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## RNA-seq of Ins<sup>+</sup> cells in human differentiation protocol across lines and isolated from human islets - Study GBCO4764

**Genomics Study Specifications**

<b>Study Name</b>	RNA-seq of Ins <sup>+</sup> cells in human differentiation protocol across lines and isolated from human islets
<b>Contact Name</b>	<a href="#">David Gifford</a> (MIT)
<b>Publication</b>	<a href="http://www.ncbi.nlm.nih.gov/pubmed/24594682">http://www.ncbi.nlm.nih.gov/pubmed/24594682</a>
<b>My Strategies</b>	<a href="#">Return to My Strategies page</a>
<b>Classification</b>	Differentiation of insulin-producing cells; Cell differentiation; Tissue expression, surveys and comparisons
<b>Links</b>	 <a href="#">Biomaterials Graph</a>  <a href="#">ArrayExpress</a>
<b>BCBC Release Date</b>	April 01, 2014
<b>Citation</b>	Hrvatn S, Deng F, O'Donnell CW, Gifford DK, Melton DA. <a href="#">MARIS: method for analyzing RNA following intracellular sorting</a> . PLoS One. 2014. 9:e89459

**Synopsis**

<b>Study Description</b>	Goals
Approaches	Results
	Conclusions
Related Studies	

The aim of this experiment was to observe the transcriptional profile of Ins<sup>+</sup> cells in human cadaveric islets and at the terminal stage of our in vitro beta-cell differentiation protocol across both the Hues8 and H1 cell lines. One polyhormonal sample (Ins<sup>+</sup>/Gcg<sup>+</sup>) was also sorted for additional comparison.

<b>Platform types</b>	Expression RNA-Seq, Expression
<b>Platforms</b>	<i>Not available</i>
<b>Study Design Type</b>	<ul style="list-style-type: none"> <li>cell_type_comparison_design</li> </ul>
<b>Study Factors</b>	<a href="#">Show study factors</a>
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**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.

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<b>Melton Lab</b>	<a href="#">Request this resource</a>	<b>Stock #:</b> <i>Not provided</i> <b>Availability Notes:</b> <i>Not provided</i>
<b>Stoekert Lab</b>		<b>Stock #:</b> <i>Not provided</i>

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