<table>
<thead>
<tr>
<th>Cell line</th>
<th>Karyotype</th>
<th>CGH</th>
</tr>
</thead>
<tbody>
<tr>
<td>TPC-1</td>
<td>49,XX,der(1)(1;10;21)(1pter→1q31::21q21→21qter), ++der(1)(1:3)(p13;p21), der(3)(3:9)(p11;q22), der(7)(q22q31), +i(8)(q10), der(10)(1:10;21)(10q11.2→10pter::1q31→1qter), +der(10)inv(10)(q11.2q21)(3:10)(q25;p11), der(21)(1;10;21) (21pter→21q21::10q21.2→10q11.2::10q21.2→10qter)[25]</td>
<td>rev ish enh(1)(q), dim(3)(p12p21), enh(3)(q24q29), dim(7)(q21q31), amp(8)(q), enh(9)(q22q34), enh(10)(q)</td>
</tr>
<tr>
<td>FB2</td>
<td>82<del>87,XXXX, der(1)(1;10;21)(1pter→1q31::21q21→21qter)x2, −i, del(7)(q22q31)x2, i(8)(q10), der(10)(1;10;21) (10pter- &gt;10q15::10q11.2→10p15::1q13→1qter)x2, 11, 14, 15, 16, 17, +20, der(21)(1;10;21) (21pter→21q21::10q21.2→10q11.2::10q21.2→10qter)x2[20]/82</del>87, idem, der(6)(1;6)(p32;q15)[5]/82~87, idem, der(11)(11;17)(q14;q24), add(12)(q24)[6]</td>
<td>rev ish dim(4), dim(7)(q22q31), dim(8)(p21p23), enh(8)(q), dim(11)(q14q25), dim(14)(q), dim(15)(q), dim(16)(p11p13), dim(16)(q), enh(20)</td>
</tr>
</tbody>
</table>

Karyotype and CGH descriptions follow the ISCN 2005 guidelines. Shared chromosomal aberrations are depicted in bold.