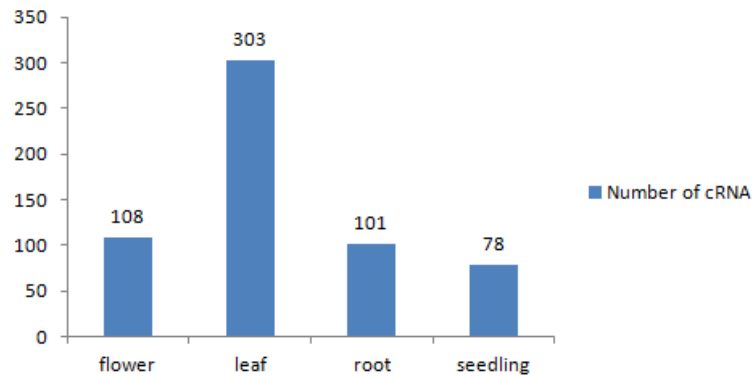
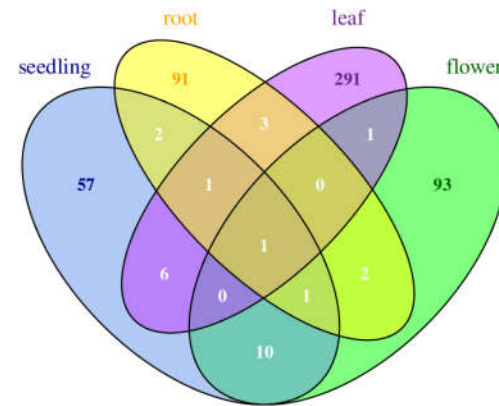


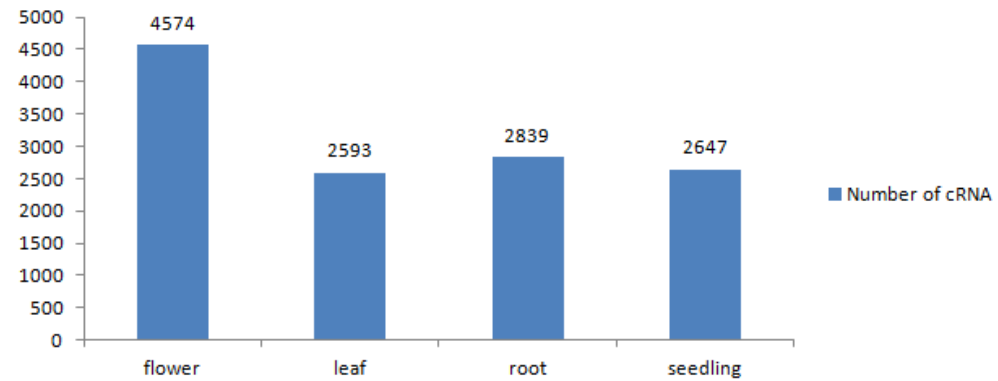
Number of cRNA



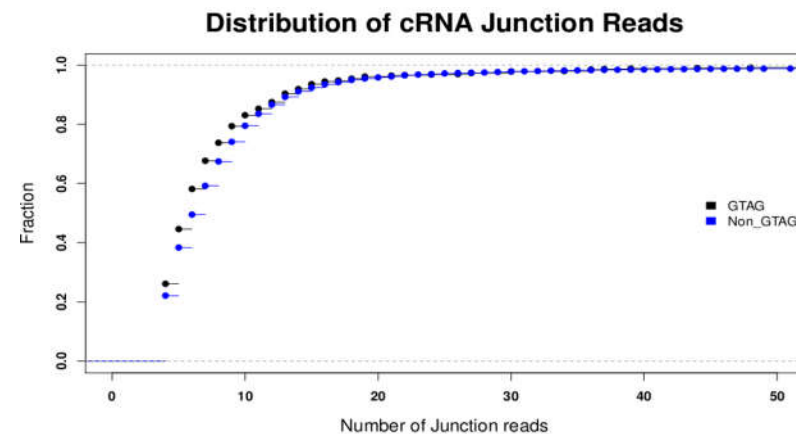
GTAG signal

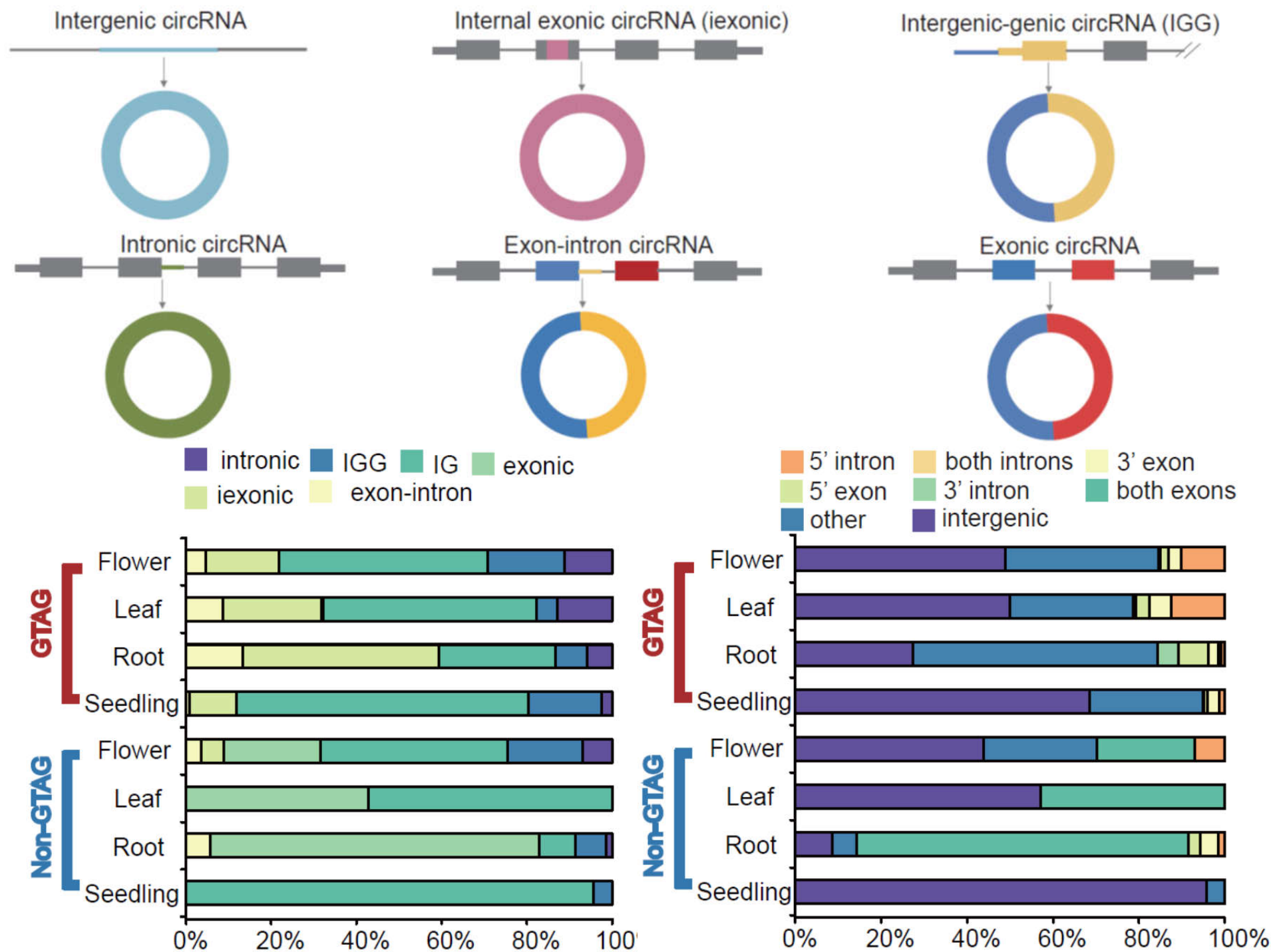


Number of cRNA



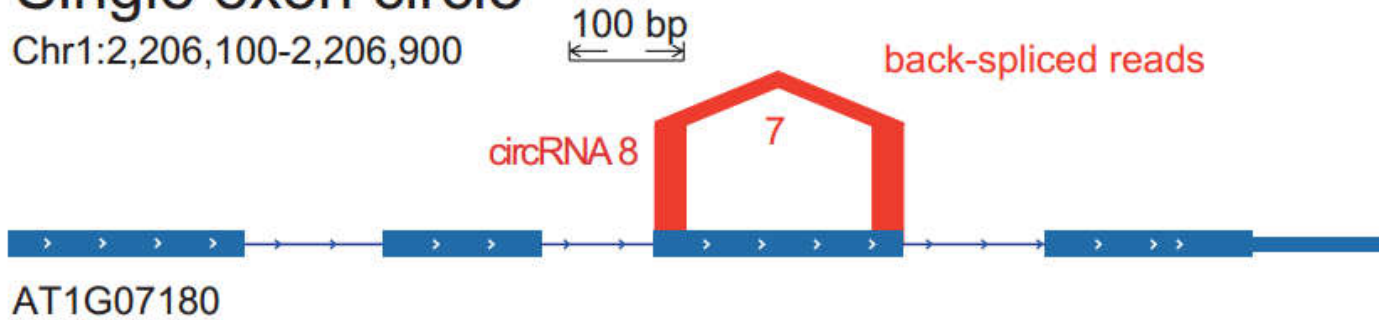
Non-GTAG signal





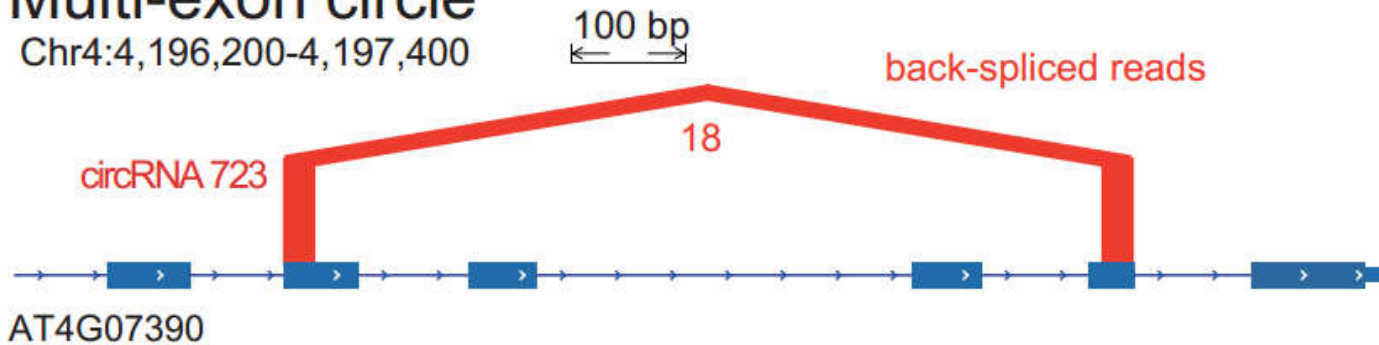
Single exon circle

Chr1:2,206,100-2,206,900



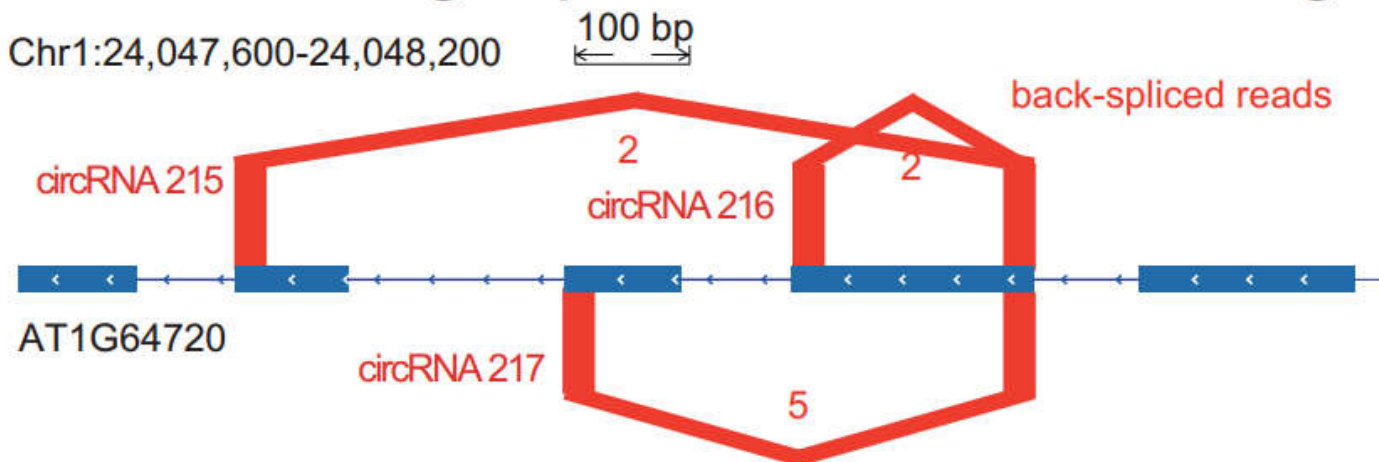
Multi-exon circle

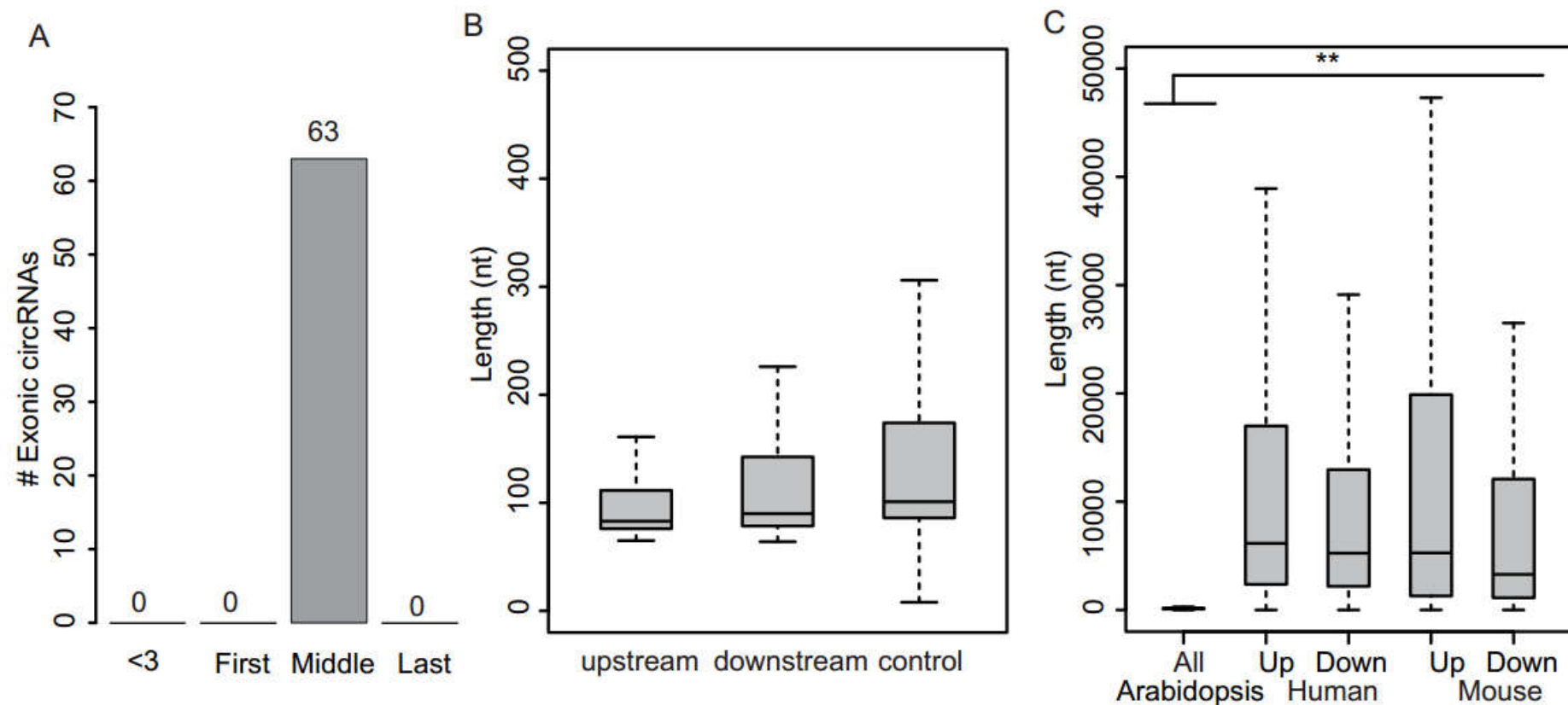
Chr4:4,196,200-4,197,400

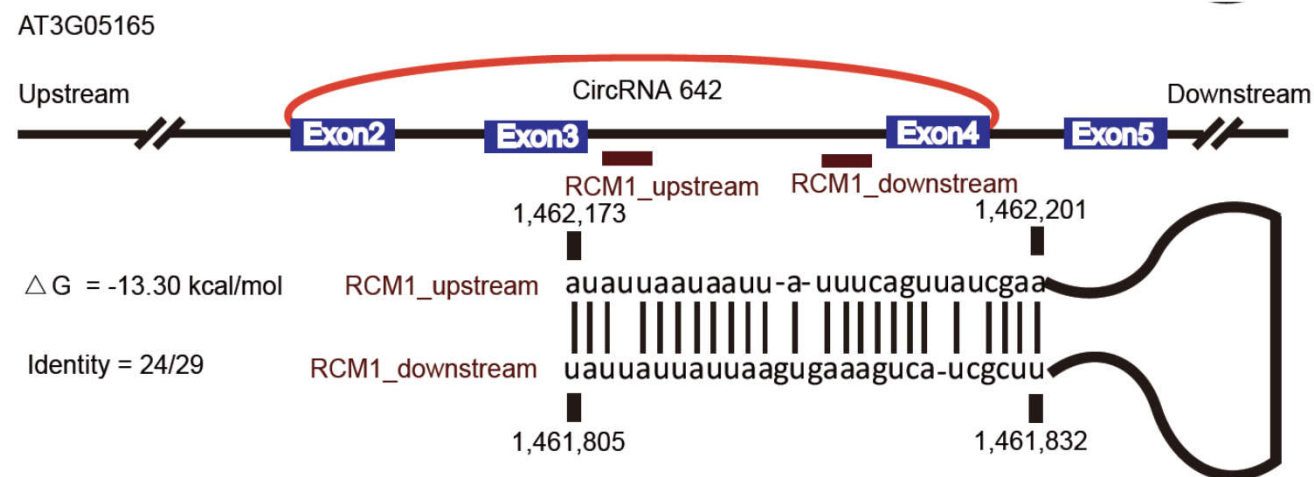
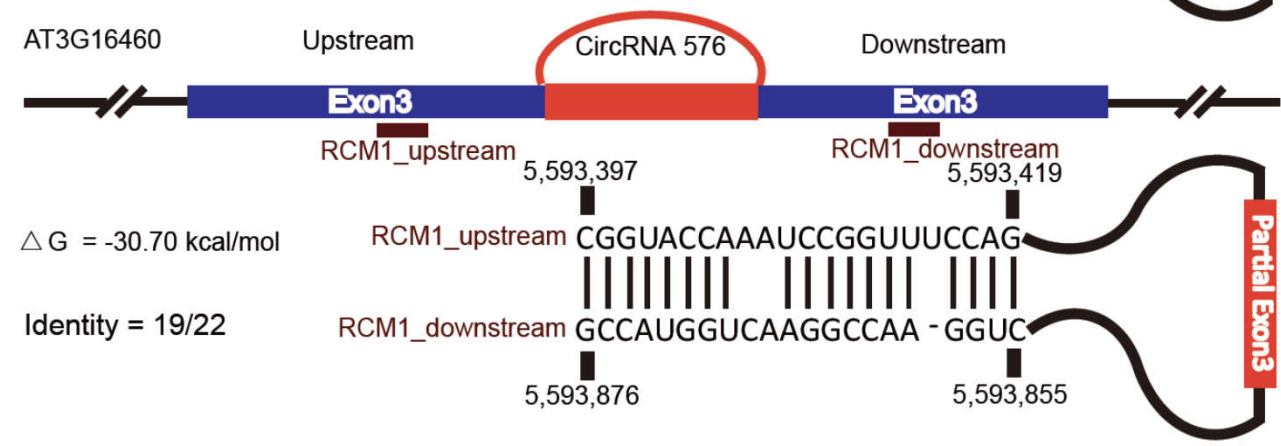
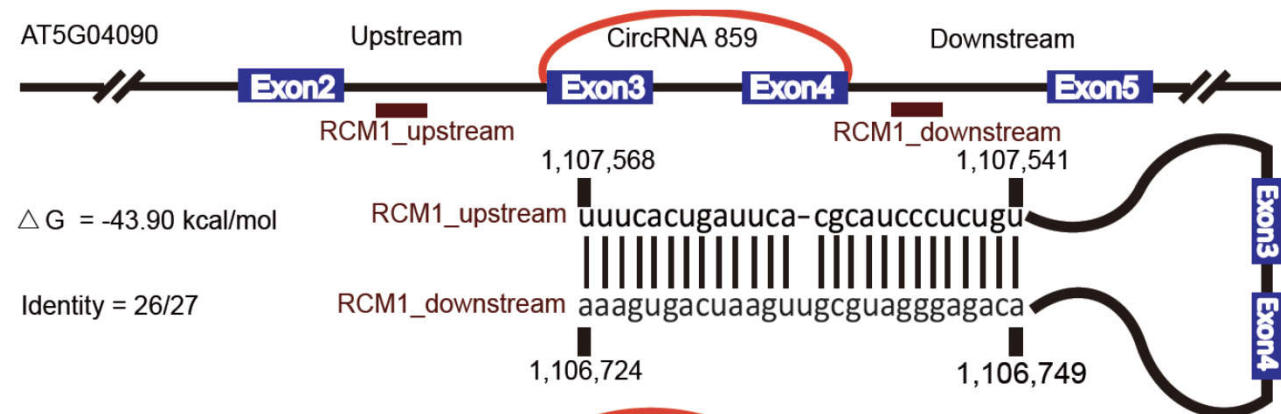


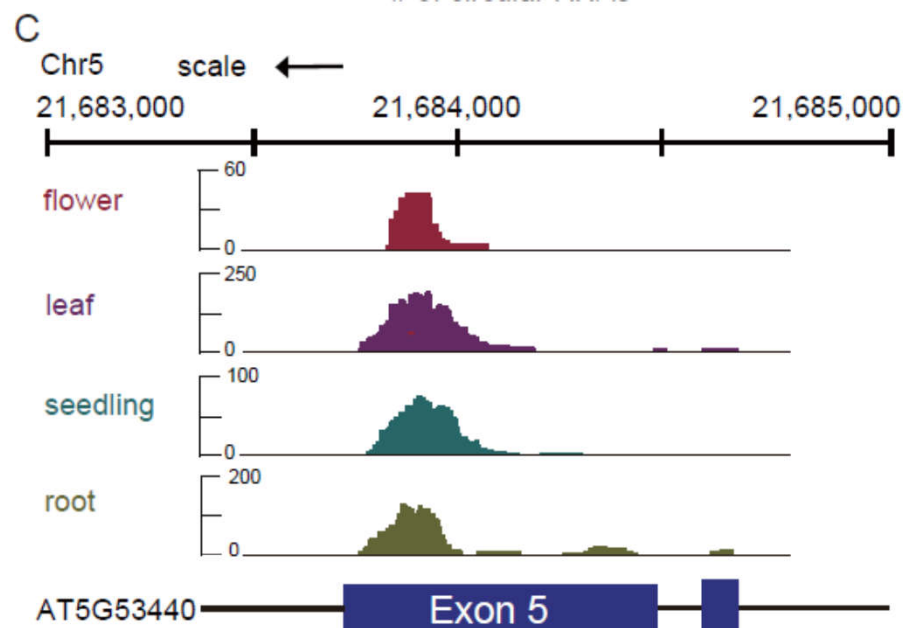
Different exon groups circularized within one gene

Chr1:24,047,600-24,048,200

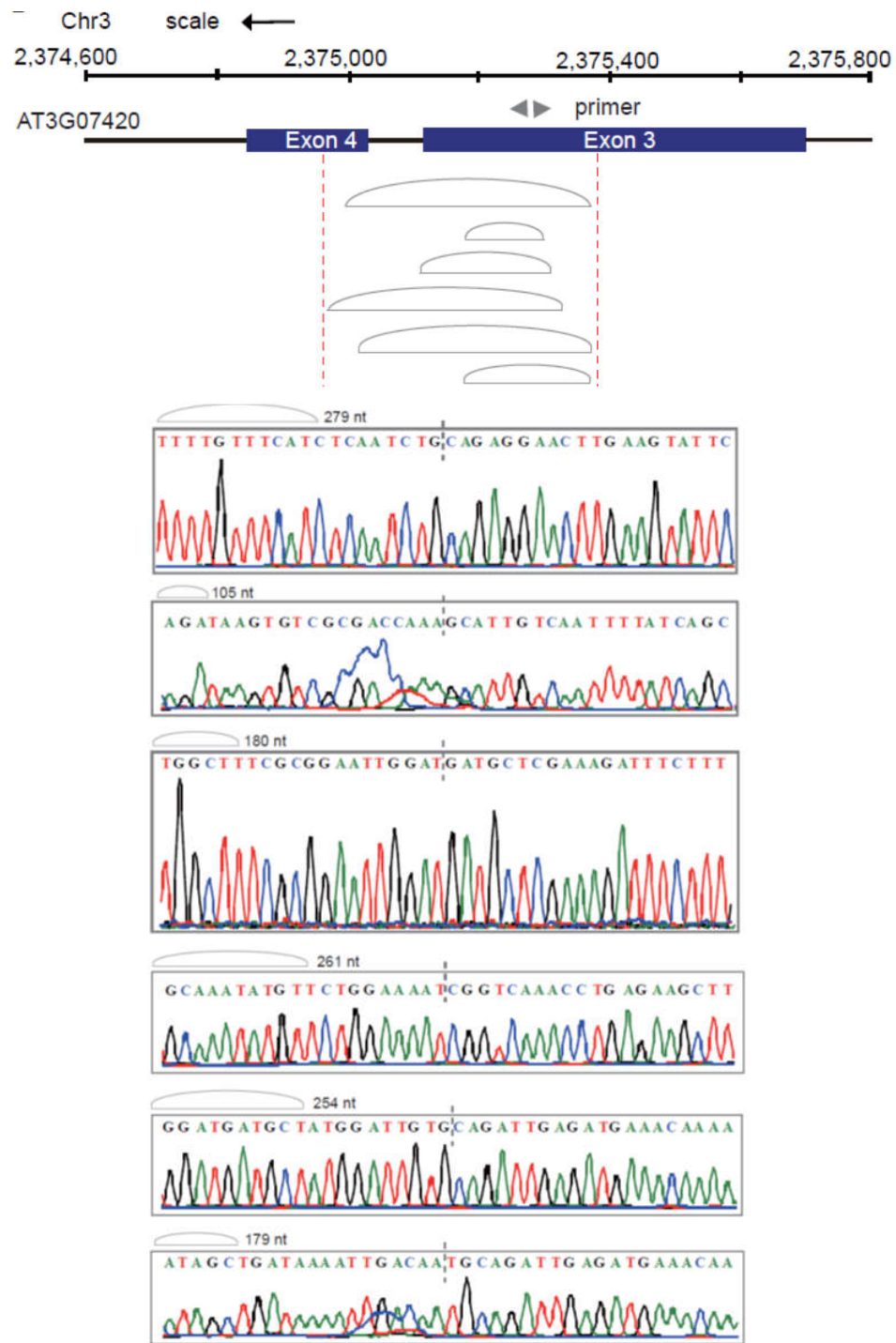




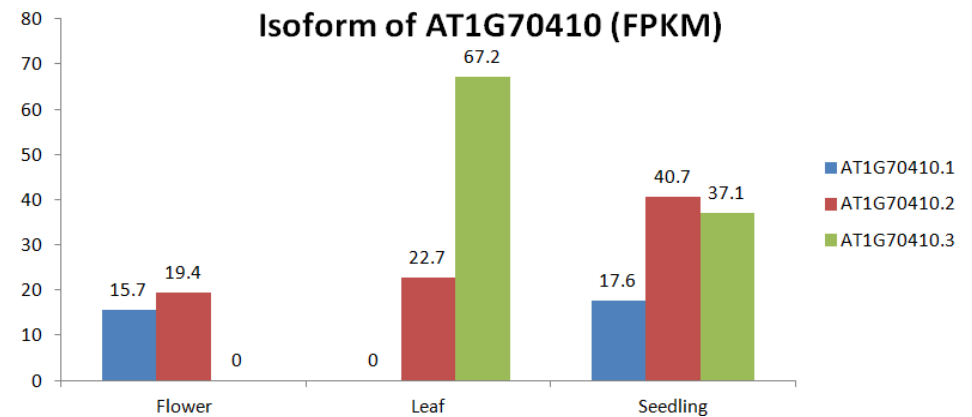
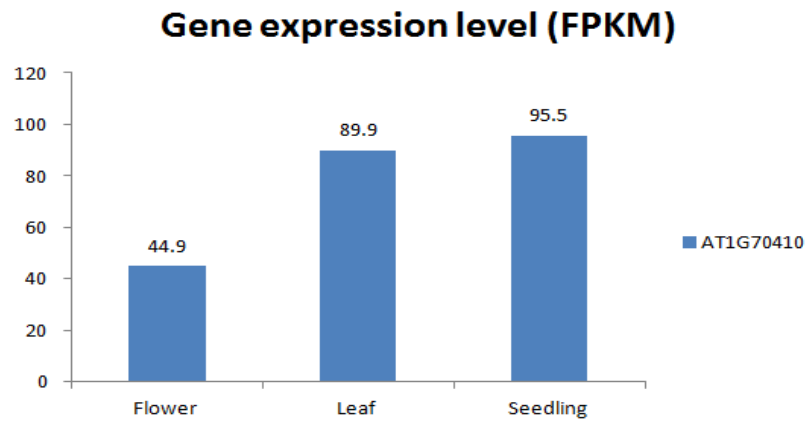
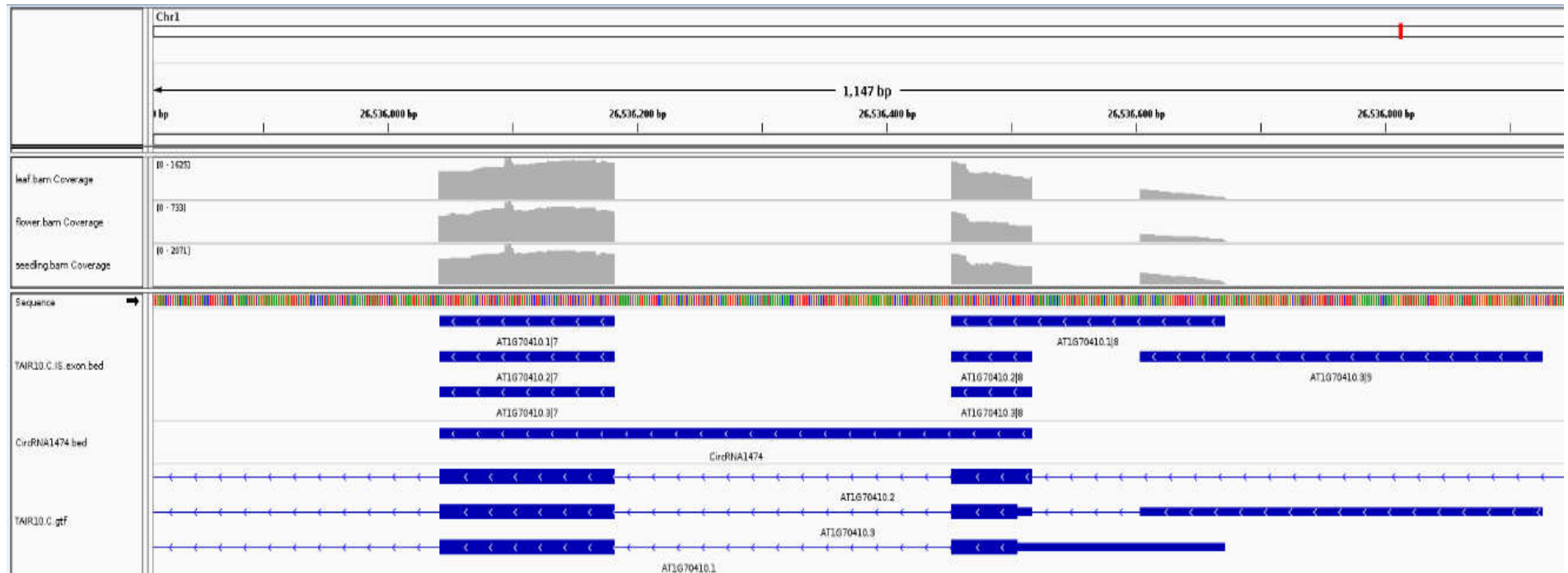




CircRNA892
CircRNA893
CircRNA894
CircRNA895
CircRNA896
CircRNA897
CircRNA898
CircRNA899
CircRNA900
CircRNA901
CircRNA902
CircRNA903
CircRNA904



Exonic circRNAs(CircRNA474) was detected in flower and Leaf but not in seedling



GO enrichment of tissue-specific circRNAs host genes

