Table S2. List of Previously Reported Candidate Genes whose Candidacy Is Further Supported by Findings in this Cohort

				Original	Original		
ID	Gene	Variant(s)	Zygosity	Report	Phenotype	Observed Phenotype	Phenotypic Comparison
					Developmental		
					delay,		
					hypotonia,		
					ataxic		
17-	ACAT	GenBank: NM_001303253.1:c.3		PMID:	movements, and chorea	Microcephaly, GDD, heterotopia,	
5507	ACAT 2	5A>G:p.Asn12Ser	Het	6150136	movements	hypomyelination, hypertonia, spasticity, and lactic acidemia	Similar
3307	2	JA>O.p.Asii125ei	пеі	0130130	Severe	lactic acidellila	Sillilai
					intellectual		
					disability,		
					microcephaly,		
					seizures, and		
		GenBank:		PMID:	some autistic	Gross motor delay, basal ganglia	
17-	ACTL	NM_016188.4:c.999T		2653989	behavioral	abnormalities, hypertonia, exaggerated startle,	
1447	6B	>A:p.Cys333Ter	Hom	1	patterns	and gastroesophageal reflux	Phenotypic expansion
				D) (ID			N/A (This is a multilocus
17-	AKAP	GenBank:		PMID: 2860077		CDD : 91 11: 1 1	phenotype; see Table S1 for a full list of variants in this
0962	6 AKAP	NM_004274.4:c.1874 A>T:p.Tyr625Phe	Het	2800077	NSID	GDD, anemia, ?hereditary spherocytosis, and ?Gilbert disease	individual.)
0902	U	A>1.p.1 y10231 lie	1101	7	Gillessen-	:Onbeit disease	ilidividuai.)
		GenBank:		PMID:	Kaesbach-	Cardiomegaly, pericardial effusion, a thick	
17-		NM_001077690.1:c.4		2596663	Nishimura	nuchal fold, oligohydramnios, hydronephrosis,	
0422	ALG9	30T>C:p.Cys144Arg	Hom	8	syndrome	and a small bladder	Similar
		GenBank:		PMID:		Periventricular heterotopia, microcephaly,	
18-		NM_001024226.1:c.3		2886815	Periventricular	failure to thrive, short stature, and dysmorphic	
1233	ARF1	79A>G:p.Lys127Glu	De novo	5	heterotopia	features	Phenotypic expansion
1.7	4.0146	GenBank:		PMID:	T 1 .		
17-	ARMC 9	NM_025139.5:c.51+5	Hom	2862550	Joubert	Louhart arm drama	Similar
6686	9	G>T	Hom	4	syndrome Autosomal-	Joubert syndrome	Similar
		GenBank:		PMID:	recessive		
17-	ATAD	NM_018188.4:c.1410		2764030	Harel-Yoon		
9070	3A	+1G>A:	Hom	7	syndrome	Intrauterine fetal death with polyhydramnios	Phenotypic expansion
		GenBank:	-				2 F F
		NM_001007026.1:c.1		PMID:	De novo		
17-		503_1508del::p.Gln50		2743129	ATN1-related		
4230	ATN1	1_Gln502del	De novo	0	GDD	GDD	Similar

		GenBank:		PMID:			
17-	ATP11	NM_173694.4:c.1675		2694447	Hemolytic		
6079	C	+2T>C:	Hemi	2	anemia	Fetus with hydrops fetalis	Phenotypic expansion
0077		GenBank:	1101111	_	unonnu	1 ords with hydrops rotalis	Thenesypre enpairs on
		NM_001001344.2:c.3		PMID:	X-linked		
17-	ATP2	542delA:p.Gln1181Ar		2291239	spinocerebellar	Failure to thrive, microcephaly, and	
5302	B3	gfsTer6	Hemi	8	ataxia	developmental regression	Phenotypic expansion
REQ	Do	GenBank:	1101111	PMID:	utu/iiu	de velopinental legiession	Thenotypic expansion
18-		NM_001127610.1:c.2		1270438	Cholestasis	2-month-old infant with cholestatic liver	Similar (different inheritance
1700	BAAT	T>A:p.?	Hom	6	(?digenic)	failure	pattern)
17-	CDK1	GenBank: NM_052988.4:c.139d elG:p.Glu47ArgfsTer		PMID: 2888634	Severe growth retardation, spine malformations, and developmental	Atrial septal defect, gross motor delay, speech delay, hypertelorism of the eye, a broad nose, hypotonia, vesicoureteral reflux, and	
5831	0	21	Hom	1	delays	gastroesophageal reflux	Phenotypic expansion
3031	U	GenBank:	Hom	1	delays	gastroesophagear renux	Thenotypic expansion
16- 2564	CHD8	NM_001170629.1:c.4 984C>T:p.Arg1662Te	De novo	PMID: 2249530	ASD	ASD	Similar
2304	CHDo	GenBank:	De novo	7	ASD	ASD	Sillilai
17- 6317	CLDN 10	NM_001160100.1:c.5 90delC:p.Pro197Leuf sTer21	Hom	PMID: 2877125	HELIX syndrome	Dry skin, no saliva, no tears, and no sweating	Similar
REQ		GenBank:		PMID:			
18-	CLHC	NM_001135598.1:c.6		2555806			
1383	1	10C>T:p.Arg204Ter	Hom	5	Myopathy	Myopathy	Similar
17- 6687	CNTN 5	GenBank: NM_014361.3:c.1300 -1G>A	Het	PMID: 2860077	Fine motor delay, speech delay, intellectual disability, learning disability, developmental regression, hypertonia, seizures tonic, spasticity, cerebellar atrophy, GDD, and epilepsy	Overgrowth, macrocephaly, speech delay, learning disability, developmental regression, ataxia, seizures, periventricular leukomalacia, and obesity	Similar
0087	J	-1U>A	пеі	<u> </u>	and epilepsy	and obesity	Similar

_	1	T	ı	ı	1	T	T
					Neurodegenera		
					tion,		
					strabismus,		
					nystagmus,		
					dysarthria,		
					muscle		
					weakness,		
					hyporeflexia,		
					and		
		GenBank:		PMID:	abnormality of		
18-	CPNE	NM_006032.3:c.546C		2743129	the cerebral	Neonatal seizures, microcephaly, GDD, failure	
0676	6	>G:p.Asn182Lys	Hom	0	white matter	to thrive, spasticity, dystonic movements	Similar
0070	0	GenBank:	Hom	PMID:	winte matter	to unive, spusicity, dystome movements	Silina
17-	CSNK	NM_022048.3:c.1099		2446388	Severe early-	Ventricular septal defect, learning disability,	
5982	1G1	C>T: p.Arg367Ter	Het	3	onset epilepsy	epilepsy, a fatty liver, and gallbladder stone	Phenotypic expansion
3984	101	C>1. p.Aig50/1er	пеі	J	Familial	ephepsy, a rany river, and gambiadder stone	i nenotypic expansion
					susceptibility		
					to myeloproliferat		
		GD1			ive and/or		
		GenBank:		DMID.			
17	DDV4	NM_016222.3:c.47_4		PMID:	lymphoprolifer	D	
17-	DDX4	8delTG:p.Val16Alafs	77.	2592068	ative	Recurrent pancytopenia and histiocytic	G: '1
6285	1	Ter12	Het	3	neoplasms	lymphohistiocytosis	Similar
					Ventricular		
					septal defect,		
					speech delay,		
					learning		
		G D I			disability, cleft		
		GenBank:		D) (ID	lip and/or		
1.0	DIAN	NM_001126057.2:c.8		PMID:	palate, skin		
18-	DMK	52_853dupCA:p.Ser2	**	2860077	laxity, joint	Congenital heart disease and a sister who died	G: '1
0998	N	85ThrfsTer74	Hom	9	laxity	of severe congenital heart disease	Similar
					Retinitis		
		GenBank:		D) (III	pigmentosa		
1.5	DWG	NM_018163.3:c.681G		PMID:	with		
17-	DNAJ	>A: r. 601_681del;		2635566	immunodeficie		a
4852	C17	p.Tyr201_Ala227del	Hom	2	ncy	Retinitis pigmentosa with immunodeficiency	Similar
1		GenBank:		PMID:			
17-	222	NM_000798.4:c.33C		1145990			
2105	DRD5	>A:p.Tyr11Ter	Het	8	Dystonia	Dystonia	Similar
		GenBank:			Chorea, speech		
		NM_022049.2:c.299_		PMID:	delay, and		
17-	GPR8	300insC:p.Glu100Asp		2712348	learning	Dystonia, dysarthria, white matter lesions, and	
3125	8	fsTer288	Hom	6	disabilities	is non-responsive to L-dopa	Similar

		GenBank:	I		Spinocerebellar		I
				PMID:	ataxia and		
17		NM_001278064.1:c.1					
17-	CDMI	568dupT:p.Cys524Va		2290194	autosomal	CDD	G: 11
5371	GRM1	lfsTer3	Hom	7	recessive 13	GDD	Similar
		GenBank:		D) (ID	Intellectual	Microcephaly, intellectual disability, seizure,	
	****	NM_005321.2:c.441d		PMID:	disability and	brain atrophy, scoliosis, hip dislocation,	
17-	HIST1	upC:p.Lys148GlnfsTe		2847585	variable .	dysmorphic, and hyperpigmented lesions in	
6787	H1E	r48	Het	7	overgrowth	the trunk starting at 2 years of age	Similar
					Developmental		
					delay,		
					intellectual		
		GenBank:		PMID:	disability, and		
17-	HIVE	NM_002114.4:c.4111		2700358	dysmorphic	Failure to thrive, microcephaly, hypotonia,	
3747	P1	A>G:p.Ile1371Val	De novo	3	features	contractures, and club foot	Similar
		GenBank:		PMID:			
18-	KCND	NM_012281.3:c.1208		2450127	ASD and	Seizure disorder, intellectual disability, muscle	
2419	2	C>G:p.Pro403Arg	De novo	8	seizures	weakness, and hypotonia	Similar
		GenBank:		PMID:			
17-	KDM5	NM_001042603.2:c.1		2193799	Intellectual		
3418	A	429T>G:p.Phe477Val	Hom	2	disability	Developmental delay and dysmorphic features	Similar
		GenBank:		PMID:	Crohn disease		
17-	LACC	NM_001128303.2:c.8		2514720	with		
8170	1	50T>C:p.Cys284Arg	Hom	3	arthropathy	Arthropathy	Similar
		GenBank:		PMID:			
18-		NM_020117.10:c.332		2260794	Infantile liver		
1546	LARS	5+1delG:	Hom	0	failure	Epileptic encephalopathy and GDD	Phenotypic expansion
					Arthrogryposis		
					multiplex		
		GenBank:			congenita,		
		NM_139284.2:c.834d	^a Het	PMID:	neurogenic,		
17-		elG:p.Ser279AlafsTer	(both	2831849	with myelin		
2109	LGI4	191	parents)	9	defect	History of multiple neonatal deaths	Similar
		GenBank:		PMID:	Intellectual		
17-		NM_001199216.1:c.7		2760634	disability and		
5632	MAG	20C>A:p.Ser240Arg	Hom	6	ataxia	Intellectual disability and ataxia	Similar
		•			Mental		
		GenBank:		PMID:	retardation and		
17-	MBOA	NM_001146056.2:c.6		2761648	autosomal	Speech delay, ataxia, and bilateral globus	
3591	<i>T7</i>	04G>A:p.Gly202Ser	Hom	0	recessive 57	pallidus involvement	Phenotypic expansion
		GenBank:			Mental		
		NM_001146056.2:c.1		PMID:	retardation and		
17-	MBOA	000delT:p.Tyr334Met		2761648	autosomal		
6540	<i>T7</i>	fsTer41	Hom	0	recessive 57	GDD, autistic features, and failure to thrive	Similar

		GenBank:		PMID:			
17-	MCID	NM_001190787.1:c.7		2504896	Primary ciliary		
1170	AS	17+2T>G:	Hom	3	dyskinesia	Primary ciliary dyskinesia	Similar
1170	710	GenBank:	110111	PMID:	Nonsyndromic	Timary chary dyskinesia	Silina
17-	NCKA	NM_013436.4:c.742-		2894009	intellectual		
2106	P1	3A>C	De novo	7	disability	Intellectual disability and epilepsy	Phenotypic expansion
2100		5.5 0	20.1070	,	Fatal infantile	interiorital disacrity and opinopsy	Thenotypic enpansion
		GenBank:			lactic acidosis		
		NM_004548.2:c.373_		PMID:	and		
17-	NDUF	375delCAG:p.Gln125		2804073	cardiomyopath		
2505	B10	del	Hom	0	V	Non-immune hydrops fetalis, died after birth	Distinct phenotype
					,	Failure to thrive, fine motor delay, gross motor	
		GenBank:		PMID:		delay, speech delay, intellectual disability,	
17-	PGAP	NM_001291726.1:c.6		2962072	Toriello-Carey	learning disability, hypotonia, and a brother	
1765	3	97C>T:p.His233Tyr	Hom	4	syndrome	with cerebellar hypoplasia	Similar
					-	Failure to thrive, fine motor delay, gross motor	
		GenBank:		PMID:		delay, intellectual disability, speech delay,	
17-	PGAP	NM_001291726.1:c.6		2962072	Toriello-Carey	cleft lip, blindness, hearing loss, tonic	
2720	3	97C>T:p.His233Tyr	Hom	4	syndrome	seizures, muscle weakness, constipation	Similar
						Macrocephaly, multiple cafe au lait spots,	
		GenBank:		PMID:		GDD, speech and cognitive delay, autistic	
17-	PIK3C	NM_006218.3:c.3103		2324628	Cowden	features, periventricular white matter changes	
0806	\boldsymbol{A}	G>A:p.Ala1035Thr	Het	8	syndrome	by MRI, and epilepsy	Similar
					Autosomal-		
					recessive,		
					severe form of		
					anterior		
		GenBank:			segment		
		NM_005029.3:c.640_		PMID:	dysgenesis and		
17-		656del:p.Ala214Argfs		2183652	microphthalmi	Gross motor delay, speech delay, cataract,	
5492	PITX3	Ter42	Hom	2	a	blindness, and spasticity	Similar
					Intellectual		
					disability,		
		GenBank:		PMID:	dysmorphic		
18-	PPP1	NM_001135629.2:c.4		2894009	facies, and	Intellectual disability, dysmorphic facies, and	
0885	R21	27C>T:p.Arg143Ter	Hom	7	hypotonia	hypotonia	Similar
1		GenBank:		PMID:			
17-	PPP2	NM_002719.3:c.398A	_	2597237		Macrocephaly, demyelination, developmental	a: ::
1250	R5C	>G:p.His133Arg	De novo	8	Overgrowth	delay, and hypotonia that improved with time	Similar
1.0		GenBank:		PMID:			
18-	PPP2	NM_178586.2:c.1343		2597237			
2160	R5C	A>G:p.Lys448Arg	Het	8	Overgrowth	Overgrowth	Similar

		1	T		1	T	I
						8-week-old boy with severe complex	
						congenital heart disease (pulmonary atresia,	
						ventricular septal defect, major	
						aortopulmonary collateral artery, patent ductus	
REO		GenBank:		PMID:		arteriosus, and hypertrophic cardiomyopathy;	
18-	PRKD	NM_002742.2:c.2554		2571311	Truncus	on prostaglandin with biventricular	
1959	1 KKD	_	Hom	0	arteriosus	hypertrophic cardiomyopathy)	Dhanatymia aymanaian
1939	1	G>T:p.Glu852Ter	пош	U		hypertropine cardiomyopathy)	Phenotypic expansion
					A recognizable		
					syndrome with		
					intellectual		
		GenBank:		PMID:	disability, heart		
16-	PUF6	NM_078480.2:c.24+1		2780495	defects, and	Dysmorphic baby with multiple congenital	
2870	0	G>C	Het	8	short stature	anomalies (cleft lip and dysplastic vertebrae)	Similar
						Failure to thrive, growth retardation,	
		GenBank:		PMID:		microcephaly, fine motor delay, gross motor	
17-	QRFP	NM_198179.2:c.373C		2743129	Intellectual	delay, speech delay, intellectual disability,	Similar (different inheritance
1020	R	>T:p.Gln125Ter	Damouo	0	disability	nystagmus, spasticity, and PMLD?	pattern)
1020	Λ	>1.p.Giii1231ei	De novo	U	-	hystaghius, spasticity, and FWLD?	pattern)
					Childhood-		
					onset		
					mitochondrial		
		GenBank:		PMID:	respiratory	Repeated episodes of encephalopathy with	
17-	QRSL	NM_018292.4:c.1042		2674149	chain complex	muscle weakness and abnormal basal ganglia	
2764	1	+3A>G:	Hom	2	deficiency	on brain MRI	Similar
		GenBank:		PMID:			
17-		NM_002872.4:c.484C		2551208		Failure to thrive, hypogammaglobulinemia,	
1228	RAC2	>T:p.Gln162Ter	Hom	1	SCID	and bronchial asthma	Similar
		GenBank:	-	PMID:	X-linked		
17-	RPS6	NM_014496.4:c.688G		1064443	intellectual		
5425	KA6	>C:p.Ala230Pro	Hemi	0	disability	GDD and hearing loss	Similar
3423	MAU	GenBank:	1101111	PMID:	Severe	ODD and hearing loss	Simiai
17	GETED						
17-	SETD	NM_014712.2:c.4812		2697495	neurodevelopm	T . 11 . 1 11 . 1 11 . 1	g: '1
1913	1A	+4A>G:	Het	0	ental disorder	Intellectual disability and epilepsy	Similar
		GenBank:					
		NM_001166496.1:c.1		PMID:	Recurrent,	Developmental delay, frequent episodes of	
17-	SLC16	079delC:p.Ala360Gly		2539074	severe	acidosis and ketotic hypoglycemia, and	
3382	A1	fsTer19	Hom	0	ketoacidosis	epilepsy	Similar
		GenBank:					
		NM_001166496.1:c.4		PMID:	Recurrent,		
17-	SLC16	2_43insC:p.Asp15Arg		2539074	severe		
4510	AI	fsTer34	Hom	0	ketoacidosis	Developmental delay, epilepsy, and ketosis	Phenotypic expansion
1310	111	1010154	110111		Complex	Bevelopmental delay, ophiopsy, and ketosis	Thenotypic expansion
		GenBank:		PMID:	partial epilepsy	Migrogaphaly fine motor dalay gross motor	
17	CI CA					Microcephaly, fine motor delay, gross motor	Db (4:ff
17-	SLC4	NM_022058.3:c.2773	**	1841348	and intellectual	delay, speech delay, intellectual disability, and	Phenotypic expansion (different
4393	A10	-2A>C	Hom	2	disability	hypotonia	inheritance pattern)

		GenBank:		PMID:	Lichtenstein-		
17-	SLC9	NM_003047.4:c.1273		2520511	Knorr	GDD, severe cerebellar dysfunction, hearing	
6065	A1	C>T:p.Arg425Cys	Hom	2	syndrome	loss, and cerebellar atrophy	Similar
DEC		G D 1		D) (ID	**	Microcephaly, gross motor delay, intellectual	
REQ		GenBank:		PMID:	Heart and brain	disability, atrial septal defect, abnormal	
18-	GMCO	NM_019108.3:c.701+	**	2701847	malformation	imaging (periventricular leukomalacia), and	g: 11
1086	SMG9	4A>G:	Hom	4	syndrome	asthma	Similar
					Severe hydranencepha		
REQ		GenBank:		PMID:	ly nearly		
18-	SPDL	NM_017785.4:c.668A		2555806	occupying the		
2221	31 DL	>G:p.Tyr223Cys	Hom	5	entire skull	Hydranencephaly and epilepsy	Similar
LLLI	1	> G.p. 1 y1223 C y s	110111	3	Congenital	Trydraneneephary and ephepsy	Similar
					myopathy,		
		GenBank:		PMID:	neuropathy,		
17-	SPTB	NM_020971.2:c.1665		2854041	and central	Speech delay, intellectual disability, ataxia,	
6278	N4	+2T>C:	Hom	3	deafness	seizures, and cerebral atrophy	Phenotypic expansion
		GenBank:		PMID:	Non-epileptic		
18-	STXB	NM_001032221.3:c.1		2136470	intellectual		
0605	P1	060T>C:p.Cys354Arg	Het	0	disability	Intellectual disability and ASD	Similar
						Previous termination of 15/32 with	
						holoprosencephaly, cyclops, cleft lip,	
1	mn ar	GenBank:		PMID:		ventricular septal defect, agenesis of the	
17-	TBC1	NM_152730.6:c.1372	**	2428556	OED IV	corpus callosum, club foot, and sandal gap (no	Di .
0889	D32	+1G>T:	Hom	6	OFD IX	sample)	Phenotypic expansion
	<i>TME M132</i>	GenBank: NM_001304438.1:c.1					N/A (This is a multilocus
	E;	483-2A > G;		PMID:			phenotype; see Table S1 for a
17-	IL12R	NM_001258216.1:c.3		2533163		Nystagmus, high arch palate, contracture, and	full list of variants in this
4280	B2	64+1G > A	Hom	8	Hearing loss	family history of similar condition	individual.)
1200	22	GenBank:	220111		110411115 1000	raming motory of billing condition	
		NM_004618.4:c.2272		PMID:			
17-	TOP3	_2273insC:p.Arg758T		3019313	Bloom-like		
3406	A	hrfsTer3	Hom	7	phenotype	Short stature and microcephaly	Similar
		GenBank:		PMID:	Microcephaly,		
17-	TRAP	NM_001079537.2:c.2		2862602	epilepsy, and	Failure to thrive, microcephaly, GDD, and	
3847	PC6B	68-2_268-1delAG:	Hom	9	autistic features	ASD	Similar
		GenBank:					
1		NM_177452.3:c.302d		PMID:			
17-	TRAP	elG:p.Gly101AlafsTer		2839783	Intellectual		
0626	PC6B	7	Hom	8	disability	GDD and microcephaly	Phenotypic expansion
REQ	TCDVI	G D 1		PMID:	Sudden infant	IUGR, swallowing difficulty, failure to thrive,	
18-	TSPYL	GenBank:	11	1527328	death with	ambiguous genitalia, history of SIDS in a	G::1
2590	1	NM_003309.3:c.725_	Hom	3	dysgenesis of	sister at age 6 months	Similar

		726delTG:p.Val242Gl ufsTer52			the testes syndrome		
		WID 1910 2			Dyskinetic movement		
		GenBank:			disorder,	Failure to thrive, microcephaly, fine motor	
		NM_001080421.2:c.4		PMID:	developmental	delay, gross motor delay, intellectual	
17-	UNC1	379C>T:p.Ala1460Va		2819236	delay, and	disability, learning disability, developmental	
3311	3A	1	De novo	9	ASD	regression, cerebral atrophy, and spasticity	Similar
						GDD, spastic quadriplegia, microcephaly,	
					Cerebellar	epilepsy, dysmorphic features, and swallowing	
		GenBank:		PMID:	ataxia with	difficulty; status post-GT-insertion: significant	
17-	VWA3	NM_144992.4:c.1191		2615703	intellectual	cerebral volume loss but no clear cerebellar	
7243	В	T>G:p.Tyr397Ter	Hom	5	disability	involvement	Phenotypic expansion
						Microcephalic failure to thrive, speech delay,	
		G D 1		D) (ID	Microcephaly	bitemporal narrowing, a long philtrum,	
1.7	IIID EU	GenBank:		PMID:	18, primary,	epicanthal folds, hypertelorism,	
17- 7303	WDFY 3	NM_014991.4:c.1014 7+2T>C:	Hot	2700854	autosomal- dominant	gastrointestinal reflux, and eventration of	Similar
/303	3	7+21>C:	Het	4	Intellectual	diaphragm	Similar
					disability,		
					motor delay,		
		GenBank:		PMID:	microcephaly,		
17-	ZBTB	NM_014415.3:c.2735		2989385	and cerebellar	GDD, microcephaly, cataract mild cerebellar	
3151	11	G>A:p.Arg912Gln	Hom	6	atrophy	atrophy, and elevated methylmalonic acid	Phenotypic expansion

Question marks denote uncertainty of the clinical feature of diagnosis as provided by the ordering physician. Abbreviations are as follows: Hom = homozygous; Het = heterozygous; Hemi = hemizygous; IUGR = intrauterine growth restriction; MRI = magnetic resonance imaging; GDD = global developmental delay; ASD = autism spectrum disorder; NSID = nonspecified intellectual disability; HELIX = hypohidrosis, electrolyte imbalance, lacrimal gland dysfunction, ichthyosis, and xerostomia; PMLD = Pelizaeus-Merzbacher-like disease; OFD = oral-facial-digital syndrome; SCID = severe combined immunodeficiency, SIDS = sudden infant death syndrome; and GT = gastronomy tube.

^aHet (both parents) infers recessive inheritance of the phenotype associated with neonatal deaths in this instance

Table S8.ª List of Couples Who Presented for Counseling with No Available DNA From the Affected Deceased Child(ren) or Fetus(es)

ID	Phenotype of the Deceased Child or Fetus	Result	Gene	Variant(s)	Zygosity	LOF
16-				GenBank:		
2873-	Previous termination because of holoprosencephaly, cyclops, cleft lip, ventricular	Ambiguous		NM_005759.5:c.991+6T	Het (both	
A	septal defect, agenesis of the corpus callosum, club foot, and a sandal gap	(VUS)	ABI2	>A	parents)	-
17-		Pathogenic or		GenBank:		
3580-		likely		NM_001177520.2:c.119	Het (both	
Α	History of a fetus with abnormal skeletal features	pathogenic	ALPL	5G>A:p.Glu399Lys	parents)	_
17-	•	Pathogenic or		GenBank:		
6127-		likely		NM_001024944.1:c.100	Het (both	
A	A deceased child with hepatomegaly and suspected metabolic disease	pathogenic	ASL	0C>T:p.Gln334Ter	parents)	y
		Pathogenic or		GenBank:		
17-		likely		NM_001024946.1:c.525-	Het (both	
4404	Neonatal death with hyperammonemia	pathogenic	ASL	83_528del	parents)	_
17-		Pathogenic or		GenBank:		
7724-		likely		NM_000049.2:c.691_69	Het (both	
В	Two deceased children with Canavan disease	pathogenic	ASPA	2insC:p.Tyr231SerfsTer5	parents)	y
				GenBank:		
				NM_001206846.1:c.113		
				8C>T:p.Gln380Ter in the		
				mother; GenBank:		
		Pathogenic or		NM_018136.5:c.4363G>		
17-		likely		T:p.Glu1455Ter in the	Het (both	
0706	Severe microcephaly	pathogenic	ASPM	father	parents)	y
		Pathogenic or		GenBank:		
17-		likely	B3GAL	NM_152490.4:c.762+1G	Het (both	
2597	A deceased child with suspected Walker-Warburg syndrome	pathogenic	NT2	>C	parents)	y
		Pathogenic or		GenBank:		
17-		likely	B3GAL	NM_152490.4:c.762+1G	Het (both	
2597	A deceased child with Walker-Warburg Syndrome.	pathogenic	NT2	>C	parents)	y
		Pathogenic or		GenBank:		
17-		likely		NM_004305.3:c.3G>T:p.	Het (both	
5685	IUFD	pathogenic	BIN1	?	parents)	у
		Pathogenic or		GenBank:		
17-		likely		NM_138793.3:c.902_90	Het (both	
1488	Two neonatal deaths with skeletal dysplasia	pathogenic	CANT1	6dup:p.Ser303Alafs*21	parents)	y
				GenBank:		
		Pathogenic or		NM_001080522.2:c.308		
18-		likely	CC2D2	4delG:p.Lys1029ArgfsTe	Het (both	
2336	Two deceased children with polycystic kidney disease and hydrocephalus	pathogenic	A	r3	parents)	_
REQ1		Ambiguous		GenBank:		
8-	History of a pregnancy with suspected Meckel-Gruber syndrome (encephalocele,	(candidate	CEP29	NM_033395.1:c.2603del	Het (both	
0971	dysplastic kidney, and no polydactyly)	gene)	5	A:p.Gln868ArgfsTer25	parents)	-

				GenBank:		
18-	A deceased child with developmental delay, speech delay, encephalopathy, delayed	Ambiguous		NM_001195139.1:c.172	Het (both	
0831	milestone, and provisional hypoxic ischemic encephalopathy	(VUS)	COG4	1C>T:p.Ser574Phe	parents)	_
REQ1	7 1 71	Pathogenic or		GenBank:	•	
8-		likely		NM_001122634.3:c.406	Het (both	
2856	Three deceased children with severe hyperammonemia	pathogenic	CPS1	C>T:p.Arg136Cys	parents)	_
		1		GenBank:	•	
17-		Ambiguous		NM_000098.2:c.451C>G	Het (both	
5536	Multiple neonatal deaths	(VUS)	CPT2	:p.Arg151Gly	parents)	_
	•			GenBank:		
18-		Ambiguous	DYNC2	NM_001080463.1:c.333	Het (both	
0769	Recurrent non-immune hydrops fetalis	(VUS)	H1	1G>A:p.Glu1111Lys	parents)	_
	• •	Pathogenic or		GenBank:		
17-		likely		NM_004092.3:c.88+5G>	Het (both	
7307	A deceased child with severe acidosis and hydrocephalus	pathogenic	ECHS1	A	parents)	_
		Ambiguous		GenBank:		
17-	A deceased child with hydrocephalous, congenital heart disease, and multiple	(candidate	EHBP1	NM_001099409.2:c.400	Het (both	
2653	congenital abnormalities	gene)	L1	4-1G>A	parents)	y
				GenBank:		
17-		Ambiguous		NM_004434.2:c.2233G>	Het (both	
3181	Four deceased children with hydrocephalus	(VUS)	EML1	A:p.Val745Ile	parents)	_
		Pathogenic or		GenBank:		
17-		likely	FAM13	NM_019000.4:c.503C>G	Het (both	
4767	A deceased child with ataxia and spasticity	pathogenic	4B	:p.Ser168Ter	parents)	_
		Ambiguous		GenBank:		
17-	History of fetal loss with arrhythmia, club foot, short limbs, hydrocephalus,	(candidate		NM_001077261.4:c.227	Het (both	
0249	encephalocele, and polyhydramnios	gene)	NCOR2	5G>A:(p.Gly759Arg)	parents)	_
		Pathogenic or		GenBank:		
17-		likely		NM_012160.4:c.1652T>	Het	
4496	Five deceased children with unexplained lactic acidosis	pathogenic	FBXL4	A:p.Ile551Asn	(carrier)	-
				GenBank:		
				NM_001039885.2:c.954		
		Pathogenic or		_955insTGCGCGCGC:p		
17-		likely		.Cys318_Leu319insCysA	Het (both	
2289	A deceased child with suspected Walker-Warburg syndrome	pathogenic	FKRP	laArg	parents)	у
REQ1		Pathogenic or		GenBank:		
8-		likely		NM_000152.4:c.1802C>	Het (both	
2239	Two deceased children with Pompe disease	pathogenic	GAA	T:p.Ser601Leu	parents)	_
		Pathogenic or		GenBank:		
17-		likely	anner-	NM_001244710.1:c.686-	Het (both	
6039	Three deceased children with suspected muscle disease	pathogenic	GFPT1	1G>A	parents)	у
		Pathogenic or				
17-		likely	HADH	NM_000183.3:c.631-	Het (both	
1160	Parents of four children who died as neonates with a suspected metabolic disease	pathogenic	В	1G>A	parents)	=

		Pathogenic or		GenBank:		
18-		likely		NM_000518.5:c.135delC	Het (both	
0957	Two deceased children with beta thalassemia major	pathogenic	HBB	:p.Phe46LeufsTer16	parents)	_
0,0,	The developed emidient with ooth that asserting in a	paarogeme	1122	GenBank:	purents)	
				NM_000518.5:c.315+1G		
		Pathogenic or	HBB	> A;		
17-	History of one deceased child with sickle cell anemia and another with severe	likely	and	NM_017668.3:c.346G >	Het (both	
4868	microcephaly	pathogenic	NDE1	T:p.Glu116Ter	parents)	v
		1 0		GenBank:		
17-		Ambiguous		NM_002161.5:c.1114-	Het (both	
1801	Ptosis and subclinical epilepsy (EEG)	(VUS)	IARS	3T>C	parents)	_
		Pathogenic or		GenBank:		
18-		likely		NM_000208.2:c.1546T>	Het (both	
1907	A deceased child with suspected Donohue syndrome	pathogenic	INSR	C:p.Trp516Arg	parents)	_
	·	Pathogenic or		GenBank:	<u> </u>	
17-		likely		NM_001101417.3:c.103	Het (both	
4689	Two deceased babies with hydrocephalus and renal anomalies	pathogenic	ISPD	6G>T:p.Glu346Ter	parents)	_
	•	Ĭ		GenBank:	ĺ	
		Pathogenic or		NM_001244193.1:c.346		
18-		likely	KIAA05	8_3469insT:p.Ile1157Tyr	Het (both	
2004	Two pregnancy losses	pathogenic	86	fsTer2	parents)	_
				GenBank:		
		Pathogenic or		NM_198129.2:c.5304-		
		likely		19C>G;		
17-		pathogenic and		NM_198129.2:c.8436+1	Het	
6683	A deceased child with dystrophic epidermolysis bullosa	VUS	LAMA3	G>A	(CMP)	y
		Pathogenic or		GenBank:		
17-		likely		NM_005562.2:c.3385C>	Het (both	
3772	One deceased baby with epidermolysis bullosa	pathogenic	LAMC2	T:p.Arg1129Ter	parents)	у
		Pathogenic or		GenBank:		
17-		likely		NM_139284.2:c.834delG	Het (both	
2109	Previous pregnancies with multiple anomalies	pathogenic	LGI4	:p.Ser279AlafsTer191	parents)	у
				GenBank:		
17-		Ambiguous	MAPK	NM_001265611.1:c.330	Het (both	
2133	Two deceased children with polycystic kidney disease	(VUS)	BP1	4C>T:p.Arg1102Cys	parents)	_
		Ambiguous		GenBank:		
17-		(candidate		NM_033540.2:c.2012+8	Het (both	
0367	Two neonatal deaths with undiagnosed sever neuromuscular disease	gene)	MFN1	G>A:	parents)	_
	Two deceased newborns with facial anomalies (bilateral, large soft tissue lesions,	Ambiguous		GenBank:		
17-	absent nasal soft tissue, and internal tubular sonolucent structures over both eyes),	(candidate	MICAL	NM_001136004.3:c.239	Het (both	
3770	and clenched hands	gene)	3	8-3T>A	parents)	_
		Pathogenic or		GenBank:		
17-		likely	MMAC	NM_015506.2:c.848G>C	Het (both	
2096	Three affected pregnancies with hydrocephalus	pathogenic	HC	:p.Ter283Serext*14	parents)	у

	Two deceased children with cardiomyopathy, failure to thrive, growth retardation,					
17-	fine and gross motor delay, speech delay, intellectual disability, learning disability,	Pathogenic or		GenBank:		
5990-	external ear malformation, recurrent fever, anemia, immunodeficiency, hypotonia,	likely	MTHF	NM_005956.4:c.517C>T	Het	
A	muscle weakness, renal agenesis, and renal tubulopathy	pathogenic	DI	:p.Arg173Cys	(carrier)	_
	masere weathless, renaragements, and renar vacuopaary	Pathogenic or	21	GenBank:	(currer)	
18-		likely	NDUFS	NM_004553.4:c.187-	Het (both	
0664	Three neonatal deaths with unexplained severe lactic acidosis	pathogenic	6	1G>A	parents)	v
0001	Three neonatal deaths with the Apathica servere factor actions	patriogenie		GenBank:	parents)	,
17-		Ambiguous		NM_004543.4:c.18696+	Het (both	
3566	Two neonatal deaths with myopathy, arthrogryposis, and dysmorphism	(VUS)	NEB	20A>G	parents)	_
3300	1 wo neonatal deaths with informity, arthrogryposis, and dysmorphism	(100)	IVLD	GenBank:	parents)	
				NM_004543.3:c.17394T		
17-		Ambiguous		>A:p.Asn5798LysAsn57	Het	
5341	Two deceased children with severe hypotonia	(VUS)	NEB	98Lys	(carrier)	_
JJ+1	1 wo deceased emidien with severe hypotolila	Pathogenic or	NED	GenBank:	(Carrier)	-
17-		likely		NM_004543.3:c.15873de	Het (both	
3501	Recurrent non-immune hydrops fetalis	pathogenic	NEB	lA:p.Val5292Serfs*8	parents)	v
3301	Recurrent non-minume nydrops fetans	Pathogenic or	IVED	GenBank:	parents)	У
18-		likely		NM_001199399.1:c.352	Het (both	
1212	Two deceased children with multiple congenital anomalies	pathogenic	NEK1	7T>A:p.Leu1176Ter	`	
1212	1 wo deceased children with multiple congenital anomalies	patriogenic	NEKI	GenBank:	parents)	У
17		A1-:				
17-	T ILIED	Ambiguous	NEE	NM_033116.5:c.2525C>	11-4	
2362	Two IUFDs with congenital anomalies	(VUS)	NEK9	T:p.Thr842Ile	Het	_
		D.d.		GenBank:		
17		Pathogenic or		NM_001293557.1:c.182	11 4 (1 - 4	
17-		likely	NODA	8_1846del:p.Asn610Serf	Het (both	
4002	Two IUFDs	pathogenic	NOD2	sTer120	parents)	У
1.7		Pathogenic or		GenBank:	II . A . A	
17-		likely	NDIID2	NM_153240.5:c.2694-	Het (both	
6935	Two neonatal deaths with renal agenesis and polydactyly	pathogenic	NPHP3	2_2694-1delAG	parents)	у
1.5		Pathogenic or		GenBank:	TT . A . I	
17-		likely		NM_153240.5:c.2694-	Het (both	
6614	Two deceased neonates with polycystic kidney disease and one with anencephaly	pathogenic	NPHP3	2_2694-1delAG	parents)	_
				GenBank:		
18-		Ambiguous		NM_001127692.2:c.323	Het (both	
2124	Static encephalopathy	(VUS)	PCCA	C>A:p.Thr108Asn	parents)	_
		Pathogenic or		GenBank:		
17-		likely		NM_000319.4:c.1554T>	Het (both	
0456	Three children with Zellweger syndrome	pathogenic	PEX5	G:p.Asn518Lys	parents)	_
				GenBank:		
17-	Three neonatal deaths with hydrocephalus, microcephaly, and cerebellum	Ambiguous		NM_012327.5:c.305T>C	Het (both	
4845	hypoplasia	(VUS)	PIGN	:p.Ile102Thr	parents)	-

				GenBank:		
		Pathogenic or		NM_138694.3:c.4325_4		
17-		likely		326insTT:p.Cys1443Serf	Het (both	
6313	Three deceased children with suspected polycystic kidney disease	pathogenic	PKHD1	sTer18	,	v
0313	Timee deceased children with suspected polycystic kidney disease	Pathogenic or	ГКПОТ	GenBank:	parents)	У
17-		likely	POMG	NM_032806.5:c.473G>	Het (both	
3043	There described the state of th	pathogenic	NT2		,	
3043	Three deceased children with congenital hydrocephalus		IVI Z	A:p.Arg158His	parents)	_
17-		Pathogenic or likely		GenBank: NM_001077366.1:c.118	11-4 (h -4h	
			DOMTI		Het (both	
7704 17-	Three deceased babies born with hydrocephalus	pathogenic	POMT1	+1G>T GenBank:	parents)	_
5785-		Pathogenic or	D 4 D 2 7	GenBank: NM_004580.4:c.598C>T	11 . 4 . 4	
		likely	RAB27		Het (both	
Α	A deceased child with suspected Griscelli syndrome	pathogenic	A	:p.Arg200Ter	parents)	У
1.7		Pathogenic or		GenBank:	TT . 0 . 1	
17-		likely	D.E.E.I	NM_052859.4:c.775+1G	Het (both	
3004	Microcephaly, GDD, epilepsy, spasticity, and brain atrophy	pathogenic	RFT1	>C	parents)	y
				GenBank:		
17-		Ambiguous		NM_004589.3:c.722A>	Het (both	
7694	Two deceased infants with brain atrophy	(VUS)	SCO1	G:p.Tyr241Cys	parents)	_
				GenBank:		
18-		Ambiguous	SLC15	NM_001170798.1:c.159	Het (both	
0170	Three deceased children with epilepsy and failure to thrive	(VUS)	A5	3-6delCTTT	parents)	_
		Pathogenic or		GenBank:		
17-	Two deceased children with two different diseases: non-immune hydrops fetalis and	likely	SLC34	NM_003052.4:c.532+2T	Het (both	
5514	nephrocalcinosis	pathogenic	A1	>C	parents)	у
		Pathogenic or		GenBank:		
17-		likely	<i>SMARC</i>	NM_001127207.1:c.445	Het (both	
1563	Four neonatal deaths with multiple anomalies	pathogenic	AL1	C>T:p.Gln149Ter	parents)	y
		Pathogenic or		GenBank:		
17-		likely		NM_018676.3:c.617G>	Het (both	
6346	A deceased neonate with history of non-immune hydrops fetalis	pathogenic	THSD1	A:p.Cys206Tyr	parents)	_
		Pathogenic or		GenBank:		
17-		likely		NM_001288951.1:c.412	Het (both	
1407	Three neonatal deaths with history of polyhydramnios and imperforate anus	pathogenic	TTC7A	C>T:p.Arg138Ter	parents)	у
		-		GenBank:		
		Pathogenic or		NM_001163809.1:c.850		
17-		likely		_851delCT:p.Leu284Val	Het (both	
8378	A deceased child with hydrocephalus	pathogenic	WDR81	fsTer9	parents)	_
	ř ř	Ambiguous		GenBank:	<u> </u>	
17-	History of a deceased child (1 year old) with club feet, clinodactyly, a small mouth,	(candidate		NM_001199144.1:c.241	Het (both	
0374	and small feet	gene)	XIRP2	9T>A:p.Phe807Ile	parents)	_
		0			,,	
18-					Het (both	
2659	IUFD with severe IUGR	gene)	ZMIZ2	T:p.Gln424Ter	parents)	v
18-		Ambiguous (candidate		GenBank: NM_174929.2:c.1270C>	Het (both	V

Abbreviations are as follows: LOF = loss of function, y = yes, Hom = homozygous, Het = heterozygous, Hemi = hemizygous, GDD = global developmental delay, IUGR = intrauterine growth restriction, IUFD = intrauterine fetal death, VUS = variant of unknown significance, EEG = electroencephalogram, and CMP = compound.

^aNegative instances can be found in Table S9.

Table S10.ª List of Individuals for Whom Flash WES Protocol was Implemented

ID	Result	Gene	Variant(s)	Zygosity	Phenotype
			GenBank:		
REQ18-	Pathogenic or	ABCA	NM_001089.3:c.3145		
1702	likely pathogenic	3	T>C:p.Ser1049Pro	Hom	NICU individual with severe unexplained lung disease
18-2213	Negative				Individual with electrolyte imbalance and suspected renal dysfunction
			GenBank:		
REQ18-		AKR1	NM_001190906.1:c.81		
3252	Ambiguous (VUS)	D1	7T>C:p.Trp273Arg	Hom	Individual with liver failure; a candidate for liver transplant
DE010	B 4		GenBank:		
REQ18-	Pathogenic or	D 4 4 T	NM_001127610.1:c.2	**	
1700	likely pathogenic	BAAT	T>A:p.?	Hom	Individual with cholestatic liver disease; a candidate for liver transplant
10 1207	D. d.		GenBank:		
18-1297-	Pathogenic or	CETD	NM_000492.3:c.579+1	11	NICH is distinct with an analysis of control lines of the same lin
В	likely pathogenic	CFTR	G>A GenBank:	Hom	NICU individual with unexplained severe lung and liver disease
DEO19	D-4h	CLCN			Individual with an arrant arrant in a great part of an arrant for the state of the
REQ18-	Pathogenic or	CLCN KB	NM_001165945.2:c.90	11	Individual with recurrent vomiting, growth retardation and/or short stature, renal
2160	likely pathogenic	KB	1G>C:p.Gly301Arg GenBank:	Hom	tubulopathy, and a history of polyhydramnios
			NM_000397.3:c.1462-		
		CYBB	1G>C:		
REQ18-	Pathogenic or	and	NM_000402.4:c.233T	Hom and	An individual with G6PD, suspected to have CGD (chronic granulomatous disease); a
1192	likely pathogenic	G6PD	> C:p.Ile78Thr;	Hemi	candidate for bone marrow transplant
1172	inkery patriogenic	GOLD	GenBank:	TICHH	Candidate for bone marrow transplant
18-0842-	Pathogenic or		NM_000531.5:c.231G		
B	likely pathogenic	OTC	>T:p.Leu77Phe	Het	Pregnant mother with history of a deceased neonate with hyperammonemia
	intery paintageme	010	GenBank:	1100	1 regimale mounts with mistory of a deceased meanage with hyperanimone man
			NM_001318859.1:c.47		
REQ18-	Pathogenic or	DGUO	8_479insTGAT:p.Asp		
1996	likely pathogenic	K	160ValfsTer3	Hom	Individual with liver failure and developmental delay; a candidate for liver transplant
			GenBank:		1
			NM_001318859.1:c.47		
REQ18-	Pathogenic or	DGUO	8_479insTGAT:p.Asp		Individual with failure to thrive, microcephaly, developmental delay, and abnormal
3659	likely pathogenic	K	160ValfsTer3	Hom	hair
			GenBank:		
REQ18-	Pathogenic or	ECHS	NM_004092.3:c.88+5	Het	
4214	likely pathogenic	1	G>A	(carrier)	Pregnant mother with history of a deceased neonate
			GenBank:		
	Ambiguous		NM_181466.2:c.132C		Individual with cholestatic jaundice, developmental delay, and dysmorphic features; a
18-2455	(candidate gene)	EIF6	>G:p.Tyr44Ter	Hom	potential candidate for liver transplant
			GenBank:		
REQ18-	Pathogenic or		NM_012160.4:c.1698		One of two children with unexplained lactic acidosis; there is an urgent need for
3341	likely pathogenic	FBXL4	A>G:p.Ile566Met	Hom	molecular diagnosis because of pregnant mother

REQ18-					NICU individual with autoimmune hemolytic anemia, thrombocytopenia,
3794	Negative				hepatosplenomegaly, and severe hypotonia
			Chr9:g.(6610356-		
REQ18-	Pathogenic or		1_6620184+1)_(66456		
2014	likely pathogenic	GLDC	92-1_6720863+1)_del	Hom	NICU individual with nonketotic hyperglycinemia
	71 0		GenBank:		71 57
			NM_001200049.3:c.75		
			02-4G>A:		
	Pathogenic or	HBB	NM_000518.5:c.20A>		
	likely pathogenic	and	T:p.Glu7Val;		
REQ18-	and ambiguous	CFAP	NM 001200049.3:c.75	Hom and	
0631	(candidate gene)	46	02-4G>A	Hom	Individual with normal GGT cholestasis
	(**************************************		GenBank:		
			NM_001289064.1:c.16		
18-0816-b	Ambiguous (VUS)	LIG1	66G>A:p.Ala556Thr	Hom	SCID features; a potential candidate for bone marrow transplant
			GenBank:		
REQ18-	Ambiguous		NM_000627.3:c.364C		
4236	(candidate gene)	LTBP1	>T:p.Gln122Ter	Hom	Individual with dysmorphic features
.200	(canaraare gene)	DIDI I	GenBank:	110111	Individual Will dybliotphie tentaleb
REQ18-	Pathogenic or		NM_004990.4:c.1066		NICU individual with anemia and/or neutropenia and/or pancytopenia, IUGR,
2652	likely pathogenic	MARS	C>T:p.Arg356Cys	Hom	elevated transaminases, cardiomyopathy, failure to thrive, and polyhydramnios
	intery puriogenie	1,11110	GenBank:	110111	ole value transmission, earlierney opanity, taitere to till ve, and polyhydraminos
REQ18-	Pathogenic or	MPV1	NM_002437.4:c.279+1		
1271	likely pathogenic	7	G>T	Hom	Individual with microcephaly, IUGR, elevated transaminases, and failure to thrive
	Final grant		GenBank:		, , , , , , , , , , , , , , , , , , , ,
	Ambiguous		NM_004536.2:c.602G		Cyclic neutropenia, arthritis, ?IBD, and suspected SCID; a candidate for bone marrow
18-2765	(candidate gene)	NAIP	>C:p.Gly201Ala	Hom	transplant
	(**** *********************************		GenBank:		
REQ18-	Pathogenic or		NM_000271.4:c.2114		
4855	likely pathogenic	NPC1	T>C:p.Leu705Pro	Hom	Individual with cholestatic liver disease; a candidate for liver transplant
			GenBank:		
18-1824-	Pathogenic or		NM_000298.6:c.1435		
В	likely pathogenic	PKLR	C>T:p.Arg479Cys	Hom	Individual with anemia and intellectual disability
			GenBank:		,
REQ18-	Pathogenic or		NM_001126131.1:c.32		
3345	likely pathogenic	POLG	86C>T:p.Arg1096Cys	Hom	Individual with liver failure and developmental delay; a candidate for liver transplant
			GenBank:		
18-1257-	Pathogenic or		NM_001083116.2:c.89		
В	likely pathogenic	PRF1	5C>T:p.Arg299Cys	Hom	Individual with immunodeficiency; a candidate for bone marrow transplant
					NICU 8-week-old boy with severe, complex congenital heart disease (pulmonary
			GenBank:		atresia, ventricular septal defect, major aortopulmonary collateral artery, patent ductus
REQ18-	Pathogenic or	PRKD	NM_002742.2:c.2554		arteriosus, and hypertrophic cardiomyopathy; on prostaglandin with biventricular
1959	likely pathogenic	1	G>T:p.Glu852Ter	Hom	hypertrophic cardiomyopathy)

			GenBank:		
			NM_000536.3:c.374_3		
REQ18-	Pathogenic or		75delCA:p.Thr125Arg		
1480	likely pathogenic	RAG2	fsTer10	Hom	Individual with immunodeficiency; a candidate for bone marrow transplant
			GenBank:		<i></i>
			NM_031229.3:c.120_1		
REQ18-	Pathogenic or	RBCK	21insTGCC:p.Leu41C		Individual with failure to thrive, gross and fine motor delay, lactic acidemia, recurrent
0419	likely pathogenic	1	vsfsTer9	Hom	fever, and immunodeficiency; a potential candidate for bone marrow transplant
	J T T T T T T T T T T T T T T T T T T T		GenBank:	-	,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,
		RNAS	NM_001286834.2:c.8		
18-0453	Ambiguous (VUS)	EH1	A>G:p.Tyr3Cys	Hom	Individual with lactic acidosis
	3		GenBank:		
17-1684-	Pathogenic or	SLC17	NM_012434.4:c.308G		NICU individual with non-immune hydrops fetalis, congenital dysmorphic features,
С	likely pathogenic	A5	>A:p.Trp103Ter	Hom	and neonatal cholestasis
	31 8		GenBank:		
REQ18-	Pathogenic or		NM_005633.3:c.1297		Individual with anemia and severe bone marrow dysplasia; a candidate for bone
4394	likely pathogenic	SOS1	G>A:p.Glu433Lys	Het	marrow transplant
	J		GenBank:		•
REQ18-	Ambiguous		NM_175610.3:c.3964		
2677	(candidate gene)	TJP1	G>A:p.Gly1322Arg	Het	NICU individual with lissencephaly and heterotopia
			GenBank:		
	Pathogenic or	TMEM	NM_001077416.2:c.82		
18-2764	likely pathogenic	231	4-11T>C	Hom	NICU individual with polydactyly and occipital encephalocele
			GenBank:		
REQ18-	Ambiguous	TOM	NM_001128917.1:c.27		
1528	(candidate gene)	M40	5A>G:p.Glu92Gly	Het	ICU adult with encephalopathy and a history of stroke-like episodes
			GenBank:		
			NM_003309.3:c.725_7		
REQ18-	Pathogenic or	TSPYL	26delTG:p.Val242Gluf		NICU case with IUGR, ambiguous genitalia, and a history of a deceased sister with
2590	likely pathogenic	1	sTer52	Hom	SIDS
			GenBank:		
			NM_001288782.1:c.10		
18-2763	Ambiguous (VUS)	TTC8	7G>A:p.Cys36Tyr	Hom	Individual with polydactyly and deteriorating renal function
			GenBank:		
REQ18-	Pathogenic or	VPS33	NM_001289148.1:c.14		Individual with microcephaly; skin, hair, and nail abnormalities; hearing loss;
2801	likely pathogenic	В	38C>T:p.Arg480Ter	Hom	hypotonia; failure to thrive; and elevated transaminases
			GenBank:		
			NM_014797.2:c.1492_		
REQ18-	Pathogenic or	ZBTB2	1493delCA:p.Gln498V		
3658	likely pathogenic	4	alfsTer15	Hom	Individual with dysmorphic features and epilepsy
			GenBank:		
REQ18-			NM_001171653.1:c.18		
4044	Ambiguous (VUS)	ZEB2	98C>T:p.Ser633Phe	Het	Individual with Hirschsprung disease

			GenBank:		
18-2346-		ZFHX	NM_024721.4:c.9271		NICU individual with tetralogy of Fallot, failure to thrive, developmental delay,
В	Ambiguous (VUS)	4	G>A:p.Gly3091Ser	Het	speech delay, facial dysmorphism, and external ear malformations

Question marks denote uncertainty of the clinical feature of diagnosis as provided by the ordering physician. Abbreviations are as follows: Hom = homozygous, Het = heterozygous, Hemi = hemizygous, IUGR = intrauterine growth restriction, NICU = neonatal intensive care unit, ICU = intensive care unit, SCID = severe combined immunodeficiency, IBD = inflammatory bowel disease, and SIDS = sudden infant death syndrome.

^aNegative instances can be found in Table S11.