



COVID-19 is an emerging, rapidly evolving situation.

Get the latest public health information from CDC: <a href="https://www.coronavirus.gov">https://www.coronavirus.gov</a>.

Get the latest research from NIH: <a href="https://www.nih.gov/coronavirus">https://www.nih.gov/coronavirus</a>.

Find NCBI SARS-CoV-2 literature, sequence, and clinical content: <a href="https://www.ncbi.nlm.nih.gov/sars-cov-2/">https://www.ncbi.nlm.nih.gov/sars-cov-2/</a>.

_	HOME   SEARCH   SITE MA	_				
GEO help: Mouse over screen elements for information.						
	Series GSE93463	Query DataSets for GSE93463				
	Status	Public on Feb 03, 2017				
	Title	polyA mRNA RNA-seq from liver (ENCSR000CHE)				
	Project	Mouse ENCODE				
	Organism	Mus musculus				
	Experiment type	Expression profiling by high throughput sequencing				
	Summary	RNA-seq on 14.5 day embryonic mouse liver				
		For data usage terms and conditions, please refer to http://www.genome.gov/27528022 and http://www.genome.gov/Pages/Research/ENCODE/ENCODE_Data_Use_Policy_for_External_Users_03-07-14.pdf				
	Overall design	https://www.encodeproject.org/ENCSR000CHE/				
	Citation(s)	ENCODE Project Consortium An integrated encyclopedia of DNA elements in the human genome. <i>Nature</i> 2012 Sep 6;489(7414):57-74. PMID: 22955616				
n.	BioProject	PRJNA66167 Mouse ENCODE transcriptome data				
	Submission date	Jan 11, 2017				
	Last update date	May 15, 2019				
	Contact name	ENCODE DCC				
	E-mail(s)	encode-help@lists.stanford.edu				
	Organization name	ENCODE DCC				
	Street address	300 Pasteur Dr				
	City	Stanford				
	State/province	CA				
	ZIP/Postal code	94305-5120				
	Country	USA				
	Platforms (1)	GPL13112 Illumina HiSeq 2000 (Mus musculus)				
	Samples (2)	GSM2453385 polyA mRNA RNA-seq from liver (ENCLB399DXG)				
		GSM2453386 polyA mRNA RNA-seq from liver (ENCLB784HFI)				
	Relations					
	Supersedes	GSM929721				
	Download family	Format				

1 of 2 5/8/21, 7:08 pm

SOFT 🖪 SOFT formatted family file(s) MINIML 🛂 MINIML formatted family file(s) Series Matrix File(s) TXT 🛂

	Supplementary file	Size	Download	File type/resource
GSE93463	_RAW.tar		(http)	TAR (of BIGWIG, TSV,
		Mb	(custom)	TXT)

Raw data provided as supplementary file Processed data provided as supplementary file

Custom GSE93463\_RAW.tar archive:

Supplementary f	ile	File size
GSM2453385_ENCFF078MDF_signal_of_	_unique_reads_mm10.bigWig	43.0 Mb
GSM2453385_ENCFF088LMD_signal_of_	_all_reads_mm10.bigWig	76.1 Mb
GSM2453385_ENCFF480SQS_transcript	_quantifications_mm10.tsv.gz	4.4 Mb
✓ GSM2453385_ENCFF668RJN_gene_qua	ntifications_mm10.tsv.gz	2.4 Mb
GSM2453385_ENCLB399DXG_README.	txt.gz	598 b
GSM2453386_ENCFF604QLO_signal_of_	_unique_reads_mm10.bigWig	41.5 Mb
GSM2453386_ENCFF837CEK_transcript	_quantifications_mm10.tsv.gz	4.4 Mb
GSM2453386_ENCFF866RWU_signal_of	_all_reads_mm10.bigWig	72.9 Mb
✓ GSM2453386_ENCFF993NNK_gene_qua	2.4 Mb	
GSM2453386_ENCLB784HFI_README.t:	602 b	
Select All	Cancel Download	2 file(s), 4.8 Mb

| NLM | NIH | GEO Help | Disclaimer | Accessibility |

5/8/21, 7:08 pm

2 of 2