

Table S1. Gene list used for filtering of WES data.

Gene associated with	Gene Symbol
a) Syndromic MAC[†]	<i>ACTB</i> ¹ , <i>ACTG1</i> ^{1,2} , <i>AHI1</i> ³ , <i>ALG3</i> ⁴ , <i>ALX1</i> ⁵ , <i>ALX3</i> ⁶ , <i>ANOS1</i> ⁷ , <i>ARX</i> ⁸ , <i>B3GALNT2</i> ⁹ , <i>B3GLCT</i> ¹⁰ , <i>BCOR</i> ¹⁰ , <i>BEST1</i> ¹⁰ , <i>BMP4</i> ¹¹ , <i>BRPF1</i> ¹² , <i>C12orf57</i> ¹⁰ , <i>CASK</i> ¹³ , <i>CC2D2A</i> ¹⁰ , <i>CCDC22</i> ¹⁴ , <i>CDK9</i> ¹⁵ , <i>CDON</i> ^{16,17} , <i>CEP290</i> ⁴ , <i>CHD7</i> ¹⁸ , <i>CPLANE1</i> ¹⁰ , <i>COL4A1</i> ^{19,20} , <i>COX7B</i> ²¹ , <i>CREBBP</i> ⁴ , <i>CRIM1</i> ²² , <i>CTU2</i> ²³ , <i>DAG1</i> ²⁴ , <i>DHX38</i> ²⁵ , <i>DPYD</i> ⁴ , <i>DSC3</i> ²⁶ , <i>EFTUD2</i> ²⁷ , <i>EP300</i> ²⁸ , <i>ERCC1</i> ²⁹ , <i>ERCC5</i> ³⁰ , <i>ERCC6</i> ³¹ , <i>EYA1</i> ¹⁰ , <i>FADD</i> ⁴ , <i>FAM111A</i> ³² , <i>FANCA</i> ¹⁰ , <i>FANCD2</i> ¹⁰ , <i>FANCE</i> ³³ , <i>FANCI</i> ³³ , <i>FANCL</i> ³³ , <i>FAT1</i> ³⁴ , <i>FBXW11</i> ³⁵ , <i>FIBP</i> ³⁶ , <i>FKRP</i> ³⁷ , <i>FKTN</i> ¹⁰ , <i>FLNA</i> ⁴ , <i>FNBP4</i> ³⁸ , <i>FOXA2</i> ³⁹ , <i>FOXL2</i> ¹⁰ , <i>FRAS1</i> ¹⁰ , <i>FREM1</i> ¹⁰ , <i>FREM2</i> ^{33,40} , <i>GDF3</i> ¹⁰ , <i>GDF6</i> ¹⁰ , <i>GJA1</i> ⁴¹ , <i>GRIP1</i> ¹⁰ , <i>GZF1</i> ⁴² , <i>HCCS</i> ¹⁰ , <i>HESX1</i> ⁴³ , <i>HDAC6</i> ³³ , <i>HMGB3</i> ¹⁰ , <i>HMX1</i> ¹⁰ , <i>HRAS</i> ⁴ , <i>IGBP1</i> ¹⁰ , <i>IKBKKG</i> ⁴⁴ , <i>INPP5E</i> ⁴⁵ , <i>INTS1</i> ⁴⁶ , <i>KCTD1</i> ⁴ , <i>KDM6A</i> ¹³ , <i>KERA</i> ¹⁰ , <i>KIAA0586</i> ⁴⁵ , <i>KIAA1109</i> ⁴⁷ , <i>KIAA1279</i> ^{4,48} , <i>KIF11</i> ⁴⁹ , <i>KIF26B</i> ⁵⁰ , <i>KMT2D</i> ⁵¹ , <i>KRAS</i> ⁴ , <i>LAMA1</i> ⁵² , <i>LRP2</i> ⁴ , <i>MAB21L2</i> ¹⁰ , <i>MAPRE2</i> ⁵³ , <i>MED12L</i> ⁵⁴ , <i>MID1</i> ⁵⁵ , <i>MITF</i> ⁴ , <i>MKS1</i> ⁴ , <i>NAA10</i> ¹⁰ , <i>NAGA</i> ⁵⁶ , <i>NDUFB11</i> ⁵⁷ , <i>NF1</i> ⁵⁸ , <i>NHS</i> ¹⁰ , <i>NRAS</i> ⁴ , <i>NTNG1</i> ⁵⁹ , <i>OFD1</i> ⁴⁵ , <i>OSGEP</i> ⁶⁰ , <i>OTX2</i> ¹⁰ , <i>PAX2</i> ⁴ , <i>PDE6D</i> ⁶¹ , <i>PIGL</i> ⁶² , <i>PITX2</i> ⁴ , <i>PITX3</i> ¹⁰ , <i>POC1B</i> ⁴⁵ , <i>POMGNT1</i> ⁶³ , <i>POMGNT2</i> ³³ , <i>POMK</i> ⁶⁴ , <i>POMT1</i> ⁶⁵ , <i>POMT2</i> ³³ , <i>PORCN</i> ⁴ , <i>PQBP1</i> ⁴ , <i>PTCH1</i> ⁴ , <i>PTPN11</i> ⁴ , <i>PUF60</i> ⁶⁶ , <i>RAB18</i> ⁶⁷ , <i>RAB3GAP1</i> ⁴ , <i>RAB3GAP2</i> ¹⁰ , <i>RARA</i> ⁶⁸ , <i>RARB</i> ¹⁰ , <i>RAX</i> ¹⁰ , <i>RERE</i> ⁶⁹ , <i>RIPK4</i> ⁷⁰ , <i>RPGRIP1L</i> ⁷¹ , <i>SALL1</i> ⁴ , <i>SALL4</i> ⁴ , <i>SEMA3E</i> ⁷² , <i>SH3PXD2B</i> ⁷³ , <i>SHH</i> ¹⁰ , <i>SIX3</i> ⁴ , <i>SLC18A2</i> ²⁶ , <i>SLC25A24</i> ⁷⁴ , <i>SLC38A8</i> ⁷⁵ , <i>SMARCA4</i> ⁷⁶ , <i>SMCHD1</i> ³³ , <i>SMG9</i> ⁷⁷ , <i>SMO</i> ⁷⁸ , <i>SMOC1</i> ¹⁰ , <i>SOX2</i> ¹⁰ , <i>SOX3</i> ⁷⁹ , <i>SOX10</i> ⁸⁰ , <i>SPINT2</i> ⁸¹ , <i>SRD5A3</i> ⁸² , <i>STRA6</i> ¹⁰ , <i>TBC1D20</i> ⁸³ , <i>TBC1D32</i> ^{26,33} , <i>TBX22</i> ⁸⁴ , <i>TCOF1</i> ¹⁰ , <i>TFAP2A</i> ⁸⁵ , <i>TMEM67</i> ⁸⁶ , <i>TMEM216</i> ⁸⁷ , <i>TMEM237</i> ⁴⁵ , <i>TUBA1A</i> ⁸⁸ , <i>TUBB</i> ⁵³ , <i>TUBGCP4</i> ⁸⁹ , <i>TWF1</i> ² , <i>VAX1</i> ⁹⁰ , <i>VSX2</i> ¹⁰ , <i>WASHC5</i> ⁹¹ , <i>WDR11</i> ⁹² , <i>WDR37</i> ⁹³ , <i>YAP1</i> ⁹⁴ , <i>ZEB2</i> ⁹⁵ , <i>ZMIZ1</i> ⁹⁶
b) Non-syndromic MAC[‡]	<i>ABCB6</i> ⁹⁷ , <i>ALDH1A3</i> ¹⁰ , <i>ATOH7</i> ¹⁰ , <i>CRYAA</i> ⁹⁸ , <i>CRYBA4</i> ¹⁰ , <i>CRYBB1</i> ³³ , <i>CRYBB2</i> ³³ , <i>CRYGC</i> ³³ , <i>FOXC1</i> ⁴ , <i>FOXE3</i> ¹⁰ , <i>GJA8</i> ^{99,100} , <i>IPO13</i> ¹⁰¹ , <i>LCP1</i> ² , <i>MAF</i> ¹⁰² , <i>MFRP</i> ³³ , <i>MYO10</i> ²⁶ , <i>NDP</i> ¹⁹ , <i>OLFM2</i> ¹⁰³ , <i>PAX6</i> ¹⁰ , <i>PIEZO2</i> ¹⁰⁴ , <i>PRSS56</i> ²⁶ , <i>PXDN</i> ¹⁰ , <i>RBP4</i> ^{105,106} , <i>SALL2</i> ¹⁰ , <i>SIX6</i> ¹⁰ , <i>TENM3</i> ¹⁰⁷ , <i>TMX3</i> ¹⁰⁸
c) Animal models with MAC	<i>ADAMTS16</i> ¹⁰⁹ , <i>ALDH7A1</i> ¹¹⁰ , <i>BCL6</i> ¹¹¹ , <i>CAP2</i> ¹¹² , <i>CDH2</i> ¹¹³ , <i>CDO1</i> ¹¹⁴ , <i>FBN2</i> ¹¹⁵ , <i>FGFR1</i> ¹¹⁶ , <i>FGFR2</i> ¹¹⁶ , <i>FOXG1</i> ¹¹⁷ , <i>FRS2</i> ¹¹⁸ , <i>GCLC</i> ¹¹⁹ , <i>HES1</i> ¹²⁰ , <i>ISPD</i> ¹²¹ , <i>JAG1</i> ¹²² , <i>LAMB1</i> ¹²³ , <i>LAMC1</i> ¹²³ , <i>LMO2</i> ¹²⁴ , <i>LMX1B</i> ⁴ , <i>MAPK8</i> ¹²⁵ , <i>MAPK9</i> ¹²⁵ , <i>NOG</i> ¹²⁵ , <i>PHACTR4</i> ¹²⁶ , <i>PPP1R12A</i> ¹²⁷ , <i>SFRP1</i> ¹²⁸ , <i>SFRP5</i> ¹²⁸ , <i>SMAD7</i> ¹²⁹ , <i>SOX4</i> ¹³⁰ , <i>SOX11</i> ^{131,132} , <i>TGFB2</i> ¹³³ , <i>TLE3</i> ¹³⁴ , <i>VAX2</i> ⁴
Candidate Genes	
d) Genes coding for proteins of the SHH signalling pathway¹³⁵	<i>CER1</i> , <i>GLI2</i> , <i>GLI3</i> , <i>GREM1</i> , <i>SUFU</i>
e) Genes coding for proteins of the WNT signalling pathway¹³⁵	<i>APC</i> , <i>CTNNB1</i> , <i>DVL1</i> , <i>FZD1</i> , <i>FZD2</i> , <i>FZD5</i> , <i>FZD6</i> , <i>FZD7</i> , <i>FZD9</i> , <i>FZD10</i> , <i>GSK3A</i> , <i>LRP5</i> , <i>LRP6</i> , <i>WNT1</i> , <i>WNT2</i> , <i>WNT2B</i> , <i>WNT3</i> , <i>WNT3A</i> , <i>WNT4</i> , <i>WNT5A</i> , <i>WNT5B</i> , <i>WNT6</i> , <i>WNT7A</i> , <i>WNT7B</i> , <i>WNT8A</i> , <i>WNT8B</i> , <i>WNT9A</i> , <i>WNT9B</i> , <i>WNT10A</i> , <i>WNT11</i> , <i>WNT16</i>

f) Additional candidate genes *AXIN2*⁴, *BMP7*¹³⁶, *BMPR1A*¹³⁷, *BOC*¹³⁸, *CDK7*¹³⁹, *CENPH*¹³⁹, *CHD2*¹³⁵, *CRX*^{135,140}, *CRYAB*¹³⁵, *CRYBA1*¹³⁵, *CRYBA2*¹³⁵, *CRYBB3*¹³⁵, *CRYGA*¹³⁵, *CRYGB*¹³⁵, *CRYGD*¹³⁵, *CRYGS*¹³⁵, *CRYZ*¹³⁵, *CYP1B1*¹⁴¹, *DIS3L2*¹⁴², *DISP1*¹⁴³, *DKK1*⁴, *DLX1*¹³⁵, *DLX2*¹³⁵, *EFNA5*¹⁴⁴, *EPHB2*¹⁴⁴, *FGF8*¹⁴⁵, *GAS1*¹⁴⁶, *GLI1*¹⁴⁷, *LHX1*¹³⁵, *NOD2*¹⁴⁸, *PAX3*¹⁴⁹, *RARG*⁶⁸, *RXYLT1*³³, *SEMA3A*¹⁵⁰, *SCLT1*¹⁵¹, *SCRIB*¹⁵², *SLC30A5*¹³⁹, *SNX3*¹⁵³, *TBX2*¹³⁵, *TBX3*⁴, *TBX5*¹³⁵, *TLN1*¹⁵⁴, *TMEM98*¹⁵⁵, *ZIC2*¹⁵⁶, *ZNF219*²⁶, *ZNF503*¹⁵⁷, *ZNF703*¹⁵⁷

† Syndromic and non-syndromic cases have been reported for some genes

‡ Only non-syndromic cases reported for all genes

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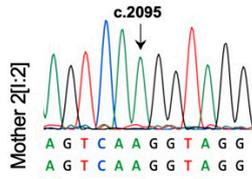
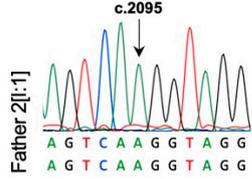
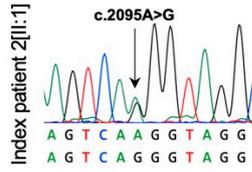
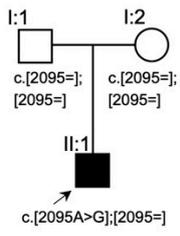
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Family 2: *CHD7*



Family 4: *PTCH1*

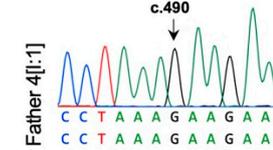
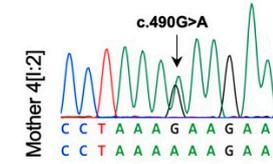
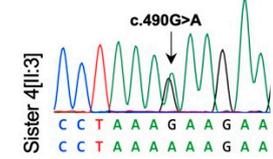
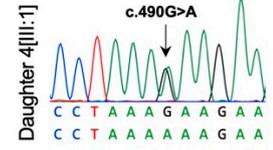
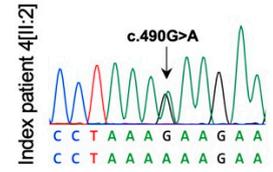
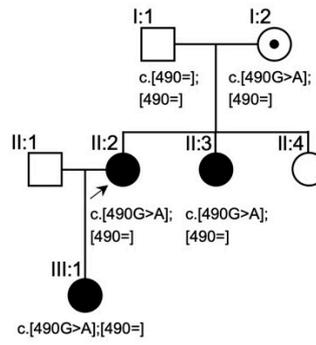
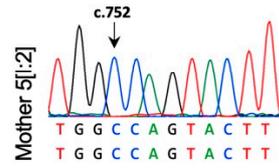
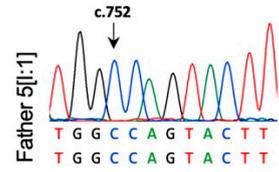
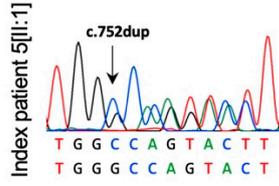
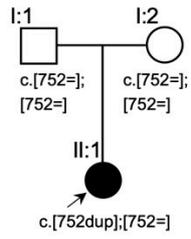
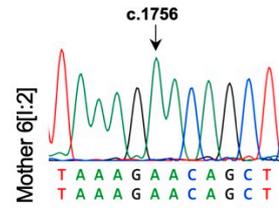
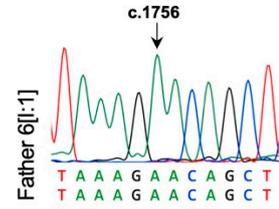
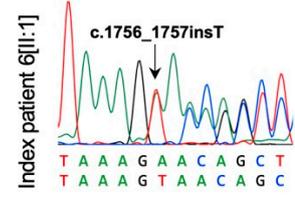
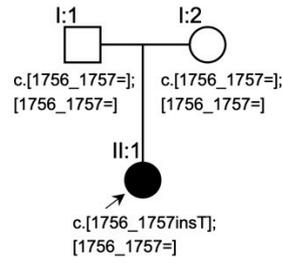


Figure S1: Pedigrees and Sanger sequencing electropherograms of families 2 and 4.

Family 5: PUF60



Family 6: BRPF1



TGFB2: Family 7

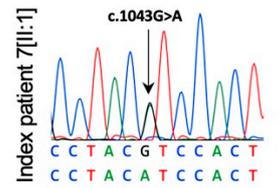
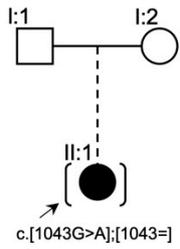


Figure S2: Pedigrees and Sanger sequencing electropherograms of families 5-7.