

SRA

Full

SRX24461561: RNAseq bud 3
1 BGISEQ (BGISEQ-500) run: 29.2M spots, 8.7G bases, 4.8Gb downloads

Design: using conventional library construction methods

Submitted by: China Agricultural University

Study: The sequencing data of the genome assembly and annotation of the Rosa hybrida 'Samantha'
[PRJNA1108167](#) • [SRP505854](#) • [All experiments](#) • [All runs](#)
[show Abstract](#)

Sample:
[SAMN41227155](#) • [SRS21214511](#) • [All experiments](#) • [All runs](#)
Organism: [Rosa hybrid cultivar](#)

Library:
Name: RNAseq-bud-3
Instrument: BGISEQ-500
Strategy: RNA-Seq
Source: TRANSCRIPTOMIC
Selection: RANDOM
Layout: PAIRED

Runs: 1 run, 29.2M spots, 8.7G bases, [4.8Gb](#)

Run	# of Spots	# of Bases	Size	Published
SRR28905042	29,223,823	8.7G	4.8Gb	2024-08-25

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