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Supplemental Data

**Heterozygous Mutations in *MAP3K7*,
Encoding TGF- β -Activated Kinase 1,
Cause Cardiospondylocarpofacial Syndrome**

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	Coverage	Reads depth (15x)	Reads depth (30x)	Total relevant variants	Variant under dominant model	After in house data analysis	Variant under recessive model	After in house data analysis
Case 1	42	57,8	46	926	27 (clinically associated 16)	1	120 (clinically associated 68)	0
Case 2	40	64	50	972				
Case 3	80	69	60	1385				
Case 5	79	68	59	1064				

Supplemental table 1 WES data