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Run Browser › **SRR28905042**

RNAseq bud 3 (SRR28905042)

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SRA archive data

SRA archive data is normalized by the SRA load process and used by the **SRA Toolkit** to read and produce formats like FASTQ, SAM, etc. The defau
by accession.

Public SRA files are now available from GCP and AWS cloud platforms as well as from NCBI. Access to most data in the cloud requires a user accoun
costs for cloud compute or to copy data outside of the specified cloud service region.

Type	Version	Created	Size	Location	Name
SRA Normalized	1	2024-05-06	4.8GB	AWS	https://sra-pub-run-odp.s3.amazonaws.com/sra/SRR28905042/SRR28905042
SRA Lite	1	2024-05-06	2.5GB	NCBI	https://sra-downloadb.be-md.ncbi.nlm.nih.gov/sos6/sra-pub-zq-40/SRR028/28905/SRR28905042/SRR28905042.lite.1
				AWS	s3://sra-pub-zq-3/SRR28905042/SRR28905042.lite.1
				GCP	gs://sra-pub-zq-103/SRR28905042/SRR28905042.lite.1

Original format

The original files submitted to SRA. These files may require specific software to open, read and interpret data.

Type	Version	Created	Size	Location	Name	Free Egress	Access Type
fastq	1	2024-05-06	3.2GB	AWS	s3://sra-pub-src-15/SRR28905042/SMT_bud_3_R1.fq.gz.1	-	Use Cloud Da
fastq	1	2024-05-06	2.7GB	AWS	s3://sra-pub-src-15/SRR28905042/SMT_bud_3_R2.fq.gz.1	-	Use Cloud Da

❓ Egress and Access: what does it mean?
❓ Why is SRA data in the cloud?
❓ What is "Cloud Data Delivery"?