



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 M.					
GEO help: Mouse over	r screen elements for information.				
Scope: Self	Format: HTML V Amount: Quick V GEO accession: GSE93464				
Series GSE9346	Query DataSets for GSE93464				
Status	Public on Feb 03, 2017				
Title	polyA mRNA RNA-seq from heart (ENCSR000CHF)				
Project	Mouse ENCODE				
Organism	Mus musculus				
Experiment type	Expression profiling by high throughput sequencing				
Summary	RNA-seq on 14.5 day embryonic mouse heart				
	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/ENCSR000CHF/				
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				
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	e 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)	GSM2453387 polyA mRNA RNA-seq from heart (ENCLB330QBQ)				
	GSM2453388 polyA mRNA RNA-seq from heart (ENCLB814UGI)				
Relations					
Supersedes	GSM929724				
Download famil	y Format				

1 of 2 5/8/21, 7:17 pm

SOFT 🖪

TXT 🛂

MINIML 🛂

SOFT formatted family file(s)
MINIML formatted family file(s)
Series Matrix File(s)

	Supplementary file	Size	Download	File type/resource
GSE93464	_RAW.tar	384.1	(http)	TAR (of BIGWIG, TSV,
		Mb	(custom)	TXT)

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

Custom GSE93464\_RAW.tar archive:

Supplementary fil	le	File size
GSM2453387_ENCFF108QKW_signal_of_	unique_reads_mm10.bigWig	72.2 Mb
GSM2453387_ENCFF261PKW_transcript_	_quantifications_mm10.tsv.gz	4.3 Mb
GSM2453387_ENCFF571QZP_signal_of_a	all_reads_mm10.bigWig	114.5 Mb
✓ GSM2453387_ENCFF662WLV_gene_quan	ntifications_mm10.tsv.gz	2.4 Mb
GSM2453387_ENCLB330QBQ_README.tx	xt.gz	600 b
GSM2453388_ENCFF139CQE_transcript_	quantifications_mm10.tsv.gz	4.3 Mb
GSM2453388_ENCFF580UUS_signal_of_ι	unique_reads_mm10.bigWig	71.5 Mb
GSM2453388_ENCFF700GCM_signal_of_a	all_reads_mm10.bigWig	112.4 Mb
✓ GSM2453388_ENCFF705YYN_gene_quan	2.4 Mb	
GSM2453388_ENCLB814UGI_README.tx	599 b	
Select All	Cancel Download	2 file(s), 4.7 Mb

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

2 of 2