



Supplemental Figure S6: UCSC Genome Browser view of ATAC-seq peaks and their NMF annotations (https://www.gigauag.uliege.be/cms/c_4791343/en/gigauag-diagnostics-software-data) around *PLCD4* (*Phospholipase C Delta 4*) gene. A recessive mutation in this gene has pleiotropic effects on multiple cow phenotypes (Reynolds *et al.*, 2021). The gene might be regulated by various combinations of shared and tissue-type-specific regulatory elements. Peak names consist of chromosome coordinate, largest NMF component and its weight. Peaks are color-coded as in the main Fig. 1A and F.