Resource Summary Report

Generated by dkNET on Apr 24, 2025

Stowers Original Data Repository

RRID:SCR_002640
Type: Tool

Proper Citation

Stowers Original Data Repository (RRID:SCR_002640)

Resource Information

URL: http://odr.stowers.org/websimr/

Proper Citation: Stowers Original Data Repository (RRID:SCR_002640)

Description: THIS RESOURCE IS NO LONGER IN SERVICE, documented January 13, 2022. Open access repository of original, unprocessed data underlying work published by Stowers researchers to allow the scientific community to validate and extend the findings made by Stowers researchers. For papers first submitted for publication after November 1, 2011, the Stowers Institute requires its members to deposit original data files into the Stowers Original Data Repository or to repositories maintained by third parties at the time of publication. Access to the Stowers Original Data Repository is free, but you will be asked to register before you can download data.

Abbreviations: Stowers ODR, ODR

Synonyms: Stowers Institute Original Data Repository

Resource Type: service resource, data repository, database, bibliography, data or information resource, storage service resource

Keywords: data set, image

Funding:

Availability: THIS RESOURCE IS NO LONGER IN SERVICE, Free, Open unspecified license, Registration required, Acknowledgement requested

Resource Name: Stowers Original Data Repository

Resource ID: SCR_002640

Alternate IDs: nlx_156067

Record Creation Time: 20220129T080214+0000

Record Last Update: 20250424T064602+0000

Ratings and Alerts

No rating or validation information has been found for Stowers Original Data Repository.

No alerts have been found for Stowers Original Data Repository.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 11 mentions in open access literature.

Listed below are recent publications. The full list is available at dkNET.

Parker HJ, et al. (2021) Analysis of lamprey meis genes reveals that conserved inputs from Hox, Meis and Pbx proteins control their expression in the hindbrain and neural tube. Developmental biology, 479, 61.

Singh NP, et al. (2021) Genome-Wide Binding Analyses of HOXB1 Revealed a Novel DNA Binding Motif Associated with Gene Repression. Journal of developmental biology, 9(1).

Prummel KD, et al. (2019) A conserved regulatory program initiates lateral plate mesoderm emergence across chordates. Nature communications, 10(1), 3857.

Parker HJ, et al. (2019) A Hox-TALE regulatory circuit for neural crest patterning is conserved across vertebrates. Nature communications, 10(1), 1189.

Chen Y, et al. (2019) Overdosage of Balanced Protein Complexes Reduces Proliferation Rate in Aneuploid Cells. Cell systems, 9(2), 129.

De Kumar B, et al. (2017) HOXA1 and TALE proteins display cross-regulatory interactions and form a combinatorial binding code on HOXA1 targets. Genome research, 27(9), 1501.

De Kumar B, et al. (2017) Hoxa1 targets signaling pathways during neural differentiation of ES cells and mouse embryogenesis. Developmental biology, 432(1), 151.

McEllin JA, et al. (2016) Analyses of fugu hoxa2 genes provide evidence for subfunctionalization of neural crest cell and rhombomere cis-regulatory modules during vertebrate evolution. Developmental biology, 409(2), 530.

De Kumar B, et al. (2015) Analysis of dynamic changes in retinoid-induced transcription and epigenetic profiles of murine Hox clusters in ES cells. Genome research, 25(8), 1229.

Chen G, et al. (2015) Targeting the adaptability of heterogeneous aneuploids. Cell, 160(4), 771.

Parker HJ, et al. (2014) A Hox regulatory network of hindbrain segmentation is conserved to the base of vertebrates. Nature, 514(7523), 490.