									Pr	edic	ted										
		DNA/CMC	DNA/MULE	DNA/Maverick	DNA/P	DNA/TcMar	DNA/hAT	LINE/CR1	LINE/I	LINE/Jockey-I	LINE/L1	LINE/R1	LINE/R2	LTR/Copia	LTR/DIRS	LTR/Gypsy	LTR/Pao	RC/Helitron			
	DNA/CMC -	3	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	Cl	Class	
	DNA/MULE -	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	Pe	rcent 100%	
	DNA/Maverick	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0			
	DNA/P-	0	0	0	10	0	0	0	0	0	0	0	0	0	0	0	0	0			
	DNA/TcMar	0	1	0	0	13	1	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		0070	
ıth	LINE/CR1 -	0	0	0	0	0	0	5	0	0	0	0	0	0	0	1	0	0			
Ground Truth	LINE/I -	1	0	0	0	0	0	1	0	57	0	0	0	0	0	2	0	0		60%	
pur	LINE/Jockey-I		0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0			
ìroı	LINE/L1 -	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		40%	
$\overline{}$	LINE/R1 -	0	1	2	0	0	0	0	0	0	0	16	0	0	0	7	0	0			
	LINE/R2 -	0	0	0	0	0	0	0	0	0	0	0	1	0	0 I	0	0	0			
	LTR/Copia		0	0	0	1	0	0	0	0	0	0	0	10	0	1	1	0			
	LTR/DIRS	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		20%	
	LTR/Gypsy -		0	0	0		1			0		0	0	0	1	216		3			
	LTR/Pao -														0	2	47	0			
	RC/Helitron -	- 0	- 0		_ 0	- 0		0	_ 0	- 0	0	_ 0	_ 0	0	0	0	0	7		0%	
Terrier (0.7) Fruit Fly Superfamily Confusion Matrix																					