	PND15	ATAC-seq	PNW21		PNW8	H3K4me3	PNW8 H3K27ac		PNW8 H3K27me3		PND7 BS	PND14 BS	PNW8 BS		
		0.6	1	1.5		3 -	2		\ 	2.74	W0-	1.5	0.2		l
Category 1 (H3K4me3)															
Category 1 (H3K27ac)															
Category 1 (No H3 mark)							1 1 1 1 1								ATAC-seq (log₂ RPKM) ■ 2
Category 2 (H3K4me3)															1 0
Category 2 (H3K27ac)				- 8-3		1			jt-		7-5		10 No.		ATAC-seq (log ₂ FC)
Category 2 (H3K27me3)						4 3			•						4
Category 2 (No H3 mark)							 								ChIP-seq (log ₂ RPKM) 5 2
Category 3 (No H3 mark)															BS (DNAme)
Category 5 (H3K27ac)		f -	-						+-3		Trans.				0.5
Category 5 (No H3 mark)															RNA-seq (log ₂ TMM FC) 4 0 -4 RNA-seq (log ₂ FC) 5 0
Category 5 (H3K27me3)					- 27										-5
Category 5 (H3K4me3)															
Category 6 (H3K4me3)		i	100	12.3	- 3	ž –			+		40.00				į
Category 6 (No H3 mark)		† † i													
		lid 1kb									Mid 1kb	-1kb Mid 1kb	-1kb Mid 1kb		
					Comp	arison (Δ'	TAC-seq) E	xpression (RNA-seq)			Comparison (RNA-seq)				
						W21 vs Pl		PND8	PND8			vs PND8			
								PND15 PNW21			PNW21 vs PND15				