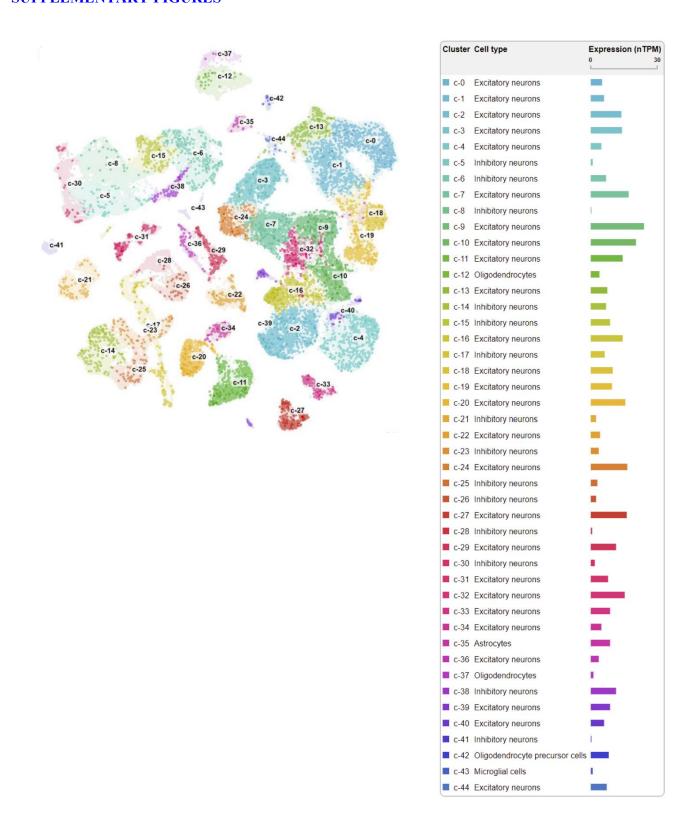
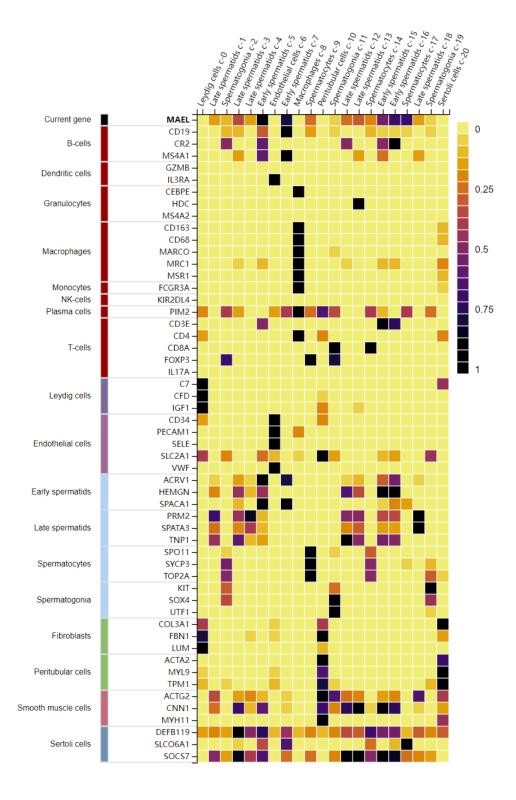
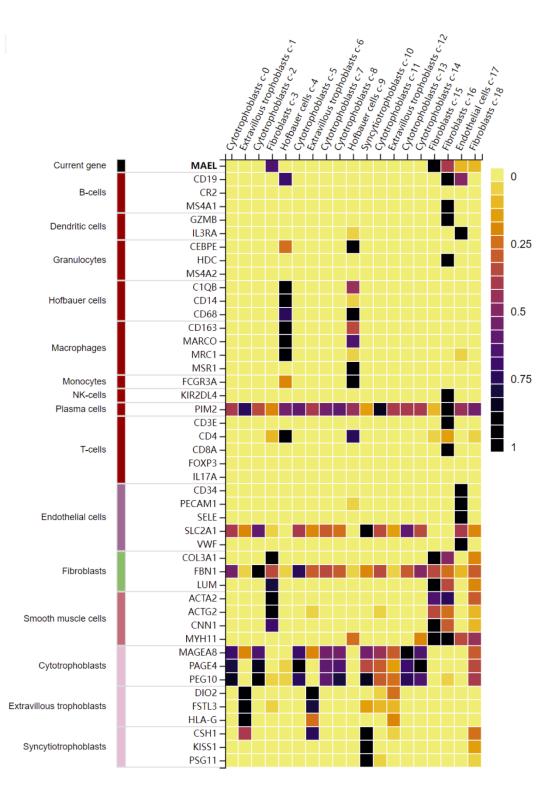
SUPPLEMENTARY FIGURES



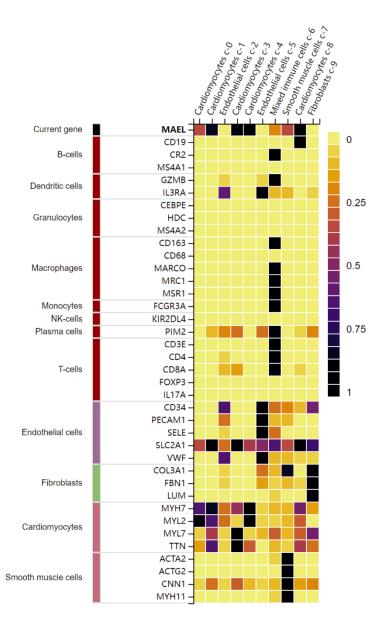
Supplementary Figure 1. UMAP plot and MAEL expression in the single cell clusters of normal brain (Allen brain map).



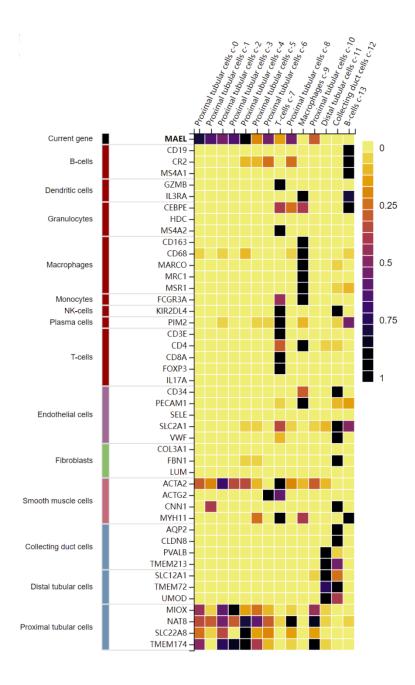
Supplementary Figure 2. mRNA expression of *MAEL* and cell type markers in different single cell type clusters of normal testis (GSE120508). Color legend: fraction of the highest expression.



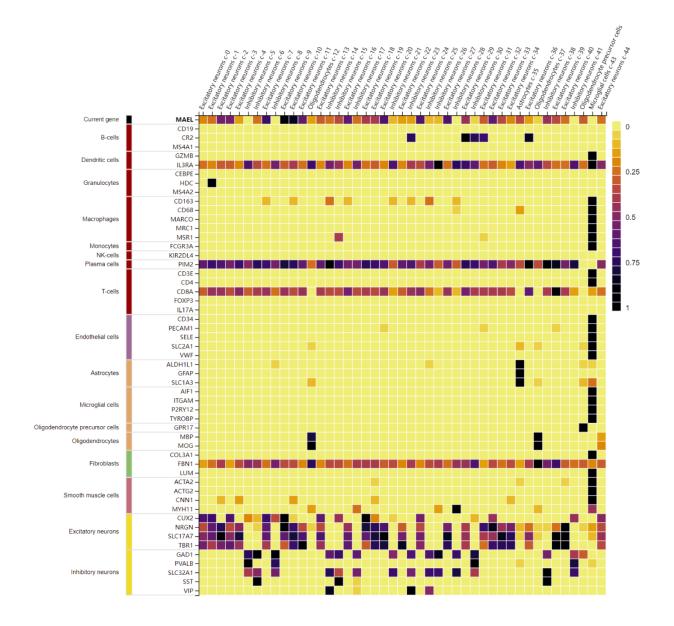
Supplementary Figure 3. mRNA expression of *MAEL* and cell type markers in different single cell type clusters of normal placenta (E-MTAB-6701). Color legend: fraction of the highest expression.



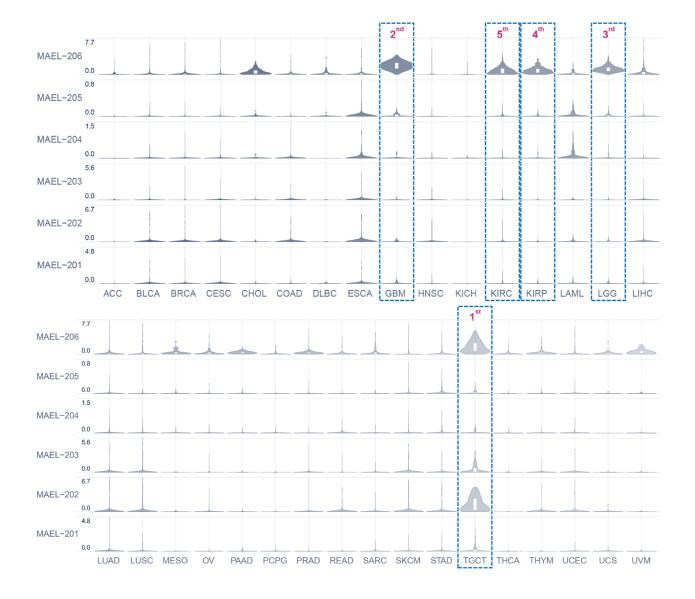
Supplementary Figure 4. mRNA expression of *MAEL* and cell type markers in different single cell type clusters of normal heart muscle (GSE109816). Color legend: fraction of the highest expression.



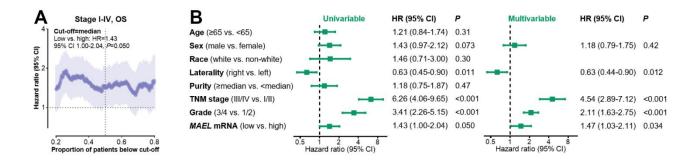
Supplementary Figure 5. mRNA expression of *MAEL* and cell type markers in different single cell type clusters of normal kidney (GSE131685). Color legend: fraction of the highest expression.



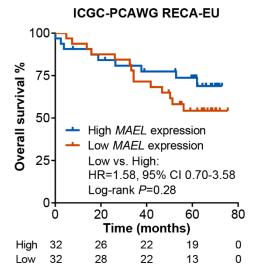
Supplementary Figure 6. mRNA expression of *MAEL* and cell type markers in different single cell type clusters of normal brain (Allen brain map). Color legend: fraction of the highest expression.



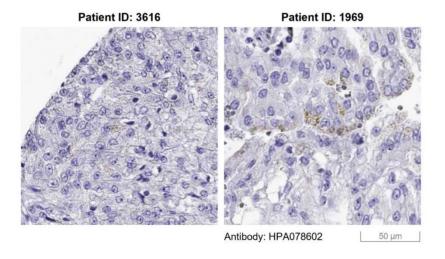
Supplementary Figure 7. mRNA expression of the six transcripts of MAEL among 33 types of cancer (unit: log2[TPM+1]).



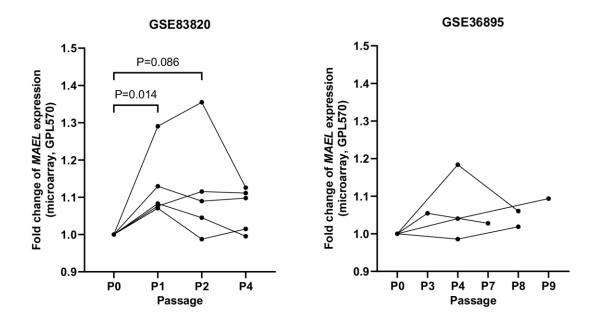
Supplementary Figure 8. Association between *MAEL* expression and recurrence-free survival in the clear cell renal cell carcinomas of the TCGA-KIRC cohort.



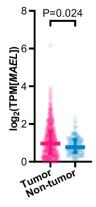
Supplementary Figure 9. Association between *MAEL* expression and overall survival in the clear cell renal cell carcinomas of the ICGC-PCAWG RECA-EU cohort.



Supplementary Figure 10. Immunohistochemical staining of *MAEL* in ccRCC samples.



Supplementary Figure 11. Expression of MAEL in ccRCC xenograft models.



Supplementary Figure 12. *MAEL* expression in clear cell renal cell carcinomas (TCGA-KIRC) and normal kidneys (TCGA-KIRC and GTEx).

E-MTAB-3267 (first-line sunitinib) Progression-free survival % 22 22 25 20 High MAEL expression Low MAEL expression Low vs. High: HR=0.56, 95% CI 0.30-1.06 Log-rank *P*=0.064 50 0 0 20 40 60 80 Time (months) High 26 10 3 0 0 Low 27 4 3 0 0

Supplementary Figure 13. High *MAEL* expression was associated with favorable progression-free survival on first-line sunitinib in advanced clear cell renal cell carcinomas (E-MTAB-3267).