Main Page

FANTOMS SSTAR (Semantic catalog of Samples, Transcription initiation And Regulators) provide a way to explore samples, transcriptional initiations, and regulators analyzed in the FANTOMS project. If you have any questions or comments, please contact fantom@gsc.riken.jp.

Search genes

Examples

Samples

Details of profiled samples (incl. primary cells, cell lines, and tissues)

- Smooth Muscle Cells
- Aortic, donor

Browse

human

RB SearchResults

Search within Gene names

Key Word: sp1

EntrezGene

Tax_id: 9606=Human, 10090=Mouse

<table>
<thead>
<tr>
<th>#</th>
<th>Symbol</th>
<th>Synonyms</th>
<th>Description</th>
<th>Tax id</th>
</tr>
</thead>
<tbody>
<tr>
<td>6688</td>
<td>sp1</td>
<td>sp1, 6688</td>
<td>spleen focus forming virus (SFFV) proviral integration oncogene sp1</td>
<td>9606</td>
</tr>
<tr>
<td>20375</td>
<td>sp1</td>
<td>sp1, 20375</td>
<td>SFFV proviral integration 1</td>
<td>10090</td>
</tr>
<tr>
<td>20700</td>
<td>Sarps1a</td>
<td>Sarps1a</td>
<td>serine (or cysteine) peptidase inhibitor, clade A, member 1A</td>
<td>10090</td>
</tr>
</tbody>
</table>