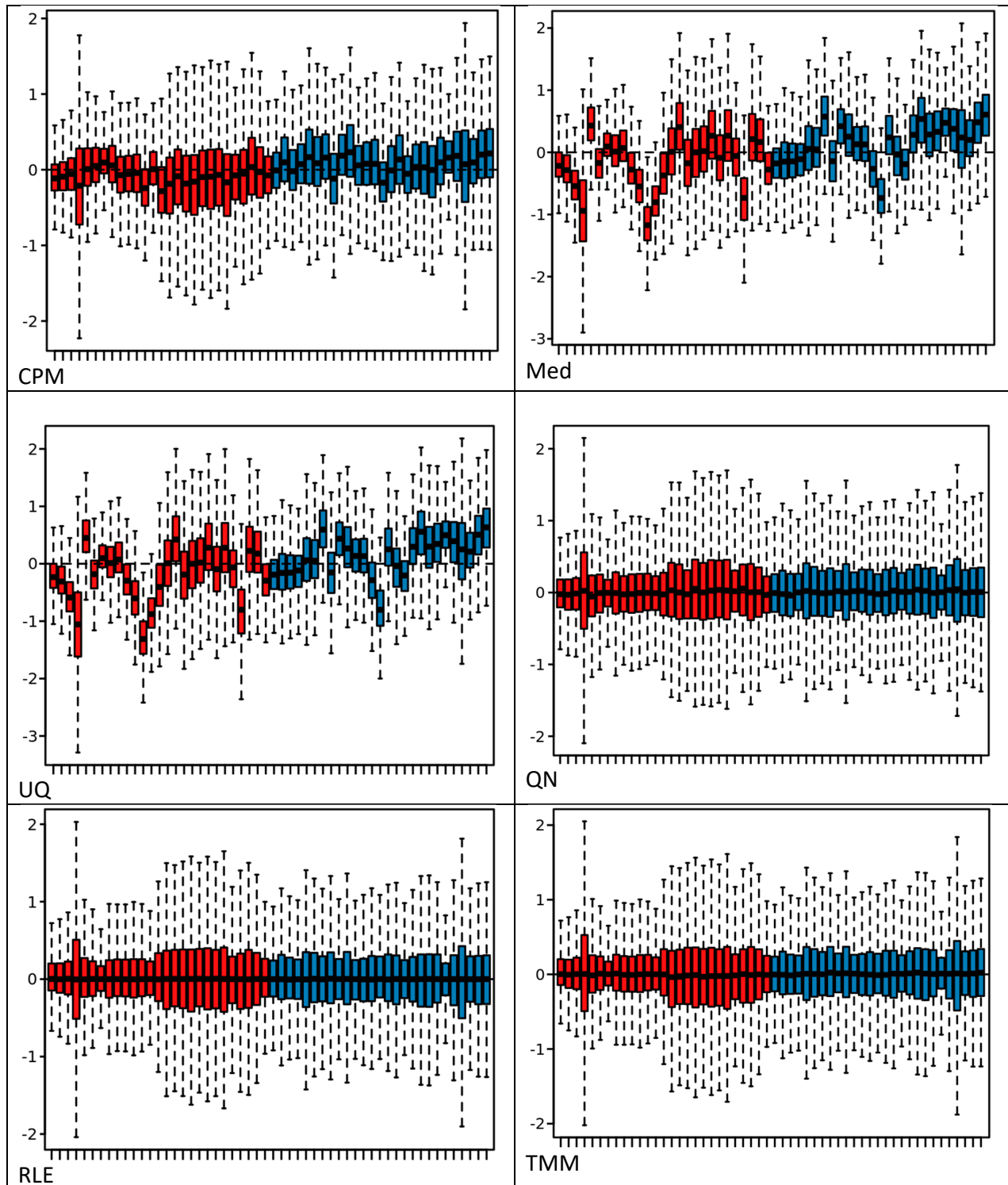
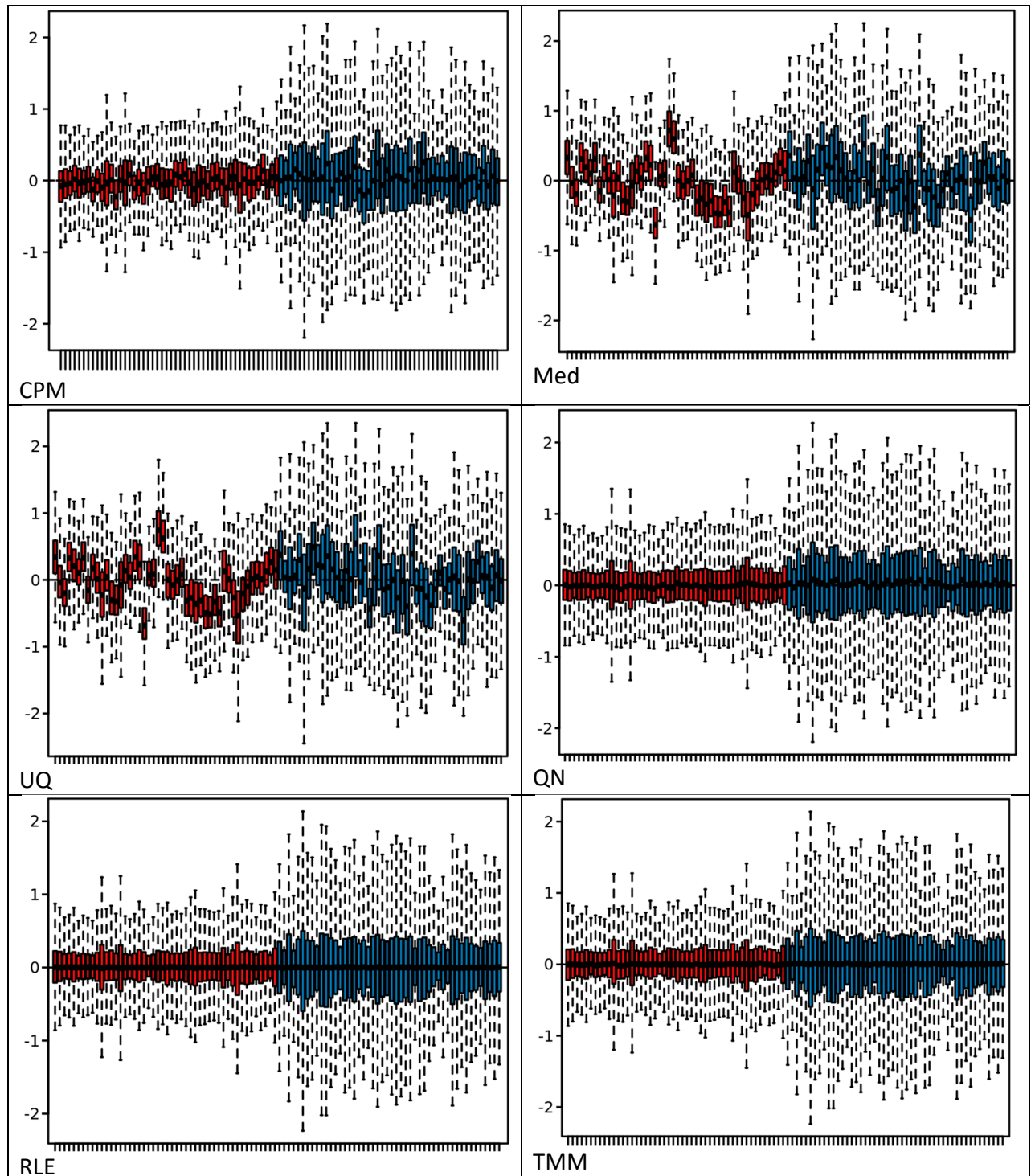


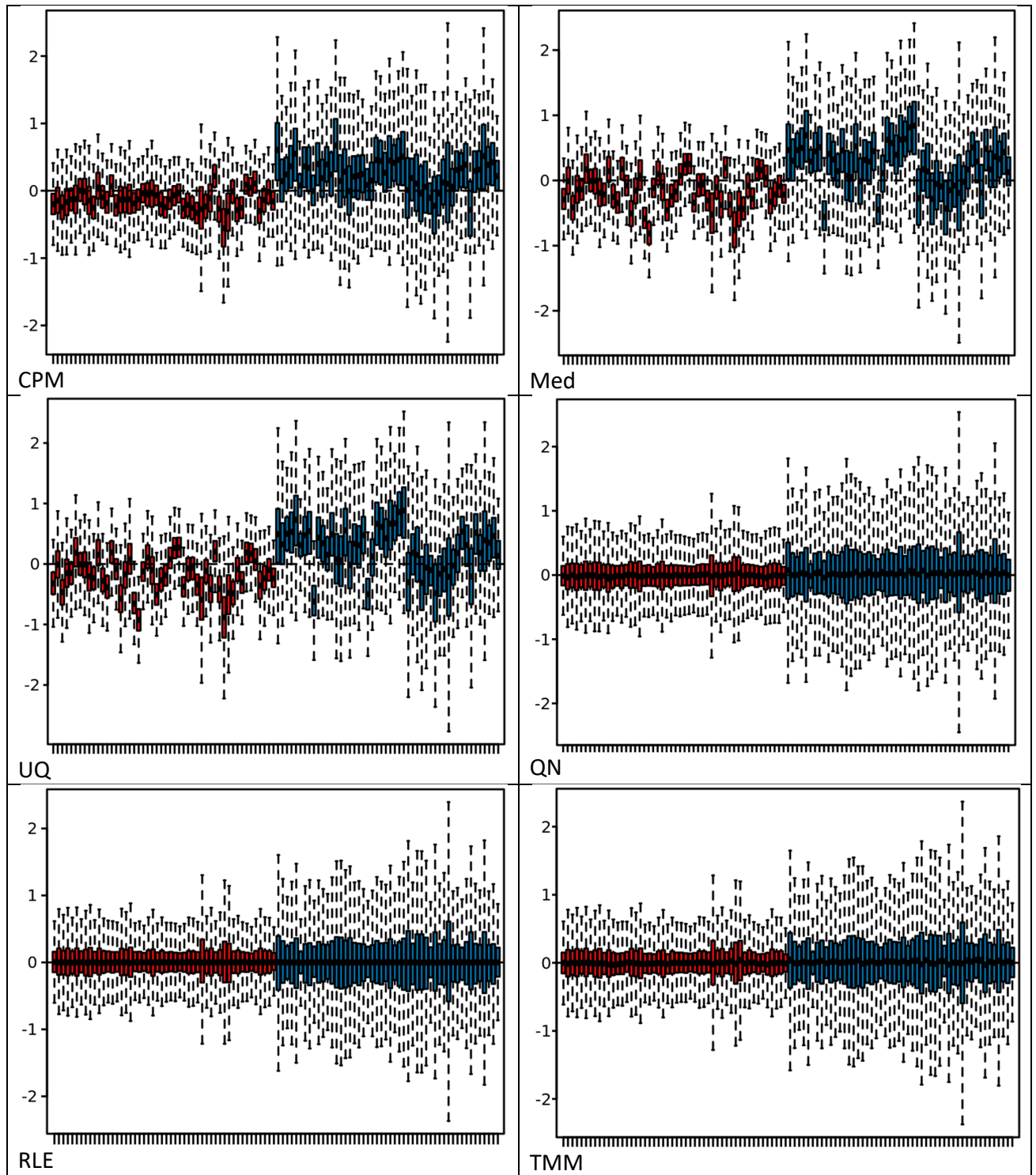
## A: STAD



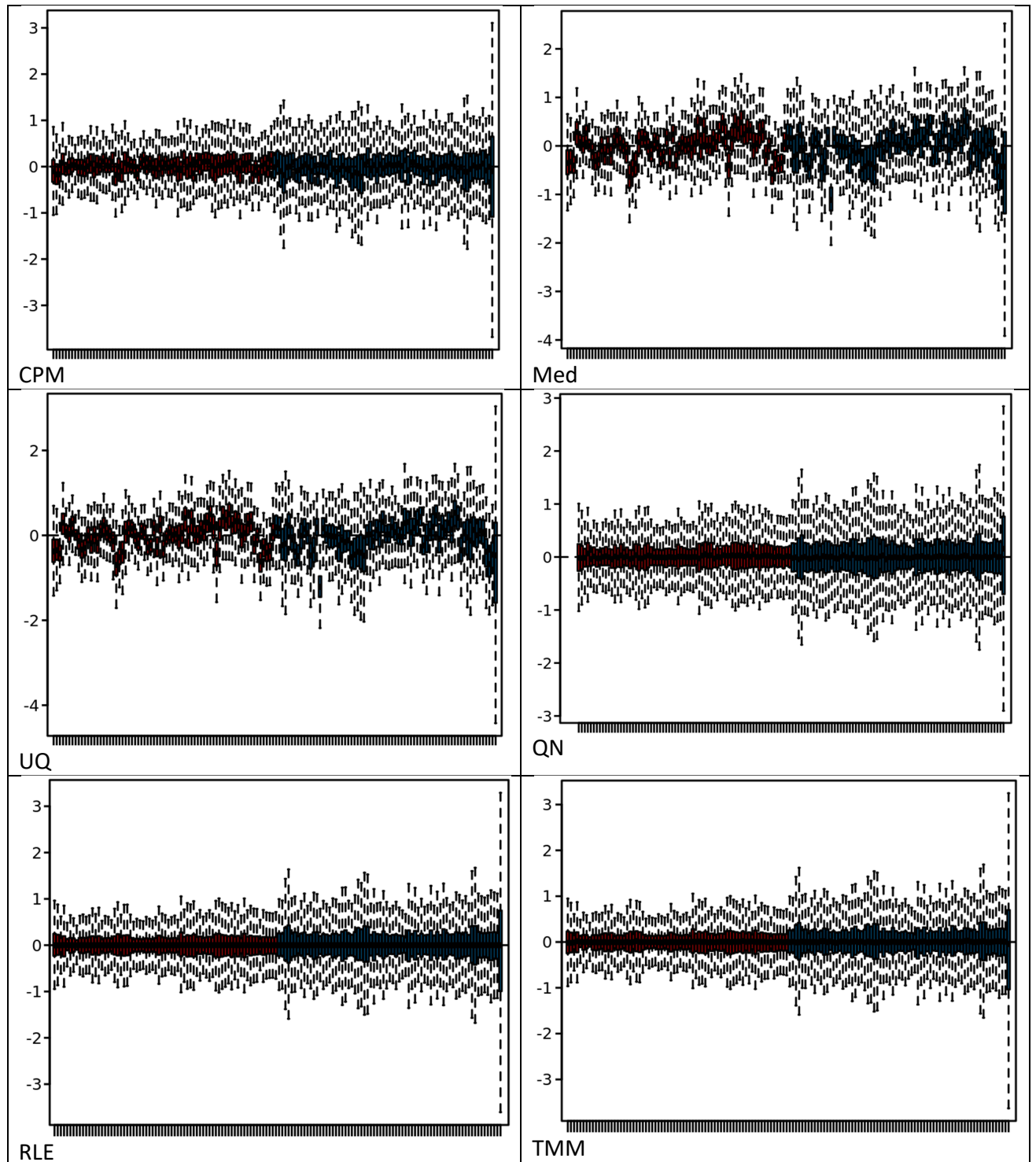
## B: LUSC



## C: LIHC

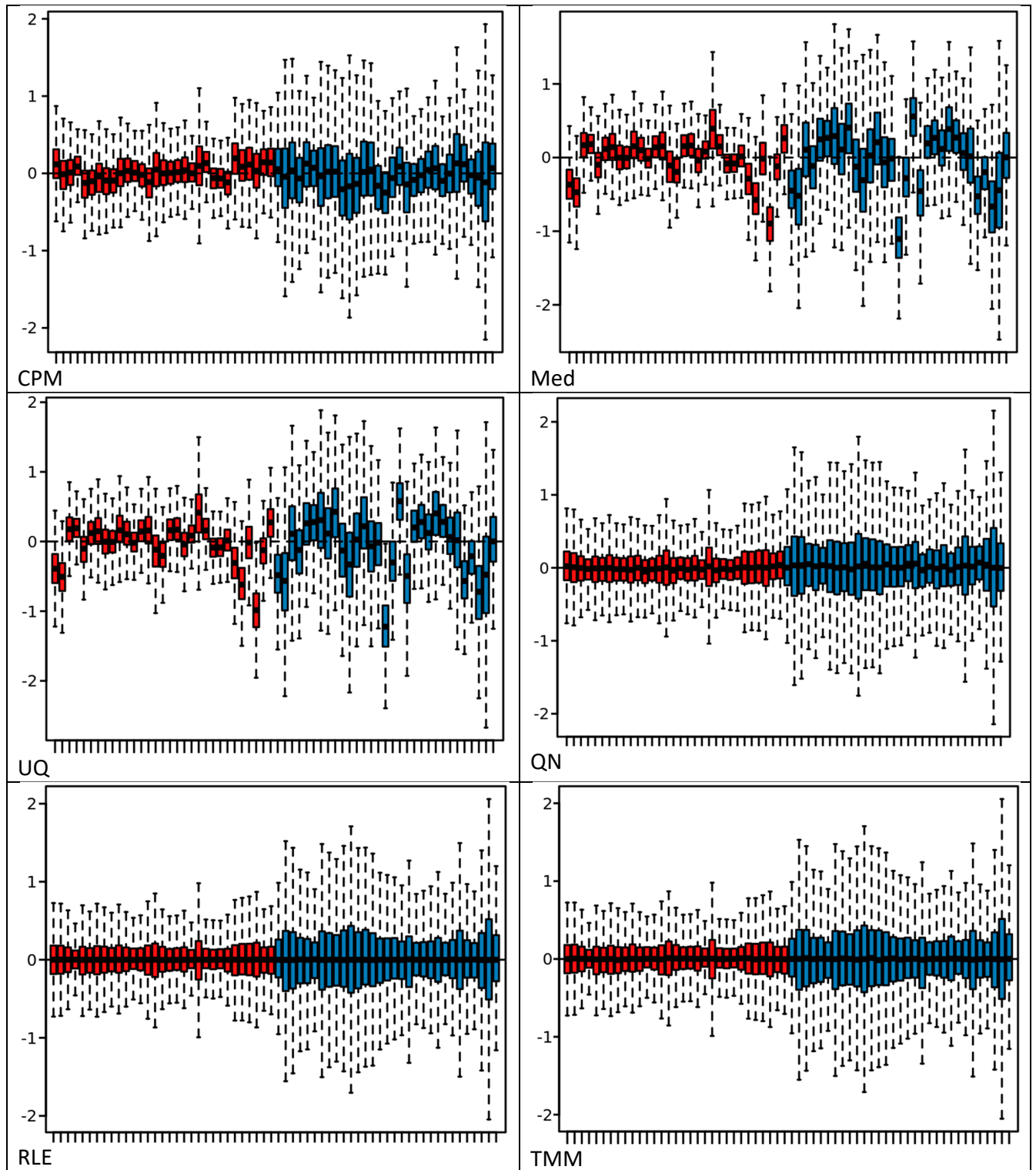


## D: KIRC

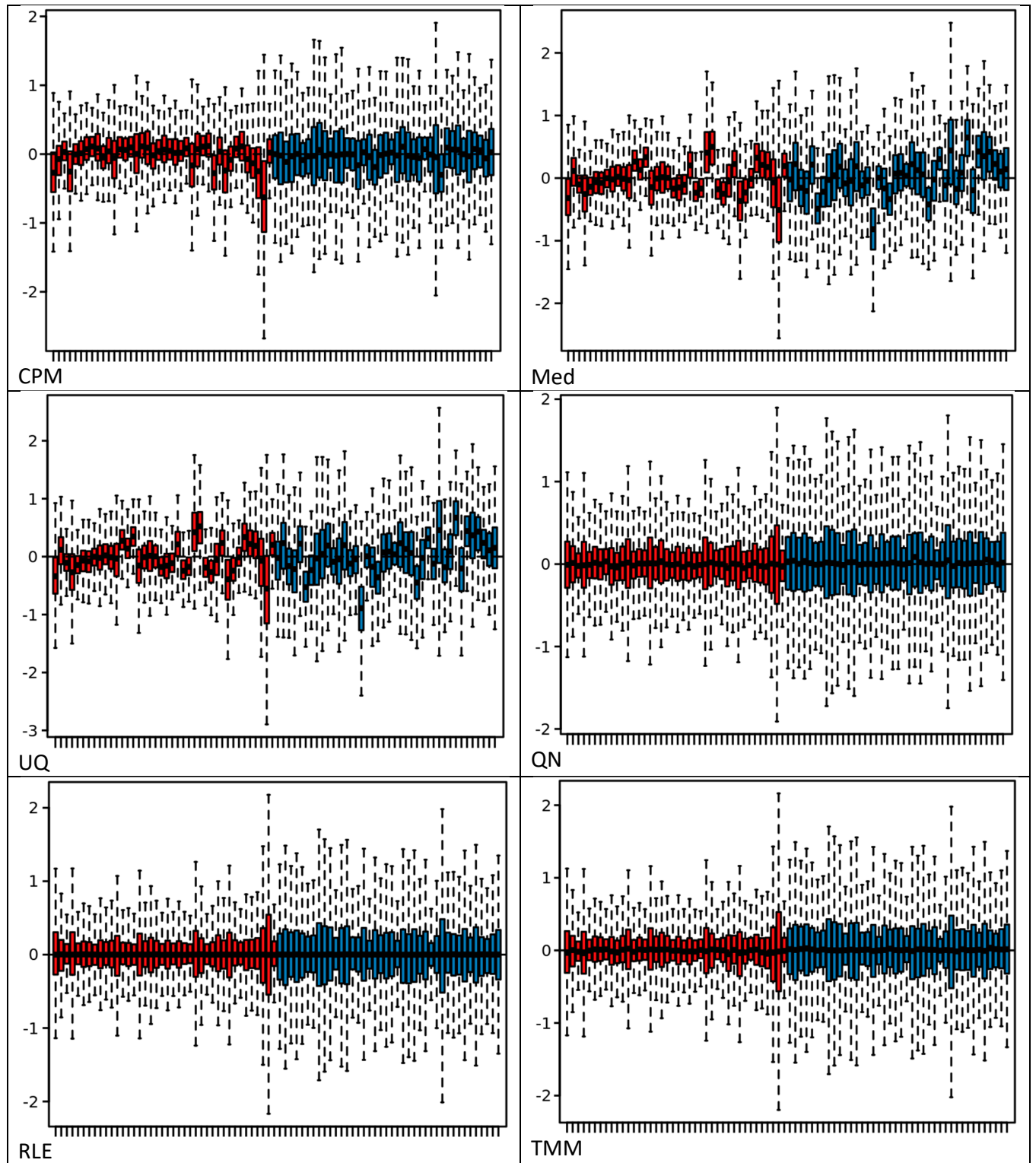




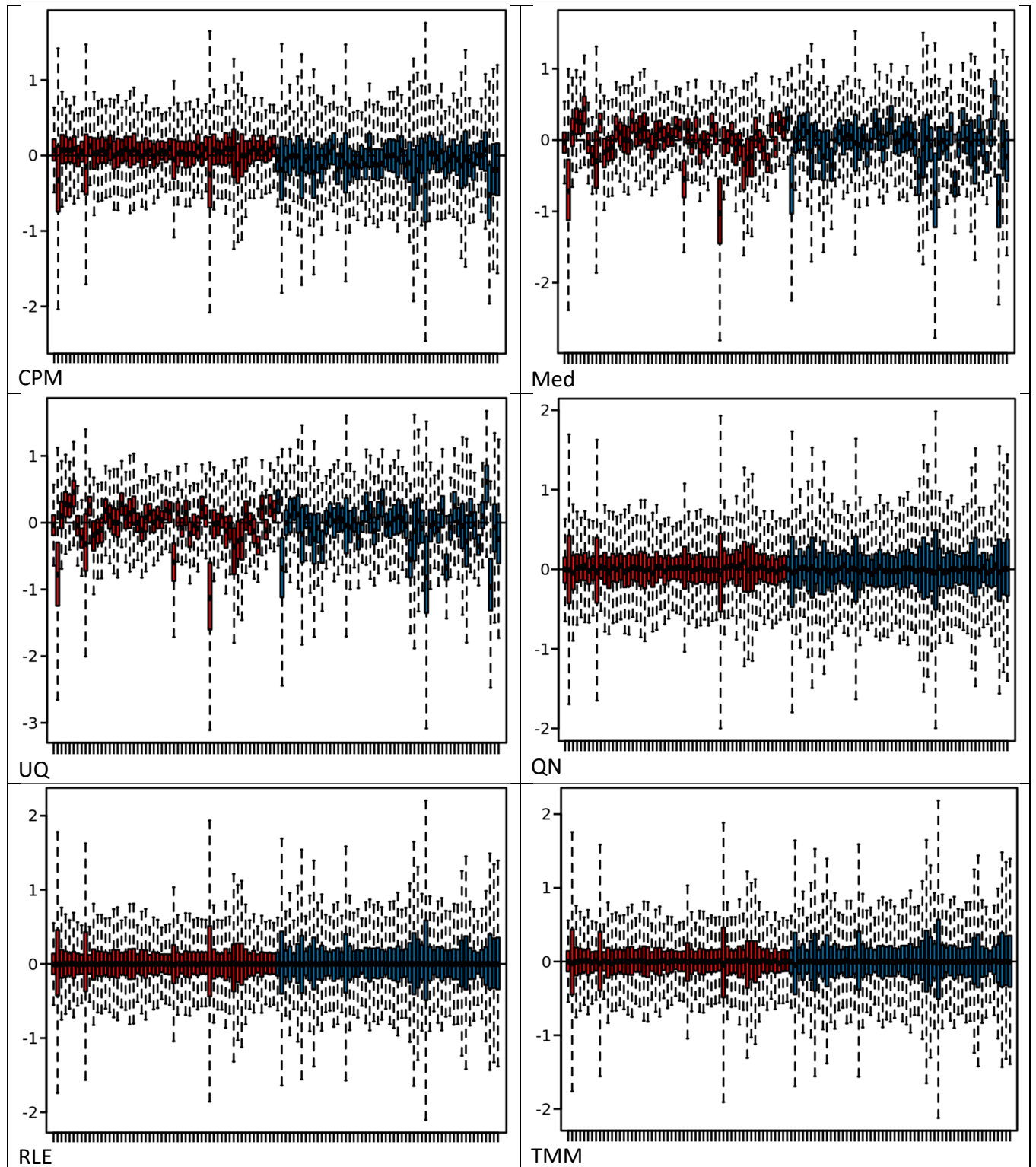
## E: KIRP



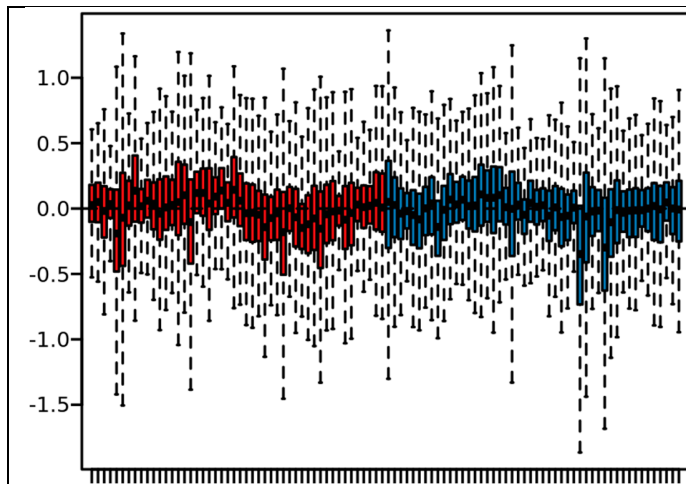
## F: BRCA



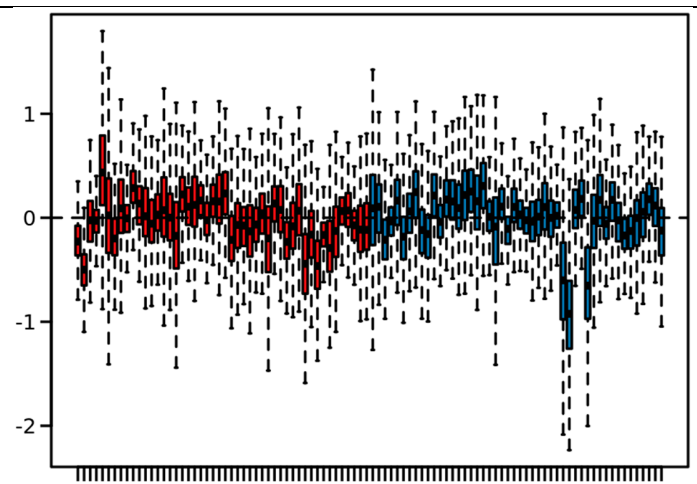
## G: THCA



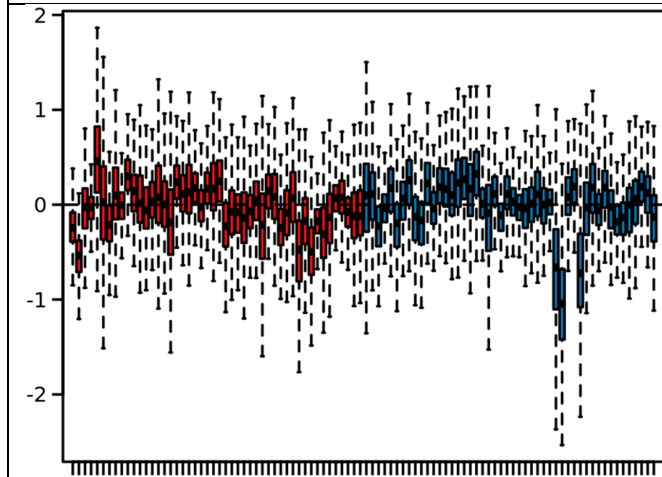
H: PRAD



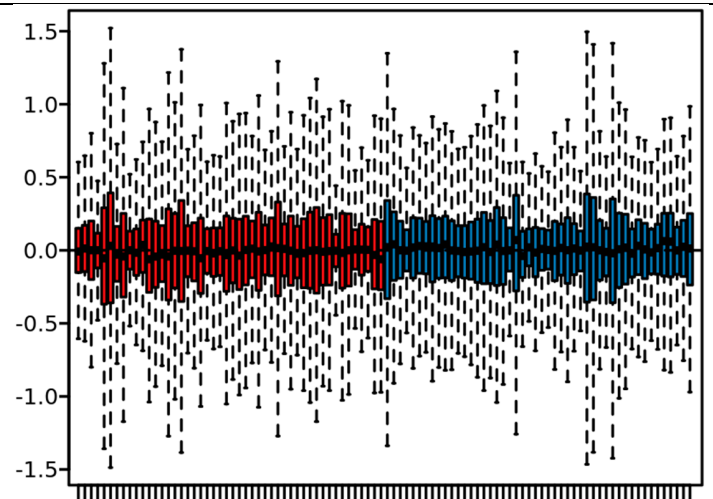
CPM



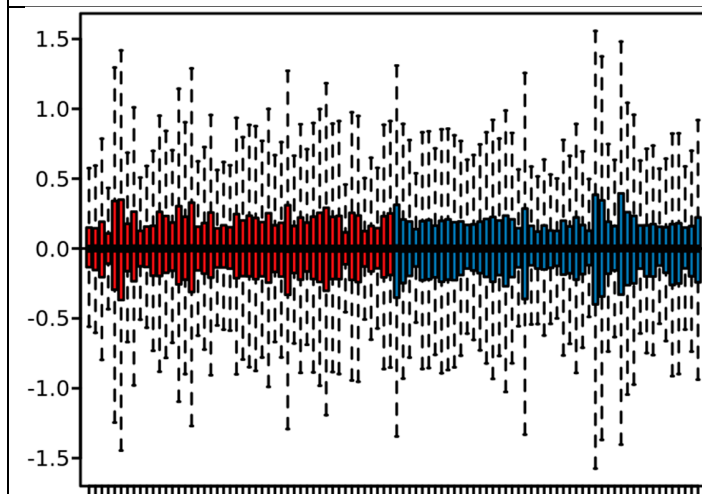
Med



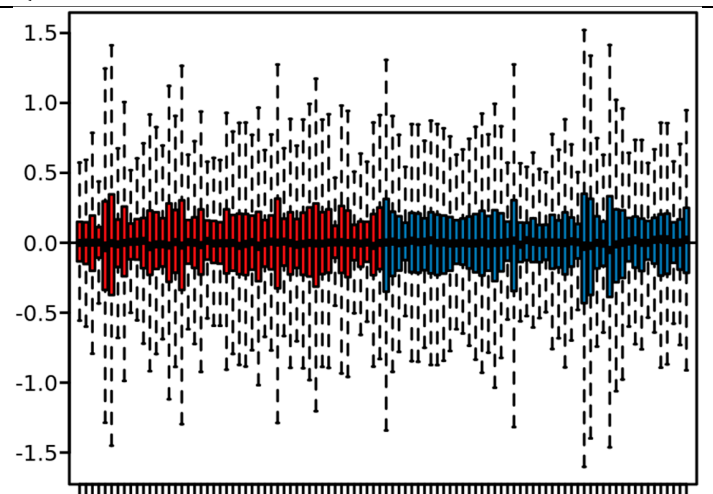
UQ



QN



RLE



TMM

**Figure S3:** RLE plots produced by NormSeq for six normalizing method (UQ, Med, CPM, RLE, QN, and TMM) for STAD (A), LUSC (B), LIHC (C), KIRC (D), KIRP (E), BRCA (F), THCA (G), and PRAD (H). Cancer samples are plotted as blue and control as red.