

Confusion Matrix of ATAC Cell Types Predicted from RNA Neighbors

Actual ATAC Cell Type	Predicted RNA Cell Type																					
	Naive CD20+ B -	Transitional B -	CD14+ Mono -	MK/E prog -	Proerythroblast -	CD4+ T naive -	CD8+ T -	Lymph prog -	ILC -	CD8+ T naive -	NK -	CD4+ T activated -	Normoblast -	cDC2 -	HSC -	CD16+ Mono -	G/M prog -	Erythroblast -	B1 B -	pDC -	Plasma cell -	ID2-hi myeloid prog -
Naive CD20+ B -	72	2	0	12	2	1	0	0	0	0	0	0	1	0	0	167	0	0	0	2	0	1
Transitional B -	0	1550	15	0	0	0	0	0	4	1	0	0	0	1	5	0	0	0	1	0	0	0
CD14+ Mono -	0	86	157	0	0	1	0	0	0	0	0	0	0	0	2	0	0	0	0	0	0	0
MK/E prog -	0	0	1	430	67	248	0	0	0	0	0	0	2	0	17	0	0	0	0	0	0	0
Proerythroblast -	0	1	1	278	258	80	0	0	0	0	0	0	0	0	2	1	0	0	0	0	0	2
CD4+ T naive -	1	2	0	51	4	1511	0	0	0	0	0	0	0	0	163	0	0	0	1	0	0	0
CD8+ T -	0	0	0	64	82	7	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1
Lymph prog -	0	3	0	2	0	1	0	569	0	0	0	0	0	0	23	0	33	0	52	0	0	0
ILC -	0	72	0	0	0	1	0	1	26	57	0	0	3	12	0	0	0	4	0	0	0	4
CD8+ T naive -	0	10	0	0	0	1	0	0	11	111	0	0	9	4	1	0	0	0	4	0	0	5
NK -	0	5	7	1	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0	0	0
CD4+ T activated -	0	0	1	0	0	17	0	0	0	0	0	0	0	0	118	0	0	0	0	0	0	0
Normoblast -	0	1	1	9	6	0	0	0	1	23	0	0	115	0	3	7	0	0	1	30	0	76
cDC2 -	0	1	0	0	0	0	0	0	5	27	0	0	0	25	1	0	0	0	45	0	0	0
HSC -	0	2	1	4	0	36	0	0	0	2	0	0	0	0	933	0	2	0	0	0	0	0
CD16+ Mono -	19	3	2	3	4	0	0	1	0	0	0	0	3	0	0	678	0	0	0	16	0	3
G/M prog -	0	0	0	2	1	0	0	107	0	0	0	0	0	0	5	0	149	0	0	0	0	0
Erythroblast -	5	2	5	13	7	1	0	0	0	0	0	0	3	0	0	8	0	0	3	0	1	0
B1 B -	0	6	0	0	0	0	0	43	1	2	0	0	0	3	10	0	0	0	275	0	0	0
pDC -	1	2	1	3	1	1	0	0	0	2	0	0	60	0	0	83	0	0	0	229	0	24
Plasma cell -	2	83	20	0	0	0	0	0	7	2	0	0	0	1	0	0	0	0	0	4	10	
ID2-hi myeloid prog -	0	8	4	4	0	0	0	0	0	9	0	0	23	2	0	1	0	0	0	3	0	137