**Supplementary Table 2.** Targeted sequencing analysis of patient

<table>
<thead>
<tr>
<th>Items</th>
<th>DATA</th>
</tr>
</thead>
<tbody>
<tr>
<td>Total yield (Mbp)</td>
<td>528.54</td>
</tr>
<tr>
<td>Mappable reads of total reads (%)</td>
<td>99.0%</td>
</tr>
<tr>
<td>Coverage of target region (≥1X)</td>
<td>99.0%</td>
</tr>
<tr>
<td>Coverage of target region (≥10X)</td>
<td>98.7%</td>
</tr>
<tr>
<td>Total number of SNPs</td>
<td>163</td>
</tr>
<tr>
<td>Number of coding SNPs</td>
<td>94</td>
</tr>
<tr>
<td>Total number of indels</td>
<td>29</td>
</tr>
<tr>
<td>Number of coding indels</td>
<td>1</td>
</tr>
</tbody>
</table>