

 An official website of the United States government

 

SRA

SRA

Search

Advanced

Help

Access

Public (2)

Source

DNA (2)

Type

genome (2)

Library Layout

paired (1)

single (1)

Platform

LS454 (2)

Strategy

Genome (2)

Data in Cloud

GS (2)

S3 (2)

File Type

sff (2)

Summary

Send to:

[Send results to Blast](#)

Links from BioProject

Items: 2

☐

[Horshecrab 454 Paired End](#)

1. 1 LS454 (454 GS FLX Titanium) run: 2.6M spots, 1.1G bases, 2.6Gb downloads
Accession: SRX202154

☐

[Horseshoe Crab Shotgun](#)

2. 9 LS454 (454 GS FLX Titanium) runs: 4.6M spots, 2.9G bases, 6.2Gb downloads
Accession: SRX202152

Filters: [Manage Filters](#)

Find related data

Database:

Select

Find items

Recent activity

Turn Off

Clear



SRA Links for BioProject (Select 178587) (2)

SRA



SRA Links for BioProject (Select 178588) (1)

SRA



Limulus polyphemus

BioProject

See more...

Clear all

[Show additional filters](#)

FOLLOW NCBI

Twitter

Facebook

LinkedIn

.cls-11,

.cls-12

/ fill:

#737373;

.cls-11 {

Connect with NLM

Twitter Facebook YouTube

National Library of Medicine

8600 Rockville Pike

Bethesda, MD 20894

Web Policies

FOIA

HHS Vulnerability Disclosure

Help

Accessibility

Careers

.cls-1{fill:#737373;}

NCBI

Insights

Blog

 An official website of the United States government



Log in

SRA

Advanced

Search

Help

Full ▼

Send to: ▼

SRX202152: Horseshoe Crab Shotgun
9 LS454 (454 GS FLX Titanium) runs: 4.6M spots, 2.9G bases, 6.2Gb downloads

Design: Followed published 454 protocols for building nebulized genomic shotgun sequencing libraries

Submitted by: UNIVERSITY OF CONNECTICUT

Study: Limulus polyphemus Genome sequencing
[PRJNA178587](#) • [SRP016906](#) • [All experiments](#) • [All runs](#)
[show Abstract](#)

Sample: Genome Reads
[SAMN01797690](#) • SRS373640 • [All experiments](#) • [All runs](#)
Organism: [Limulus polyphemus](#)

Library:
Instrument: 454 GS FLX Titanium
Strategy: WGS
Source: GENOMIC
Selection: RANDOM
Layout: SINGLE

Spot descriptor:

5 forward

Related information 
[BioProject](#)
[BioSample](#)
[Taxonomy](#)

Recent activity 
[Turn Off](#) [Clear](#)

 A Brief History of NCBI's Formation and Growth - The

 Limulus polyphemus

BioProject

 Limulus polyphemus, whole genome shotgun

Nucleotide

 Sample from Limulus polyphemus

biosample

 Horseshoe Crab Transcriptome

biosample

See more...

Runs: 9 runs, 4.6M spots, 2.9G bases, [6.2Gb](#)

Run	# of Spots	# of Bases	Size	Published
SRR610294	424,445	246.9M	537.3Mb	2013-10-31
SRR616995	507,175	321.5M	672.6Mb	2013-10-31
SRR616996	981,591	559.4M	1.2Gb	2013-10-31
SRR616997	846,004	548.9M	1.1Gb	2013-10-31
SRR616999	698,717	332.6M	784.7Mb	2013-10-31
SRR617000	27,222	12M	29Mb	2013-10-31
SRR617001	403,776	231M	510.4Mb	2013-10-31
SRR617003	456,253	533.2M	1.1Gb	2013-10-31
SRR617004	219,255	140.4M	292.4Mb	2013-10-31

ID: 269021

https://www.ncbi.nlm.nih.gov/sra/SRX202152[accn][11/18/2025 7:29:26 PM]

Seikagaku Corporation Exhibit 2075 - Page 3 of 6
Charles River Laboratories, Inc. v. Seikagaku Corporation
PGR2025-00023

FOLLOW NCBI

TwitterFacebookLinkedIn

Connect with NLM

TwitterFacebookYouTube

National Library of Medicine

8600 Rockville Pike

Bethesda, MD 20894

Web Policies

FOIA

HHS Vulnerability Disclosure

Help

Accessibility

Careers

NLM | NIH | HHS | USA.gov

 An official website of the United States government

?

Log in

SRA

SRA

Advanced

Search

Help

Full

Send to

SRX202154: Horshecrab 454 Paired End
1 LS454 (454 GS FLX Titanium) run: 2.6M spots, 1.1G bases, 2.6Gb downloads

Design: Published protocols for 454 paired-end 3KB sequencing
Submitted by: UNIVERSITY OF CONNECTICUT
Study: Limulus polyphemus Genome sequencing
[PRJNA178587](#) • [SRP016906](#) • [All experiments](#) • [All runs](#)
[show Abstract](#)

Sample: Genome Reads
[SAMN01797690](#) • SRS373640 • [All experiments](#) • [All runs](#)
Organism: [Limulus polyphemus](#)

Library:
Instrument: 454 GS FLX Titanium
Strategy: WGS
Source: GENOMIC
Selection: RANDOM
Layout: PAIRED

Spot descriptor:

5

forward

forward

Runs: 1 run, 2.6M spots, 1.1G bases, [2.6Gb](#)

Run	# of Spots	# of Bases	Size	Published
SRR610297	2,581,149	1.1G	2.6Gb	2013-10-31

Related information

[BioProject](#)
[BioSample](#)
[Taxonomy](#)

Recent activity

[Turn Off](#) [Clear](#)

 A Brief History of NCBI's Formation and Growth - The

 Limulus polyphemus BioProject

 Limulus polyphemus, whole genome shotgun Nucleotide

 Sample from Limulus polyphemus biosample

 Horseshoe Crab Transcriptome biosample

[See more...](#)

ID: 269023

FOLLOW NCBI

Twitter

Facebook

LinkedIn

.cls-11, .cls-12 { fill: #737373; }

.cls-1{fill:#737373; } Insights Blog

Connect with NLM

Twitter Facebook Youtube

National Library of Medicine

8600 Rockville Pike

Bethesda, MD 20894

Web Policies

FOIA

HHS Vulnerability Disclosure

Help

Accessibility

Careers

NLM

NIH

HHS

USA.gov