

Overview of methylation differences for sample replicates

A **All methylation sites**

Sample replicates	# sites $\Delta\beta > 0.1$	# sites $\Delta\beta > 0.2$	# sites $\Delta\beta > 0.3$	# sites $\Delta\beta > 0.4$
MBC3	7,696 (1.635%)	253 (0.054%)	32	8
MBC6	11,573 (2.458%)	304 (0.066%)	32	11
CNOR3	15,616 (3.317%)	307 (0.065%)	22	10
CNOR5	19,347 (4.11%)	1,171 (0.249%)	165	36

B **Unmethylated sites (avg. beta < 0.2 for MBC or CNOR sample groups)**

Sample replicates	# sites $\Delta\beta > 0.1$	# sites $\Delta\beta > 0.2$	# sites $\Delta\beta > 0.3$	# sites $\Delta\beta > 0.4$
MBC3	498 (0.295%)	25 (0.015%)	5	1
MBC6	760 (0.45%)	41 (0.024%)	6	1
CNOR3	718 (0.437%)	12 (0.007%)	3	1
CNOR5	1,820 (1.108%)	106 (0.065%)	19	6