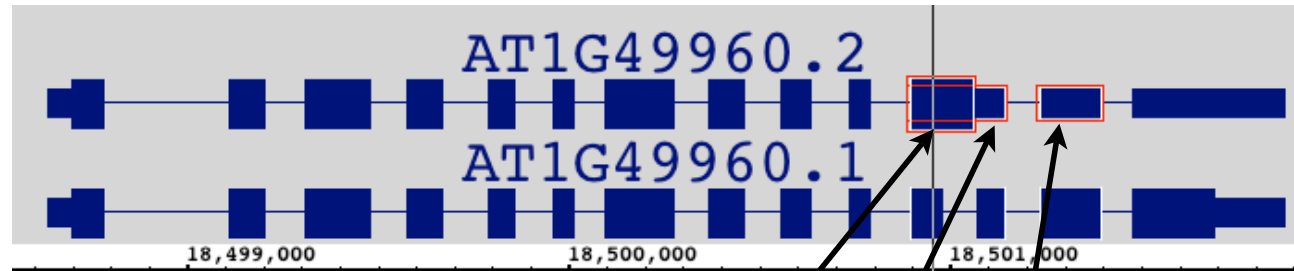


testMeasureEejDistancePlusMultipleExons



A

translated part of
the last coding
region exon
(including stop
codon)

B

last coding region
exon

C

second to last
UTR exon

property	no ID	Selection Info	Search	Slic
name	AT1G49960.2	AT1G49960.2	AT1G49960.2	50.2
id	AT1G49960.2	AT1G49960.2	AT1G49960.2	60.2
chromosome	chr1	chr1	chr1	60.2
start	18,500,896	18,500,896	18,501,234	
end	18,501,062	18,501,147	18,501,395	
length	166	251	161	
type		TAIR9 ALL	TAIR9 ALL	
cds min		18498699	18498699	
cds max		18501062	18501062	
forward		true	true	
score		0.0	0.0	

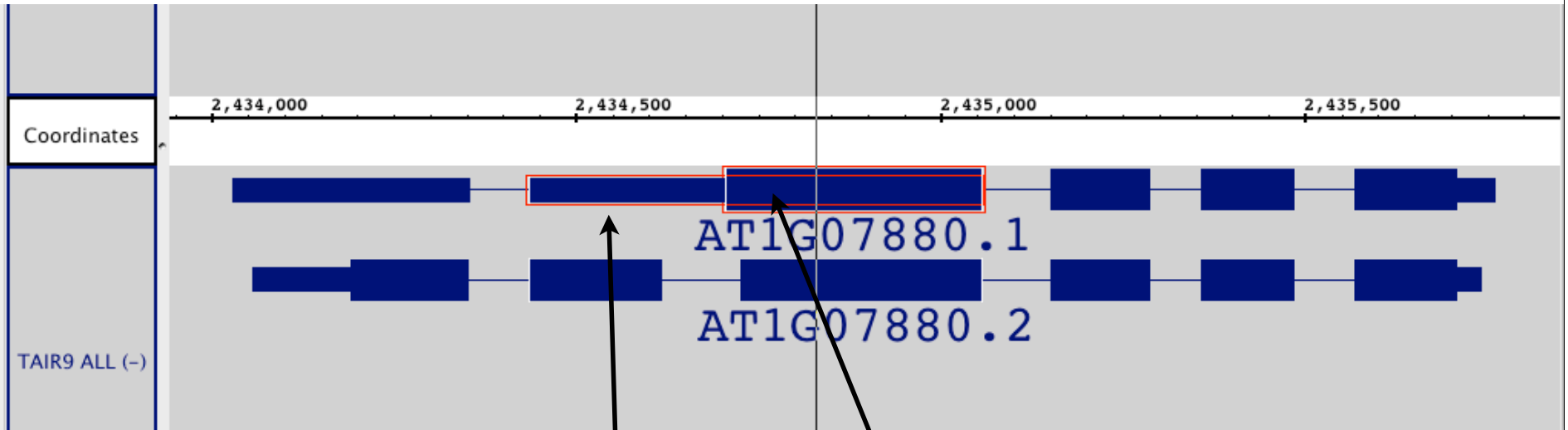
$$\text{EEJ distance} = 251 - 166 + 161$$

B

A

C

testMeasureEejDistanceMinus



A translated part of the last coding region exon (including stop codon)

B the second to last exon (also last translated exon)

Data Access		Selection Info	
property	AT1G07880.1	no ID	
name	AT1G07880.1		
id	AT1G07880.1		
chromosome	chr1	chr1	
start	2,435,059	2,434,705	
end	2,434,437	2,435,059	
length	622	354	
type	TAIR9 ALL		
cds min	2434705		
cds max	2435712		
forward	false		
score	0.0		

2 Selections

B **A**

$$\text{EEJ distance} = 622 - 354$$