		Predicted																
		DNA/P	DNA/TcMar	DNA/Transib	DNA/hAT	LINE/Jockey	LINE/L1_L2	LINE/R1	LINE/R1-LOA	LINE/RTE	LINEJockey	LTR/Copia	LTR/Gypsy	LTR/Pao	RC/Helitron			
Ground Truth	DNA/P -	8	0	0	0	0	0	0	0	0	0	0	0	0	0	Cla	Class Percent	
	DNA/TcMar -	2	14	0	0	0	0	0	0	0	0	0	0	0	0	Pe		
	DNA/Transib -	0	0	6	0	0	0	0	0	0	0	0	0	0	0			
	DNA/hAT -	0	0	0	1	0	0	0	0	0	0	0	0	0	0			
	LINE/Jockey -	0	0	0	0	44	1	0	0	2	0	0	2	0	1		80%	
	LINE/L1_L2 -	0	0	0	0	0	0	0	0	0	0	0	0	0	0			
	LINE/R1 -	0	0	0	0	1	0	0	0	2	0	2	2	0	0		60%	
	LINE/R1-LOA -	0	0	0	0	0	0	0	0	4	0	0	0	0	0			
	LINE/RTE -	0	0	0	0	0	0	0	0	0	0	0	0	0	0		4007	
	LINEJockey -	0	0	0	0	9	0	0	0	0	0	0	0	0	0		40%	
	LTR/Copia -	0	2	0	0	0	0	0	0	0	0	6	0	1	1		20%	
	LTR/Gypsy-	2	6	0	2	8	5	0	0	2	0	2	119	3	0			
	LTR/Pao-	0	2	0	0	1	0	0	0	1	0	0	0	25	1			
	RC/Helitron -	0	0	0	0	0	0	0	0	0	0	0	0	0	10		0%	
TEclass2 (0.9) Fruit Fly Superfamily Confusion Matrix										U / 0								