Accessory Publication

Fig. S1. Representative electropherograms of DNA from normal and resorbed embryos from

control and tamoxifen treated group following bisulfite sequencing indicating the changes in

methylation patterns at H19 DMR. Blue and red arrows represent methylated and unmethylated

CpGs respectively. (A, B) Sequencing result of paternally derived allele with methylated CpG

sites of the normal embryo from control and tamoxifen treated group. (C, D) Electropherograms

of the paternally derived clone of the resorbed embryo from control and tamoxifen treated group

showing scattered methylation errors. (E) Maternally derived clones that were unmethylated at all

CpG sites in normal as well as resorbed embryo from control and tamoxifen treated group.

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