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<!DOCTYPE HTML PUBLIC "-//W3C//DTD HTML 4.0 Transitional//EN">
<html>

  <head>
    <title>Mouse Medaka MHC - Vertebrate Genomics</title>
    <meta http-equiv="Content-Type" content="text/html;
charset=iso-8859-1">
    <meta http-equiv="content-language" content="en">
    <meta http-equiv="Reply-to"
content="wwwadm@molgen.mpg.de">
    <meta name="robots" content="index, follow">
    <meta name="allow-search" content="yes">
    <meta http-equiv="Content-Script-Type"
content="text/javascript">
    <meta http-equiv="Content-Style-Type" content="text/css">
    <link rel=STYLESheet type="text/css" href="ags2001.css">
    <script language="JavaScript" src="mainjs2001.js"
type="text/javascript">
<!--
    self.defaultStatus = document.title;
// -->
</script>
  </head>

  <body bgcolor="#FFFFFF" text="#000000" link="#007B70" alink="#
990000" vlink="#2E90B4" leftmargin="0" topmargin="0" marginwidth="0"
marginheight="0">
    <table cellpadding="0" cellspacing="0" border="0"
width="100%" height="120">
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        <td rowspan="2" width="7%" valign="middle"
align="center"></td>
        <td align="left" rowspan="2" valign="middle"
width="83%">
          <div align="left">
            <font
face="Verdana,Arial,Helvetica,sans-serif" size="+2">MAX PLANCK
INSTITUTE for MOLECULAR GENETICS</font></div>
          </td>
        <td align="left" rowspan="2" valign="middle"
width="10%"><a href="http://www.molgen.mpg.de/"><font size="-3">home
</font></a>
          <p><a
href="http://www.molgen.mpg.de/contact.html"><font size="-3">contact
</font></a></p>
          <p><a
href="http://www.molgen.mpg.de/search.html"><font size="-3">search
</font></a></p>
        </td>
      </tr>
      <tr valign="top" align="left"></tr>
      <tr valign="top" align="left">
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bgcolor="#E3E4E0" background="mhead2001_brownline_back.gif">
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border="0" height="36">
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valign="middle" height="36" nowrap background="mhead2001
_titlesmall_blank.gif"><font class="catsmalltitle"
face="Verdana,Arial,Helvetica,sans-serif" color="#FFFFFF" size="-2">
&nbsp;<a href="http://www.molgen.mpg.de/research/lehrach/"></a>Vertebrate Genomics</font></td>
    <td height="36"
valign="middle" nowrap align="left"><font class="catsubtitle"
face="Verdana,Arial,Helvetica,sans-serif" color="#000000" size="+1">
Mouse - Medaka - MHC</font></td>
    <td height="36"
valign="middle" nowrap width="10" align="left"></td>
    <td height="36"
valign="middle" nowrap align="left"><font size="+1">/Projects/Medaka
transcriptome analysis</font></td>
    </tr>
  </table>
</td>
  <td height="36" align="left"
bgcolor="#E3E4E0" background="mhead2001_brownline_back.gif"
width="10%"></td>
</tr>
</table>
<br>
<br>
<table cellpadding="0" cellspacing="0" border="0">
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border="0">
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        <td><a href="people.html"
class="leftm">People</a></td>
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      <tr valign="top" align="left">
        <td colspan="2"><br>
        </td>
      </tr>
      <tr valign="top" align="left">
        <td width="32"
align="right"><a href="projects.html"></a></td>
        <td><a
href="projects.html" class="leftm">Projects</a></td>
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      <tr valign="top" align="left">
        <td colspan="2"><br>
        </td>
      </tr>
      <tr valign="top" align="left">
        <td width="32"
align="right"><a href="publications.html"></a></td>
        <td><a
href="publications.html" class="leftm">Publications</a></td>
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    </table>
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</table>
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        </tr>
        <tr valign="top" align="left">
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align="right"><a href="protocols.html"></a></td>
            <td><a
href="protocols.html" class="leftm">Protocols</a></td>
        </tr>
    </table>
</td>
    <td class="tdmain"><!-- START OF CONTENTS -->
    <p class="maintext">
    <table border="0" width="800">
    <tr><td>For research on the medaka
transcriptome we
fingerprinting
redundancy of
in our
from different
(ovary,
From a reduced
obtained
are used as a resource
expression analysis.
started with the oligonucleotide
approach (OFP) as a method to reduce
our medaka cDNA libraries. We included
analysis in total 119,040 cDNA clones
adult organs and developmental stages
gastrula, neurula and organogenesis).
redundancy set we generated 5' ESTs and
10,016 unique sequences. These clones
for further experiments, including
    </td></tr>
    <tr><td>See <a
href="CloneInfo_mod.html"><b><big>EST
database</big></b></a> for details on
clones and
sequences.</td></tr>
    <tr><td>
    <tr align="left"><td></td></tr>
    <tr align="left"><td></td></tr>
    </table>
    </p>
    </td>
    </tr>
</table>
<!-- PAGE FOOTER -->
</body>
</html>

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