Method:
DNA isolation was carried out from cell pellet (cell layer).
Genetic characteristics were determined by PCR-single-locus-technology.
16 independent PCR-systems D8S1179, D21S11, D7S820, CSF1PO, D3S1358, TH01, D13S317, D16S539, D2S1338, AMEL, D5S818, FGA, D19S433, vWA, TPOX and D18S51 were investigated. 
(Thermo Fisher, AmpFISTR® Identifier® Plus PCR Amplification Kit)
In parallel, positive and negative controls were carried out yielding correct results.

Result:

<table>
<thead>
<tr>
<th>Client Sample Name</th>
<th>Sample Code</th>
<th>D8S1179</th>
<th>D21S11</th>
<th>D7S820</th>
<th>CSF1PO</th>
<th>D3S1358</th>
<th>TH01</th>
<th>D13S317</th>
<th>D16S539</th>
<th>D2S1338</th>
<th>AMEL</th>
<th>D5S818</th>
<th>FGA</th>
</tr>
</thead>
<tbody>
<tr>
<td>8</td>
<td>CL00000354</td>
<td>10,13</td>
<td>29,29</td>
<td>9,11</td>
<td>12,13</td>
<td>13,15</td>
<td>7,7</td>
<td>8,10</td>
<td>9,11</td>
<td>19,23</td>
<td></td>
<td>11,12</td>
<td>20,21</td>
</tr>
</tbody>
</table>

The table shows the result of the cell line analysis and the comparison with the online database of the DSMZ (http://www.dsmz.de/de/service/services-human-and-animal-cell) and the Cellosaurus database (https://web.expasy.org/cellosaurus). Please note that only the PCR-systems according to ANSI/ATCC standard ASN-0002 were aligned (DSS818, D13S317, D7S820, D16S539, VWA, TH01, TPOX, CSF1PO, AMEL).

This report was created automatically and is therefore valid without a signature.