### Clustered Data Summary, Contigs and Best Annotation Hits

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**Datasets sub-menu**
- Annotation display options: show contigs where automatic annotation produced or did not produce results
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#### CLUSTERED DATA SUMMARY

**Sort by annotation results**
- Contigs can be displayed in groups pre-sorted by number of reads and length of sequence or sorted by entering range values

<table>
<thead>
<tr>
<th>Contigs sorted by the number of reads</th>
<th>Contigs sorted by length</th>
</tr>
</thead>
<tbody>
<tr>
<td>Number of reads</td>
<td>Number of contigs</td>
</tr>
<tr>
<td>-----------------</td>
<td>-------------------</td>
</tr>
<tr>
<td>1-300</td>
<td>12</td>
</tr>
<tr>
<td>91-300</td>
<td>142</td>
</tr>
<tr>
<td>31-40</td>
<td>501</td>
</tr>
<tr>
<td>5-20</td>
<td>212</td>
</tr>
<tr>
<td>5-7</td>
<td>219</td>
</tr>
<tr>
<td>4</td>
<td>4</td>
</tr>
<tr>
<td>3</td>
<td>4</td>
</tr>
<tr>
<td>2</td>
<td>4</td>
</tr>
</tbody>
</table>

Range (e.g. 100-200) or single value:
- [ ]

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#### CLUSTERED DATA SUMMARY, CONTIGS AND BEST ANNOTATION HITS

<table>
<thead>
<tr>
<th>Contig name</th>
<th>Length</th>
<th>Contig length</th>
<th>Number of reads</th>
<th>Number of hits from annotation</th>
</tr>
</thead>
<tbody>
<tr>
<td>Contig7593974</td>
<td>1584</td>
<td>1584</td>
<td>12</td>
<td>12</td>
</tr>
</tbody>
</table>

**Best hits selected from annotation**
- *Salmo salar hypoxia-inducible factor 1 (S Paran et al.)*
  - Database: SGP-Sal
  - Annotation: Hypoxia-inducible factor 1 (Salmo salar) [SGP] annotation

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**Access contig and reads sequences**

**Access detailed automatic annotation results**