



Additional file 3: C+G content in oak and *Arabidopsis*.

Distribution of C+G content for certain populations of nucleotide sequences (oak solid lines, *Arabidopsis* dotted lines): coding sequences from all gene models (green, TAIR file TAIR10_cds_20110103_representative_gene_model_updated), 5'-UTRs (blue, TAIR file TAIR10_5_utr_20101028), 3'-UTRs (magenta, TAIR file TAIR10_3_utr_20101028), and entire contigs for oak contigs having no gene models (solid red) or all *Arabidopsis* intron sequences (dotted red, TAIR file TAIR10_intron_20101028).