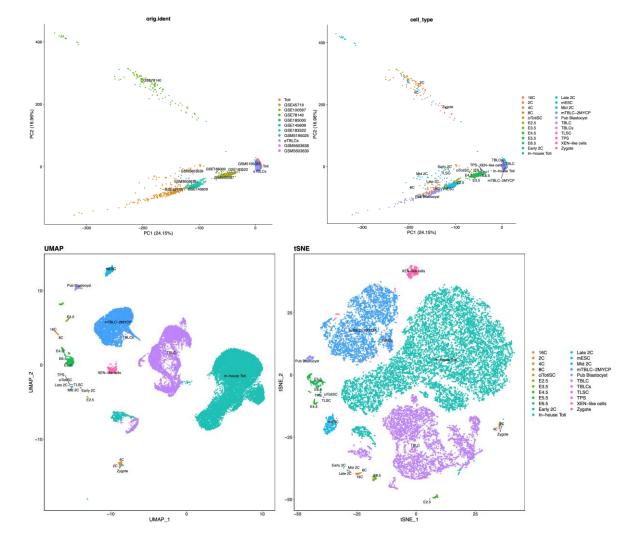


过滤前细胞数: 14244 过滤后细胞数: 13936

过滤比例:2%

分析目的: 我们in house toti cells (10X) 是否其他细胞相比, 其整体表达谱更类似于小鼠2C胚胎

1.UMAP/tSNE/PCA分析, 胚胎公共数据: Zygote, Early 2C, Middle 2C, Late 2C, 2C, 4C, 8C, 16C(GSE45719), E3.5, E4.5, E6.5(GSE100597), E6.75, 对照: mESC(GSE78140), 其他已经published 细胞公共数据: ciTotiSC (GSE:), L-EPSC (GSE145609), D-EPSC (GSE145609), TBLC (GSM5195025), 2MYCP-TBLC (GSM7798458), TLSC(GSM5603638, GSM5603639), TPS (GSE183522)。

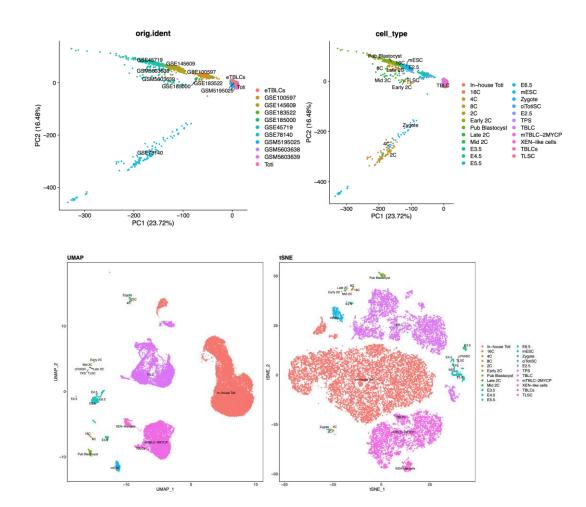


Merge data

Correction Method: ComBat_seq

分析目的: 我们in house toti cells (10X) 是否其他细胞相比, 其整体表达谱更类似于小鼠2C胚胎

1.UMAP/tSNE/PCA分析, 胚胎公共数据: Zygote, Early 2C, Middle 2C, Late 2C, 2C, 4C, 8C, 16C(GSE45719), E3.5, E4.5, E6.5(GSE100597), E6.75, 对照: mESC(GSE78140), 其他已经published 细胞公共数据: ciTotiSC (GSE:), L-EPSC (GSE145609), D-EPSC (GSE145609), TBLC (GSM5195025), 2MYCP-TBLC (GSM7798458), TLSC(GSM5603638, GSM5603639), TPS (GSE183522)。

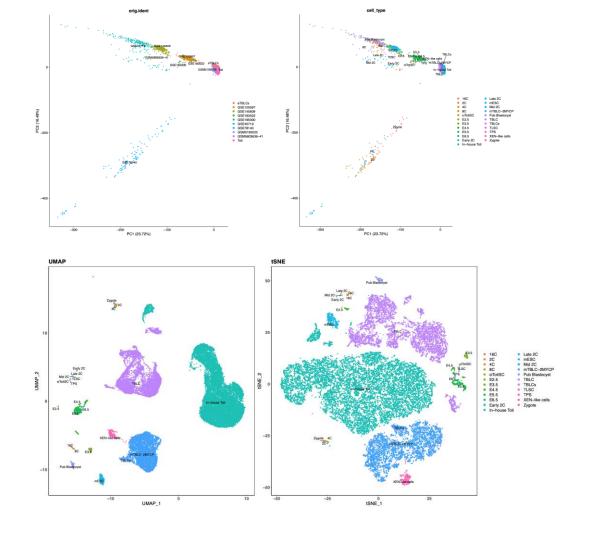


Merge data

Correction Method: Harmony

分析目的: 我们in house toti cells (10X) 是否其他细胞相比, 其整体表达谱更类似于小鼠2C胚胎

1.UMAP/tSNE/PCA分析, 胚胎公共数据: Zygote, Early 2C, Middle 2C, Late 2C, 2C, 4C, 8C, 16C(GSE45719), E3.5, E4.5, E6.5(GSE100597), E6.75, 对照: mESC(GSE78140), 其他已经published 细胞公共数据: ciTotiSC (GSE:), L-EPSC (GSE145609), D-EPSC (GSE145609), TBLC (GSM5195025), 2MYCP-TBLC (GSM7798458), TLSC(GSM5603638, GSM5603639), TPS (GSE183522)。



Raw Merge data

Correction Method: None

2. Specific genes 表达: totipotent genes, 2C specific genes及maternal及ZGA基因 在zygote/E2C/M2C/L2C高表达的基 因在我们in house细胞中也表达,在mESC中高表达的pluripotent基因在我们的细胞中低表达。

这些基因set有对应的gene list

Merge data **Correction Method:** ComBat_seq corrected marker heatm

ap_ComBat_seq.pdf

Merge data

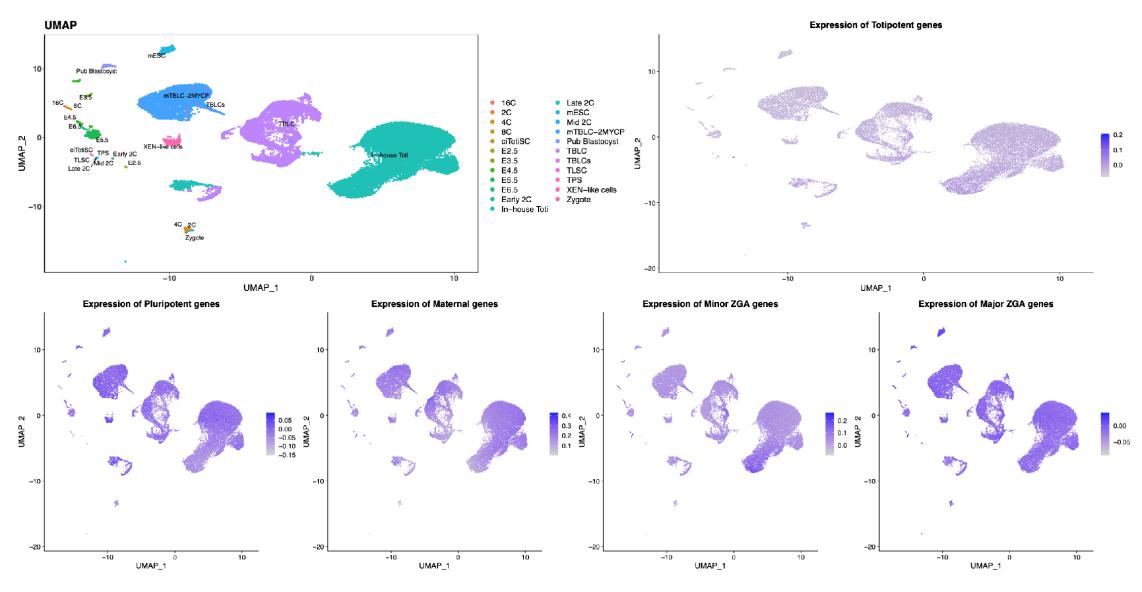
Correction Method: Harmony

corrected_marker_heatm ap_Harmony.pdf

Raw Merge data Correction Method: None

corrected_marker_heatm ap_raw_merge_data.pdf

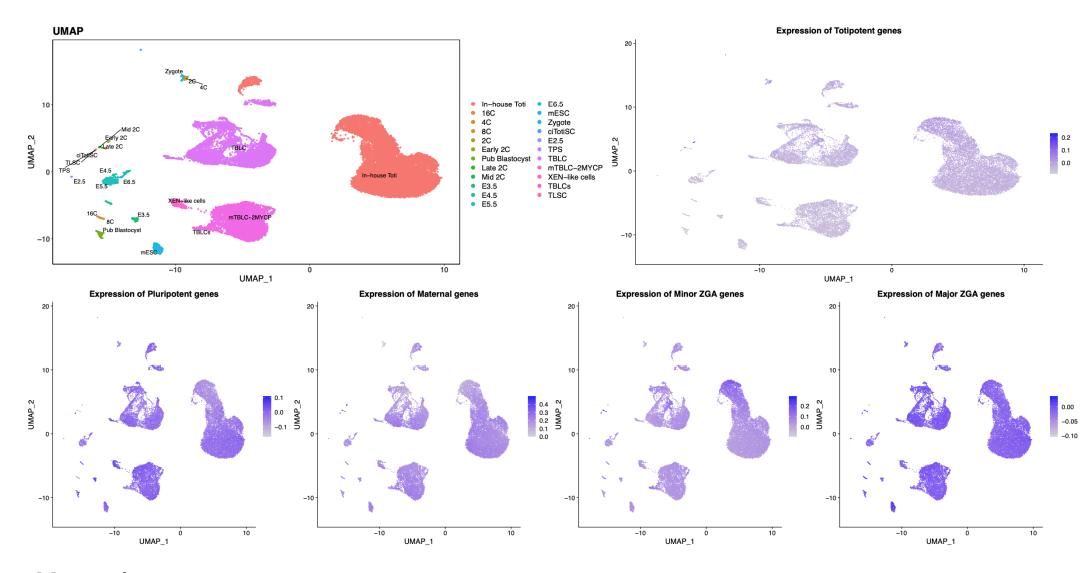
2. UMAP或者tSNE分析 totipotent genes 及pluripotent genes 表达情况



Merge data
Correction Method: ComBat_seq

specific_feature_plot_ComBat_seq.pdf

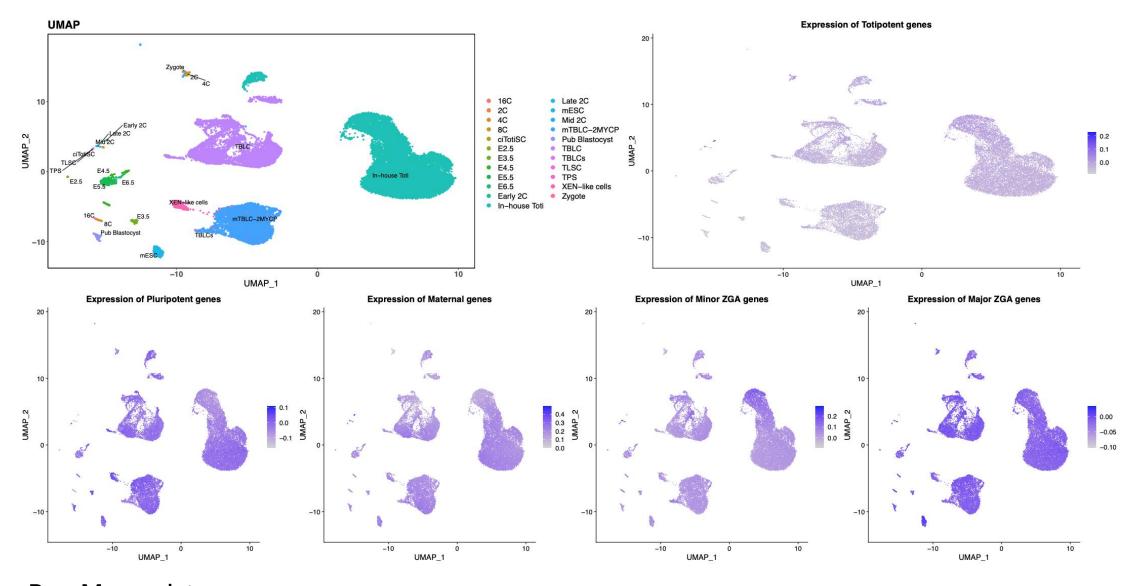
2. UMAP或者tSNE分析 totipotent genes 及pluripotent genes 表达情况



Merge data Correction Method: Harmony

 $specific_feature_plot_Harmony.pdf$

2. UMAP或者tSNE分析 totipotent genes 及pluripotent genes 表达情况



Raw Merge data Correction Method: None

specific_feature_plot_raw_merge_data.pdf