

Transcriptome analysis of apical meristem enriched bud samples for size dependent flowering commitment in *Crocus sativus* reveal role of sugar and auxin signalling

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Table S1: SSRs Statistics of transcripts

Total number	18796
Total number of SSR containing sequence	13905
Number of sequence >1 SSR	3125
Number of SSR in compound formation	2641

Table S2: SSRs Distribution of transcripts

Unit size	Number
2	9194
3	8767
4	508
5	175
6	152

Table S3: Annotation statistics

Source	Number of Annotated Transcripts
NR-Uniprot	74766
PlantTFDB	40463
KEGG	5325
GO	56230
Pfam	22571
PRINTS	1107
ProDom	146
ProSitePatterns	1693
ProSiteProfiles	7905
SMART	2746
SUPERFAMILY	21716
TIGRFAM	787

