

Name	Chromosome	Fragment Size	Forward	Reverse
JAG1	Chr20	125bp	CTCCTTGAGGCATTGA	TGTCATGCAGAACGTGAA
GATA6	Chr18	145bp	ATCTCTTCCTCGTCCTC	GATGCGAAGCGTAGGAACT
NOTCH1	Chr9	142bp	AACGAGGTCGGCTCCTA	CACACTCGTGGGTGACG
TBX2	Chr17	135bp	CTGTAGCCCAGTCCCAATAC	GCGGCTACAATCTCCATCA
NKX2-1	Chr14	101bp	TGGTCTACGTGTCTGTCAGTCTGT	CGCTCGGATTCTCTCCGGTA
OFD1 (control for normalization)	ChrX	137bp	AGGTGTTCTGCTGCTGAGATGGAA	TCCCTTTGTGCCAGATGAAGAGA

Supplemental Table 4. Summary of Chromosomal Distribution of CNVs

Chromosome	Duplicated CNVs	Deleted CNVs	Total CNVs
1	1	24	25
2	2	21	23
3	11	32	43
4	26	15	41
5	41	24	65
6	14	44	58
7	4	14	18
8	12	30	42
9	1	12	13
10	10	9	19
11	2	9	11
12	5	5	10
13	3	6	9
14	7	34	41
15	2	32	34
16	8	11	19
17	22	39	61
18	0	6	6
19	8	3	11
20	0	8	8
21	0	7	7
22	27	7	34
X	5	10	15

bp, base pairs; CGH, comparative genomic hybridization; CNVs, copy number variants.