High-resolution DNase-seq profiles of targeted loci in mES-derived primitive ectoderm - Study GBCO4773

Genomics Study Specifications

Study Name: High-resolution DNase-seq profiles of targeted loci in mES-derived primitive ectoderm
Contact Name: David Gifford (MIT)
Publication: Not provided
My Strategies: Return to My Strategies page
Classification: Tissue expression, surveys and comparisons

Links
- Biomaterials Graph
- ArrayExpress

BCBC Release Date: April 01, 2014
Citation: unavailable
Synopsis

The aim of this experiment was to profile DNase-I accessibility at a subset of genomic regions in extremely high coverage. After DNase-I treatment, DNA fragments from specific loci were targeted using bead capture, amplified, and sequenced.

Platform types
- Open chromatin DNase-Seq

Platforms
- Not available

Study Design Type
- compound_treatment_design

Study Factors
- Show study factors

Study Assays
- Show study assays

Access to Study Data

To access the Study Data you must "Request this Resource" (below) and the supplier must fill your Request. Then Beta Cell Genomics will contact you with details on how to access the data.

Gene List(s)

To access this study's gene list(s) you must "Request this Resource" (below) and the supplier must fill your Request.

Repositories

Melton Lab
- Request this resource
  - Stock #: Not provided
  - Availability Notes: Not provided

Stoeckert Lab
- Request this resource
  - Stock #: Not provided
  - Availability Notes: Not provided

Comments
There are no comments for this entry.