	PND15	ATAC-seq	PNW21		PNW8 H3K4me		PNW8 H3K27me3	PND7 BS	PND14 BS	PNW8 BS		
	Zin	0.8	XXXX	1.5 -	3	2.5	white was	0.5	1.5 -	1 -		
		04		0.5		VA GINGO			V0	0.2		l
Catagony 1 (H3K4ma2)			鬟		季草				- =			
Category 1 (H3K4me3)		i										
Category 1 (H3K27ac)						1						
	- 1											
Category 1 (No H3 mark)		-				1-						
		-				1						ATAC-seq (log ₂ RPKM)
												2
Category 2 (H3K4me3)			亖									1
												0
Category 2 (H3K27ac)										*		ATAC-seq (log ₂ FC)
Category 2 (H3K27me3)		-										4
		F				l I						-0
		<u> </u>				1 1						ChIP-seq (log ₂ RPKM)
Category 2 (No H3 mark)				-					+ 2			 5
	= 1											2
		Ī										0
Category 3 (No H3 mark)		<u> </u>										BS (DNAme)
Category 5 (H3K27ac)	-	F	-					reas Indian				0.5
												0
												RNA-seq (log ₂ TMM FC)
		Ť. – –		F.		i				House, and		4
Category 5 (No H3 mark)		1				1						0
		1		1								= -4
		Ī										RNA-seq (log ₂ FC)
		1										0
Category 5 (H3K27me3)												- 5
]
Category 5 (H3K4me3)		l		 		1						
Category 6 (H3K4me3)												
		<u> </u>				i						
Category 6 (No H3 mark)		i i			-1-2	i						
	-1kb M	lid 1kb		lid 1kb	-1kb Mid 1k		o -1kb Mid 1kb		-1kh Mid 1kh	-1kh Mid 1kh		J
	- IND IV	IIG IKD	- IND IV	iid IND	- IND IVIIG IF	D-IND WING IKE	J-IND IVIIG TKD	- IND WING TRD	- IND WING TRD	- IND WING TRD		
	Comparison (ATAC-seq)				ATAC-seq) Ex	Expression (RNA-seq) Comparisor		on (RNA-seq)				
	PNW21 vs PND15				PND15	PND8 PND15 PNW2 ²		15 vs PND8 21 vs PND15				
						PNU15 PNW21	■ PNW21	VS PNU15				