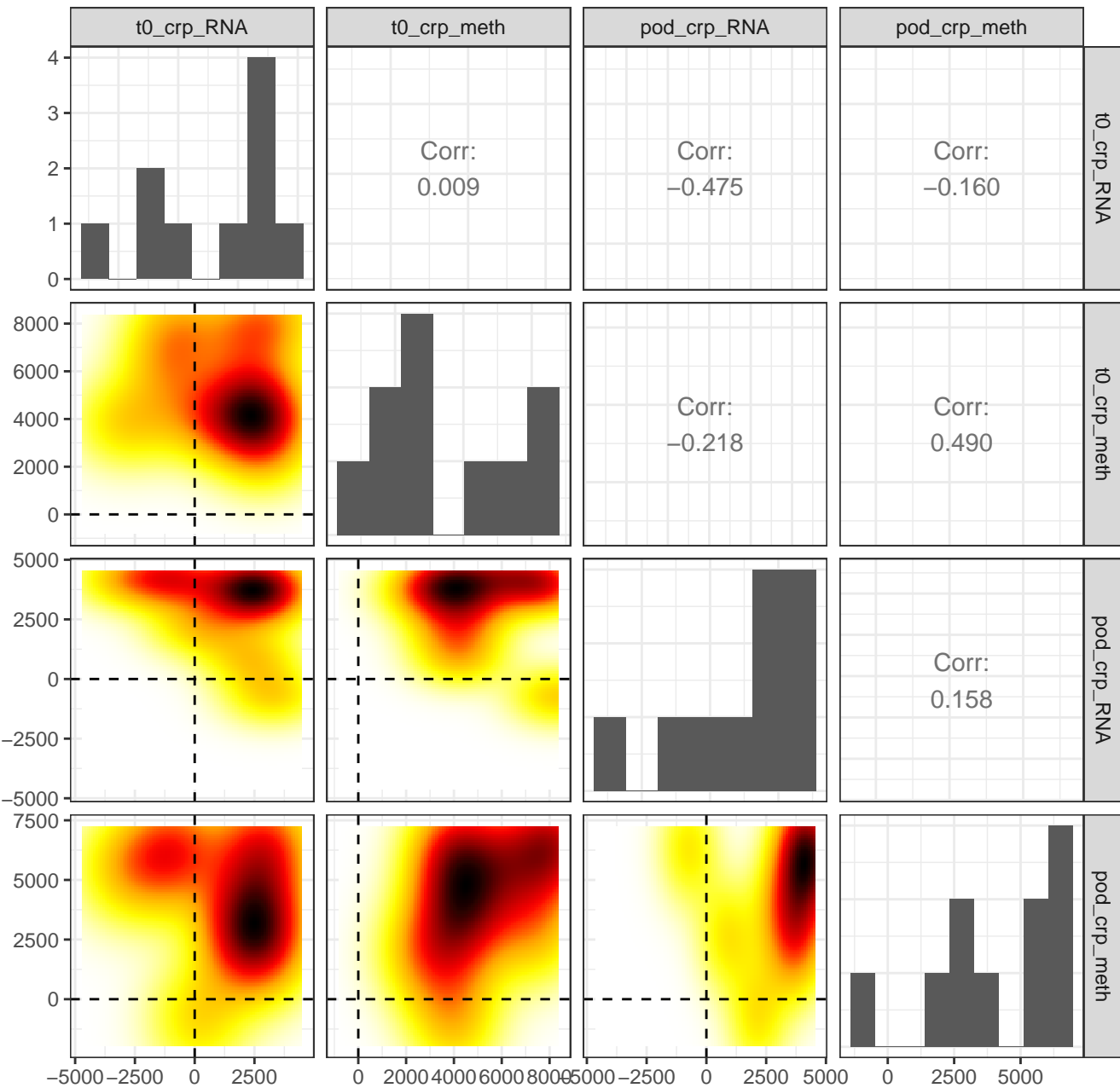
Platelet activation, signaling and aggregation



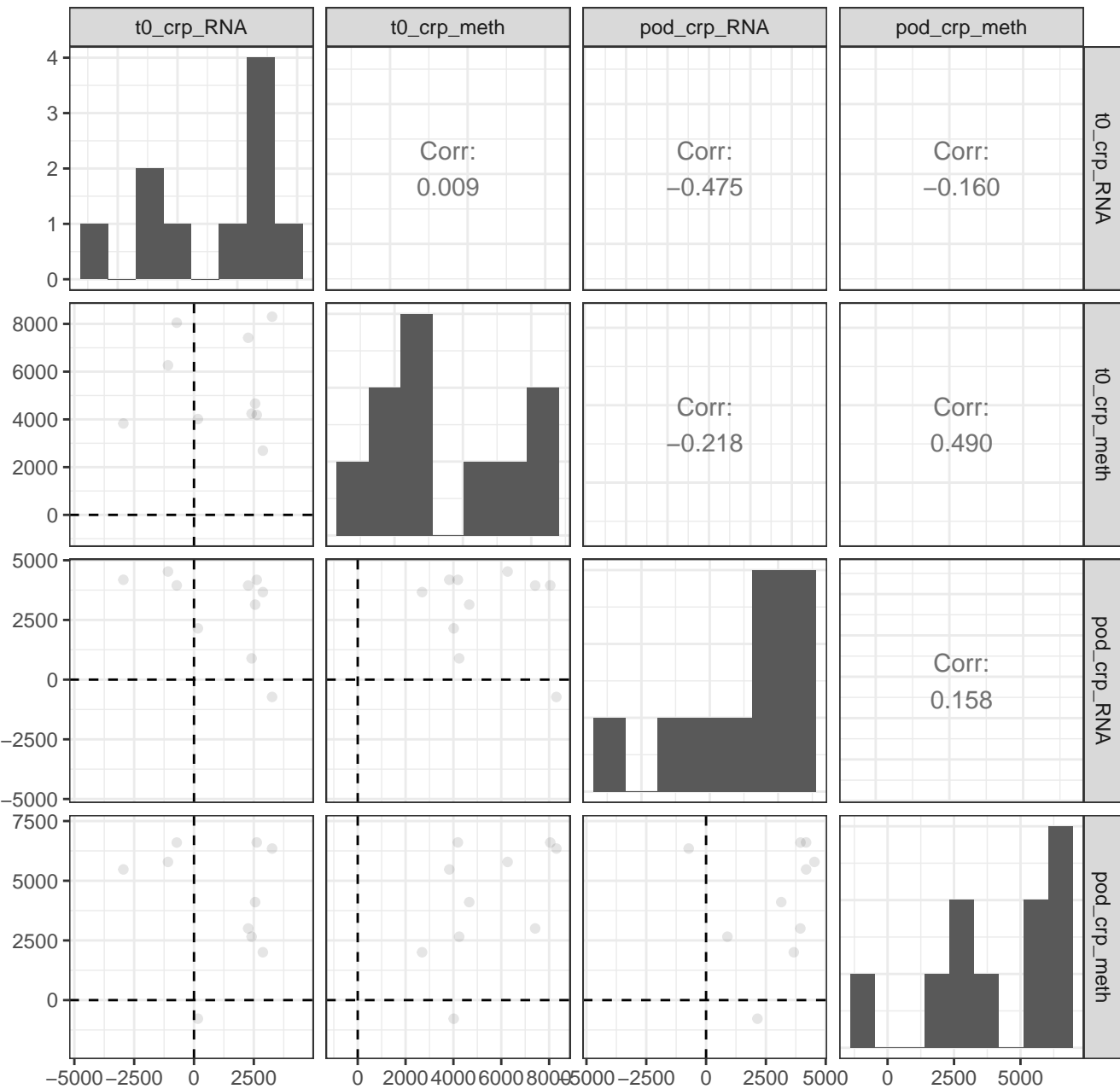
Platelet activation, signaling and aggregation



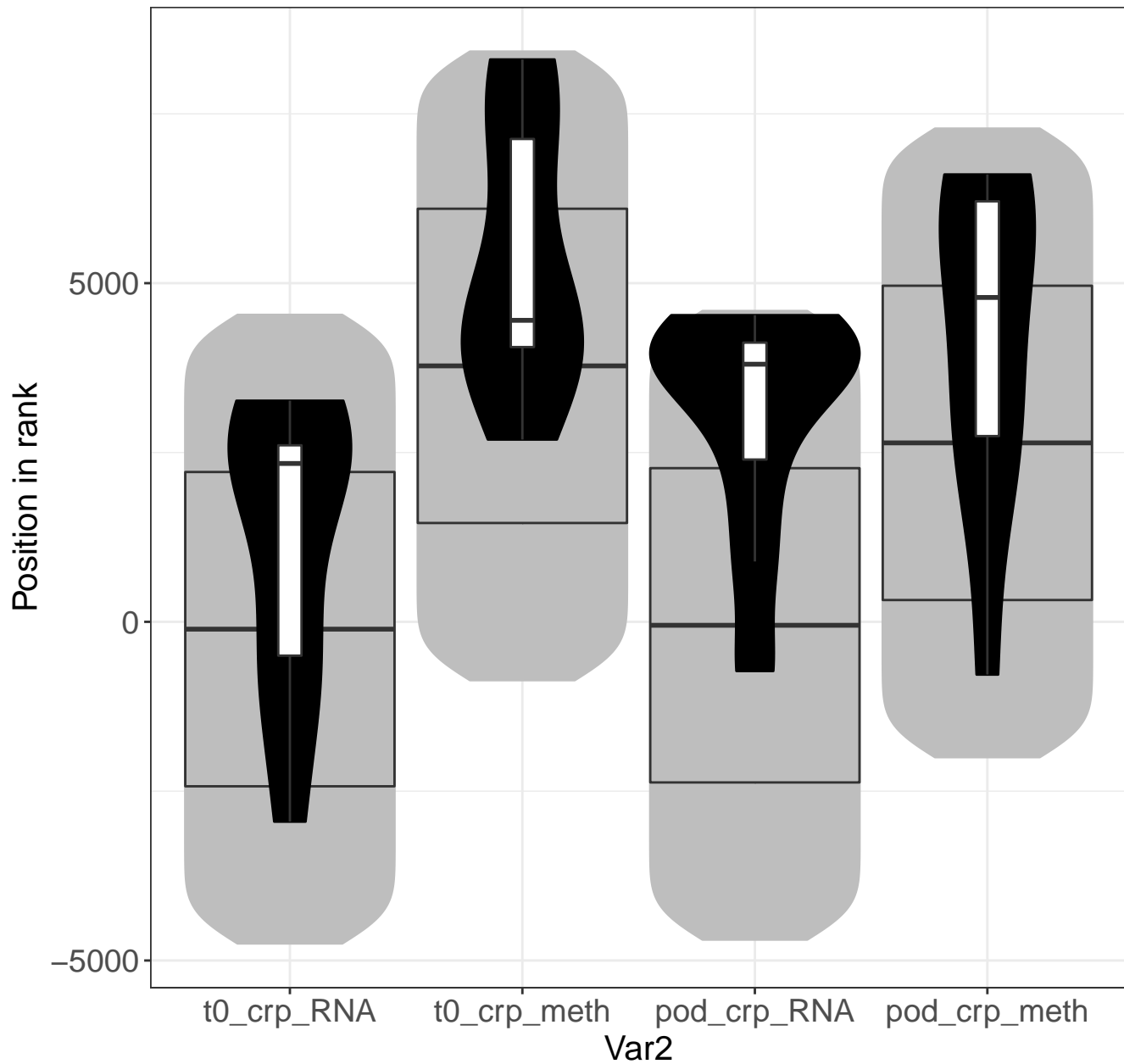
alpha-linolenic acid (ALA) metabolism



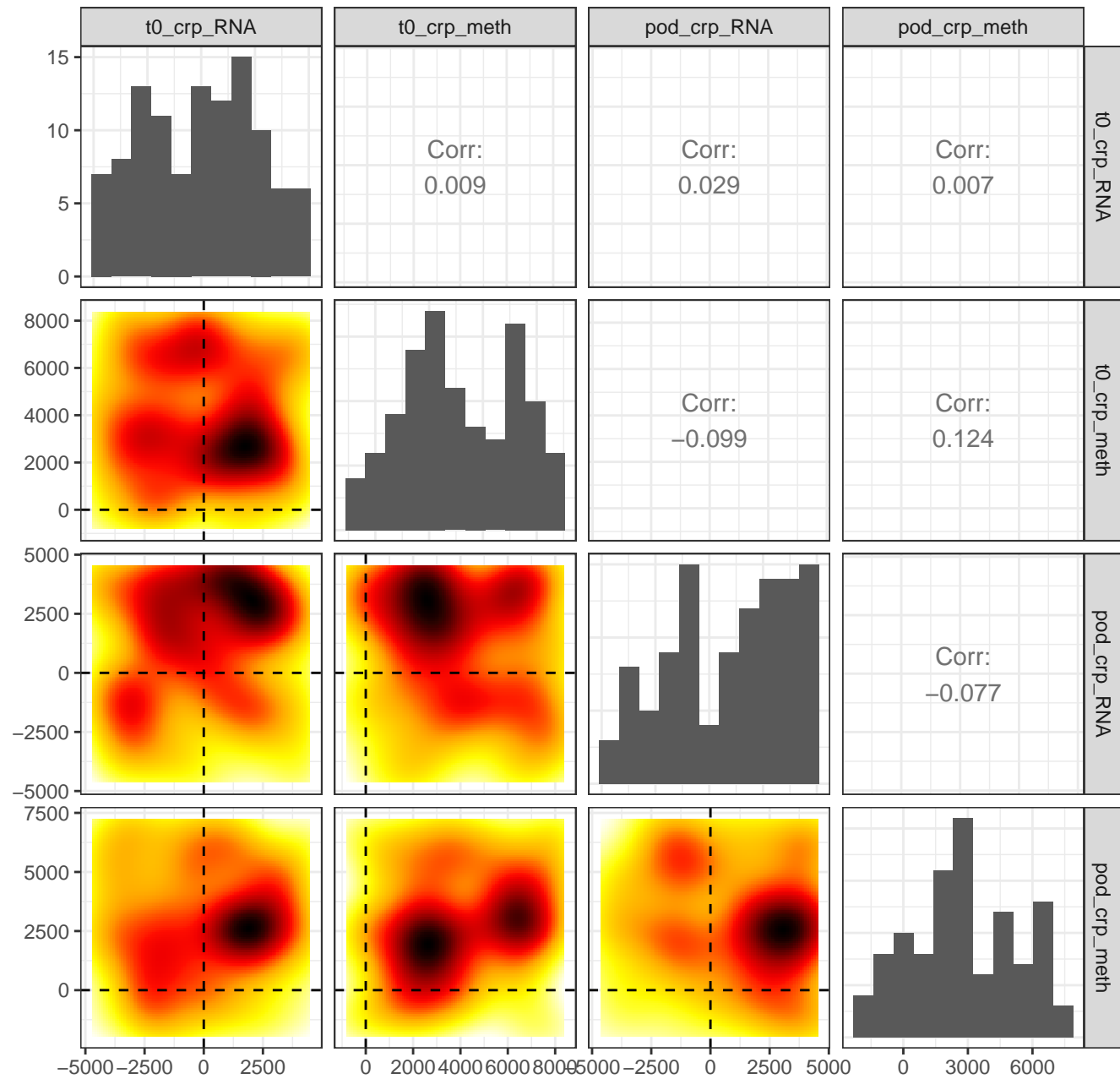
alpha-linolenic acid (ALA) metabolism



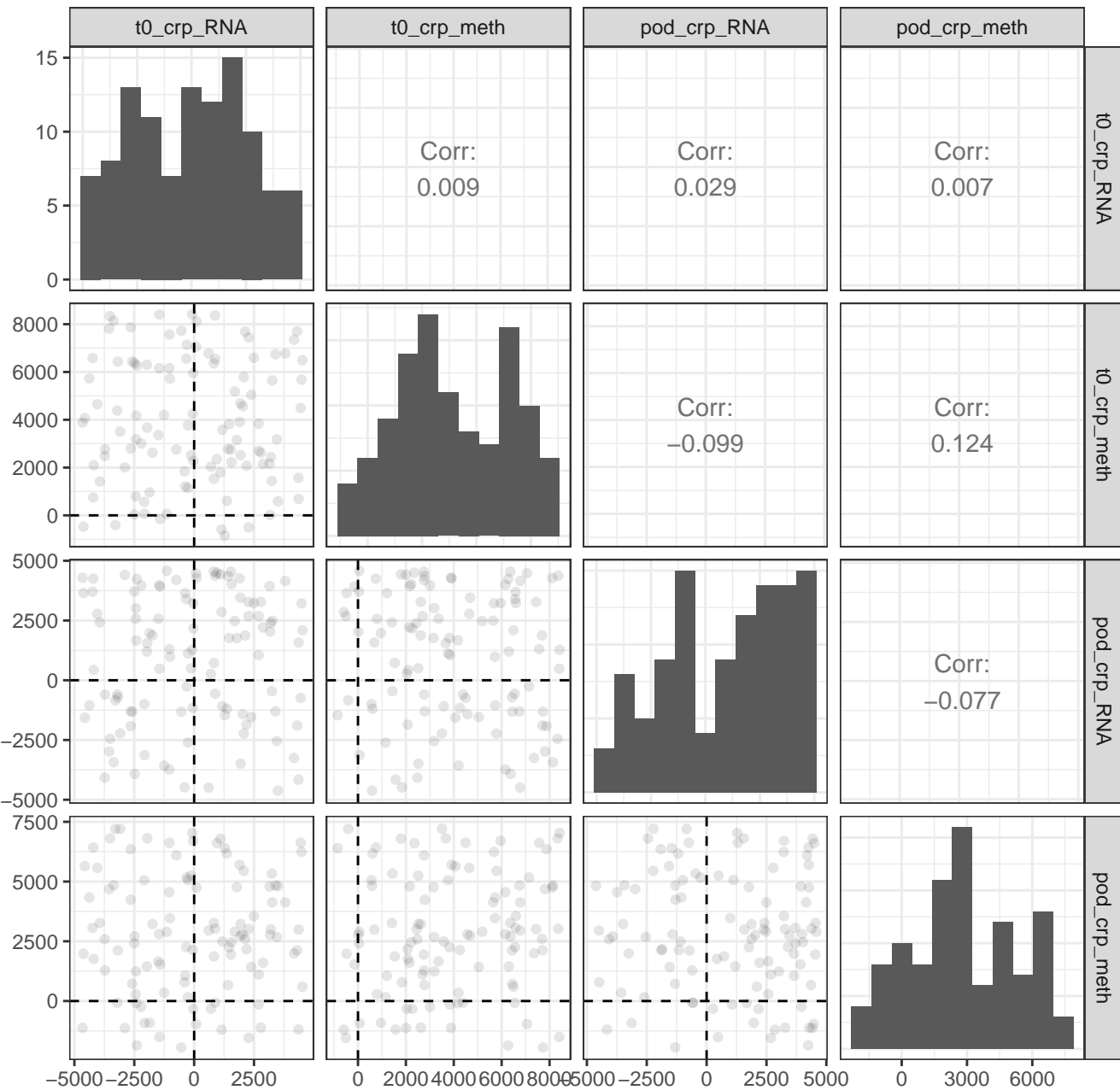
alpha-linolenic acid (ALA) metabolism



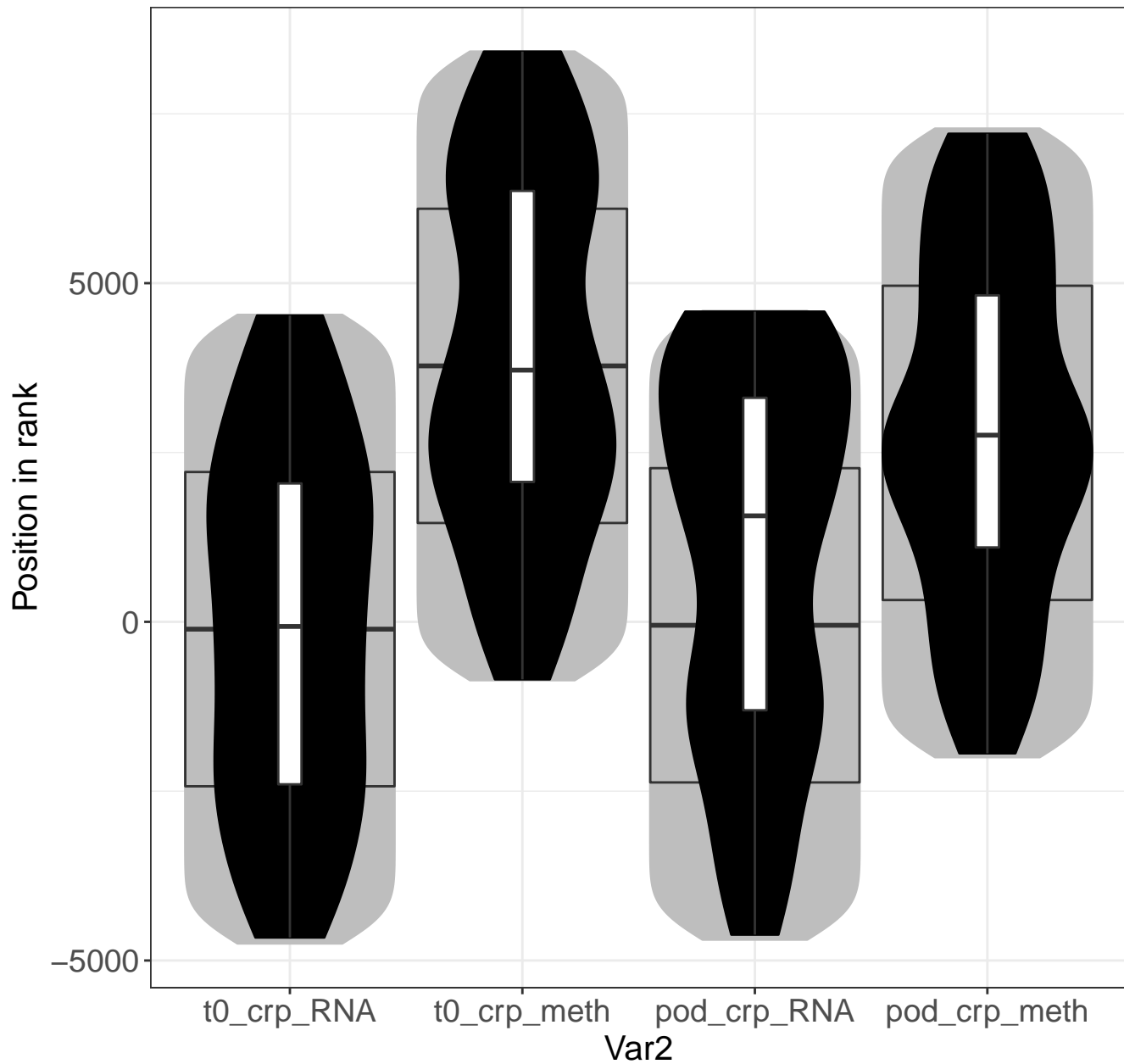
Toll-like Receptor Cascades



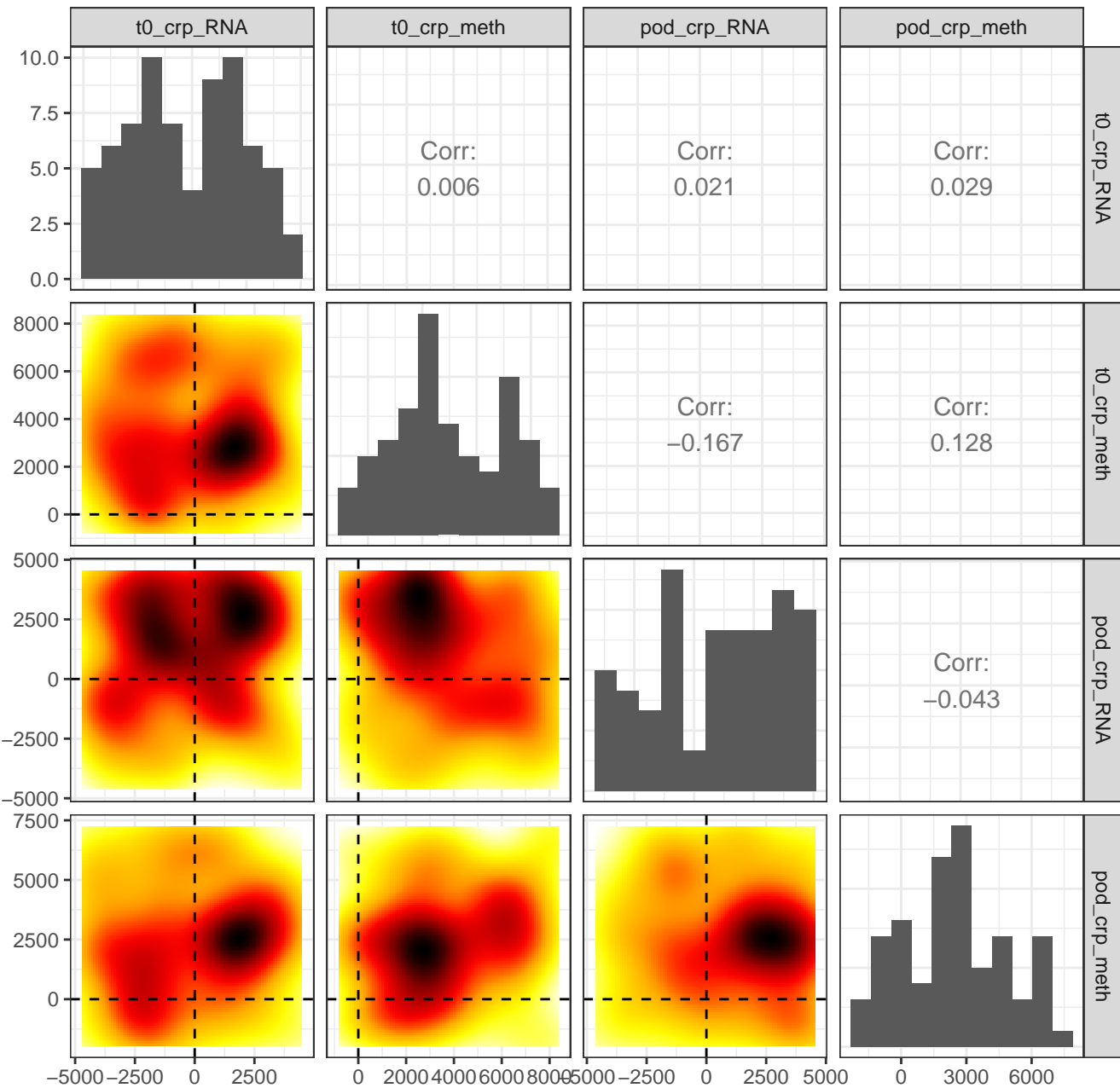
Toll-like Receptor Cascades



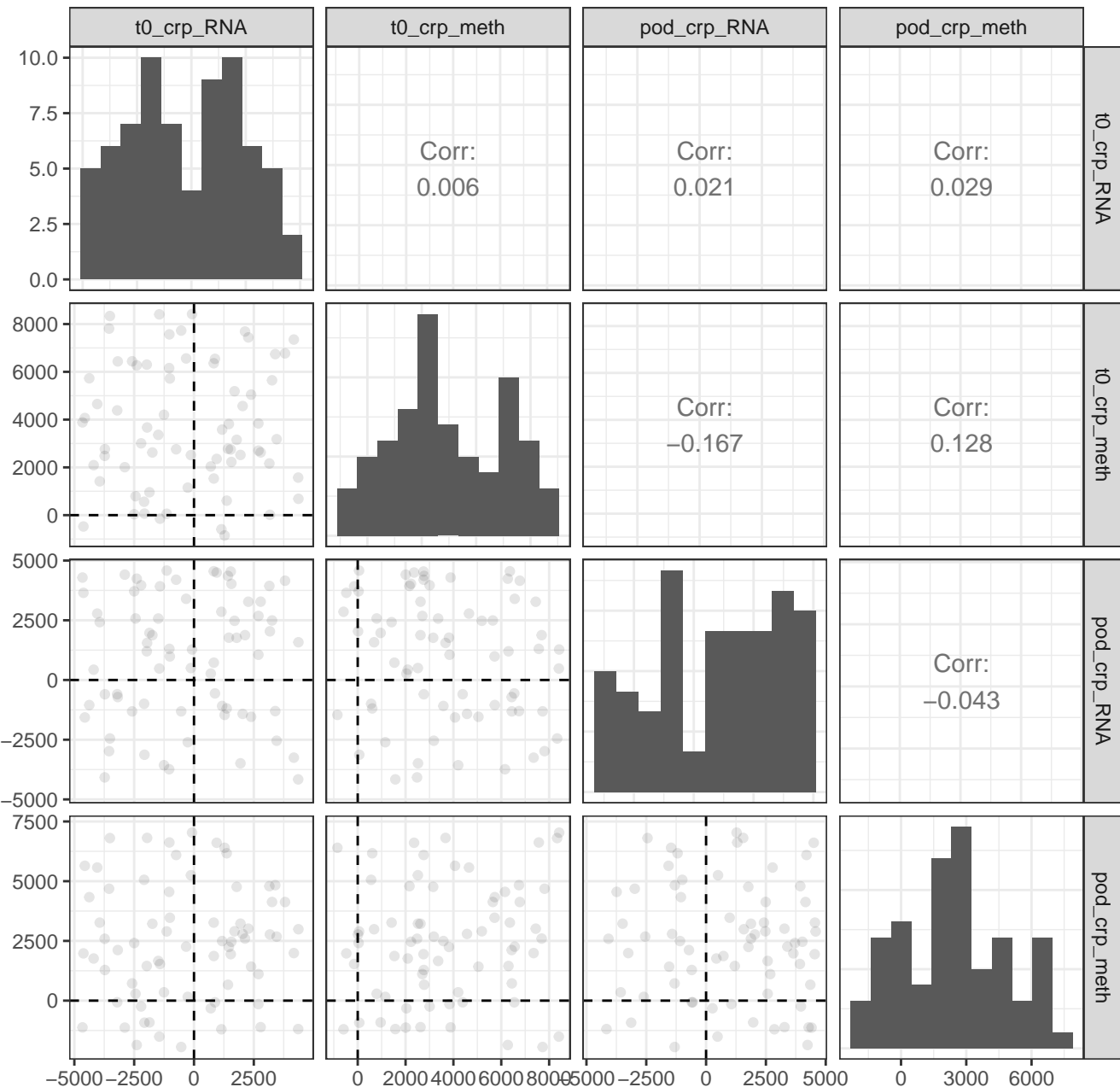
Toll-like Receptor Cascades



Toll Like Receptor 2 (TLR2) Cascade



Toll Like Receptor 2 (TLR2) Cascade



Toll Like Receptor 2 (TLR2) Cascade

