Supplemental Material for "Coding DNA repeated throughout intergenic regions of the Arabidopsis thaliana genome: Evolutionary footprints of RNA silencing"

Supplemental Table S1. Nucleotide (nt) and Record Number Statistics for Each File.

	3'UTR	5'UTR	Intergenic	Intron	CR		Modified 3'UTR	Modified 5'UTR	Modified Intron	Modified CR
	(TAIR8 3 utr 20080228)	(TAIR8 5 utr 20080228)	(TAIR8 intergenic 20080229)	(TAIR8 intron 20080228)	(TAIR8 cds 20080412)	total				
A	1781005	1051305	17154467	6956962	11356253	38299992	1334617	772103	5355876	9372541
C	928128	768970	7793572	3944349	8041458	21476477	685470	558409	3006940	6636208
G	1076869	612553	7762361	4319539	9444647	23215969	797754	428752	3289810	7786499
T	2233102	1181958	17133876	10204684	10746155	41499775	1692779	843727	7804427	8878292
D	0	0	1	0	0	1	0	0	0	0
K	0	0	63	0	0	63	0	0	0	0
M	0	0	81	0	1	82	0	0	0	0
N	0	0	225278	0	0	225278	0	0	2	3
R	0	0	47	0	0	47	0	0	0	0
S	0	0	33	1	0	34	0	0	0	0
W	0	0	143	0	1	144	0	0	0	0
Y	0	0	93	1	1	95	0	0	0	0
Non-ACGT total	0	0	225739	2	3	225744	0	0	0	0
Total nt	6019104	3614786	50070015	25425536	39588516	124717957	4510620	2602991	1945705	32673540
Records	25273	24267	31353	154240	32825	-	19820	18873	118333	26990