a

Total percent of reads with indel split by guide RNA – sperm vs fin-clip samples

<table>
<thead>
<tr>
<th>Guide RNA</th>
<th>Sperm</th>
<th>Fin Clip</th>
</tr>
</thead>
<tbody>
<tr>
<td>guide1</td>
<td>0%</td>
<td>25%</td>
</tr>
<tr>
<td>guide2</td>
<td>25%</td>
<td>50%</td>
</tr>
<tr>
<td>guide3</td>
<td>50%</td>
<td>75%</td>
</tr>
<tr>
<td>guide4</td>
<td>0%</td>
<td>25%</td>
</tr>
<tr>
<td>guide5</td>
<td>25%</td>
<td>50%</td>
</tr>
<tr>
<td>guide6</td>
<td>50%</td>
<td>75%</td>
</tr>
<tr>
<td>guide7</td>
<td>0%</td>
<td>25%</td>
</tr>
<tr>
<td>guide8</td>
<td>25%</td>
<td>50%</td>
</tr>
<tr>
<td>guide9</td>
<td>50%</td>
<td>75%</td>
</tr>
</tbody>
</table>

b

Frequencies of individual variants in sperm vs fin clip

Indel frequency – fin clip (%) vs Indel frequency – sperm (%)