	 			n TE insertions	 	 	 	 Promodomoja UC2 2:PPD7/PPD0 Pro	
								Bromodomain	
								Bromodomain HG3.10:KAIZA/KAIZB Bromodomain HG3.10:KAIZA/KAIZB Gene presence Bromodomain HG3.10:KAIZA/KAIZB Gene presence Bromodomain HG3.10:KAIZA/KAIZB Gene presence Bromodomain HG3.10:KAIZA/KAIZB GENE CRITTIES Bromodomain HG3.10:KAIZA/KAIZB GENE GENE GENE GENE GENE GENE GENE GEN	
								Bromodomain. HG3. 10:KAI 2B/KD18 RWD3/PHIP Bromodomain. HG7. 1:BRWD1/BRWD3/PHIP Bromo	sence main C2H ip WD40 G_3
								Biomodomain 165,2 BA21 ABA21 BBA22/AB. BHO BIOMODOMAIN 165,2 BA21 ABA21 BBA22/AB. BHO BIOMODOMAIN 165,2 BA21 ABA21 BBA22/AB. BHO BIOMODOMAIN 165,2 BA21 ABA21 BBA22/AB. WILLIAM BIOMODOMAIN 165,2 BA21 ABA21 BBA22/AB. MISS BIOMODOMAIN 165,2 BA21 ABA21 BBA22/AB. FLORE BIOMODOMAIN 165,2 BA21 ABA21 BBA22/AB. GOOD BIOMODOMAIN 165,2 BA21 ABA21 BBA22/AB. LAMB BIOMODOMAIN 165,2 BA21 ABA21 BBA22/AB. BAB21	odomain 11 _Acf1_DNA_bd
								Bromodomain HG1 16 like BRD2/BRD3/BRD. Brome Bromodomain HG1 fil6 like BRD2/BRD3/BRD. Gene Bromodomain HG1 fil6 like BRD2/BRD3/BRD. BE1 Bromodomain HG1 fil6 like BRD2/BRD3/BRD. Ark 2 Bromodomain HG1 16 like BRD2/BRD3/BRD. HAT F Bromodomain HG1 16 like BRD2/BRD3/BRD. Integr	domain resence AT11 se_H2C2 aseH_2
								Bromodomain.HG3.7cECR2 Gene presence Bromodomain.HG3.7cECR2 Gene presence Bromodomain.HG3.7cECR2 Bromodomain.BG3.7cECR2 Bromodomain.BG3.7cECR2 Bromodomain.BG3.7cECR2 WHIMI Bromodomain.HG3.7cECR2 HDD Bromodomain.HG3.7cECR2 Bromodo	
								Bromodomain.HG4.7:like:SMARCA2/SMARCA Gen Bromodomain.HG4.7:like:SMARCA2/SMARCA Bron Bromodomain.HG4.7:like:SMARCA2/SMARCA Heli Bromodomain.HG4.7:like:SMARCA2/SMARCA Chro Bromodomain.HG4.7:like:SMARCA2/SMARCA SNE Bromodomain.HG4.7:like:SMARCA2/SMARCA SNE	presence odomain ase_C no 2_N C5HC2H 1_box