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ChIP-Seq analysis of enhancer related histone modifications during hESC differentiation towards pancreatic endoderm. - Study GBCO4574

Genomics Study Specifications

Study Name	ChIP-Seq analysis of enhancer related histone modifications during hESC differentiation towards pancreatic endoderm.
Contact Name	Maïke Sander (University of California, San Diego)
Publication	Not provided
My Strategies	Return to My Strategies page
Classification	Cell differentiation; Tissue expression, surveys and comparisons
Links	 Biomaterials Graph  GEO
BCBC Release Date	February 12, 2013
Citation	<i>unavailable</i>

Synopsis

Study Description	Goals	
Approaches	Results	Conclusions
Related Studies		

The objective of these experiments was to employ ChIP-Seq to map changes in enhancer related chromatin modifications during CyT49 hESC differentiation towards the pancreatic lineage. Such data will be used to identify enhancer elements during pancreatic lineage specification of hESCs.

Platform types	Epigenomic, Histone modification ChIP-Seq
Platforms	<i>Not available</i>
Study Design Type	<ul style="list-style-type: none"> development_or_differentiation_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

Sander 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>

Access Status

 This resource is publicly viewable.

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Primary contributor: [Sander Lab](#)

Resource Tags

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Resource History & Actions

Approved on Feb 12, 2013
Last modified on Feb 26, 2013

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