

**Table S4**  
**Copy number determination by KIR MLPA technique.**

<b>Gene detected</b>	<b>Determining CNV</b>
2DL1	Average of all probes
2DL2	Average of all probes
2DL3	Average of #2 in Mix 2 and 3
2DL4	Average of all probes
2DL5 *	2DL5 #2 in Mix 2
2DS1	Average of #2 multiplied by 2 in Mix 2 and 3
2DS2	2DS2 in Mix 2
2DS3	Average 2DS3 #2 in Mix 2 and 2DS3 #1 in Mix 3
2DS4 All	2DS4 All and independently copies of WT + trunc
2DS4 WT	2DS4 WT
2DS4 truncated	2DS4 trunc multiplied by 2
2DS5	Average of all probes
2DP1	Average of all probes
3DL1	Average of 3DL1 #1 in Mix 1 and 3
3DL2	Average of all probes
3DL3	Average of all probes
3DS1	Average of all probes
3DP1	3DP1 #1 in Mix1

\* Detects both 2DL5A and B