<table>
<thead>
<tr>
<th>project</th>
<th>description</th>
</tr>
</thead>
<tbody>
<tr>
<td>Mouse - Medaka - MHC</td>
<td>Mouse - Medaka - MHC</td>
</tr>
<tr>
<td>Projects/Medaka transcriptome analysis</td>
<td>Projects/Medaka transcriptome analysis</td>
</tr>
</tbody>
</table>

<form action="http://www.molgen.mpg.de/~rodent/bin/MedakaDB_new.cgi" method="post">
<br>
<table width = "80%">
<tr><td align = "right"><h3><font size = "5">Search in Medaka ESTdb</font></h3></td><td align = "right"><a href="http://www.molgen.mpg.de/~rodent/MedakaDB_help.html"><font color = "red" size = "4">&nbsp;&nbsp;HELP</font></a></td></tr>
<h4>Search for clone or cluster information</h4>
<p><select name="search_crit">
<option value="Clone_id">Clone_id</option>
<option value="UniqueSequence_id">UniqueSequence_id</option>
<option value="Rearray1_id">Rearray1_id</option>
<option value="Rearray2_id">Rearray2_id</option>
<option value="OFPcluster_id">OFPcluster_id</option>
<option value="Trace_id">Trace_id</option>
<option value="AnnotationKeyword">AnnotationKeyword</option>
</select>
<input type=text name="search_what" size=30>
<p><input type=submit name="Query" value="Search"></p>
</h4></form>