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SRX202160: Horseshoe Crab Transcriptome

1 LS454 (454 GS FLX Titanium) run: 772,309 spots, 188.2M bases, 131.6Mb downloads

**Design:** Normalized polyA selected RNA using Evrogen Trimmer Kit and used broken T primer approach for cDNA construction and amplification.

**Submitted by:** UNIVERSITY OF CONNECTICUT

**Study:** Limulus polyphemus Transcriptome or Gene expression

[PRJNA178588](#) • [SRP016911](#) • [All experiments](#) • [All runs](#)

[hide Abstract](#)

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

**Sample:** Horseshoe Crab Transcriptome

[SAMN01797697](#) • SRS373647 • [All experiments](#) • [All runs](#)

**Organism:** [Limulus polyphemus](#)

Library:

**Instrument:** 454 GS FLX Titanium

**Strategy:** WGS

**Source:** TRANSCRIPTOMIC

**Selection:** cDNA

**Layout:** SINGLE

Spot descriptor:



**Runs:** 1 run, 772,309 spots, 188.2M bases, [131.6Mb](#)

| Run                       | # of Spots | # of Bases | Size    | Published  |
|---------------------------|------------|------------|---------|------------|
| <a href="#">SRR610305</a> | 772,309    | 188.2M     | 131.6Mb | 2013-10-31 |

ID: 269029