

Description of Additional Supplementary Files

File name: Supplementary Data 1

Description: The source data behind the graphs in the paper.

File name: Supplementary Data 2

Description: Enrichr analysis of genes expressed in NPE and PE tumors that correlate with those expressed in ERMS ($P \leq 0.05$).

File name: Supplementary Data 3

Description: Up- and downregulated genes in NPE vs PE RMS ($\log_2FC \leq -0.5$ and ≥ 0.5) $P \leq 0.5$.

File name: Supplementary Data 4

Description: Enrichr analysis of genes upregulated in NPE RMS versus those in PE RMS ($P < 0.05$).

File name: Supplementary Data 5

Description: Enrichr analysis of genes downregulated in NPE RMS versus those in PE RMS ($P < 0.05$).

File name: Supplementary Data 6

Description: GSEA of top ranked differentially expressed genes in NPE versus PE RMS.

File name: Supplementary Data 7

Description: Comparison of gene expression profiles in RMS patients with gene expression of MSigDB pathways. Results are ranked by ADIPOQ expression.