

Additional file 6: CpG methylation relative to Ctf and DNaseI hypersensitivity.

Locations of Ds-TR CpG methylation profiles relative to ChIP-seq data for Ctf (Kim et al., 2007, *Cell* 128: 1231-145) and DNaseI hypersensitivity (Sabo et al., 2006, *Nature Methods* 3:511-518). Region shown corresponds to 72,917,882-72,920,900 of the mouse X chromosome (mm9).

