



a\_embryo Tang\_embryo\_D6\_PE  
a\_embryo Tang\_embryo\_D12\_EPI  
colon\_Adult Goblet\_2  
colon\_Adult OLFM4\_High  
a\_embryo Tang\_embryo\_D6\_EPI  
colon\_Adult Endocrine  
colon\_Adult Enter\_2  
colon\_Adult Goblet\_1  
colon\_Adult Enter\_1  
G\_Sintes\_7 Secretory Progenitor  
G\_Sintes\_9 Enteroendocrine  
im\_embryo Pre\_EmbryoPrelineage  
im\_embryo Pre\_Embryo\_trophectoderm\_mural  
im\_embryo Pre\_Embryo\_trophectoderm\_polar  
im\_embryo Pre\_Embryoepiblast  
im\_embryo Pre\_Embryoprimitive\_endoderm  
a\_embryo Tang\_embryo\_D10\_TE  
a\_embryo Tang\_embryo\_D12\_TE  
a\_embryo Tang\_embryo\_D6\_TE  
a\_embryo Tang\_embryo\_D8\_TE  
a\_embryo Tang\_embryo\_D8\_PE  
a\_embryo Tang\_embryo\_D8\_EPI  
a\_embryo Tang\_embryo\_D10\_EPI  
colon\_Adult Immune  
colon\_Adult Mesenchymal  
colon\_Adult Enter\_3  
colon\_Adult MKI67\_High  
G\_Sintes\_15 LGR5+ Stem Cells  
G\_Sintes\_10 Enterocyte Cell  
G\_Sintes\_11 Paneth-Like Cell  
G\_Sintes\_4 PROM1Low Progenitor  
G\_Stomach\_4 Parietal Progenitor  
G\_Stomach\_1 PROCRHigh Progenitor  
G\_Stomach\_3 HES1High Progenitor  
G\_Sintes\_5 Enterocyte  
G\_Sintes\_1 Tuft Progenitor  
G\_Sintes\_2 PROCR+ Progenitor  
G\_Sintes\_8 LGR5+ Stem Cells  
G\_Sintes\_6 Secretory Progenitor  
G\_Esophagus\_1 Ciliated Epithelial  
G\_Stomach\_5 Pit Progenitor  
G\_Stomach\_6 Pit Cell  
a\_embryo Tang\_embryo\_D10\_PE  
a\_embryo Tang\_embryo\_D12\_PE  
G\_Esophagus\_2 KRT6B+ Secretory Progenitor  
G\_Esophagus\_3 FGFR1LowNME5- Epithelial  
G\_Esophagus\_4 FGFR1HighNME5- Epithelial  
G\_Sintes\_7 Goblet Progenitor  
G\_Sintes\_8 Goblet Cell  
G\_Sintes\_6 Goblet Cell  
G\_Sintes\_3 Enterocyte Progenitor -1  
G\_Sintes\_4 Enterocyte Progenitor -2  
G\_Stomach\_2 Progenitor at Neck  
G\_Sintes\_2 MKI67+ Progenitor  
G\_Sintes\_9 Enteroendocrine  
G\_Sintes\_1 DCLK1+ Progenitor  
G\_Sintes\_3 PROM1High Progenitor



### CellType

Normal cells  
Tumor cells

### Cell\_info

log2.NM.epithelial.COUNT.normal.uniGene  
log2.Tang.colon.cancer.FPKM.500  
log2.Tang.colon.cancer.TPM.688  
log2.Tumor.epithelial.COUNT.normal.uniGene