

Supplemental Data**Duplications Involving a Conserved Regulatory Element****Downstream of *BMP2* Are Associated****with Brachydactyly Type A2**

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Table S1. Primer Information

Primer	Primer Sequence (5'-3')		Amplicon position on chr. 20 [bp]	
	Forward	Reverse	Start	Stop
P1	TGGTAATCCAAACCAAGAGGAA	CCACTTAAAATTCAAGAGGTAGTAGCA	6,807,799	6,807,870
P2	ATGAACCCTATGGAGCGTTTC	AAACCCAGCCTGACTTCTG	6,809,794	6,809,888
P3	GATGCTGTTGAGGGTGTCCA	ACAGAGCACATGCCCTGTT	6,811,685	6,811,770
P4	CAGCTCATTTGAACCTTATCTGAAC	AAGCCTTCTTGAGCCACAC	6,812,767	6,812,863
P5	TGAAGATCTGAAGTGGCTGTTCTG	CTGATGTTGAAGTTACCTGCCAATT	6,814,403	6,814,488
Primer Sequence (5'-3') for amplification and sequencing of the junction fragment				Primer position on chr. 20 [bp]
P6	GAGAAAGTAAAGTTGTTGATG		6,813,701	6,813,722
P6		GTGCAACTGTATAAGCCTAT	6,808,654	6,808,672

Amplicons used to confirm the duplicated region by quantitative real-time PCR (P1-P5) and primers for sequencing of the junction fragment (P6 Forward 6,813,701-6,813,722, P6 Reverse 6,808,654-6,808,672). Nucleotide positions corresponding to Ensembl release 50, July 2008.