

Supplementary Material

Supplementary table 1. Primers sequences used in long-range PCR for breakpoints mapping.

Deletion coordinates	Primers spanning the putative breakpoints
chr16:67193822-67387415	5'-AAAAGTATTAAGAGTAAAATGTACATGGAAACAAAAGGGG 5'-GAAATTGCCATCATGCCACACGGAAC
chr16:67324886-67330557	5'-GACCCCACCAGAGCCCATCCACAGAC 5'-GGGGAAACAGTACCTGTATCAAGATGTTTTG
chr16:67328695-67328844	5'-CAGCCAATCAGCGGTACGGG 5'-CGGGCGGGGGCGGGCGGGCCGGCGTCCCTCGCAAGTCAGG
chr16:67416845-67424923	5'-CAAATGATTATCACAGCAAATGAGAGCAAGTCAATTAG 5'-CTTTTGGACATCACCACCATGTAAAGAGTGATG
chr16:67424298-67425126	5'-CTTAAGTTATGTTCTCTAGTAACTTGCTTGG 5'-CTTTTGGACATCACCACCATGTAAAGAGTGATG

Electronic-Database Information

The URLs for data presented herein are as follows:

National Center for Biotechnology Information (NCIB), <http://www.ncbi.nlm.nih.gov/>

Online Mendelian Inheritance in Man (OMIM) for CADHERIN 1, *CDH1* (+192090),
<http://www.ncbi.nlm.nih.gov/Omim/>

The genome database, <http://www.gdb.org>

MRC Holland, <http://www.mlpa.com/pages/indexpag.html>

Ensembl *Homo sapiens* core database, http://www.ensembl.org/Homo_sapiens/Info/Index

UCSC genome browser, <http://biodev.cbm.fvg.it/remo/DEL.txt>

UCSC genome browser Human Mar, <http://genome.ucsc.edu/cgi-bin/hgGateway>