

Supplementary information, Table S11

Table S11 Summary of whole genome bisulfite sequencing (WGSB)

Sample	Library index	Raw reads (M)	Raw bases (G)	Clean reads (M)	Clean bases (G)	Clean Rate (%)	Unique mapped reads (M)	Unique mapped ratio (%)	Bisulfite conversion rate (%)
<i>Crygc</i> ^{-/-} -4 W-SSCs	21	484.9	49.1	460.9	46.4	94.5	137.4	59.68	99.40
Line-HDR 1-SC-8	10	115.01	11.62	112.37	11.35	97.70	37.89	66.97	99.57
Line-HDR 1-SC-8	11	101.5	10.3	99.0	10.0	97.5	32.6	65.63	99.55