

Run

Legend ?

Experiment

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Biosample

Bioproject

Bioproject	SRA Study	Title
PRJNA178588	SRP016911	Limulus polyphemus Transcriptome or Gene expression

Abstract:
Normalized 454 cDNA amoebocyte transcriptome. An individual sampled from coastal waters near Cape Cod (in Woods Hole, MA).

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National Library of Medicine

8600 Rockville Pike

Bethesda, MD 20894

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COMMIT_INFO

Horseshoe Crab Transcriptome (SRR610305)

- Metadata
- Analysis
- Reads
- Data access**
- FASTA/FASTQ download

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 default toolkit configuration enables it to find and retrieve SRA runs 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 account with the cloud service provider. The user's account will incur costs for cloud compute or to copy data outside of the specified cloud service region.

Type	Version	Created	Size	Location	Name	Free Egress	Access Type
SRA Normalized	1	2012-10-31	131.6MB	AWS	https://sra-pub-run-odp.s3.amazonaws.com/sra/SRR610305/SRR610305	worldwide	anonymous
SRA Lite	1	2022-06-06	49.4MB	NCBI	https://sra-downloadb.be-md.ncbi.nlm.nih.gov/sos5/sra-pub-zq-11/SRR000/610/SRR610305/SRR610305.lite.1	worldwide	anonymous
				AWS	s3://sra-pub-zq-9/SRR610305/SRR610305.lite.1	s3.us-east-1	aws identity
				GCP	gs://sra-pub-zq-109/SRR610305/SRR610305.lite.1	gs.us-east1	gcp identity

- [Egress and Access: what does it mean?](#)
- [Why is SRA data in the cloud?](#)
- [What is "Cloud Data Delivery"?](#)

