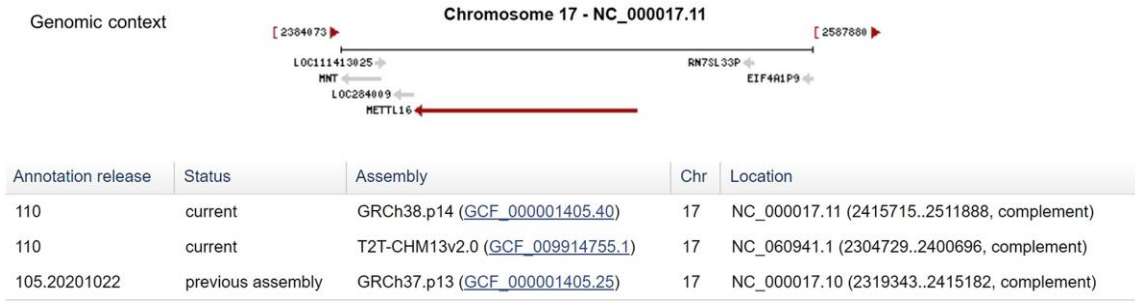
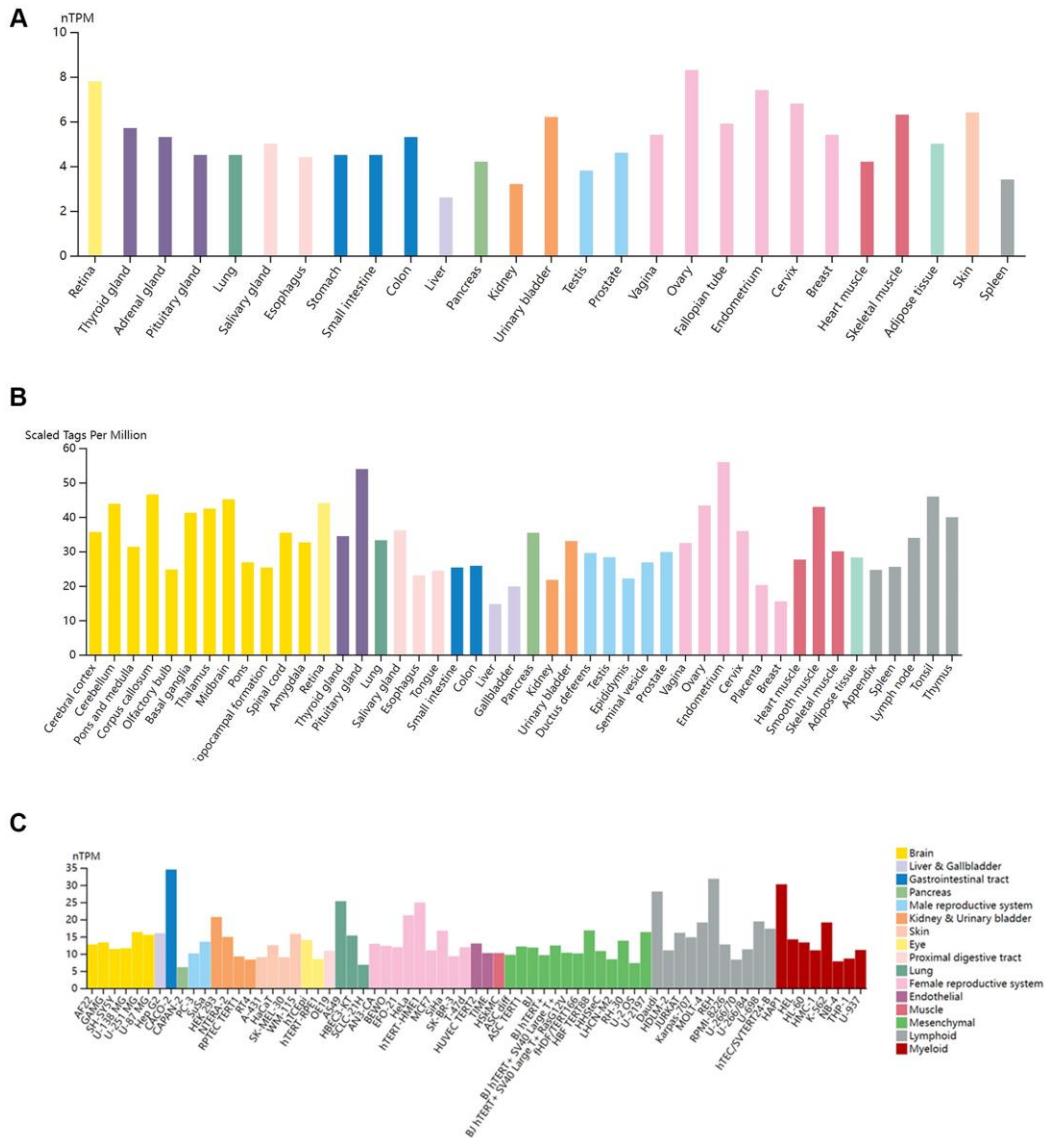


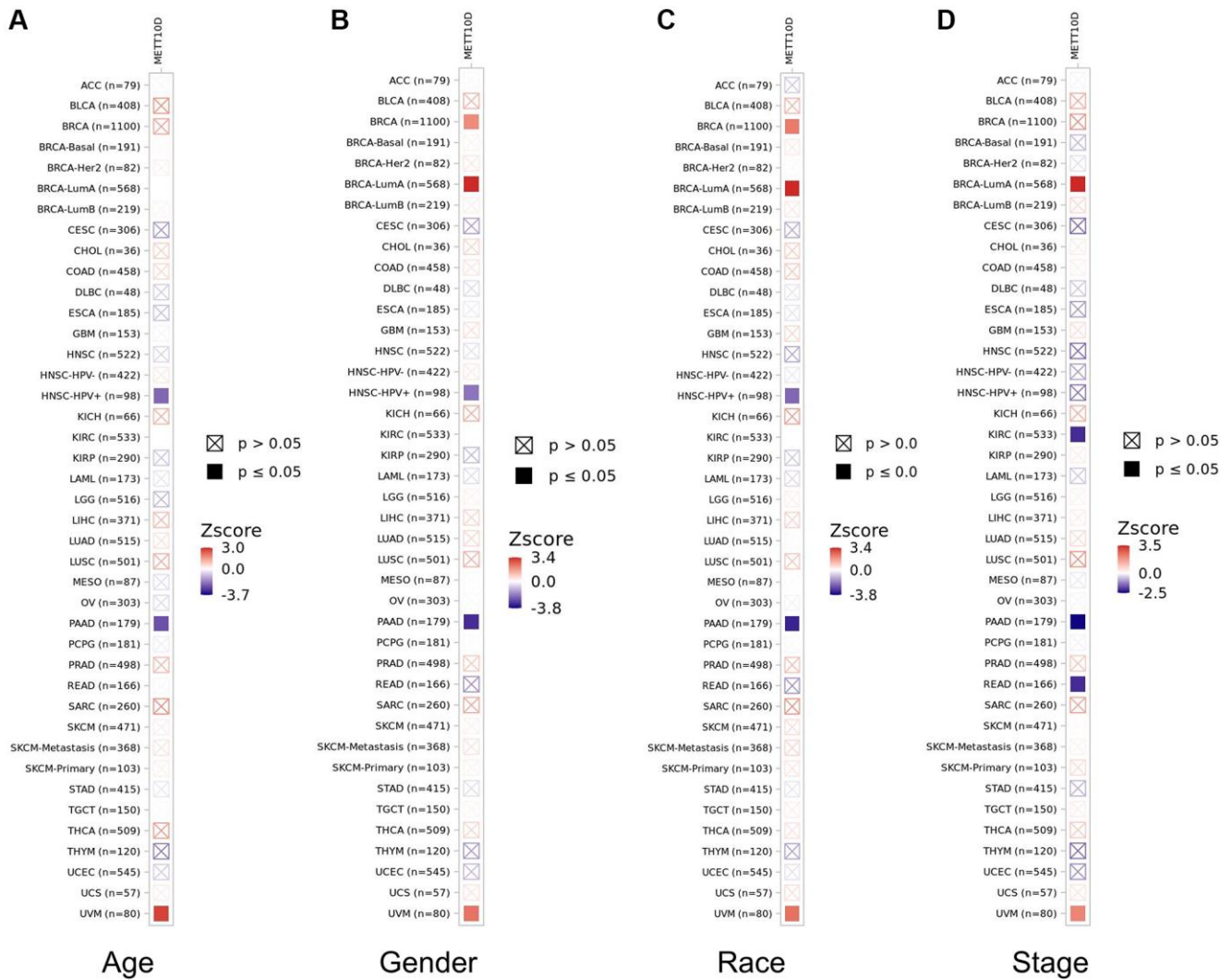
SUPPLEMENTARY FIGURES



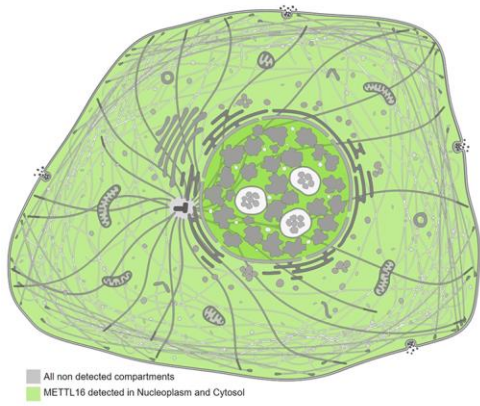
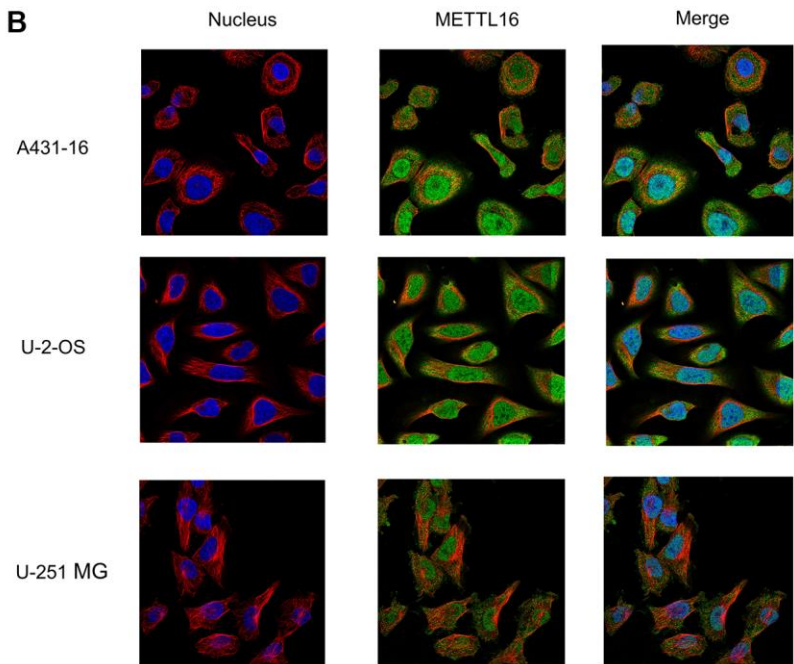
Supplementary Figure 1. Structural characteristics of METTL16.



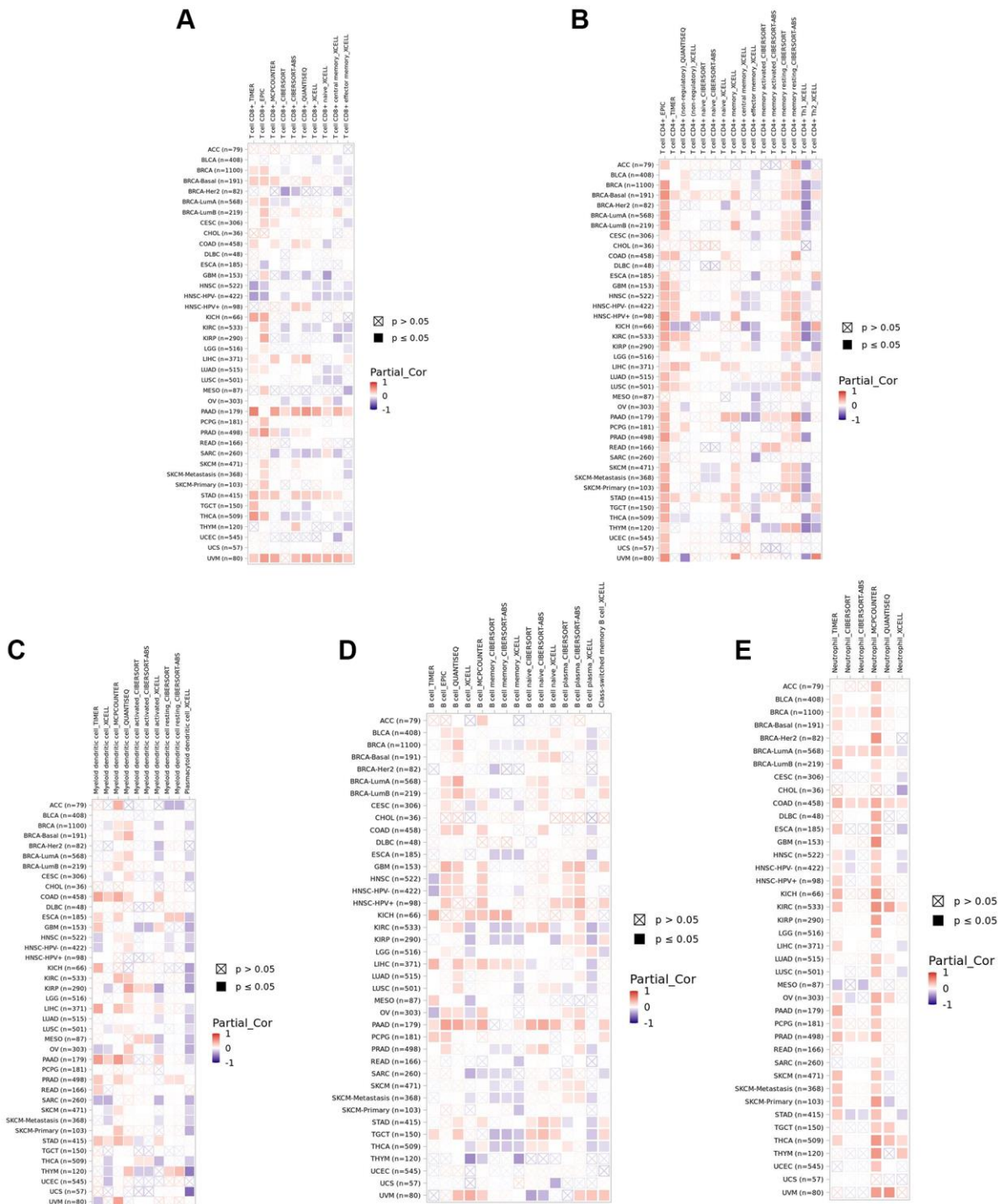
Supplementary Figure 2. Expression level of METTL16 in different cells and tissues. (A) HPA tissue dataset; (B) Tissue consensus dataset; (C) Cell line dataset.



Supplementary Figure 3. The relationship between METTL16 gene expression with age (A), gender (B), race (C), and stage (D).

A**B**

Supplementary Figure 4. The subcellular distribution of METTL16 (A) and the nucleus are marked in blue, the METTL16 marked in green, the microtubules are marked in red (B).



Supplementary Figure 5. Correlation analysis between METTL16 expression levels and the expression degree of immune cells, including CD8+ T cell (A), CD4+ T cell (B), Myeloid (C), B cell (D), Neutrophil (E).