								Pr	edic	ted									
		DNA/CACTA	DNA/Mutator	DNA/P	DNA/PIF	DNA/Tc1	DNA/hAT	LINE/CR1	LINE/I	LINE/L1	LINE/R1	LTR/Bel-Pao	LTR/Copia	LTR/ERV	LTR/Gypsy	RC/Helitron			
	DNA/CACTA	3	1	0	0	1	4	0	0	0	0	1	0	0	0	0	Cl	Class Percent	
	DNA/Mutator -	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	Pe		
	DNA/P-	1	3	0	1	0	8	0	0	0	0	1	0	0	0	0			
	DNA/PIF-	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		80%	
	DNA/Tc1-	3	0	0	1	4	6	0	0	1	0	1	0	1	1	0			
h	DNA/hAT -	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0			
Ground Truth	LINE/CR1 -	0	0	0	0	1	3	0	0	0	0	0	0	0	1	0		60%	
	LINE/I -	12	3	0	2	3	9	0	0	17	0	6	1	0	15	0		60%	
	LINE/L1 -	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		40%	
	LINE/R1-	6	1	0	0	0	2	0	0	2	0	0	0	1	19	0			
	LTR/Bel-Pao	7	2	0	1	6	5	0	0	2	0	9	1	1	16	0			
	LTR/Copia -	4	1	0	0	0	3	0	0	1	0	2	0	0	1	0		20%	
	LTR/ERV -	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0			
	LTR/Gypsy-	45	17	0	1	10	33	0	0	6	0	43	3	0	56	0			
	RC/Helitron	1	0	0	0	0	9	0	0	0	0	0	0	0	3	0		0%	
	TERL Fruit Fly Superfamily Confusion Matrix								U / 0										