

**Table S1.** Genes with a recognizable association with macrocephaly, retrieved from OMIM and the scientific literature.

Gene	Gene/Locus name	Cytogenetic location (hg38)
<i>ABCC9</i>	ATP binding cassette subfamily C member 9	12p12.1
<i>ACAN</i>	Aggrecan	15q26.1
<i>ACER3</i>	Alkaline ceramidase 3	11q13.5
<i>ADK</i>	Adenosine kinase	10q22.2
<i>ADNP</i>	Activity dependent neuroprotector homeobox	20q13.13
<i>AKT1</i>	AKT serine/threonine kinase 1	14q32.33
<i>AKT3</i>	AKT serine/threonine kinase 3	1q43-q44
<i>ALKBH8</i>	AlkB homolog 8, tRNA methyltransferase	11q22.3
<i>AMER1</i>	APC membrane recruitment protein 1	Xq11.2
<i>AMPD1</i>	Adenosine monophosphate deaminase 1	1p13.2
<i>ANKH</i>	ANKH inorganic pyrophosphate transport regulator	5p15.2
<i>ANKRD17</i>	Ankyrin repeat domain 17	4q13.3
<i>ANTXR1</i>	Anthrax toxin receptor 1	2p13.3
<i>APC2</i>	APC regulator of WNT signaling pathway 2	19p13.3
<i>APOC2</i>	Apolipoprotein C-II	19q13.32
<i>ARSB</i>	Arylsulfatase B	5q14.1
<i>ARSK</i>	Arylsulfatase family member K	5q15
<i>ASH1L</i>	ASH1 like histone lysine methyltransferase	1q22
<i>ASPA</i>	aspartoacylase	17p13.2
<i>ASXL2</i>	ASXL transcriptional regulator 2	2p23.3
<i>AUTS2</i>	Activator of transcription and developmental regulator AUTS2	7q11.22
<i>B3GLCT</i>	Beta 3-glucosyltransferase	13q12.3
<i>B4GALT1</i>	Beta-1,4-galactosyltransferase 1	9p21.1
<i>B4GALT7</i>	Beta-1,4-galactosyltransferase 7	5q35.3
<i>BAP1</i>	BRCA1 associated protein 1	3p21.1
<i>BGN</i>	Biglycan	Xq28
<i>BICD2</i>	BICD cargo adaptor 2	9q22.31
<i>BICRA</i>	BRD4 interacting chromatin remodeling complex associated protein	19q13.33
<i>BLTP1</i>	Bridge-like lipid transfer protein family member 1	4q27
<i>BRAF</i>	B-Raf proto-oncogene, serine/threonine kinase	7q34
<i>BRWD3</i>	Bromodomain and WD repeat domain containing 3	Xq21.1
<i>C12orf57</i>	Chromosome 12 open reading frame 57	12p13.31
<i>CACNA1E</i>	Calcium voltage-gated channel subunit alpha1 E	1q25.3
<i>CAMK2G</i>	Calcium/calmodulin dependent protein kinase II gamma	10q22.2
<i>CANT1</i>	Calcium activated nucleotidase 1	17q25.3
<i>CASK</i>	Calcium/calmodulin dependent serine protein kinase	Xp11.4
<i>CBL</i>	Cbl proto-oncogene	11q23.3
<i>CCND2</i>	Cyclin D2	12p13.32
<i>CCNK</i>	Cyclin K	14q32.2
<i>CDC42BPB</i>	CDC42 binding protein kinase beta	14q32.32
<i>CDH2</i>	Cadherin 2	18q12.1
<i>CDKN1C</i>	Cyclin-dependent kinase inhibitor 1C	11p15.4
<i>CEP120</i>	Centrosomal protein, 120kD	5q23.2
<i>CHD1</i>	Chromodomain helicase DNA binding protein-1	5q15-q21.1
<i>CHD3</i>	Chromodomain helicase DNA binding protein-3	17p13.1
<i>CHD4</i>	Chromodomain helicase DNA-binding protein-4	12p13.31
<i>CHD8</i>	Chromodomain helicase DNA-binding protein 8	14q11.2
<i>COL2A1</i>	Collagen II, alpha-1 polypeptide	12q13.11
<i>COL3A1</i>	Collagen III, alpha-1 polypeptide	2q32.2
<i>CPE</i>	Carboxypeptidase E	4q32.3
<i>CRPPA</i>	CDP-L-ribitol pyrophosphorylase A	7p21.2
<i>CSF1R</i>	Colony stimulating factor 1 receptor	5q32

<b>5GALNACT1</b>	Chondroitin sulfate N-acetylgalactosaminyltransferase 1	8p21.3
<b>CSNK2A1</b>	Casein kinase 2 alpha 1	20p13
<b>CUL4B</b>	Cullin 4B	Xq24
<b>CUL7</b>	Cullin 7	6p21.1
<b>CWC27</b>	CWC27 spliceosome associated cyclophilin	5q12.3
<b>D2HGDH</b>	D-2-hydroxyglutarate dehydrogenase	2q37.3
<b>DAG1</b>	Dystroglycan 1	3p21.31
<b>DDX3X</b>	DEAD-box helicase 3 X-linked	Xp11.4
<b>DEAF1</b>	DEAF1 transcription factor	11p15.5
<b>DEPDC5</b>	DEP domain containing 5	22q12.2-q12.3
<b>DHCR24</b>	24-dehydrocholesterol reductase	1p32.3
<b>DICER1</b>	Dicer 1, ribonuclease III	14q32.13
<b>DIS3L2</b>	DIS3 like 3'-5' exoribonuclease 2	2q37.1
<b>DLX3</b>	Distal-less homeobox 3	17q21.33
<b>DMXL2</b>	Dmx like 2	15q21.2
<b>DNAJC21</b>	DnaJ heat shock protein family (Hsp40) member C21	5p13.2
<b>DNM1</b>	Dynamitin 1	9q34.11
<b>DNMT3A</b>	DNA methyltransferase 3 alpha	2p23.3
<b>DOCK3</b>	Dedicator of cytokinesis 3	3p21.2
<b>DOCK6</b>	Dedicator of cytokinesis 6	19p13.2
<b>DPYSL5</b>	Dihydropyrimidinase-like 5	2p23.3
<b>DVL1</b>	Dishevelled segment polarity protein 1	1p36.33
<b>DVL3</b>	Dishevelled segment polarity protein 3	3q27.1
<b>DYNC2I1</b>	Dynein 2 intermediate chain 1	7q36.3
<b>DYNC2LI1</b>	Dynein cytoplasmic 2 light intermediate chain 1	2p21
<b>EBP</b>	EBP cholesterol delta-isomerase	Xp11.23
<b>EED</b>	Embryonic ectoderm development	11q14.2
<b>EHMT1</b>	Euchromatic histone lysine methyltransferase 1	9q34.3
<b>EIF2B1</b>	Eukaryotic translation initiation factor 2B subunit alpha	12q24.31
<b>EIF2B2</b>	Eukaryotic translation initiation factor 2B subunit beta	14q24.3
<b>EIF2B3</b>	Eukaryotic translation initiation factor 2B subunit gamma	1p34.1
<b>EIF2B4</b>	Eukaryotic translation initiation factor 2B subunit delta	2p23.3
<b>EIF2B5</b>	Eukaryotic translation initiation factor 2B subunit epsilon	3q27.1
<b>EML1</b>	EMAP like 1	14q32.2
<b>ERF</b>	ETS2 repressor factor	19q13.2
<b>ETFA</b>	Electron transfer flavoprotein subunit alpha	15q24.2-q24.3
<b>ETFB</b>	Electron transfer flavoprotein subunit beta	19q13.41
<b>ETFDH</b>	Electron transfer flavoprotein dehydrogenase	4q32.1
<b>EXT2</b>	Exostosin glycosyltransferase 2	11p11.2
<b>EZH2</b>	Enhancer of zeste 2 polycomb repressive complex 2 subunit	7q36.1
<b>FAM111A</b>	FAM111 trypsin like peptidase A	11q12.1
<b>FAM149B1</b>	Family with sequence similarity 149 member B1	10q22.2
<b>FAM20C</b>	FAM20C golgi associated secretory pathway kinase	7p22.3
<b>FAR1</b>	Fatty acyl-CoA reductase 1	11p15.3
<b>FBN1</b>	Fibrillin 1	15q21.1
<b>FBXW11</b>	F-box and WD repeat domain containing 11	5q35.1
<b>FGFR1</b>	Fibroblast growth factor receptor 1	8p11.23
<b>FGFR2</b>	Fibroblast growth factor receptor 2	10q26.13
<b>FGFR3</b>	Fibroblast growth factor receptor 3	4p16.3
<b>FH</b>	Fumarate hydratase	1q43
<b>FIBP</b>	FGF1 intracellular binding protein	11q13.1
<b>FKTN</b>	Fukutin	9q31.2
<b>FLNA</b>	Filamin A	Xq28
<b>FMR1</b>	Fragile X messenger ribonucleoprotein 1	Xq27.3
<b>FOXP1</b>	Forkhead box P1	3p13
<b>GABBR2</b>	Gamma-aminobutyric acid type B receptor subunit 2	9q22.33
<b>GATAD2B</b>	GATA zinc finger domain containing 2B	1q21.3
<b>GCDH</b>	Glutaryl-coa dehydrogenase	19p13.13

<b>GDF6</b>	Growth differentiation factor 6	8q22.1
<b>GFAP</b>	Glial fibrillary acidic protein	17q21.31
<b>GJA1</b>	Gap junction protein alpha 1	6q22.31
<b>GLI2</b>	GLI family zinc finger 2	2q14.2
<b>GLI3</b>	GLI family zinc finger 3	7p14.1
<b>GNAI1</b>	G protein subunit alpha i1	7q21.11
<b>GNAI3</b>	G protein subunit alpha i3	1p13.3
<b>GNAQ</b>	G protein subunit alpha q	9q21.2
<b>GNE</b>	Glucosamine (UDP-N-acetyl)-2-epimerase/N-acetylmannosamine kinase	9p13.3
<b>GPC3</b>	Glypican 3	Xq26.2
<b>GPC4</b>	Glypican 4	Xq26.2
<b>GPSM2</b>	G protein signaling modulator 2	1p13.3
<b>GRIA3</b>	Glutamate ionotropic receptor AMPA type subunit 3	Xq25
<b>GRIN2B</b>	Glutamate ionotropic receptor NMDA type subunit 2B	12p13.1
<b>GUSB</b>	Glucuronidase beta	7q11.21
<b>H1-4</b>	H1.4 linker histone, cluster member	6p22.2
<b>H3-3A</b>	H3.3 histone A	1q42.12
<b>H3-3B</b>	H3.3 histone B	17q25.1
<b>HADHA</b>	3-oxoacyl-coa dehydrogenase trifunctional multienzyme complex subunit alpha	2p23.3
<b>HDAC6</b>	Histone deacetylase 6	Xp11.23
<b>HECW2</b>	HECT, C2 and WW domain containing E3 ubiquitin protein ligase 2	2q32.3
<b>HEPACAM</b>	Hepatic and glial cell adhesion molecule	11q24.2
<b>HERC1</b>	CT and RLD domain containing E3 ubiquitin protein ligase family member 1	15q22.31
<b>HEXB</b>	Hexosaminidase subunit beta	5q13.3
<b>HMG A2</b>	High mobility group AT-hook 2	12q14.3
<b>HRAS</b>	HRas proto-oncogene, GTPase	11p15.5
<b>HSD17B4</b>	Hydroxysteroid 17-beta dehydrogenase 4	5q23.1
<b>HUWE1</b>	HECT, UBA and WWE domain containing E3 ubiquitin protein ligase 1	Xp11.22
<b>HYLS1</b>	HYLS1 centriolar and ciliogenesis associated	11q24.2
<b>IDS</b>	Iduronate 2-sulfatase	Xq28
<b>IDUA</b>	Alpha-L-iduronidase	4p16.3
<b>IFITM5</b>	Interferon induced transmembrane protein 5	11p15.5
<b>IFT43</b>	Intraflagellar transport 43	14q24.3
<b>IFT74</b>	Intraflagellar transport 74	9p21.2
<b>IFT81</b>	Intraflagellar transport 81	12q24.11
<b>IGBP1</b>	Immunoglobulin binding protein 1	Xq13.1
<b>IGF2</b>	Insulin like growth factor 2	11p15.5
<b>IL6ST</b>	Interleukin 6 cytokine family signal transducer	5q11.2
<b>INPP5E</b>	Inositol polyphosphate-5-phosphatase E	9q34.3
<b>INPPL1</b>	Inositol polyphosphate phosphatase like 1	11q13.4
<b>ITCH</b>	Itchy E3 ubiquitin protein ligase	20q11.22
<b>KCNH1</b>	Potassium voltage-gated channel subfamily H member 1	1q32.2
<b>KCNJ8</b>	Potassium inwardly rectifying channel subfamily J member 8	12p12.1
<b>KCTD13</b>	Potassium channel tetramerization domain containing 13	16p11.2
<b>KDM1A</b>	Lysine demethylase 1A	1p36.12
<b>KDM5B</b>	Lysine demethylase 5B	1q32.1
<b>KDM5C</b>	Lysine demethylase 5C	Xp11.22
<b>KIAA0753</b>	KIAA0753	17p13.1
<b>KIDINS220</b>	Kinase D interacting substrate 220	2p25.1
<b>KIF22</b>	Kinesin family member 22	16p11.2
<b>KIF7</b>	Kinesin family member 7	15q26.1
<b>KMT2E</b>	Lysine methyltransferase 2E (inactive)	7q22.3
<b>KMT5B</b>	Lysine methyltransferase 5B	11q13.2
<b>KPTN</b>	Kaptein, actin binding protein	19q13.32
<b>KRAS</b>	KRAS proto-oncogene, GTPase	12p12.1
<b>L1CAM</b>	L1 cell adhesion molecule	Xq28
<b>L2HGDH</b>	L-2-hydroxyglutarate dehydrogenase	14q21.3
<b>LAMB1</b>	Laminin subunit beta 1	7q31.1

<b>LBR</b>	Lamin B receptor	1q42.12
<b>LRP2</b>	LDL receptor related protein 2	2q31.1
<b>MAB21L2</b>	mab-21 like 2	4q31.3
<b>MADD</b>	MAP kinase activating death domain	11p11.2
<b>MAN1B1</b>	Mannosidase alpha class 1B member 1	9q34.3
<b>MAN2B1</b>	Mannosidase alpha class 2B member 1	19p13.13
<b>MAN2C1</b>	Mannosidase alpha class 2C member 1	15q24.2
<b>MAP2K1</b>	Mitogen-activated protein kinase kinase 1	15q22.31
<b>MAP2K2</b>	Mitogen-activated protein kinase kinase 2	19p13.3
<b>MCCC2</b>	Methylcrotonyl-coa carboxylase subunit 2	5q13.2
<b>MCEE</b>	Methylmalonyl-coa epimerase	2p13.3
<b>MECP2</b>	Methyl-cpg binding protein 2	Xq28
<b>MED12</b>	Mediator complex subunit 12	Xq13.1
<b>MESD</b>	Mesoderm development LRP chaperone	15q25.1
<b>MGAT2</b>	alpha-1,6-mannosyl-glycoprotein 2-beta-N-acetylglucosaminyltransferase	14q21.3
<b>MITF</b>	Melanocyte inducing transcription factor	3p13
<b>MLC1</b>	Modulator of VRAC current 1	22q13.33
<b>MOCS1</b>	Molybdenum cofactor synthesis 1	6p21.2
<b>MOCS2</b>	Molybdenum cofactor synthesis 2	5q11.2
<b>MPDZ</b>	Multiple PDZ domain crumbs cell polarity complex component	9p23
<b>MRAS</b>	Muscle RAS oncogene homolog	3q22.3
<b>MSL3</b>	MSL complex subunit 3	Xp22.2
<b>MTOR</b>	Mechanistic target of rapamycin kinase	1p36.22
<b>MTM1</b>	Myotubularin 1	Xq28
<b>MUSK</b>	Muscle associated receptor tyrosine kinase	9q31.3
<b>MYCN</b>	MYCN proto-oncogene, bHLH transcription factor	2p24.3
<b>MYH8</b>	Myosin heavy chain 8	17p13.1
<b>MYMK</b>	Myomaker, myoblast fusion factor	9q34.2
<b>NBPF1</b>	NBPF member 1	1p36.13
<b>NDN</b>	Necdin, MAGE family member	15q11.2
<b>NDUFAF3</b>	NADH:ubiquinone oxidoreductase complex assembly factor 3	3p21.31
<b>NDUFAF5</b>	NADH:ubiquinone oxidoreductase complex assembly factor 5	20p12.1
<b>NDUFS4</b>	NADH:ubiquinone oxidoreductase subunit S4	5q11.2
<b>NDUFV1</b>	NADH:ubiquinone oxidoreductase core subunit V1	11q13.2
<b>NEK1</b>	NIMA related kinase 1	4q33
<b>NF1</b>	Neurofibromin 1	17q11.2
<b>NFIA</b>	Nuclear factor I A	1p31.3
<b>NFIB</b>	Nuclear factor I B	9p23-p22.3
<b>NFIX</b>	Nuclear factor I X	19p13.13
<b>NKX3-2</b>	NK3 homeobox 2	4p15.33
<b>NONO</b>	Non-POU domain containing octamer binding	Xq13.1
<b>NR2F1</b>	Nuclear receptor subfamily 2 group F member 1	5q15
<b>NRAS</b>	NRAS proto-oncogene, GTPase	1p13.2
<b>NSD1</b>	Nuclear receptor binding SET domain protein 1	5q35.3
<b>NXN</b>	Nucleoredoxin	17p13.3
<b>OBSL1</b>	Obscurin like cytoskeletal adaptor 1	2q35
<b>ODC1</b>	Ornithine decarboxylase 1	2p25.1
<b>OFD1</b>	OFD1 centriole and centriolar satellite protein	Xp22.2
<b>OPHN1</b>	Oligophrenin 1	Xq12
<b>OTUD5</b>	OTU deubiquitinase 5	Xp11.23
<b>P4HB</b>	Prolyl 4-hydroxylase subunit beta	17q25.3
<b>PAK1</b>	P21 (RAC1) activated kinase 1	11q13.5-q14.1
<b>PAK3</b>	P21 (RAC1) activated kinase 3	Xq23
<b>PAM16</b>	Presequence translocase associated motor 16	16p13.3
<b>PC</b>	Pyruvate carboxylase	11q13.2
<b>PCGF2</b>	Polycomb group ring finger 2	17q12
<b>PDSS1</b>	Decaprenyl diphosphate synthase subunit 1	10p12.1
<b>PEX1</b>	Peroxisomal biogenesis factor 1	7q21.2

<b>PHF21A</b>	PHD finger protein 21A	<b>11p11.2</b>
<b>PHF6</b>	PHD finger protein 6	<b>Xq26.2</b>
<b>PIGA</b>	Phosphatidylinositol glycan anchor biosynthesis class A	<b>Xp22.2</b>
<b>PIGM</b>	Phosphatidylinositol glycan anchor biosynthesis class M	<b>1q23.2</b>
<b>PIGN</b>	Phosphatidylinositol glycan anchor biosynthesis class N	<b>18q21.33</b>
<b>PIGT</b>	Phosphatidylinositol glycan anchor biosynthesis class T	<b>20q13.12</b>
<b>PIGV</b>	Phosphatidylinositol glycan anchor biosynthesis class V	<b>1p36.11</b>
<b>PIK3CA</b>	Phosphatidylinositol-4,5-bisphosphate 3-kinase catalytic subunit alpha	<b>3q26.32</b>
<b>PIK3R1</b>	Phosphoinositide-3-kinase regulatory subunit 1	<b>5q13.1</b>
<b>PIK3R2</b>	Phosphoinositide-3-kinase regulatory subunit 2	<b>19p13.11</b>
<b>PKDCC</b>	Protein kinase domain containing, cytoplasmic	<b>2p21</b>
<b>PLAG1</b>	PLAG1 zinc finger	<b>8q12.1</b>
<b>PLCB4</b>	Phospholipase C beta 4	<b>20p12.3-p12.2</b>
<b>PLCH1</b>	Phospholipase C eta 1	<b>3q25.31</b>
<b>PLG</b>	Plasminogen	<b>6q26</b>
<b>POC1A</b>	POC1 centriolar protein A	<b>3p21.2</b>
<b>POLE</b>	DNA polymerase epsilon, catalytic subunit	<b>12q24.33</b>
<b>POLR3A</b>	RNA polymerase III subunit A	<b>10q22.3</b>
<b>POMK</b>	Protein O-mannose kinase	<b>8p11.21</b>
<b>POMT1</b>	Protein O-mannosyltransferase 1	<b>9q34.13</b>
<b>POP1</b>	POP1 homolog, ribonuclease P/MRP subunit	<b>8q22.2</b>
<b>PPP1CB</b>	Protein phosphatase 1 catalytic subunit beta	<b>2p23.2</b>
<b>PPP1R12A</b>	Protein phosphatase 1 regulatory subunit 12A	<b>12q21.2-q21.31</b>
<b>PPP2CA</b>	Protein phosphatase 2 catalytic subunit alpha	<b>5q31.1</b>
<b>PPP2R5D</b>	Protein phosphatase 2 regulatory subunit b'delta	<b>6p21.1</b>
<b>PTCH1</b>	Patched 1	<b>9q22.32</b>
<b>PTCH2</b>	Patched 2	<b>1p34.1</b>
<b>PTEN</b>	Phosphatase and tensin homolog	<b>10q23.31</b>
<b>PTPN11</b>	Protein tyrosine phosphatase non-receptor type 11	<b>12q24.13</b>
<b>PUF60</b>	Poly(U) binding splicing factor 60	<b>8q24.3</b>
<b>RAB39A</b>	RAB39A, member RAS oncogene family	<b>11q22.3</b>
<b>RAB39B</b>	RAB39B, member RAS oncogene family	<b>Xq28</b>
<b>RAC1</b>	Rac family small GTPase 1	<b>7p22.1</b>
<b>RAF1</b>	Raf-1 proto-oncogene, serine/threonine kinase	<b>3p25.2</b>
<b>RALA</b>	RAS like proto-oncogene A	<b>7p14.1</b>
<b>RHEB</b>	Ras homolog, mTORC1 binding	<b>7q36.1</b>
<b>RIN2</b>	Ras and Rab interactor 2	<b>20p11.23</b>
<b>RIT1</b>	Ras like without CAAX 1	<b>1q22</b>
<b>RNF125</b>	Ring finger protein 125	<b>18q12.1</b>
<b>RNF135</b>	Ring finger protein 135	<b>17q11.2</b>
<b>ROR2</b>	Receptor tyrosine kinase like orphan receptor 2	<b>9q22.31</b>
<b>RPS6KA3</b>	Ribosomal protein S6 kinase A3	<b>Xp22.12</b>
<b>RRAS2</b>	RAS related 2	<b>11p15.2</b>
<b>RXYLT1</b>	Ribitol xylosyltransferase 1	<b>12q14.2</b>
<b>SATB2</b>	SATB homeobox 2	<b>2q33.1</b>
<b>SEC23A</b>	SEC23 homolog A, COPII coat complex component	<b>14q21.1</b>
<b>SEC23B</b>	SEC23 homolog B, COPII coat complex component	<b>20p11.23</b>
<b>SEC24D</b>	SEC24 homolog D, COPII coat complex component	<b>4q26</b>
<b>SERPINH1</b>	serpin family H member 1	<b>11q13.5</b>
<b>SETD2</b>	SET domain containing 2, histone lysine methyltransferase	<b>3p21.31</b>
<b>SGSH</b>	N-sulfoglucosamine sulfohydrolase	<b>17q25.3</b>
<b>SHANK3</b>	SH3 and multiple ankyrin repeat domains 3	<b>22q13.33</b>
<b>SHOC2</b>	SHOC2 leucine rich repeat scaffold protein	<b>10q25.2</b>
<b>SIX3</b>	SIX homeobox 3	<b>2p21</b>
<b>SLC25A1</b>	Solute carrier family 25 member 1	<b>22q11.21</b>
<b>SLC2A1</b>	Solute carrier family 2 member 1	<b>1p34.2</b>
<b>SLC2A10</b>	Solute carrier family 2 member 10	<b>20q13.12</b>
<b>SLC44A1</b>	Solute carrier family 44 member 1	<b>9q31.1-q31.2</b>

<b>SMAD4</b>	SMAD family member 4	18q21.2
<b>SMO</b>	Smoothened, frizzled class receptor	7q32.1
<b>SNRPN</b>	Small nuclear ribonucleoprotein polypeptide N	15q11.2
<b>SNX10</b>	Sorting nexin 10	7p15.2
<b>SNX14</b>	Sorting nexin 14	6q14.3
<b>SON</b>	SON DNA and RNA binding protein	21q22.11
<b>SOS1</b>	SOS Ras/Rac guanine nucleotide exchange factor 1	2p22.1
<b>SOST</b>	Sclerostin	17q21.31
<b>SOX9</b>	SRY-box transcription factor 9	17q24.3
<b>SPINT2</b>	Serine peptidase inhibitor, Kunitz type 2	19q13.2
<b>SPOP</b>	Speckle type BTB/POZ protein	17q21.33
<b>SPRED1</b>	Sprouty related EVH1 domain containing 1	15q14
<b>SPRED2</b>	Sprouty related EVH1 domain containing 2	2p14
<b>STRADA</b>	STE20 related adaptor alpha	17q23.3
<b>STT3A</b>	STT3 oligosaccharyltransferase complex catalytic subunit A	11q24.2
<b>SUFU</b>	SUFU negative regulator of hedgehog signaling	10q24.32
<b>SUZ12</b>	SUZ12 polycomb repressive complex 2 subunit	17q11.2
<b>SYN1</b>	Synapsin I	Xp11.3-p11.23
<b>SZT2</b>	SZT2 subunit of KICSTOR complex	1p34.2
<b>TAOK1</b>	TAO kinase 1	17q11.2
<b>TBC1D2B</b>	TBC1 domain family member 2B	15q24.3-q25.1
<b>TBC1D7</b>	TBC1 domain family member 7	6p24.1
<b>TBCK</b>	TBC1 domain containing kinase	4q24
<b>TBL1X</b>	Transducin beta like 1 X-linked	Xp22.31-p22.2
<b>TCF20</b>	Transcription factor 20	22q13.2
<b>TCIRG1</b>	T cell immune regulator 1, ATPase H <sup>+</sup> transporting V0 subunit a3	11q13.2
<b>TET3</b>	Tet methylcytosine dioxygenase 3	2p13.1
<b>THRA</b>	Thyroid hormone receptor alpha	17q21.1
<b>TMCO1</b>	Transmembrane and coiled-coil domains 1	1q24.1
<b>TMEM53</b>	Transmembrane protein 53	1p34.1
<b>TMEM165</b>	Transmembrane protein 165	4q12
<b>TMEM216</b>	Transmembrane protein 216	11q12.2
<b>TNFRSF11B</b>	TNF receptor superfamily member 11b	8q24.12
<b>TONSL</b>	Tonsoku like, DNA repair protein	8q24.3
<b>TPI1</b>	Triosephosphate isomerase 1	12p13.31
<b>TRIM37</b>	Tripartite motif containing 37	17q22
<b>TRIO</b>	Trio Rho guanine nucleotide exchange factor	5p15.2
<b>TRIP11</b>	Thyroid hormone receptor interactor 11	14q32.12
<b>TRIP12</b>	Thyroid hormone receptor interactor 12	2q36.3
<b>TSC1</b>	TSC complex subunit 1	9q34.13
<b>TSC2</b>	TSC complex subunit 2	16p13.3
<b>UPF3B</b>	UPF3B regulator of nonsense mediated mRNA decay	Xq24
<b>USP9X</b>	ubiquitin specific peptidase 9 X-linked	Xp11.4
<b>VPS35L</b>	VPS35 endosomal protein sorting factor like	16p12.3
<b>VPS4A</b>	Vacuolar protein sorting 4 homolog A	16q22.1
<b>WASHC5</b>	WASH complex subunit 5	8q24.13
<b>WDR81</b>	WD repeat domain 81	17p13.3
<b>WNT5A</b>	Wnt family member 5A	3p14.3
<b>XYLT1</b>	Xylosyltransferase 1	16p12.3
<b>YME1L1</b>	YME1 like 1 ATPase	10p12.1
<b>ZBTB20</b>	Zinc finger and BTB domain containing 20	3q13.31
<b>ZBTB42</b>	Zinc finger and BTB domain containing 42	14q32.33
<b>ZBTB7A</b>	Zinc finger and BTB domain containing 7A	19p13.3
<b>ZDHHC9</b>	Zinc finger DHHC-type palmitoyltransferase 9	Xq26.1
<b>ZIC2</b>	Zic family member 2	13q32.3
<b>ZNF469</b>	Zinc finger protein 469	16q24.2
<b>ZSWIM6</b>	Zinc finger SWIM-type containing 6	5q12.1

**Figure S1.** Enriched biological processes of the gene set related to macrocephaly

