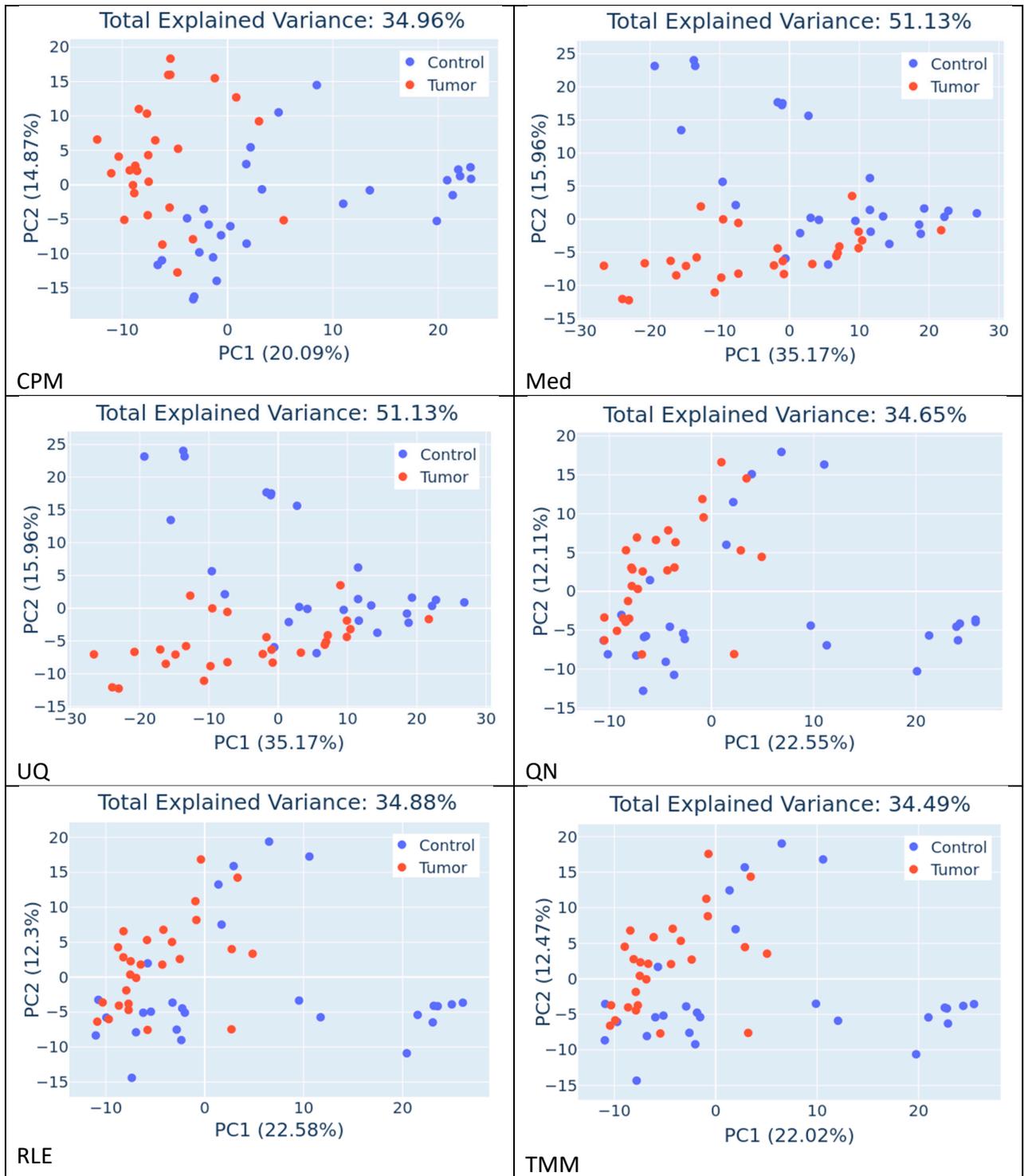
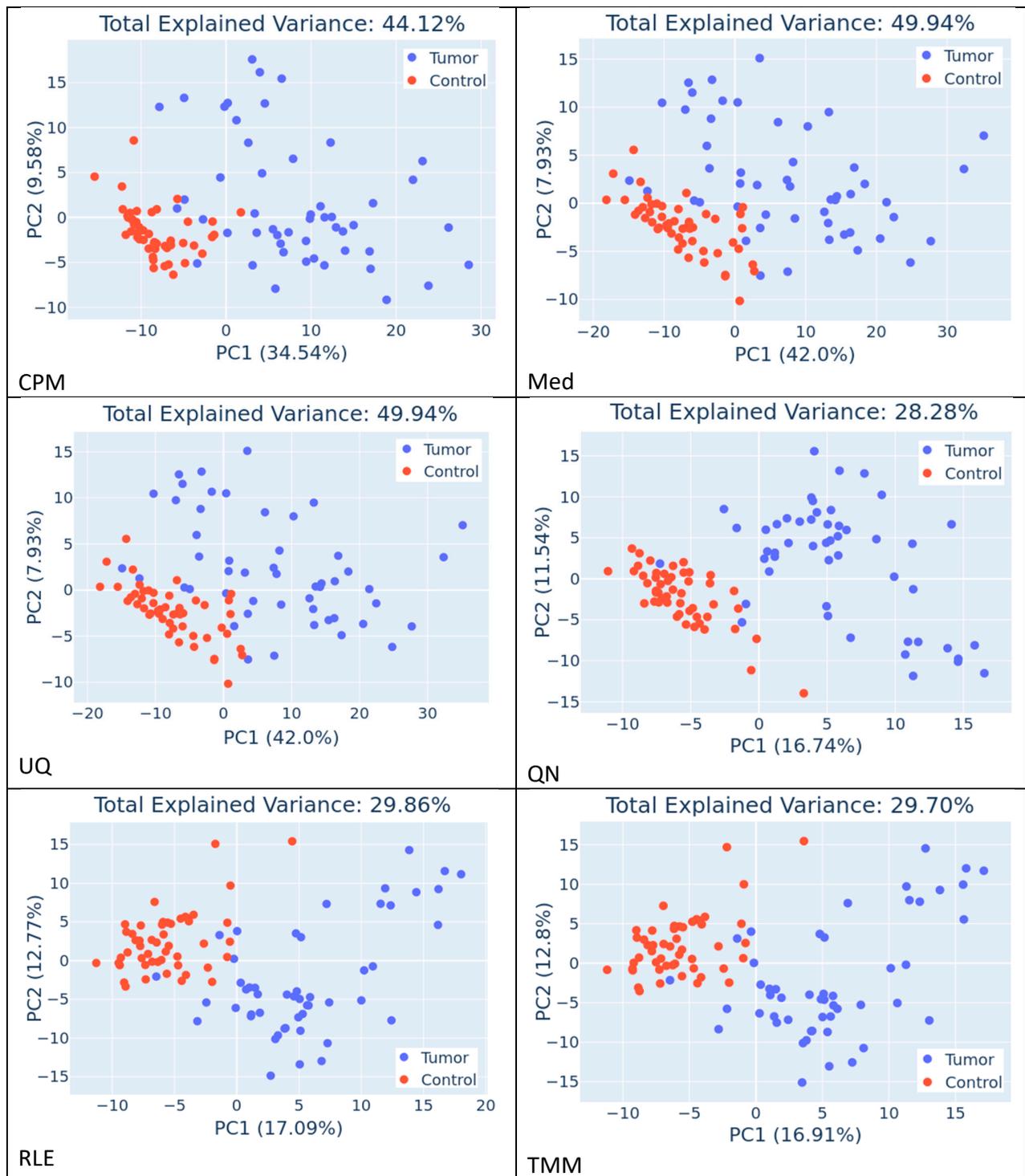


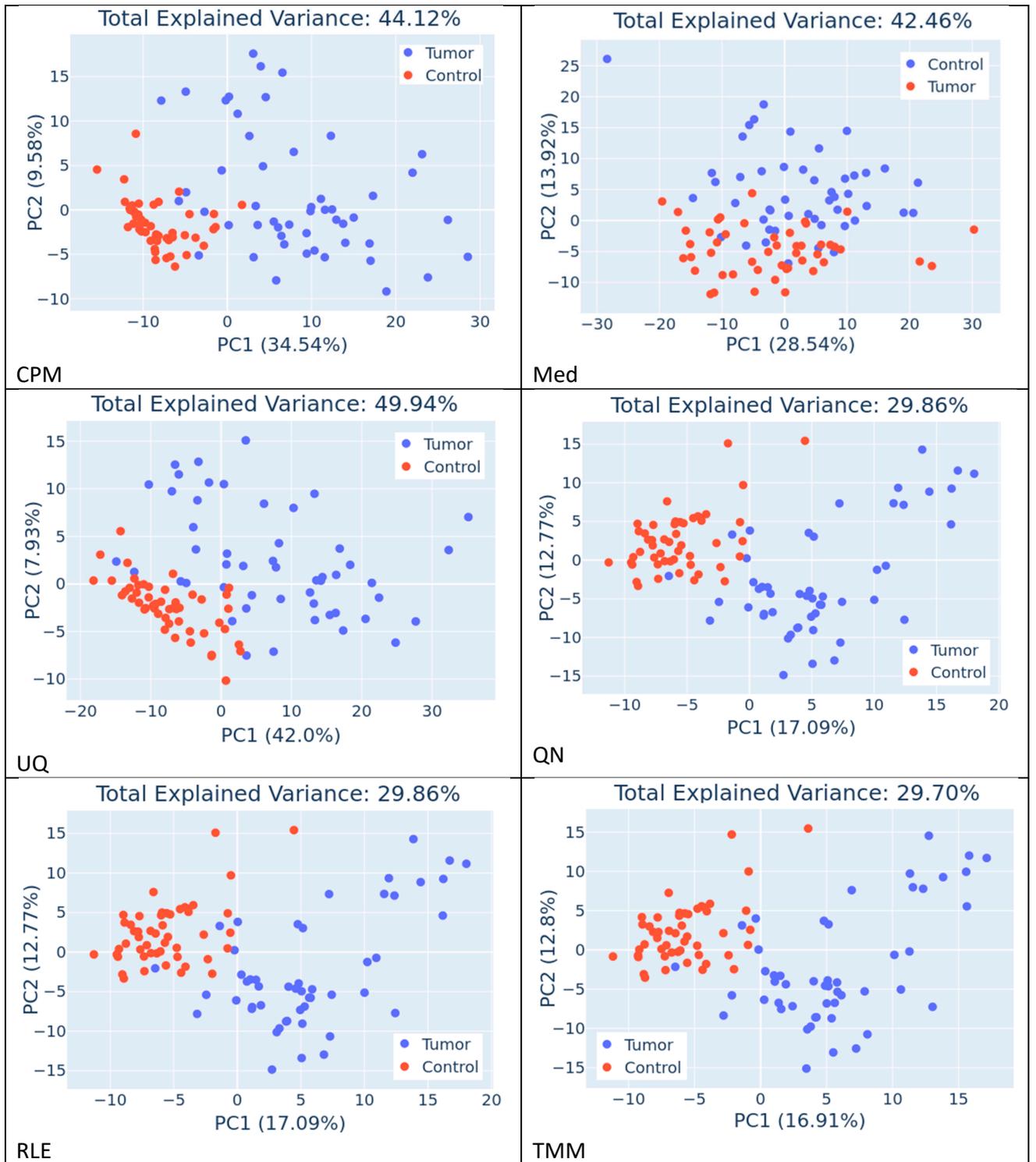
A: STAD



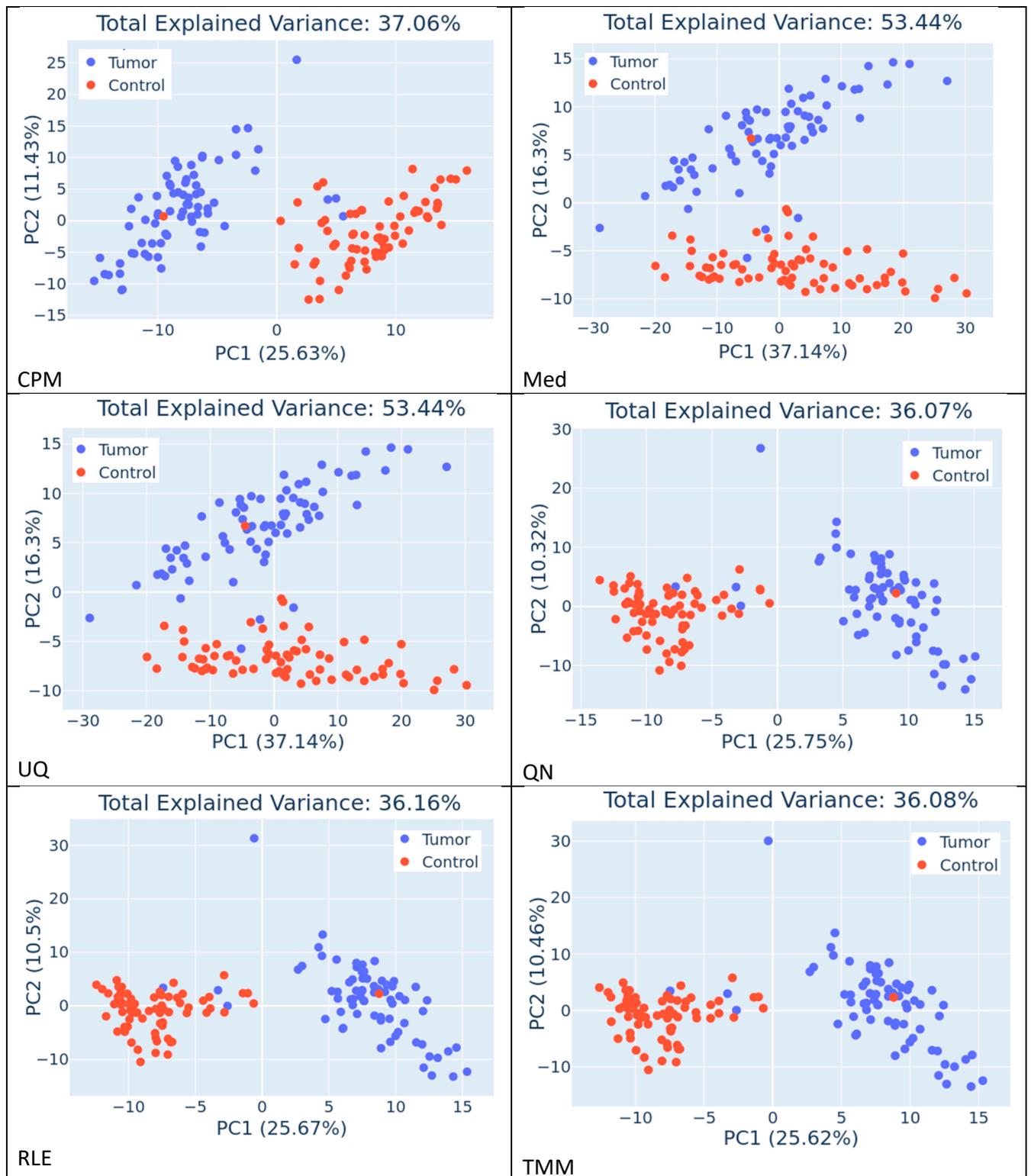
B: LUSC



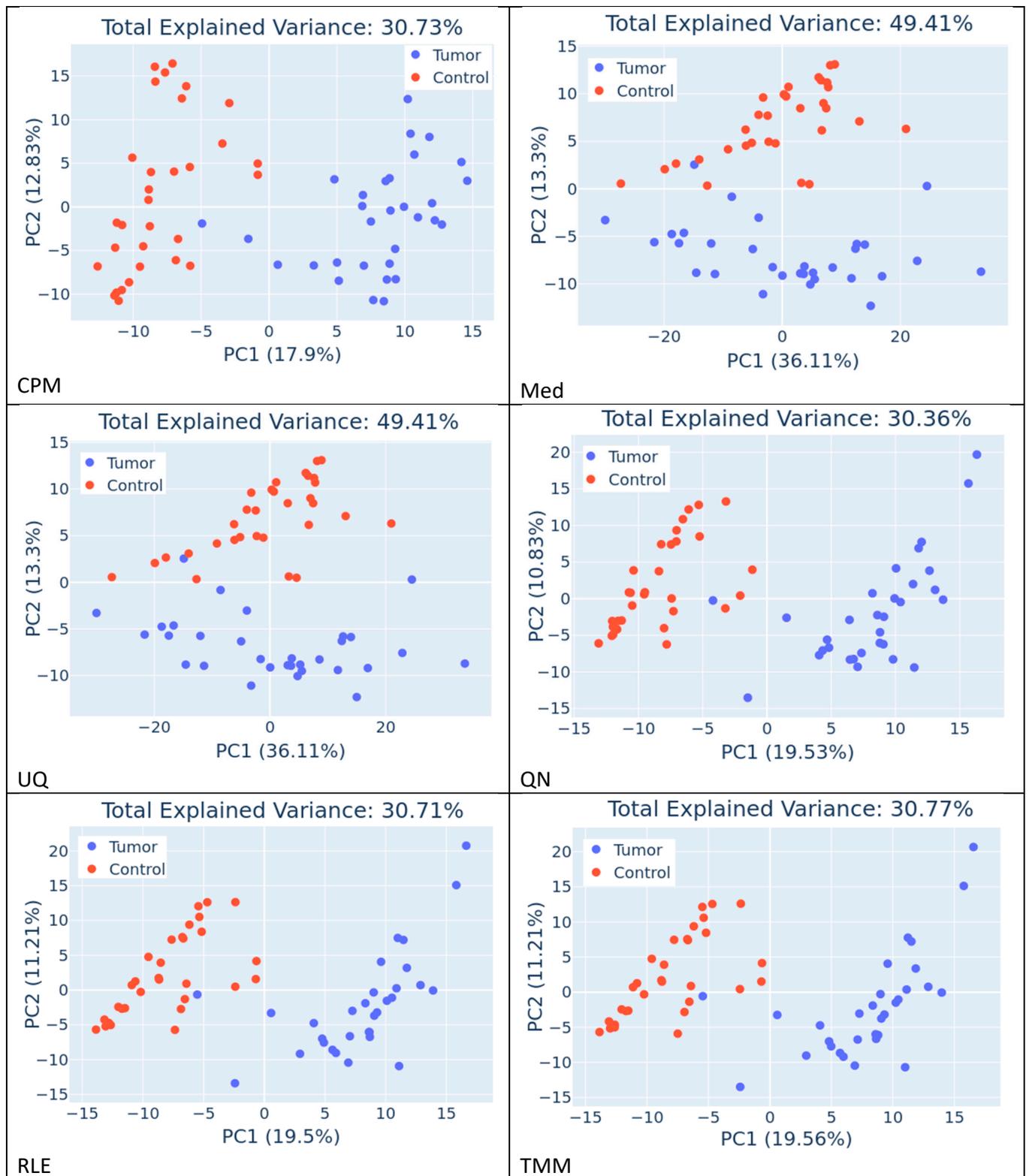
C: LIHC



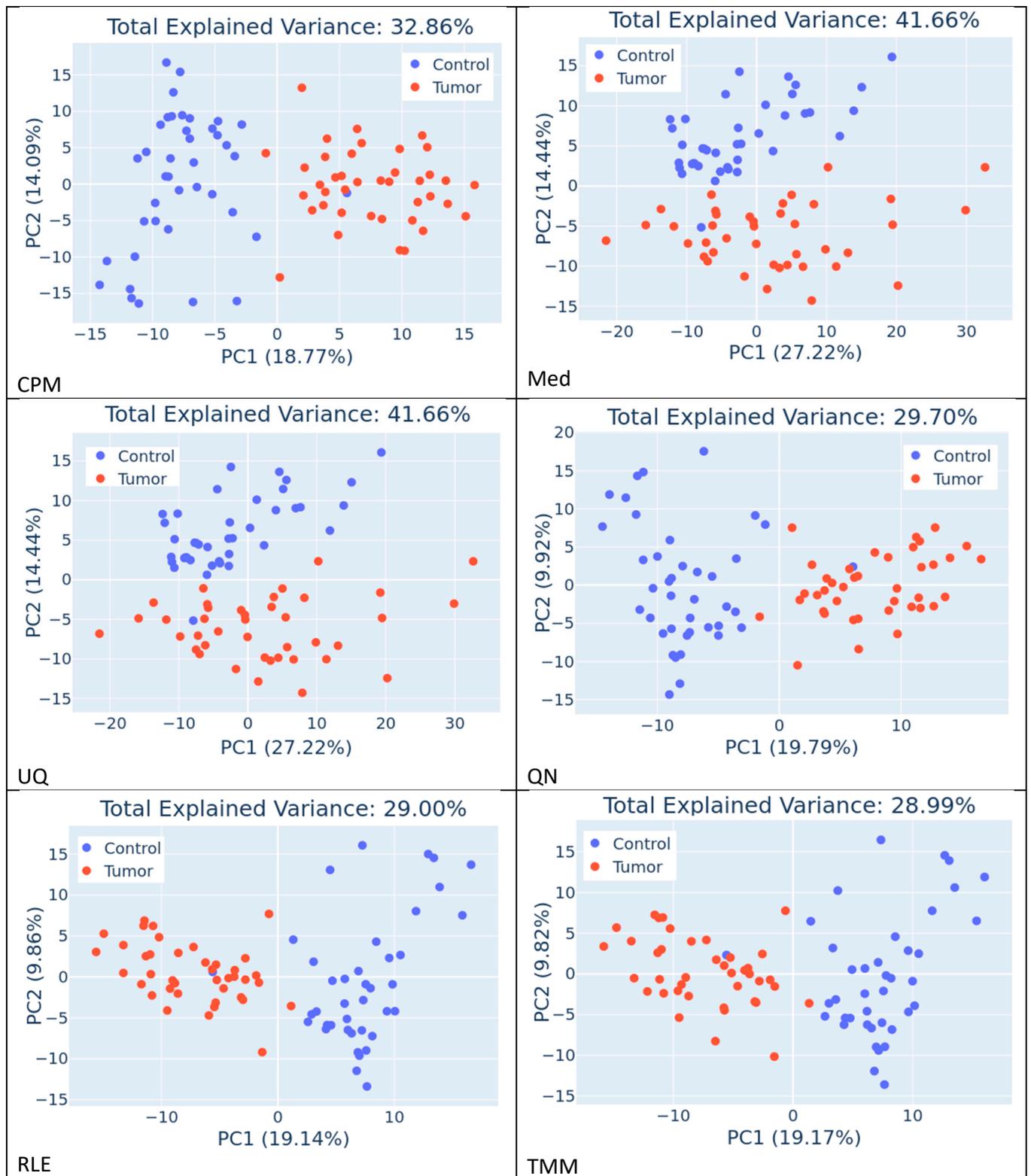
D: KIRC



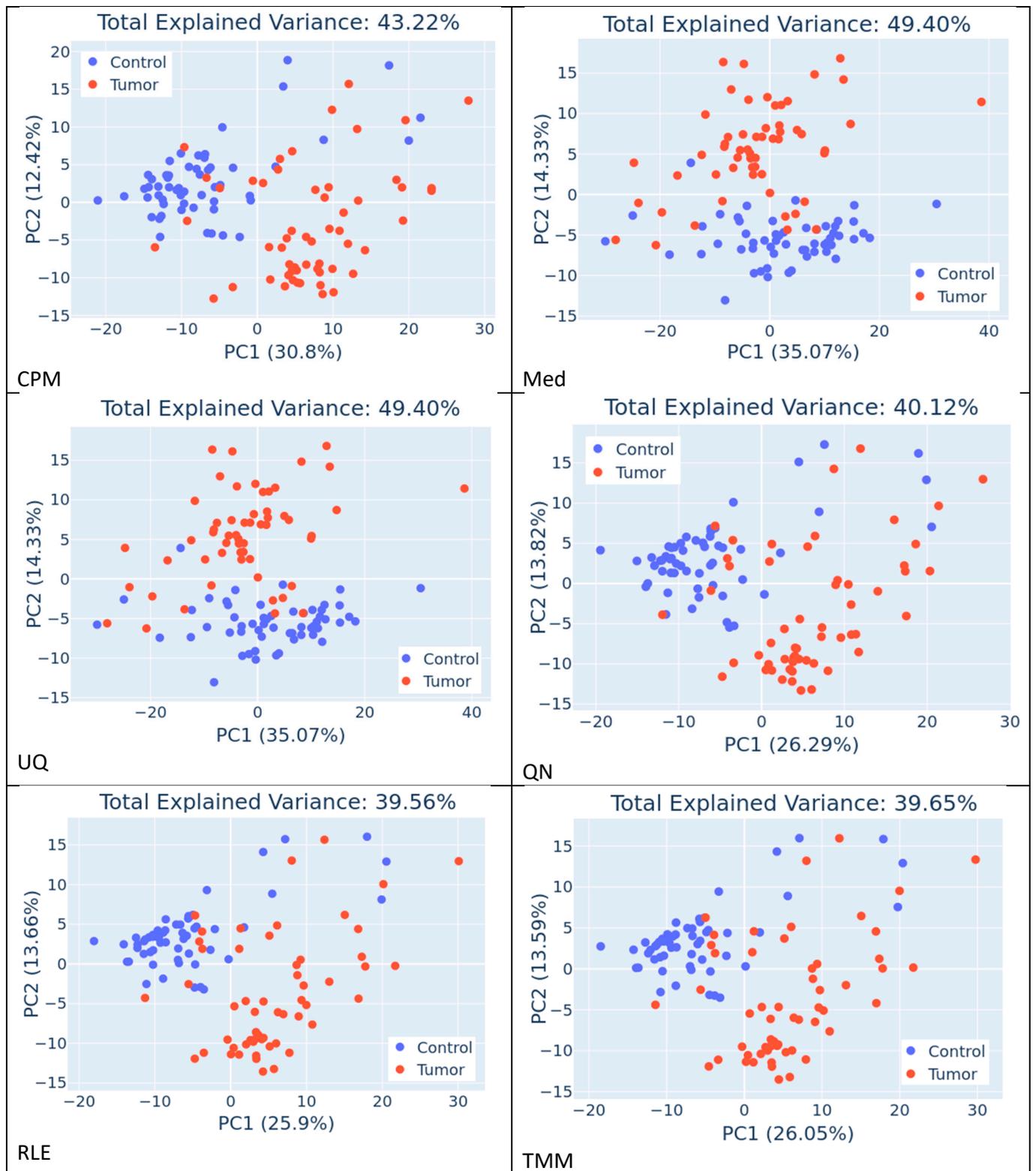
E: KIRP



F: BRCA



G: THCA



H: PRAD

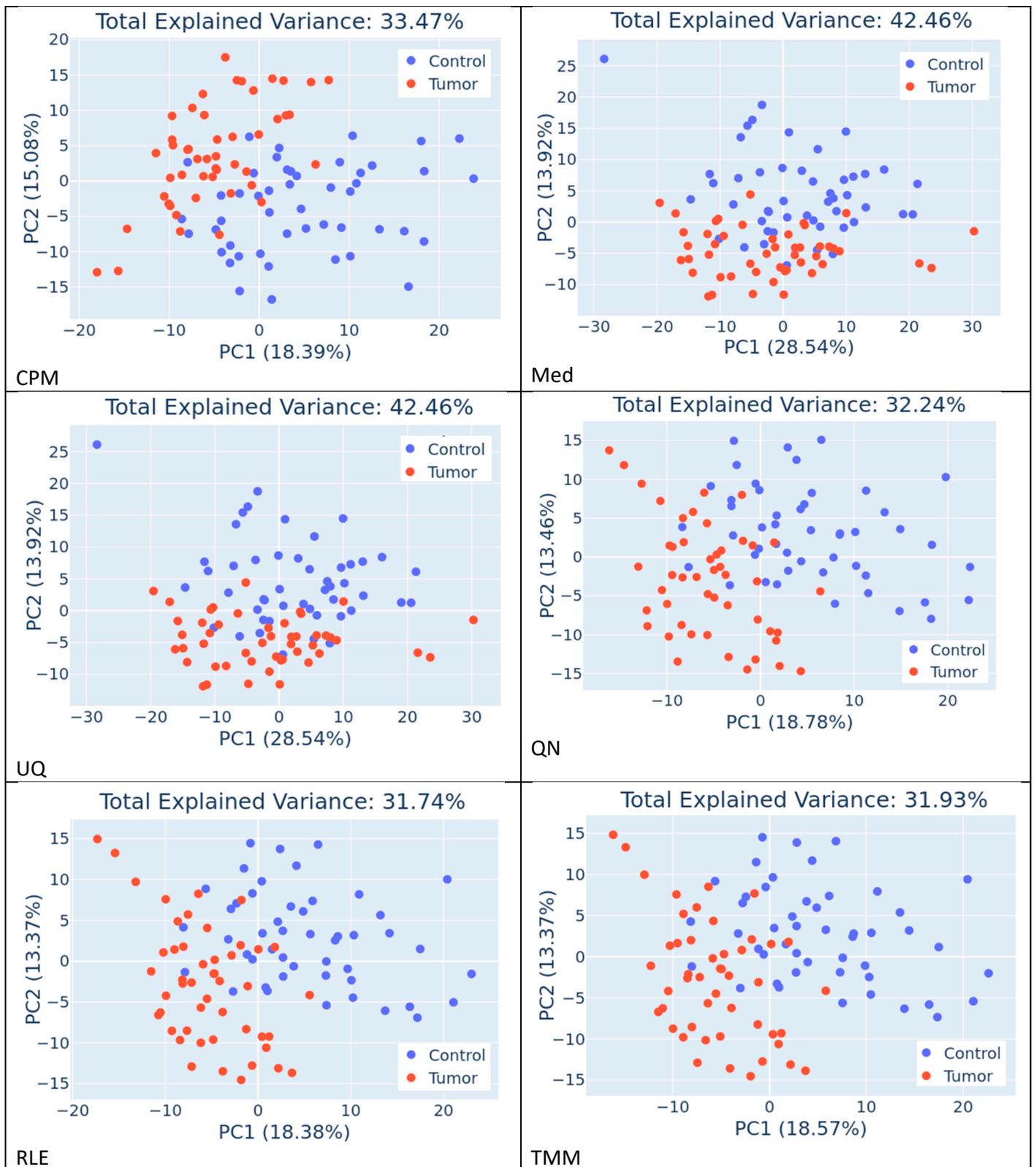


Figure S4: PCA plots produced by NormSeq for six normalizing method (UQ, Med, CPM, RLE, QN, and TMM) for STAD (A), LUSC (B), LIHC (C), KIRC (D), KIRP (E), BRCA (F), THCA (G), and PRAD (H).