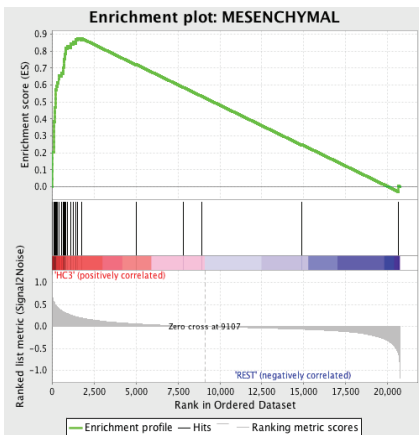
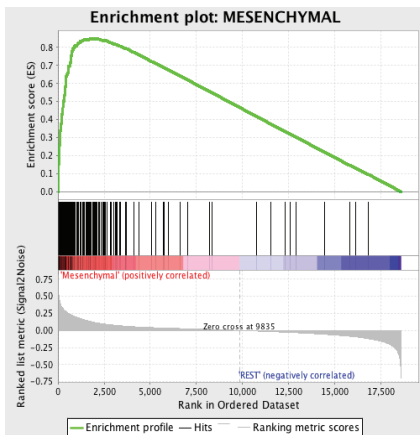


**A****B**

**Figure S1. Comparison of gene expression across different mesenchymal tumor subtypes. (A)** Significant enrichment of the top 50 genes from the Verhaak [7] mesenchymal signature in the Paugh [11] HC3/mesenchymal subtype. **(B)** Significant enrichment of the Paugh HC3/mesenchymal signature in the Verhaak mesenchymal subtype.