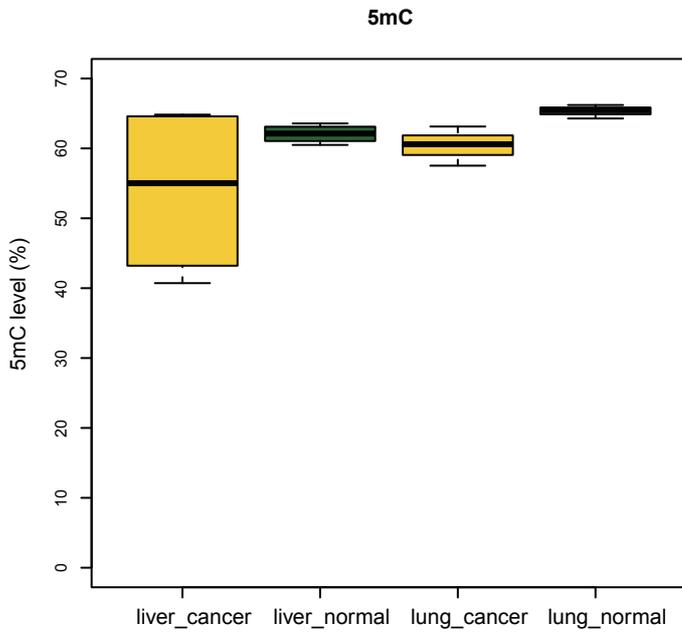
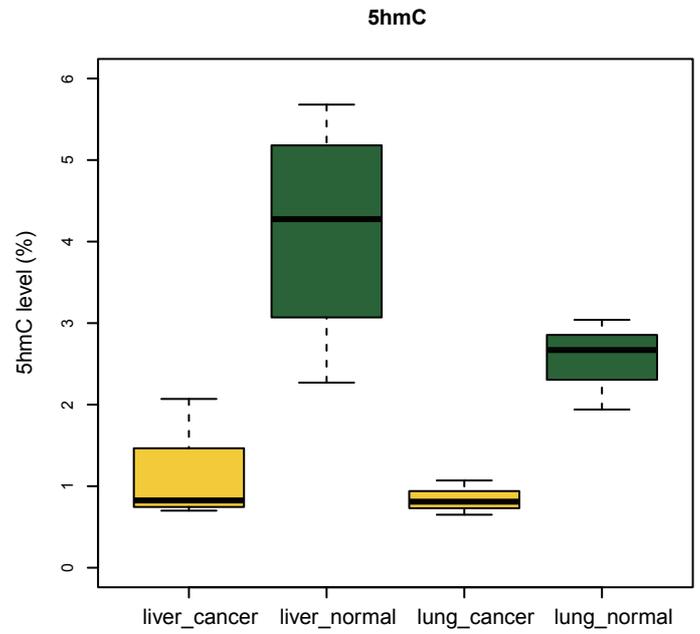


A



B



Supplemental Figure 1. Global methylation and hydroxymethylation levels in samples used in this study. (A) methylation (B) hydroxymethylation. Smoothed methylation and hydroxymethylation levels of all CpG sites across genome was averaged as global 5mC and 5hmC levels.