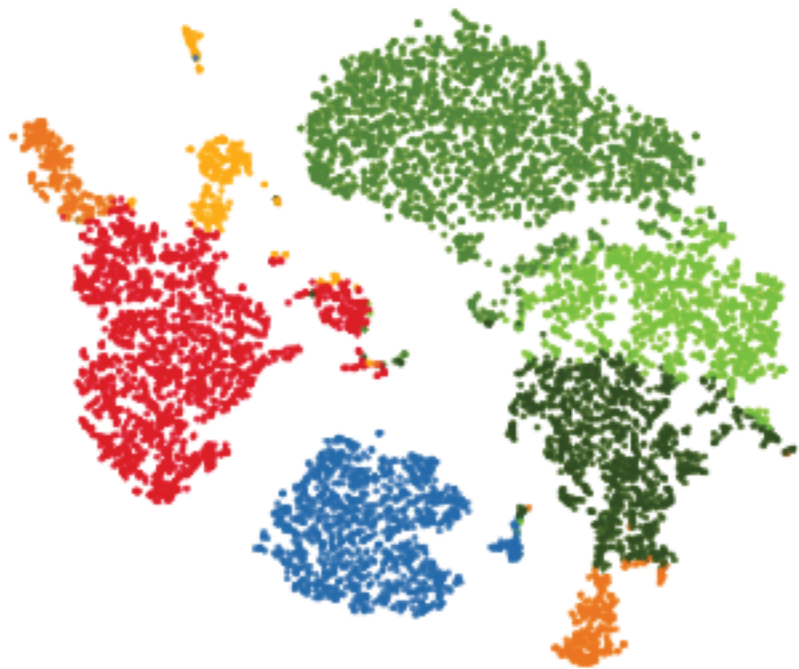






The usual alternative manual labor



Cluster  
cells

CD14



Find specific  
genes



Read papers  
about those genes



## Cell Type Assignments



File Edit View Insert Format

Undo Redo Print Paste 100% \$ % .0

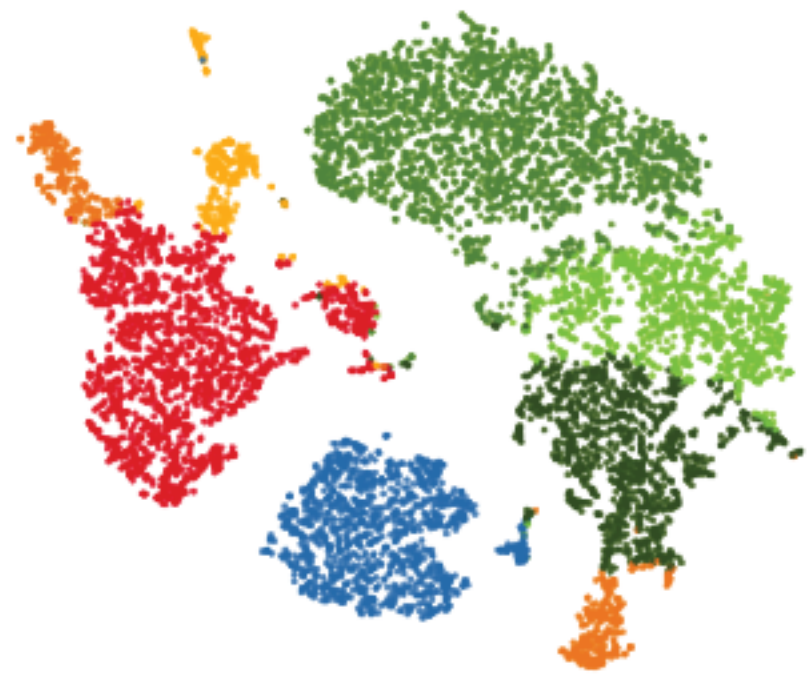
*fx*

	A	B	C
1	cluster	subset_cl	predicted_cell_type
2	1	1	Endothelial cells
3	1	2	Unknown
4	1	3	Endothelial cells
5	2	1	T cells
6	3	1	Hepatocytes
7	4	1	B cells
8	4	2	B cells
9	4	3	B cells

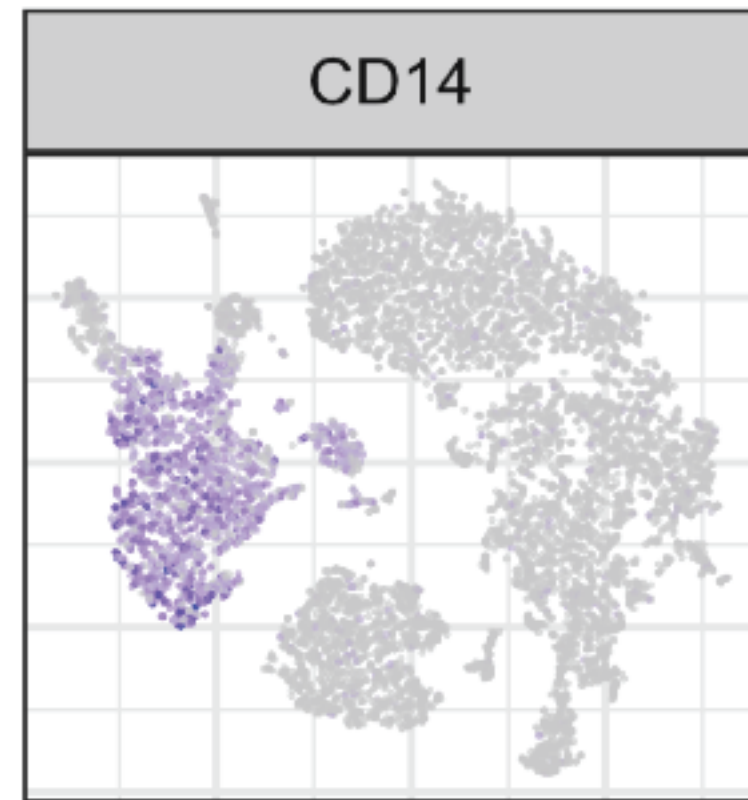
Take your best  
guess for each cluster

# The usual cell annotation workflow involves lots of manual labor

---



Cluster  
cells



Find specific  
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Read papers  
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Cell Type Assignments			
File Edit View Insert Format			
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Take your best  
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# sci-RNA-seq resolves nearly all the cell types in the worm

