Supplementary Figure 2. Genomic distribution of differentially methylated CpG sites in each hormonal treatment, by gain or loss methylation, in normal (NUP), stage I (Endo I) and stage IV (Endo IV) eSF. Only enrichments with p<0.05 by z-test are shown. Orange line: gain of methylation; Blue line: loss of methylation; black line: total interrogated in the HM450 platform.