

Supplementary table 3. Definition of the deletion breakpoints by custom array-CGH in seven cases with CCRs.*

case	Last present probe	First deleted probe	Size (bp)	Last deleted probe	First present probe	Size (bp)
PROXIMAL BREAKPOINT				DISTAL BREAKPOINT		
42 chr.13	38,424,805	38,425,739	934	41,584,272	41,584,776	504
43 chr.4	144,739,260	144,739,715	455	147,230,945	147,231,355	410
45 chr.18	37,449,354	37,449,869	515	38,502,229	38,502,488	259
47 chr.4	168,151,549	168,152,975	1426	169,559,872	169,560,813	941
47 chr.11	46,262,486	46,262,997	511	46,894,644	46,895,443	799
49 chr.4	173,348,140	173,348,630	490	175,649,430	175,650,960	1530
49 chr.4	179,161,593	179,163,446	1853	179,979,486	179,980,450	964
50 chr.6	108,228,956	108,232,380	3424	111,694,448	111,694,811	363
50 chr.11	88,287,991	88,299,437	11446	94,140,125	94,140,467	342
51 chr.1	103,146,735	103,147,223	488	106,498,911	106,499,367	456
51 chr.2	181,978,746	181,979,546	800	184,990,778	184,991,262	484

*The table shows the precise chromosomal position of the two probes, one present and one deleted, flanking the proximal and the distal breakpoints for each deletion, and the genomic distance that separates them. Map positions refer to the Genome Assembly May 2004 (BUILD 35).