



Prenatal Targeted Oligo + SNP 60k Array CGH Disease Regions Screened

Associated Condition	Gene	Chr/Band	OMIM #	Associated Condition	Gene	Chr/Band	OMIM #
1p32-31 Deletion	DIRAS3	1p32-p31	613735	16q11.2-q12.2 Deletion	Multiple	16q11.2-q12.2	
1p36 Deletion	Multiple	1p36	607872	17p13.1 Deletion	Multiple	17p13.1	613776
1q21.1 Deletion (1.35 Mb)	Multiple	1q21.1	612474	17p13.3 Distal deletion	Multiple	17p13.3	
1q21.1 Deletion (200 Kb)	Multiple	1q21.1	274000	17p13.3 Distal duplication, not including PAFAH1B1	Multiple	17p13.3	613215
1q41-42 Deletion	Multiple	1q41-q42	612530	17p13.3 Proximal duplication, including PAFAH1B1	Multiple	17p13.3	
1q43-44 Deletion	AKT3	1q43-q44	612337	17q11.2 Deletion	NF1	17q11.2	613675
2p16.1-p15 Deletion	Multiple	2p16.1-15	612513	17q12 Deletion	Multiple	17q12	614527
2p16.3 Deletion	NRXN1	2p16.3	600565	17q12 Duplication	Multiple	17q12	614526
2p21 Deletion	Multiple	2p21	606407	17q21.31 Deletion	Multiple	17q21.31	610443
2q23.1 Deletion	EPC2,	2q23.1	611000,	17q21.31 Duplication	Multiple	17q21.31	613533
	MBD5		611472	17q23.1-q23.2 Deletion	Multiple	17q23.1-q23.2	613355
2q32.2-q33 Deletion	SATB2	2q32.2q33	119540	18q Deletion	Multiple	18q	601808
2q37.3 Deletion	HDAC4	2q37.3	600430	18q21.1 Deletion	TCEB3B	18q21.1	609522
3q29 Deletion	PAK2	3q29	609425	19q13.11 Deletion	Multiple	19q13.11	613026
3q29 Duplication	Multiple	3q29	611936	22q11.2 Distal deletion	Multiple	22q11.2	611867
5p13.2 Duplication	NIPBL	5p13.2	613174	22q11.2 Duplication	Multiple	22q11.2	608363
5q35.2-q35.3 Duplication	NSD1	5q35.2-q35.3	606681	22q13.3 Deletion	Multiple	22q13.3	606232
	FOXC1	6p24	612582	Xp11.23-p11.22 Duplication	Multiple	Xp11.23-p11.22	300801
6pter-p24 Deletion	Multiple	6q24.3	612863				
6q24.3 Deletion	Multiple	6q25.2-q25.3	612863	18q21.3 Deletion	Multiple	18q21.3	300578
6q25.2-q25.3 Deletion	Multiple	7q11.23	609757	Xp21.2-p11.4 Deletion	Multiple	Xp21.2-p11.4	
7q11.23 Duplication	CTSB	8p23.1	116810				
8p23.1 Deletion	Multiple	8p23.1	222400	Xq28 Duplication	Multiple	Xq28	300815
8p23.1 Deletion / Congenital diaphragmatic hernia 2 (CDH2)	Multiple	8p23.1		Aarskog-Scott syndrome	FGD1	Xp11.22	305400
8p23.1 Duplication	CHD7	8q12.1-q12.2	608892	Adrenal hypoplasia congenital	NR0B1	Xp21.2	300200
8q12 Duplication	Multiple	9p	158170	Agammaglobulinemia, X-linked	BTK	Xq22.1	300755
9p Deletion	GABBR2,	9q22.32-	607340,	Alagille	JAG1	20p12.2	118450
9q22.32-q22.33 Deletion	TGFBR1	q22.33	190181	Albright hereditary osteodystrophy-like syndrome/2q37 deletion syndrome	HDAC4	2q37.3	600430
	EHMT1	9q34.3	610253	Alpha-thalassemia mental retardation	HBA1, HBA2, SOX8	16p13.3	141750
9q34.3 Deletion / Kleefstra	Multiple	10q22-q23		Alpha-thalassemia mental retardation, X-linked	ATRX	Xq21.1	300032
10q22-23 Deletion	GRIP1,	12q14.1-q15	604597,	Alport plus diffuse leiomyomatosis, X-linked	COL4A6	Xq22.3	308940
12q14.1-q15 Deletion	HMGA2,	HMGA2,	600698,	Alport, X-linked	COL4A5	Xq22.3	301050
	LEMD3		607844	Androgen insensitivity	AR	Xq12	300068
12q14.3-Related primordial dwarfism / Russell-Silver-like	HMGA2	12q14.3	600698	Angelman / Prader-Willi	MAGEL2,	15q11.2	105830,
12q24.21-q24.23 Duplication	Multiple	12q24.21-q24.23			MKRN3,	NDN,	176270,
					NIPA1,	NIPA2,	105830
13q14 Deletion	Multiple	13q14	613884		PWS IC,	SNORD	
14q11-q22 Deletion	Multiple	14q11-q22	613457		107/64/108,	SNORD	
14q12 Duplication	FOGX1	14q12	164874		109B,	SNORD	
14q22-q23 Deletion	Multiple	14q22-q23			116 cluster,		
14q32.2 Deletion causing UPD(14) maternal phenotype	DLK1,	14q32.2	176290,		SnoRNA,		
	RTL1		611896		SNRPN,		
14q32.2 Deletion causing UPD(14) paternal phenotype	MEG3,	14q32.2	605636,		SNURF,		
	MEG8,		613648,		UBE3A		
	RTL1		611896				
15q11-q13 Duplication	Multiple	15q11-q13	608636				
15q13.3 Deletion	Multiple	15q13.3	612001	Aniridia	PAX6	11p13	106210
15q24 Deletion	Multiple	15q24	613406	ANKRD11 haploinsufficiency	ANKRD11	16q24.3	611192
15q24.1-q24.2 Duplication	Multiple	15q24.1-q24.2		Atrial septal defect 7	NKX2-5	5q35.1	108900
16p11.2 Deletion	Multiple	16p11.2	611913				
16p12.1 Deletion	Multiple	16p12.1	136570				
16p12.2-p11.2 Deletion	Multiple	16p12.2-p11.2	613604				
16p13.11 Deletion	Multiple	16p13.11					
16p13.3 Deletion	Multiple	16p13.3	610543				
16p13.3 Duplication	CREBBP	16p13.3	613458				

Associated Condition	Gene	Chr/Band	OMIM #	Associated Condition	Gene	Chr/Band	OMIM #
Autism, SHANK2-related	SHANK2	11q13.3-q13.4	613436	Duane radial ray	SALL4	20q13.2	607323
Autistic Features, X-linked, susceptibility	NLGN4	Xp22.32-p22.31		Duchenne muscular dystrophy	DMD	Xp21.2-p21.1	310200
AZF Regions	Multiple	Yq11.21	415000	Early-onset ataxia with oculomotor apraxia and hypoalbuminemia	APTX	9p21.1	208920
Bannayan-Riley-Ruvalcaba	PTEN	10q23.31	153480	Emery-Dreifuss muscular dystrophy 1, X-linked (EDMD1)	EMD	Xq28	300384
Bartter 1	SLC12A1	15q21.1	601678	EPHA7 haploinsufficiency	EPHA7	6q16.1	602190
Bartter 2	KCNJ1	11q24.3	241200	Epileptic encephalopathy, early infantile	PCDH19	Xq22.1	300088
Bartter 3	CLCNKB	1p36.13	607364	Familial adenomatous polyposis 1	APC	5q22.2	175100
Bartter 4A	BSND	1p32.3	602522	Feingold	MYCN	2p24.3	164280
Bartter 4B	CLCNKA	1p36.13	613090	Focal dermal hypoplasia	PORCN	Xp11.23	305600
Bartter with autosomal dominant hypocalcemia	CASR	3q21.1	601199	Fragile X	AFF2, FMR1	Xq28, Xq27.3	300624
Basal cell nevus	PTCH1	9q22.32	109400	Fryns	DISP1	1q41	229850
Beckwith-Wiedemann	CDKN1C, H19, H19	11p15.4, 11p15.5	130650	Gitelman	SLC12A3	16q13	263800
	IC, IGF2, INS,			GLUT1 deficiency	SLC2A1	1p34.2	606777
	KCNQ1 IC, PHLDA2,			Glycerol kinase deficiency	GK	Xp21.2	307030
	SLC22A18			Gonadal Dysgenesis	SRY	Yp11.31	480000
Benign familial neonatal seizures	KCNQ2	20q13.33	121200	Greig cephalopolysyndactyly	GLI3	7p14.1	175700
Beta thalassemia	HBB	11p15.4	613985	Hemophilia A	F8	Xq28	306700
Bilateral frontoparietal polymicrogyria	GPR56	16q21	606854	Hemophilia B	F9	Xq27.1	306900
Blepharophimosis	FOXL2	3q22.3	110100	Hereditary hemorrhagic telangiectasia	SMAD4	18q21.2	175050
Branchiootic syndrome 3	SIX1	14q23.1	608389	Hereditary hemorrhagic telangiectasia 2	ACVR1L	12q13.13	600376
Branchiootorenal	EYA1	8q13.3	113650	Heterotaxy, visceral 1	ZIC3	Xq26.3	306955
Campomelic dysplasia	SOX9	17q24.3	114290	Hirschsprung disease, susceptibility to, 1	RET	10q11.21	142623
Cat-eye syndrome	ATP6V1E1	22q11	115470	Holoprosencephaly 1	LSS, TRAPP10	21q22.3	236100
Cerebellar ataxia, mental retardation, and dysequilibrium	VLDLR	9p24.2	224050	Holoprosencephaly 2	SIX3	2p21	157170
Cerebral cavernous malformations 1	KRIT1	7q21.2	116860	Holoprosencephaly 3	SHH	7q36.3	142945
Cerebral cavernous malformations 2	CCM2	7p13	603284	Holoprosencephaly 4	TGIF1	18p11.31	142946
Cerebral cavernous malformations 3	PDCD10	3q26.1	603285	Holoprosencephaly 5	ZIC2	13q32.3	609637
Charcot Marie Tooth Type 1A	PMP22	17p12	118220	Holoprosencephaly 6	Multiple	2q37.1-q37.3	605934
CHARGE	CHD7	8q12.1-q12.2	214800	Holoprosencephaly 7	PTCH1	9q22.32	610828
Chondrodysplasia punctata 1, X-linked	ARSE	Xp22.33	302950	Holoprosencephaly 8	Multiple	14q13.1-q13.2	609408
Choroideremia	CHM	Xq21.2	303100	Holoprosencephaly 9	GLI2	2q14.2	610829
Chronic granulomatous disease	CYBB	Xp11.4	306400	Holoprosencephaly and preaxial polydactyly	FBXW11	5q35.1	264480
Cleidocranial dysplasia	RUNX2	6p21.1	119600	Holt-Oram	TBX5	12q24.21	142900
Cohen syndrome	VPS13B	8q22.2	216550	Hydrocephalus and nephrogenic diabetes insipidus	AVPR2, L1CAM	Xq28	
Congenital diaphragmatic hernia	Multiple	15q26.1	142340	Hyper-IgE recurrent infection	DOCK8	9p24.3	243700
Congenital hemidysplasia with ichthyosiform erythroderma and limb defects (CHILD)	NSDHL	Xq28	308050	Hypohidrotic ectodermal dysplasia (XHED)	EDA	Xq13.1	305100
Cornelia de Lange 1	NIPBL	5p13.2	122470	Hypoparathyroidism, sensorineural deafness, renal disease	GATA3	10p14	146255
Cornelia de Lange 2	SMC1A	Xp11.22	300590	Hypospadius 2	MAMLD1	Xq28	300758
Cortical dysplasia-focal epilepsy	CNTNAP2	7q35-q36	610042	Hypotonia-cystinuria	PREPL, SLC3A1	2p21	606407
Cowden disease	PTEN	10q23.31	158350	Ichthyosis, X-linked	STS, VCX3A	Xp22.31	308100
Craniofrontonasal	EFNB1	Xq13.1	304110	Idiopathic short stature (ISSX)	SHOX	Xp22.33, Yp11.32	300582
Craniosynostosis type 2	MSX2	5q35.2	604757	Incontinentia Pigmenti	IKBKG	Xq28	308300
Cri-du-Chat	TERT	5p15.2	123450	Infantile hyperinsulinism with enteropathy and deafness	ABCC8, KCNJ11, USH1C	11p15-p14	606528
Curarino	MNX1	7q36.3	176450	Infantile spasms, CDKL5-related / Atypical Rett	CDKL5	Xp22.13	300672
Dandy-Walker	Multiple	3q22-q24	220200	Infantile spasms, MAGI2-related	MAGI2	7q21.11	606382
Deafness, autosomal recessive 22	OTOA	16p12.2	607039				
DiGeorge 1	CRKL, HIRA, TBX1	22q11.21	188400				
DiGeorge 2	Multiple	10p14-p13	601362				
Down syndrome critical region	Multiple	21q22.3	190685				
Dravet	SCN1A	2q24.3	607208				



Prenatal Targeted Oligo + SNP 60k Array CGH Disease Regions Screened (continued)

Associated Condition	Gene	Chr/Band	OMIM #	Associated Condition	Gene	Chr/Band	OMIM #
Intellectual disability with cerebellar hypoplasia and distinctive facial appearance	OPHN1	Xq12	300486	Microphthalmia, isolated, with cataract 2 Miller-Dieker	SIX6 ABR, PAFAH1B1, PITPNAs, YWHAE	14q23.1 17p13.3	212550 247200
Intellectual disability with language impairment and autistic features	FOXP1	3p13	613670	Mohr-Tranebaerg	TIMM8A	Xq22.1	304700
Intellectual disability with microcephaly with pontine and cerebellar hypoplasia	CASK	Xp11.4	300749	Mowat-Wilson	ZEB2	2q22.3	235730
Intellectual disability with panhypopituitarism	SOX3	Xq27.1	300123	Muscular Dystrophy-Dystroglycanopathy A6	LARGE	22q12.3	613154
Intellectual disability with stereotypical movements, epilepsy, and/or cerebral malformations	MEF2C	5q14.3	613443	Myoclonus dystonia	SGCE	7q21.3	159900
Jacobsen	Multiple	11q23	147791	Myotubular myopathy 1	MTM1	Xq28	310400
Joubert 3	AHI1	6q23.3	608629	Nablus mask-like facial	Multiple	8q22.1	608156
Joubert 4	NPHP1	2q13	609583	Nail-patella (NPS)	LMX1B	9q33.3	161200
Joubert 5	CEP290	12q21.32	610188	Nephronophthisis 1	NPHP1	2q13	256100
Juvenile polyposis, BMPR1A-related	BMPR1A	10q23.2	174900	Nephropathic cystinosis	CTNS	17p13.2	219800
Juvenile polyposis, SMAD4-related	SMAD4	18q21.2	174900	Neurofibromatosis 1	JJAZ1, NF1	17q11.2	162200
Kallmann 1	KAL1	Xp22.31	308700	Neurofibromatosis 2	NF2	22q12.2	101000
Langer Mesomelic Dysplasia	SHOX	Xp22.33, Yp11.32	249700	Neurosensory deafness (DFNB1)	GJB6	13q12.11	220290
Langer-Giedion	Multiple	8q24.11-q24.13	150230	NFIA haploinsufficiency	NFIA	1p31.3	600727
Laron syndrome	GHR	5p13-p12	262500	Noonan 1	PTPN11	12q24.13	163950
Leri-Weill Dyschondrosteosis	SHOX	Xp22.33, Yp11.32	127300	Noonan 4	SOS1	2p22.1	610733
Lesch-Nyhan	HPRT1	Xq26.2-q26.3	300322	Noonan 5	RAF1	3p25.2	611553
Leukodystrophy	Multiple	11q14.2-q14.3		Norrie disease	NDP	Xp11.3	310600
Li-Fraumeni 1	TP53	17p13.1	151623	Oculocutaneous albinism 2 (OCA2)	OCA2	15q12-q13	203200
Lissencephaly 1	PAFAH1B1	17p13.3	607432	Opitz G	MID1	Xp22.2	300000
Lissencephaly 2	RELN	7q22.1	257320	Ornithine transcarbamylase deficiency	OTC	Xp11.4	311250
Lissencephaly, X-linked	DCX	Xq23	300067	Orofacial Cleft 11	BMP4	14q22.2	600625
Loeys-Dietz, type 1A	TGFBR1	9q22.33	609192	Orofaciodigital 1	OFD1	Xp22.2	311200
Loeys-Dietz, type 2B	TGFBR2	3p24.1	610380	Osteogenesis Imperfata	COL1A1	17q21.33	166200
Long QT	KCNQ1	11p15.5-p15.4	192500	Osteopathia striata with cranial sclerosis	FAM123B	Xq11.2	300373
Lowe	OCRL	Xq25-q26	309000	Otodontal dysplasia	FGF3	11q13	166750
Lymphoproliferative, X-linked	SH2D1A	Xq25	308240	Pallister-Killian	Multiple	12p	601803
Macrocephaly/Autism	PTEN	10q23.31	605309	Paragangliomas 1	SDHD	11q23.1	168000
Marfan 1	FBN1	15q21.1	154700	Paragangliomas 4	SDHB	1p36.13	115310
McLeod	XK	Xp21.1	300842	Pelizaeus-Merzbacher disease	PLP1	Xq22.2	312080
Menkes	ATP7A	Xq21.1	309400	Peutz-Jeghers	STK11	19p13.3	175200
Mental retardation 1	MBD5	2q23.1	156200	Pitt-Hopkins	TCF4	18q21.2	610954
Mental retardation 21	IL1RAPL1	Xp21.3-p21.2	300143	Polycystic kidney disease 1	PKD1	16p13.3	601313
Mental retardation 5	SYNGAP1	6p21.32	612621	Polycystic kidney disease 2	PKD2	4q22.1	613095
Mental retardation 6	GRIK2	6q16.3	611092	Potocki-Lupski	Multiple	17p11.2	610883
Mental retardation 7	TUSC3	8p22	611093	Potocki-Shaffer	Multiple	11p11.2	601224
Mental retardation 9	FTSJ1	Xp11.23	309549	Prader-Willi-like phenotype	SIM1	6q16.3	176270
Mental retardation 94	GRIA3	Xq25	300699	Renal cysts and diabetes	HNF1B	17q12	137920
Mesomelic dysplasia, Kantaputra type	HOXD	2q24-q32 gene cluster	156232	Retinoblastoma	RB1	13q14.2	180200
Metachromatic leukodystrophy	ARSA	22q13.33	250100	Rett Syndrome	MECP2	Xq28	312750
Microphthalmia 2	BCOR	Xp11.4	300166	Rett syndrome, congenital variant	FOXG1	14q12	613454
Microphthalmia 3	SOX2	3q26.33	206900	Rieger	PITX2	4q25	180500
Microphthalmia 5	OTX2	14q22.3	610125	Rubinstein-Taybi	CREBBP, DNASE1	16p13	180849
Microphthalmia 7	HCCS	Xp22.2	309801	Russell-Silver	CPA4, GRB1	7q32.2, 7p11.2	180860
				Saethre-Chotzen	TWIST1	7p21.1	101400
				Short stature	LHX4	1q25.2	262700
				SHOX	SHOX	Xp22.33	312865
				Siderius type intellectual disability	PHF8	Xp11.22	300263
				Simpson-Golabi-Behmel 1	GPC3	Xq26.2	312870
				Smith-Lemli-Optiz	DHCR7	11q13.4	270400
				Smith-Magenis	RAI1	17p11.2	182290

Associated Condition	Gene	Chr/Band	OMIM #	Associated Condition	Gene	Chr/Band	OMIM #
Sotos	NSD1	5q35.2-q35.3	117550	XY Sex Reversal 2	NROB1	Xp21.2	300018
Speech-language disorder 1	FOXP2	7q31.1	602081	XY Sex Reversal 3	NR5A1	9q33.3	612965
Split-hand / foot malformation 1	SHFM1	7q21.2-q21.3	183600	XY Sex Reversal 4	DMRT1	9p24.3	
Split-hand / foot malformation 1 with deafness	DLX5	7q21.3	220600	Unique pericentromeric regions	Multiple	43 sites	
Split-hand / foot malformation 3	BTRC, FBXW4, LBX1, POLL	10q24	246560	Unique subtelomeric regions	Multiple	41 sites	
Split-hand / foot malformation 4	TP63	3q28	605289	Aneuploidy	Multiple	Any individual chromosome	
Split-hand / foot malformation 5	DLX1, DLX2	2q31	606708	Uniparental Disomy (UPD)	Multiple	Chromosomes 7,11,14,15, 16 & 20	
Stickler 1	COL2A1	12q13.11	108300				
Stickler 2	COL11A1	1p21.1	604841				
Synpolydactyly	EVX2, HOXD13, HOXD9	2q31.1	186000				
Timothy Syndrome	CACNA1C	12p13.33	601005				
Toe syndactyly, telecanthus, anogenital and renal malformations (STAR)	FAM58A	Xq28	300707				
Townes-Brocks	SALL1	16q12.1	107480				
Treacher-Collins	TCOF1	5q32	154500				
Tuberous sclerosis 1	TSC1	9q34.13	191100				
Tuberous sclerosis 2	TSC2	16p13.3	613254				
Ulnar-mammary	TBX3	12q24.21	181450				
Usher IIC	GPR98	5q14.3	605472				
Van der Woude	IRF6	1q32.2	119300				
Vascular endothelial growth factor disorders	VEGFA	6p21.1	192240				
Velocardiofacial	TBX1	22q11.21	192430				
Von Hippel-Lindau	VHL	3p25.3	193300				
Waardenburg I	PAX3	2q36.1	193500				
Waardenburg IIA	MITF	3p14.1-p13	193510				
WAGR	PAX6, WT1	11p13 WT1	194072				
Williams-Beuren	ELN, GTF2I, GTF2IRD1, LIMK1, NCF1	7q11.23	194050				
Wilms Tumor 1	WT1	11p13	194070				
Wolf-Hirschhorn	LETM1, WHSC1, WHSC2	4p16.3	194190				
X-linked juvenile retinoschisis	RS1	Xp22.13	312700				
X-linked lymphoproliferative type 2	XIAP	Xq25	300635				
X-linked mental retardation 30	PAK3	Xq23	300558				
X-linked mental retardation, ARX-related	ARX	Xp21.3	300419				
Xp11.22 X-linked mental retardation 17	Multiple	Xp11.22	300705				
XX Sex Reversal 2	SOX9	17q24.3	278850				
XY Sex Reversal 1	SRY	Yp11.31	400044				

NOTE: The Prenatal Targeted oligonucleotide array offered by the Genetics Center is a diagnostic screening tool designed to evaluate targeted childhood-onset disease regions of the genome for gains or losses of genomic material. This specific array will also detect whole chromosome loss of heterozygosity (LOH) for chromosomes 7, 11, 14, 15, 16, and 20. As of March 2013, the above listed genetic conditions have been associated with the indicated regions being evaluated by the Prenatal Targeted array. Copy number losses of more than 2 Mb thorough out the genome will also be reported. This listing is not all-inclusive and is subject to change as more is discovered about the associations between syndromes, genes, and the targeted regions. The Prenatal Targeted array may detect copy number variations of uncertain clinical significance, or changes associated with cancer predisposing disorders that may manifest later in life and be unrelated to prenatal ultrasound findings. Also, imbalances in highly repetitive regions (pericentromeric and telomeric regions) may not be detected. For certain conditions, additional testing may be warranted. Please contact the Genetics Center to discuss patient samples if there are any questions.