Alternative splicing of flowering time gene *FT* is associated with halving of time to flowering in coconut

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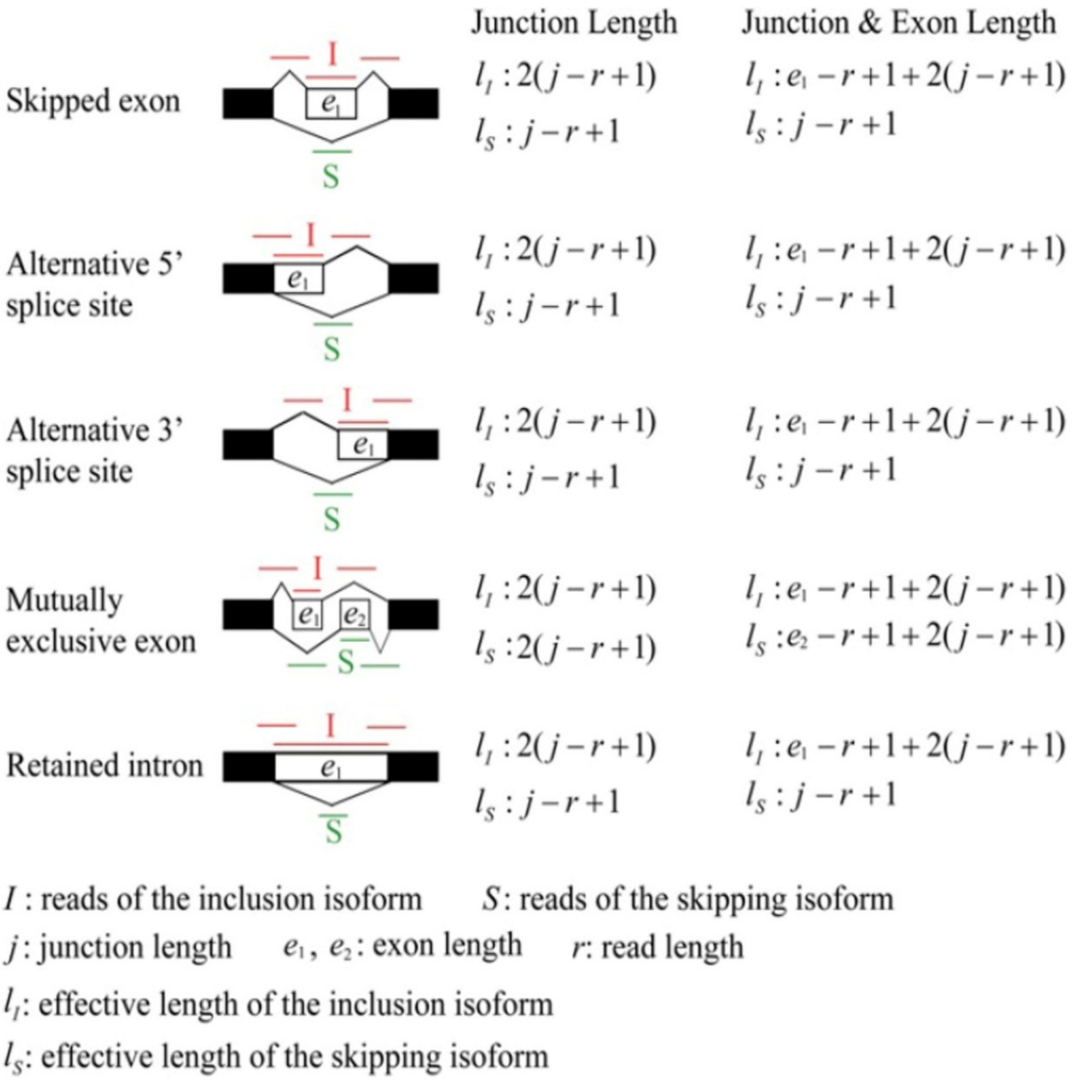

	Junction Length	Junction & Exon Length
Skipped exon	$l_I : 2(j-r+1)$ $l_S : j-r+1$	$l_I : e_1-r+1+2(j-r+1)$ $l_S : j-r+1$
Alternative 5' splice site	$l_I : 2(j-r+1)$ $l_S : j-r+1$	$l_I : e_1-r+1+2(j-r+1)$ $l_S : j-r+1$
Alternative 3' splice site	$l_I : 2(j-r+1)$ $l_S : j-r+1$	$l_I : e_1-r+1+2(j-r+1)$ $l_S : j-r+1$
Mutually exclusive exon	$l_I : 2(j-r+1)$ $l_S : 2(j-r+1)$	$l_I : e_1-r+1+2(j-r+1)$ $l_S : e_2-r+1+2(j-r+1)$
Retained intron	$l_I : 2(j-r+1)$ $l_S : j-r+1$	$l_I : e_1-r+1+2(j-r+1)$ $l_S : j-r+1$

I : reads of the inclusion isoform S: reads of the skipping isoform

j: junction length e_1, e_2: exon length r: read length

l_I: effective length of the inclusion isoform

l_S: effective length of the skipping isoform

Supplementary figure 1 Relative abundance calculation of different isoforms produced by alternative splicing