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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	<i>unavailable</i>
Synopsis	<div style="border: 1px solid gray; padding: 5px;"> <p>Study Description Goals</p> <p>Approaches Results Conclusions</p> <p>Related Studies</p> <p>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.</p> </div>
Platform types	Open chromatin DNase-Seq
Platforms	<i>Not available</i>
Study Design Type	<ul style="list-style-type: none"> 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 #: <i>Not provided</i> Availability Notes: <i>Not provided</i>
Stoekert Lab	Request this resource	Stock #: <i>Not provided</i> Availability Notes: <i>Not provided</i>

Comments

Access Status

 This resource is publicly viewable.

Request this Resource

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Primary contributor: [Melton Lab](#)

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Resource History & Actions

Approved on Apr 01, 2014
Last modified on Apr 15, 2014

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