	Predicted																					
		DNA/CACTA	DNA/Maverick	DNA/P	DNA/Tc1	DNA/Transib	DNA/hAT	LINE/CR1	LINE/I	LINE/L1_L2	LINE/R1	LINE/R2	LINE/RTE	LTR/Bel-Pao	LTR/Copia	LTR/ERV	LTR/Gypsy	RC/Helitron				
Ground Truth	DNA/CACTA -	0	0	0	0	7	0	0	0	0	0	0	0	0	0	0	1	0	Cl	lass		
	DNA/Maverick	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	Pe	rcent		
	DNA/P-	0	0	8	0	0	0	0	0	0	0	0	1	0	0	0	0	1		100%		
	DNA/Tc1	0	0	2	14	0	0	0	0	1	0	0	0	0	0	0	0	0				
	DNA/Transib	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		80%		
	DNA/hAT -	0	0	0	0	0	1	0	0	0	0	0	0	0	0	0	0	0				
	LINE/CR1		0	0	0	0	0	0	0	1	0	0	0	0	0	0	0	0				
	LINE/I -	0	0	0	1	0	0	0	62	2	0	0	4	0	0	1	2	3		60%		
	LINE/L1_L2 -	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		40%		
	LINE/R1	0	0	0	0	0	0	0	2	0	0	0	7	1	3	0	6	0				
	LINE/R2 -	0	0	0	0	0	0	0	0	0	0	0	1	0	0	0	0	0				
	LINE/RTE		0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0				
	LTR/Bel-Pao		0	0	4	0	0	0	2	1	0	0	2	31	0	0	3	2		20%		
	LTR/Copia		0	0	2	0	0	0	0	0	0	0	0	1	7	0	2	1				
	LTR/ERV -		0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0				
	LTR/Gypsy -														3	2						
	RC/Helitron -														0		0	13		0%		
	TEclass2 (0.7) Fruit Fly Superfamily Confusion Matrix																					