

Additional file 7: Fig. S3

Phasing of methylome and genome variation.

Screenshots of the visualization of EPI_13's Oxford Nanopore long read sequencing BAM files in IGV [19] at the *AR* gene locus. We observe a paternally inherited SNV in *cis* with unmethylation of the promotor region of the gene, confirming the inactivation of the maternal allele showed in Fig. 4.

