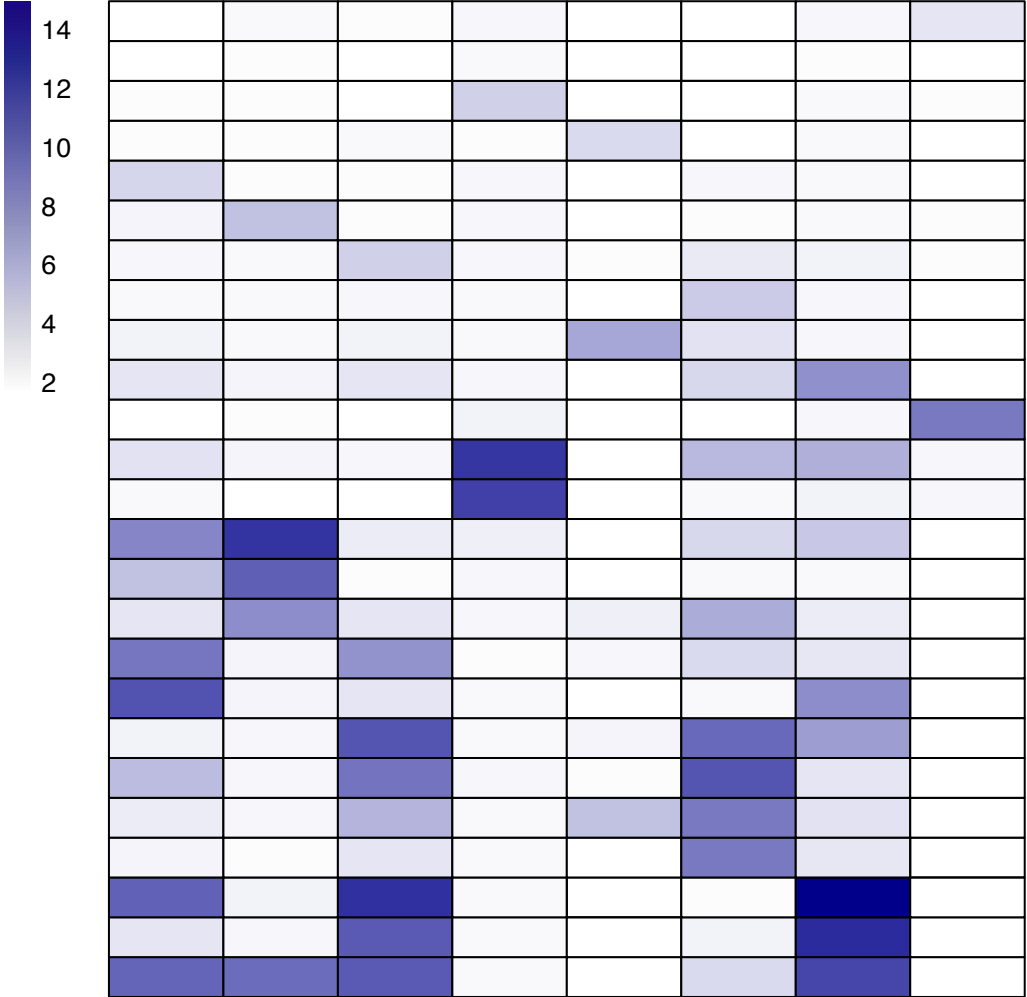


25 Joint Epigenetic States



	H1	1 (3.79%)	1 (4.98%)
	Q	0 (83.89%)	0 (79.27%)
	Pc1	3 (2.4%)	3 (2.49%)
	T1	2 (4.71%)	2 (6.41%)
	N	5 (0.9%)	5 (0.47%)
	C	9 (0.21%)	9 (0.87%)
	A	10 (0.1%)	10 (0.39%)
	E1	4 (1.29%)	4 (2.12%)
	T	8 (0.27%)	8 (0.48%)
	PE	12 (0.15%)	12 (0.09%)
	H	7 (0.39%)	7 (0.36%)
	BPE	18 (0.13%)	18 (0.09%)
	Pc	11 (0.36%)	11 (0.22%)
	PENC	24 (0.03%)	24 (0.03%)
	CN	13 (0.15%)	13 (0.15%)
	EC	22 (0.01%)	22 (0.04%)
	ENA	21 (0.06%)	21 (0.06%)
	PN	19 (0.08%)	19 (0.12%)
	PEA	20 (0.05%)	20 (0.1%)
	ENA	16 (0.08%)	16 (0.12%)
	ETA	17 (0.05%)	17 (0.11%)
	E	6 (0.44%)	6 (0.63%)
	PNA	15 (0.19%)	15 (0.12%)
	PA	14 (0.22%)	14 (0.27%)
	PNCA	23 (0.04%)	23 (0.03%)

Human

Mouse

ATAC  
CTCF  
H3K27ac  
H3K27me3  
H3K36me3  
H3K4me1  
H3K4me3  
H3K9me3

P = Promoter like  
E = Enhancer like  
N = Nuclease accessible  
H = Heterochromatin  
Q = Quiescent  
A = Active  
C = CTCF  
B = Bivalent  
T = Transcribed