

Supplementary information, Figure S3

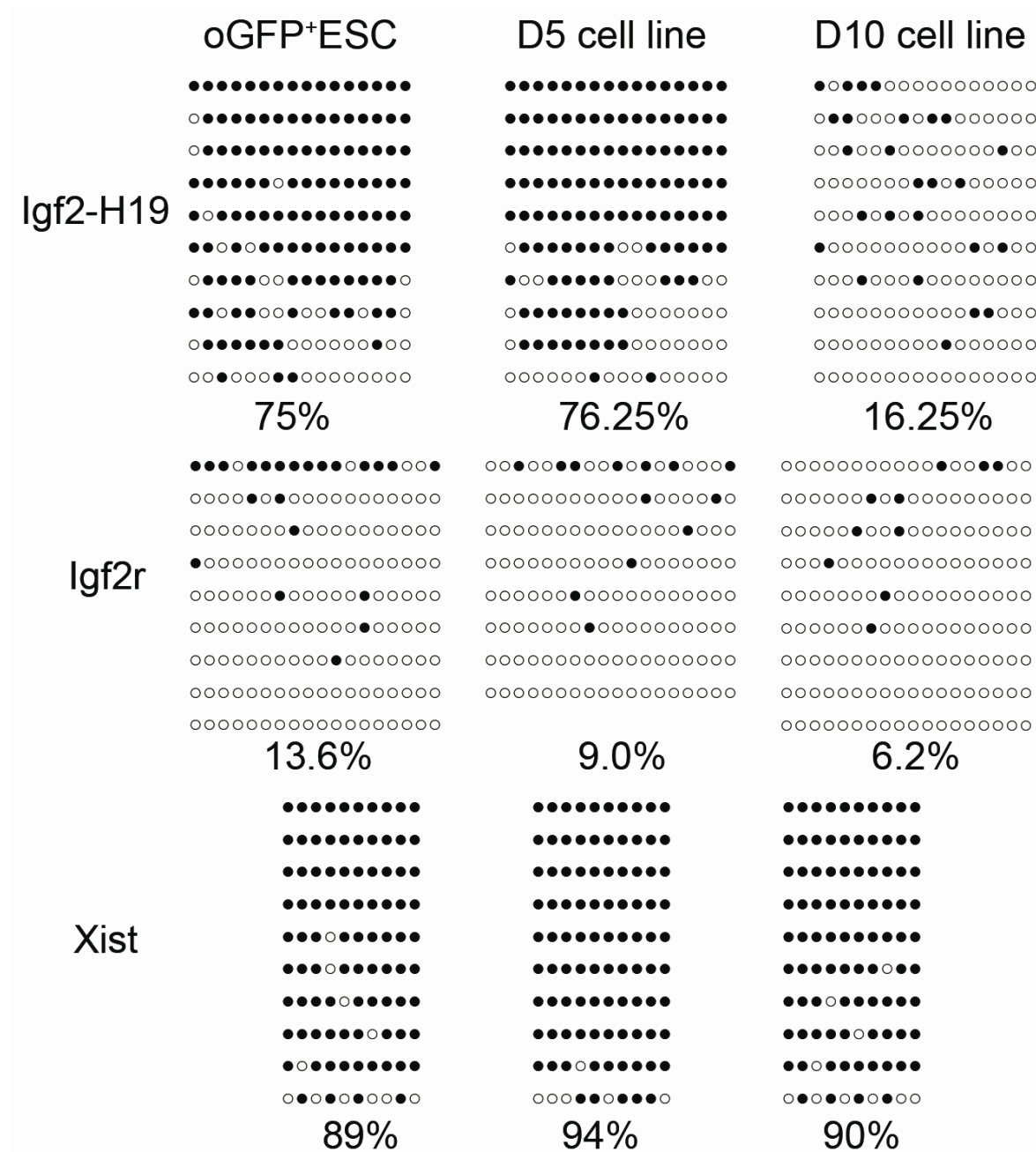


Figure S3 Methylation status of imprinted genes. Methylation status of imprinted genes in oGFP⁺ ESCs, D5 and D10 cell lines was determined using bisulphate sequencing. The D10 cell lines showed significant hypomethylation (16.25%) of the differently methylated regions for the *Igf2-H19* locus, but the methylation patterns of the differently methylated regions for *Igf2r* and *Xist* in the D5 and D10 cell lines are similar to the oGFP⁺ ESCs.