



Supplemental Figure S2. Number of expressed genes across 91 tissues and cell types in cattle. Counts of expressed genes (all annotated Ensembl genes in UMD3.1.1; Fragments Per Kilobase per Million mapped reads (FPKM) > 0) across all 91 tissues and cell types. For tissues and cell types with multiple samples, we calculate the mean of expressed genes among all these samples.