

Table 2 Statistics of differentially expressed small RNA sequence reads in two different cotton accessions under two different water regimes

Description	Differentially expressed reads in			
	KC3-C <i>Vs.</i> KC3-WS	KC3-C <i>Vs.</i> Suvin-C	KC3-WS <i>Vs.</i> Suvin-WS	Suvin-C <i>Vs.</i> Suvin-WS
Total number*	5138	5341	7494	8469
Match genome [#]	789 (15.35%)	759 (14.21%)	932 (12.43%)	1250 (14.75%)
rRNA	4	6	6	7
snRNA	1	1	5	3
snoRNA	11	10	15	10
tRNA	20	19	28	25
tmRNA	0	0	0	1

* Total number of distinct reads found in each comparison

[#]number of reads that match with *G. raimondii* genome