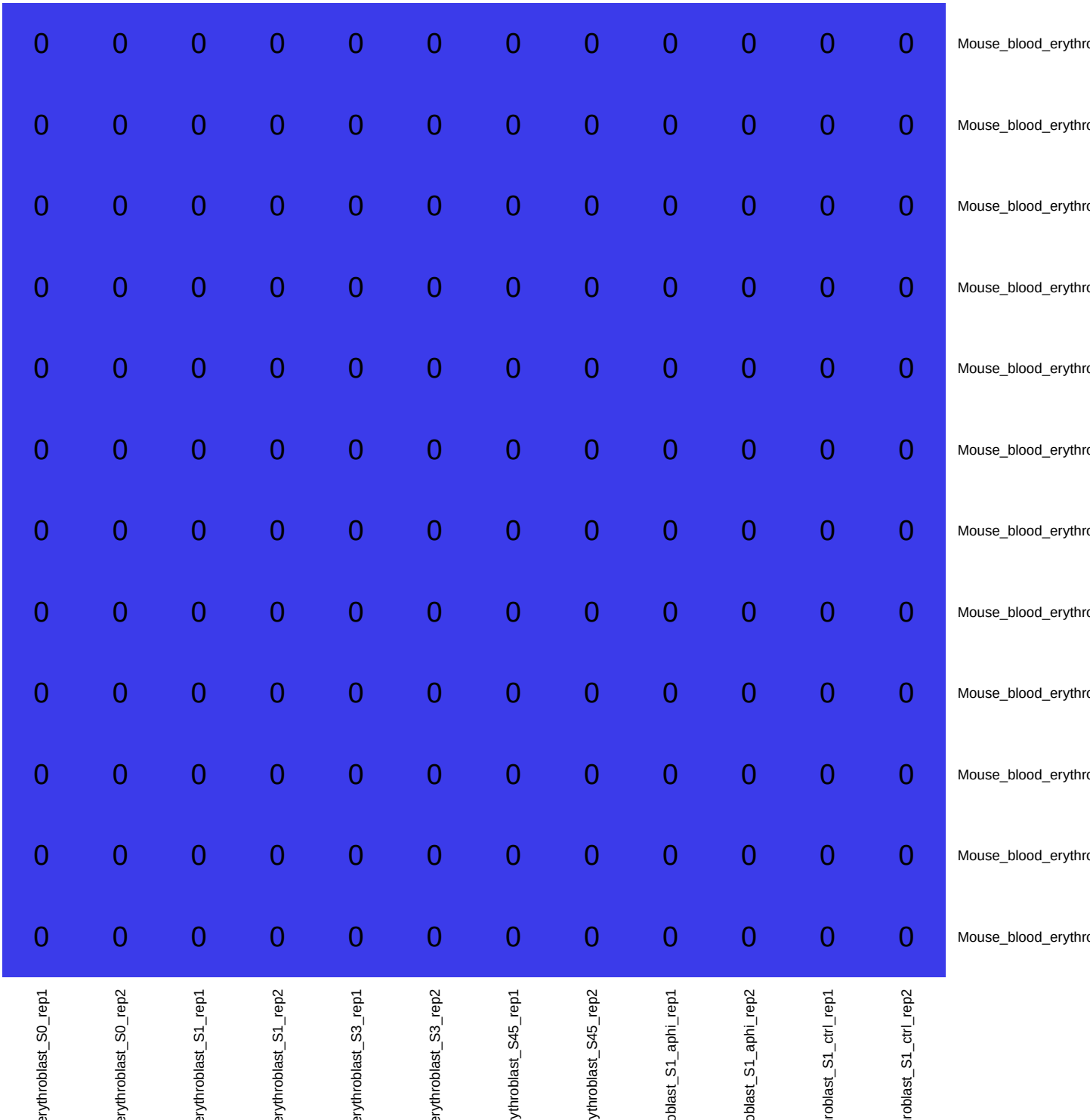
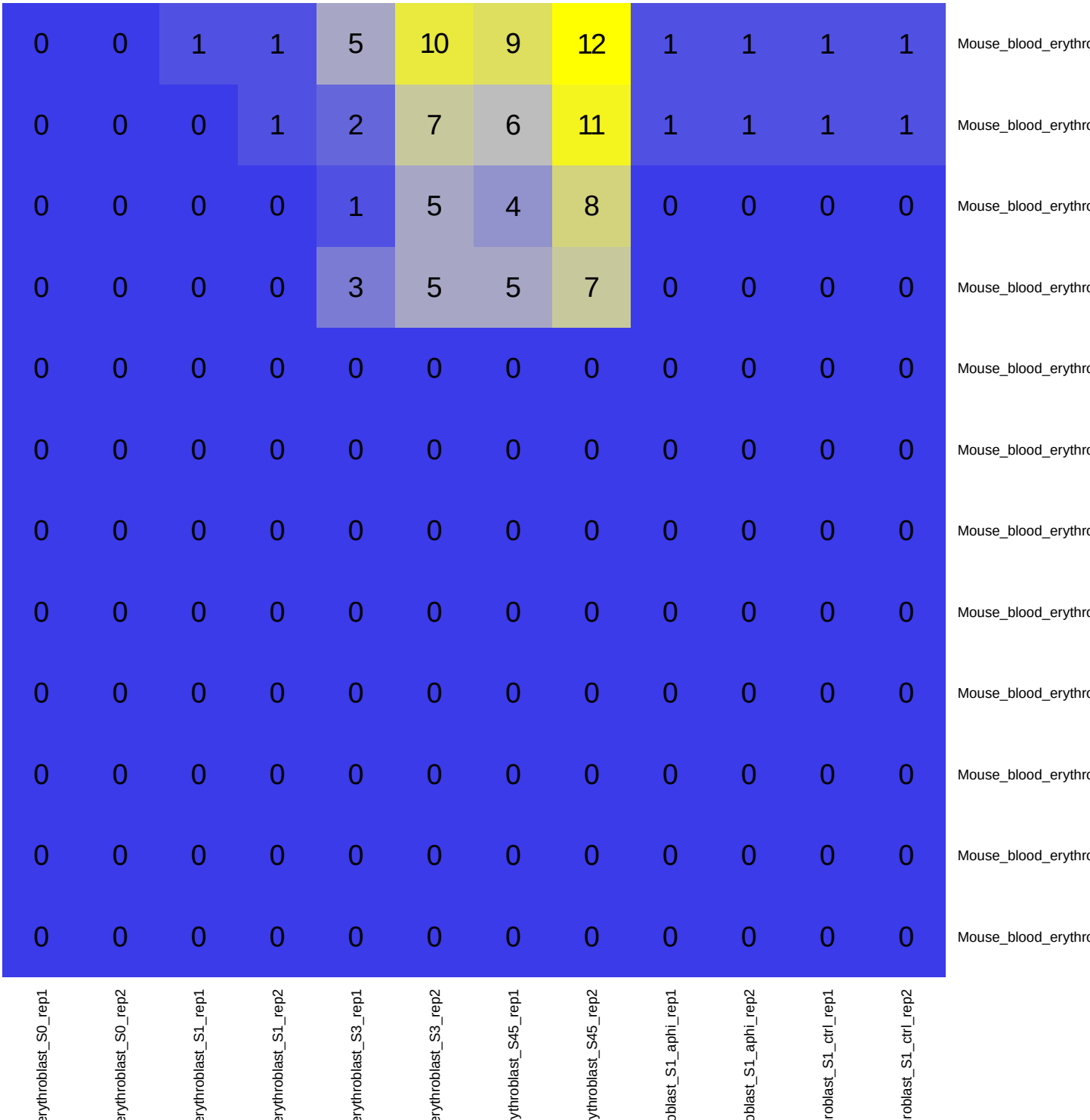
CancerGene_Promoter (#regions: 586) − RRBS regions classified as: stronglyHypermeth

(should be read as <row> is stronglyHypermeth when compared to <column>)



CancerGene_Promoter (#regions: 586) − RRBS regions classified as: hypermethylated

(should be read as <row> is hypermethylated when compared to <column>)



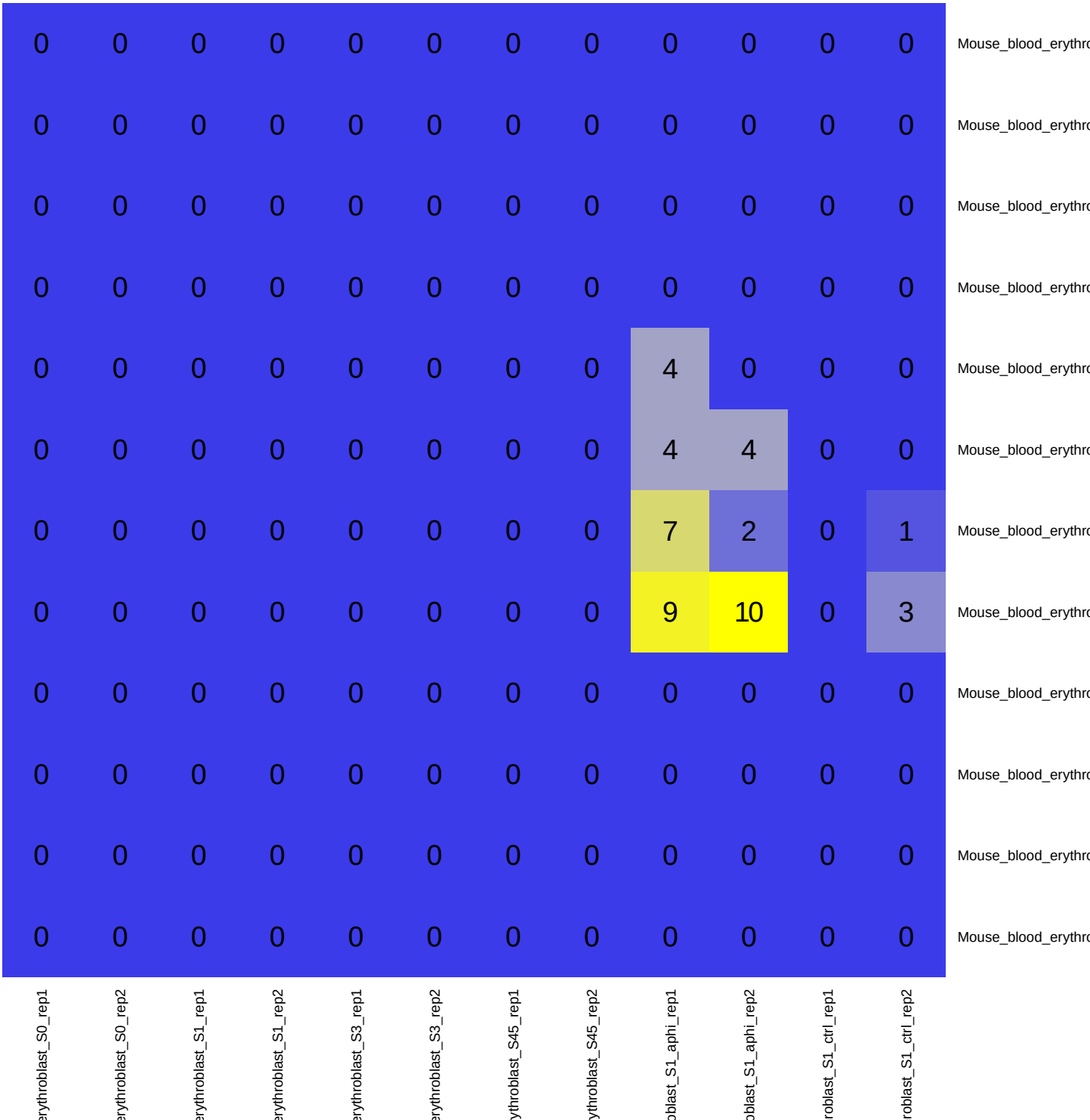
CancerGene_Promoter (#regions: 586) − RRBS regions classified as: insignificant

(should be read as <row> is insignificant when compared to <column>)

0	576	575	577	567	565	561	557	577	577	573	578	Mouse_blood_erythro
0	0	576	577	574	563	570	563	577	577	577	577	Mouse_blood_erythro
0	0	0	579	576	570	572	567	580	580	580	582	Mouse_blood_erythro
0	0	0	0	575	572	576	571	581	581	581	581	Mouse_blood_erythro
0	0	0	0	0	577	580	580	574	580	580	580	Mouse_blood_erythro
0	0	0	0	0	0	580	579	572	572	578	576	Mouse_blood_erythro
0	0	0	0	0	0	0	582	573	571	581	581	Mouse_blood_erythro
0	0	0	0	0	0	0	0	571	565	579	576	Mouse_blood_erythro
0	0	0	0	0	0	0	0	0	583	584	584	Mouse_blood_erythro
0	0	0	0	0	0	0	0	0	0	583	583	Mouse_blood_erythro
0	0	0	0	0	0	0	0	0	0	0	584	Mouse_blood_erythro
0	0	0	0	0	0	0	0	0	0	0	0	Mouse_blood_erythro
erythroblast_S0_rep1	erythroblast_S0_rep2	erythroblast_S1_rep1	erythroblast_S1_rep2	erythroblast_S3_rep1	erythroblast_S3_rep2	ythroblast_S45_rep1	ythroblast_S45_rep2	oblast_S1_aphi_rep1	oblast_S1_aphi_rep2	roblast_S1_ctrl_rep1	roblast_S1_ctrl_rep2	

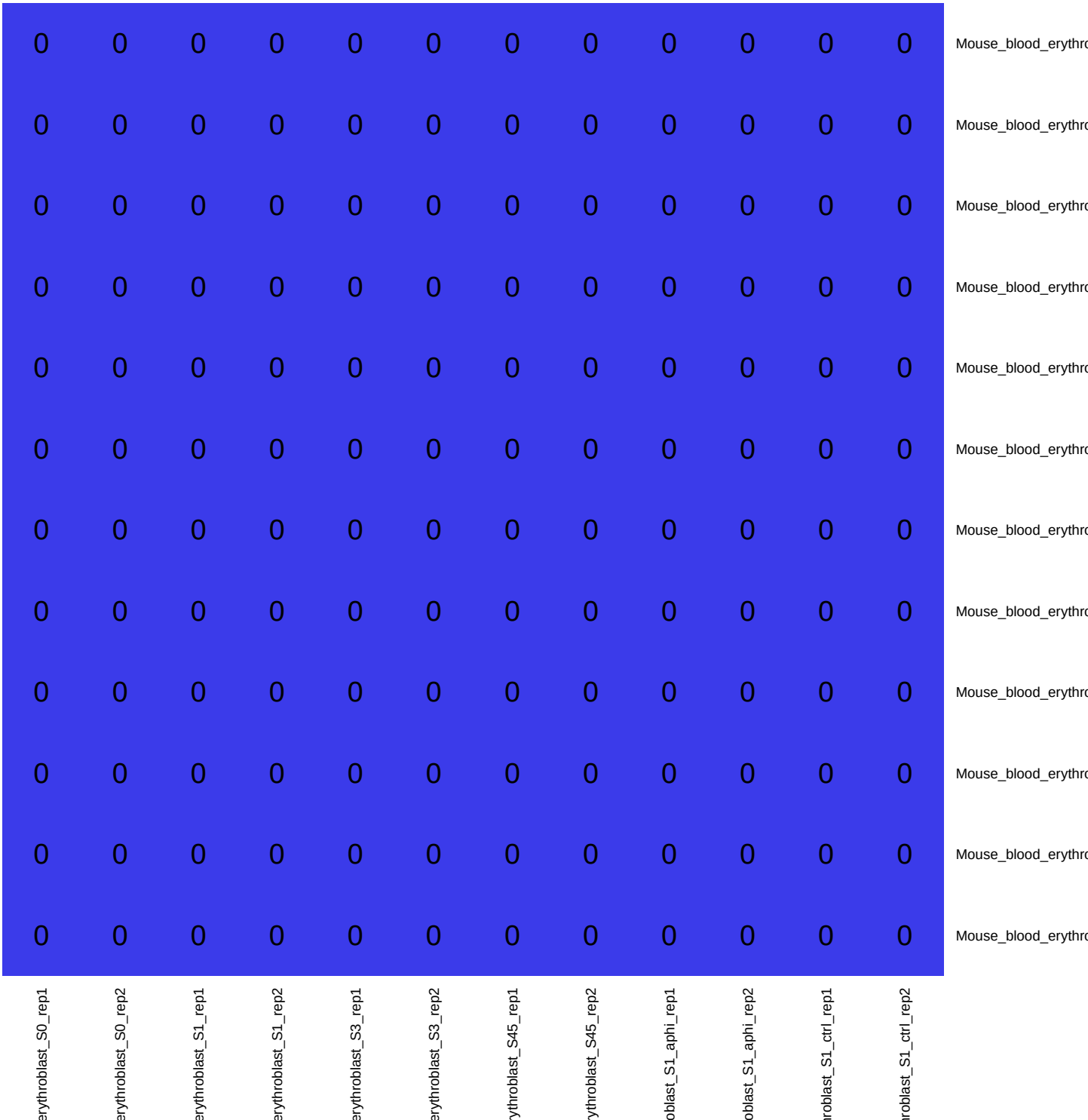
CancerGene_Promoter (#regions: 586) − RRBS regions classified as: hypomethylated

(should be read as <row> is hypomethylated when compared to <column>)



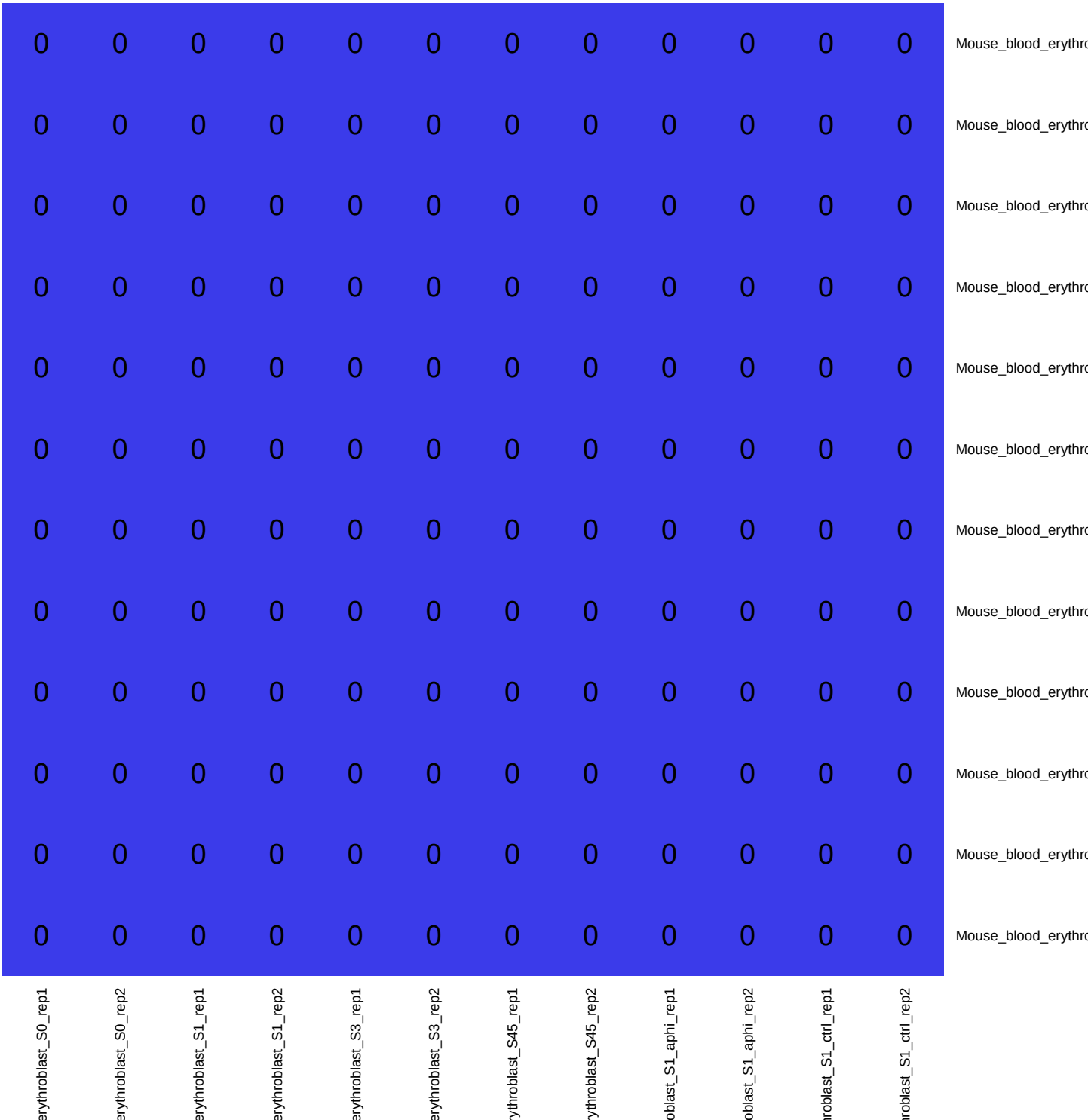
CancerGene_Promoter (#regions: 586) − RRBS regions classified as: stronglyHypometh

(should be read as <row> is stronglyHypometh when compared to <column>)



CancerGene_Promoter (#regions: 586) − RRBS regions classified as: inconclusive

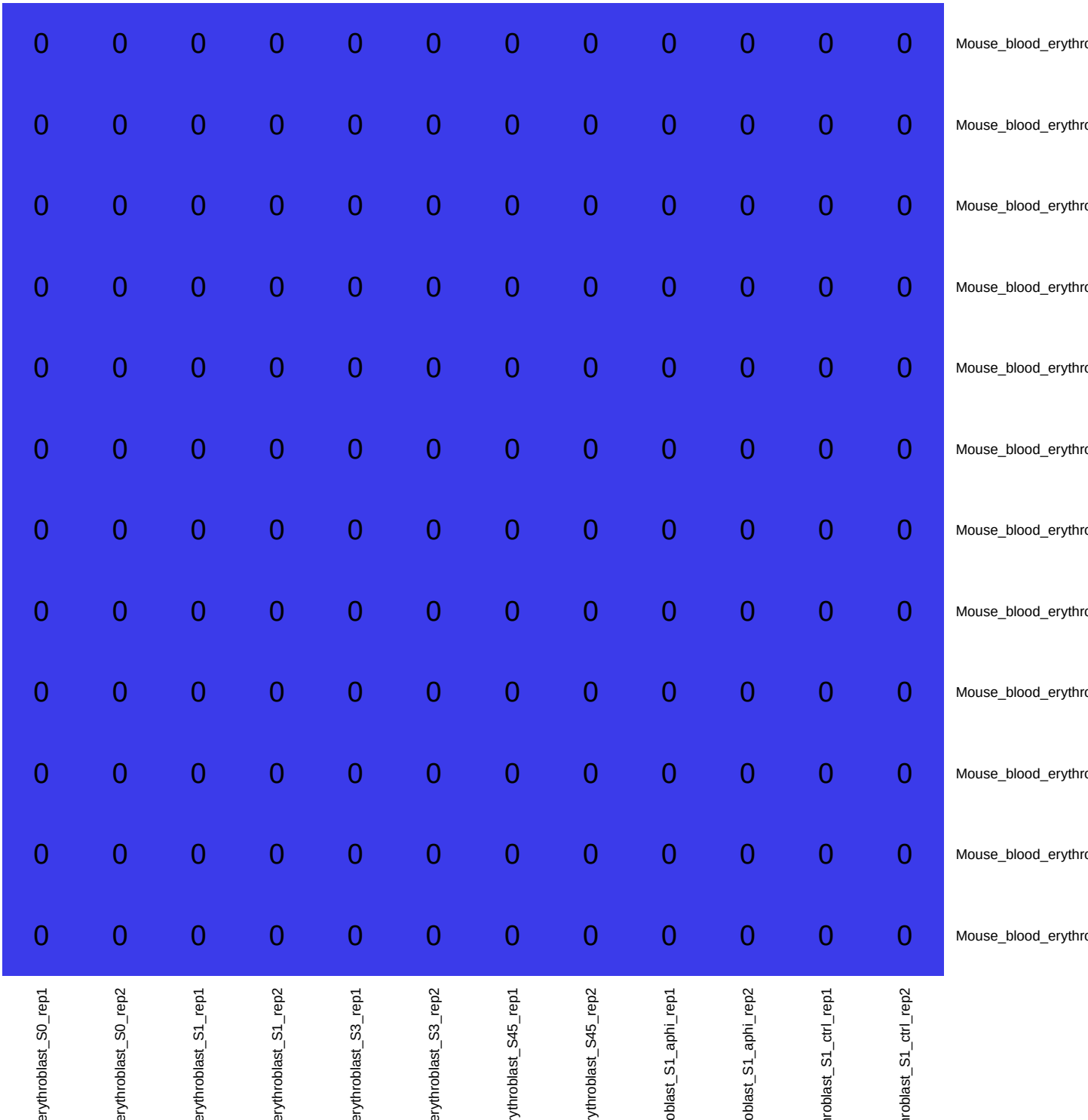
(should be read as <row> is inconclusive when compared to <column>)



CancerGene_Promoter (#regions: 586) − RRBS regions classified as: lowCoverage

(should be read as <row> is lowCoverage when compared to <column>)

	erythroblast_S0_rep1	erythroblast_S0_rep2	erythroblast_S1_rep1	erythroblast_S1_rep2	erythroblast_S3_rep1	erythroblast_S3_rep2	ythroblast_S45_rep1	ythroblast_S45_rep2	oblast_S1_aphi_rep1	oblast_S1_aphi_rep2	roblast_S1_ctrl_rep1	roblast_S1_ctrl_rep2
Mouse_blood_erythro	0	10	9	8	9	7	8	8	8	8	8	7
Mouse_blood_erythro	0	0	10	8	9	9	8	8	8	8	8	8
Mouse_blood_erythro	0	0	0	7	9	7	5	6	6	6	6	4
Mouse_blood_erythro	0	0	0	0	8	8	5	5	5	5	5	5
Mouse_blood_erythro	0	0	0	0	0	9	6	6	7	6	6	6
Mouse_blood_erythro	0	0	0	0	0	0	6	7	7	7	7	5
Mouse_blood_erythro	0	0	0	0	0	0	0	4	4	4	4	3
Mouse_blood_erythro	0	0	0	0	0	0	0	0	3	3	3	3
Mouse_blood_erythro	0	0	0	0	0	0	0	0	0	3	2	2
Mouse_blood_erythro	0	0	0	0	0	0	0	0	0	0	3	3
Mouse_blood_erythro	0	0	0	0	0	0	0	0	0	0	0	2
Mouse_blood_erythro	0	0	0	0	0	0	0	0	0	0	0	0

CancerGene_Promoter (#regions: 586) − RRBS regions classified as: noCoverage

(should be read as <row> is noCoverage when compared to <column>)