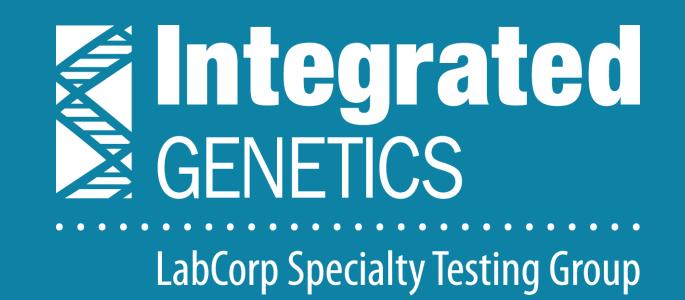
Comparison of pan-ethnic and ethnic-based carrier screening panels for individuals of Ashkenazi Jewish descent



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I. Introduction

The intent of carrier screening is to identify individuals at risk for a child with a genetic disease. ACMG guidelines currently recommend that individuals of Ashkenazi Jewish descent be screened for carrier status for nine diseases¹. A joint statement from ACMG, ACOG, NSGC, SMFM, and the Perinatal Quality Foundation acknowledges benefits of screening for more than nine diseases (expanded carrier screening)². Here we analyze detection rates for Ashkenazi Jewish individuals screened by panels with different numbers of diseases, to assess the benefit of disease panels targeted to the Ashkenazi Jewish population.

II. Materials and Methods

Array-based hybridization and allele-specific primer extension with a custom Illumina Infinium ™ array (IGv1.1) were used to detect 434 mutations in 87 genes that cause 87 diseases or a subset of 147 mutations in 18 genes that cause 18 diseases. Mutations were confirmed by Sanger sequencing. The 87-gene panel was intended for pan-ethnic carrier screening (pan-ethnic panel, Table 1), and the 18-gene panel was intended for Ashkenazi Jewish carrier screening (AJ panel).

The study sample comprised individuals self-identified as Ashkenazi Jewish and their indication for testing was carrier screening with no personal or family history of a genetic disorder. A total of 1150 individuals were tested in the pan-ethnic panel and 1248 individuals were tested in the AJ panel.

To compare pan-ethnic and AJ-based panels, positive findings for the AJ individuals tested in the pan-ethnic panel were re-analyzed with the 18 genes in the AJ panel and with the 9 genes recommended by ACMG. Similarly, positive findings for the AJ individuals tested in the AJ panel were re-analyzed with the 9 genes recommended by ACMG.

Table 1. The 87 genes and diseases included in the pan-ethnic panel.

Gene	Disease	Gene	Disease	Gene	Disease
ABCC8	Familial Hyperinsulinism, ABCC8-Related	DPYD	Dihydropyrimidine Dehydrogenase Deficiency	MEFV	Familial Mediterranean Fever
ACADM	MCAD Deficiency	ETHE1	Ethylmalonic Encephalopathy	MMAA	Methylmalonic Acidemia, MMAA-Related
ADA	Adenosine Deaminase Deficiency	FAH	Tyrosinemia Type 1	MMAB	Methylmalonic Acidemia, MMAB-Related
AGA	Aspartylglucosaminuria	FANCC	Fanconi Anemia Group C	ММАСНС	Cobalamin C Disease (Methylmalonic Aciduria and Homocystinuria Type cblC)
AGL	Glycogen Storage Disease Type IIIa and IIIb	FKTN	Walker-Warburg Syndrome, FKTN-Related	MUT	Methylmalonic Acidemia, MUT-Related
AGXT	Primary Hyperoxaluria Type 1	G6PC	Glycogen Storage Disease Type Ia	NBN	Nijmegen Breakage Syndrome
ALDH3A2	Sjogren-Larsson Syndrome	GAA	Pompe Disease (Glycogen Storage Disease Type II)	NEB	Nemaline Myopathy, NEB-Related
ALDOB	Hereditary Fructose Intolerance	GALC	Krabbe Disease	NPC1	Niemann-Pick Disease Type C, NPC1-Related
ARSA	Metachromatic Leukodystrophy	GALT	Galactosemia, GALT-Related	NPC2	Niemann-Pick Disease Type C, NPC2-Related
ASL	Argininosuccinic Aciduria	GBA	Gaucher Disease	NPHS1	Nephrotic Syndrome, NPHS1-Related (Congenital Finnish Nephrosis)
ASPA	Canavan Disease	GCDH	Glutaric Acidemia Type 1	NPHS2	Nephrotic Syndrome, NPHS2-Related
ASS1	Citrullinemia Type I	GLDC	Glycine Encephalopathy, GLDC-Related (Non-Ketotic Hyperglycinemia, GLDC-Related)	PAH	Phenylalanine Hydroxylase Deficiency (including PKU)
ATM	Ataxia-Telangiectasia	GRHPR		PCCA	Propionic Acidemia, PCCA-Related
АТР7В	Wilson Disease	GSS	Glutathione Synthetase Deficiency	РССВ	Propionic Acidemia, PCCB-Related
BBS1	Bardet-Biedl Syndrome, BBS1 Related	HADHA	LCHAD Deficiency	PCDH15	Usher Syndrome Type IF
BBS10	Bardet-Biedl Syndrome, BBS10 Related	НВВ	Beta Hemoglobinopathy, including Sickle Cell Disease	PEX1	Zellweger Syndrome Spectrum, PEX1-Related
BCKDHA	Maple Syrup Urine Disease Type 1A	HEXA	Tay-Sachs Disease	PEX7	Rhizomelic Chondrodysplasia Punctata Type 1
BCKDHB	Maple Syrup Urine Disease Type 1B	HEXB	Sandhoff Disease	PKHD1	Polycystic Kidney Disease, Autosomal Recessive
BCS1L	GRACILE Syndrome	HMGCL	HMG-CoA Lyase Deficiency	PMM2	Congenital Disorder of Glycosylation Type 1a
BLM	Bloom Syndrome	HLCS	Holocarboxylase Synthetase Deficiency	PPT1	Neuronal Ceroid-Lipofuscinosis, PPT1-Related
CBS	Homocystinuria, CBS-Related	HSD17B4	D-Bifunctional Protein Deficiency	RMRP	Cartilage-Hair Hypoplasia
CFTR	Cystic Fibrosis	IDUA	Mucopolysaccharidosis Type I (Hurler Syndrome)	SACS	Autosomal Recessive Spastic Ataxia of Charlevoix- Saguenay (ARSACS)
CLN3	Neuronal Ceroid-Lipofuscinosis, CLN3-Related	IKBKAP	Familial Dysautonomia	SLC12A6	Andermann Syndrome
CLN5	Neuronal Ceroid-Lipofuscinosis, CLN5-Related	LAMA3	Junctional Epidermolysis Bullosa, LAMA3-Related	SLC17A5	Salla Disease
CLN8	Neuronal Ceroid-Lipofuscinosis, CLN8-Related	LAMB3	Junctional Epidermolysis Bullosa, LAMB3-Related	SLC26A2	Sulfate Transporter-Related Osteochondrodysplasias
CLRN1	Usher Syndrome Type III	LAMC2	Junctional Epidermolysis Bullosa, LAMC2-Related	SLC37A4	Glycogen Storage Disease Type Ib
CTNS	Cystinosis	LRPPRC	Leigh Syndrome, French-Canadian Type	SMPD1	Niemann-Pick Disease Type A/B
DHCR7	Smith-Lemli-Opitz Syndrome	MAN2B1	Alpha-Mannosidosis	TMEM216	Joubert Syndrome 2
DLD	Dihydrolipoamide Dehydrogenase Deficiency (MSUD Type 3)	MCOLN1	Mucolipidosis Type IV	TTP1	Neuronal Ceroid-Lipofuscinosis, TTP1-Related

Blue: the pan-ethnic panel genes that are not also AJ or ACMG Green plus yellow: the subset of 18 AJ panel genes Yellow: the subset of 9 ACMG recommended AJ genes

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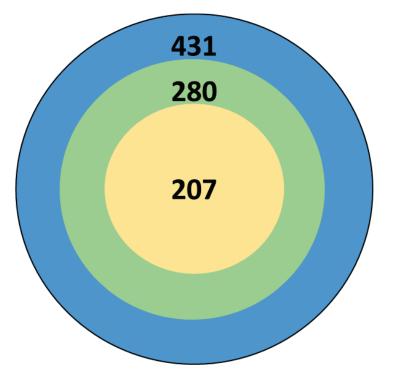


III. Results and Discussion

Table 2. Comparison of positive results for Ashkenazi Jewish individuals tested in the pan-ethnic and AJ panels.

Test	Number of genes assessed	individuals	Number of individuals positive:				Total number	%
icst			1 gene	2 genes	3 genes	4 genes	positive	Positive
Pan-ethnic panel	87	1150	344	72	13	2	431	37.5
	18		246	30	4	0	280	24.3
	9		186	19	2	0	207	18.0
AJ panel	18	1248	259	53	5	2	319	25.5
	9		203	33	1	0	237	18.9

Figure 1. Companion diagram for Table 2.

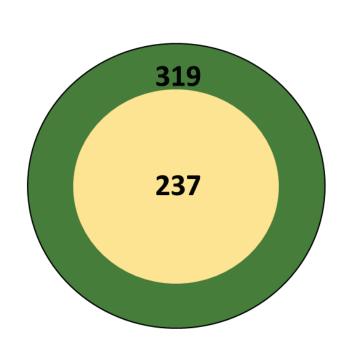


Pan-ethnic panel

recommended genes

- N = 1150 AJ individuals tested
- Individuals who were positive for at least one of the 87 pan-ethnic genes
- Individuals who would have been positive if screening was limited to the 18 AJ genes Individuals who would have been positive if

screening was limited to the 9 ACMG



AJ panel

N=1248 AJ individuals tested

- Individuals who were positive for at least one of the 18 AJ genes
- Individuals who would have been positive if screening was limited to the 9 ACMG recommended genes

DISCUSSION: In the pan-ethnic panel 431/1150 (37.5%) AJ individuals were carriers of at least one disease. If these individuals were tested in the AJ panel, the detection rate would be 280/1150 (24.3%). If these individuals were tested for the nine ACMG recommended diseases, the detection rate would be 207/1150 (18.0%). The validity of this extrapolation is demonstrated by the equivalent findings that the detection rate is 25.5% for AJ individuals tested with the AJ panel and 18.9% for the ACMG recommended diseases. (Table 2 and Figure 1)

Table 3. Twenty-one diseases not included in the ethnic-specific AJ panel, but identified in Ashkenazi Jewish carriers tested in the pan-ethnic panel.

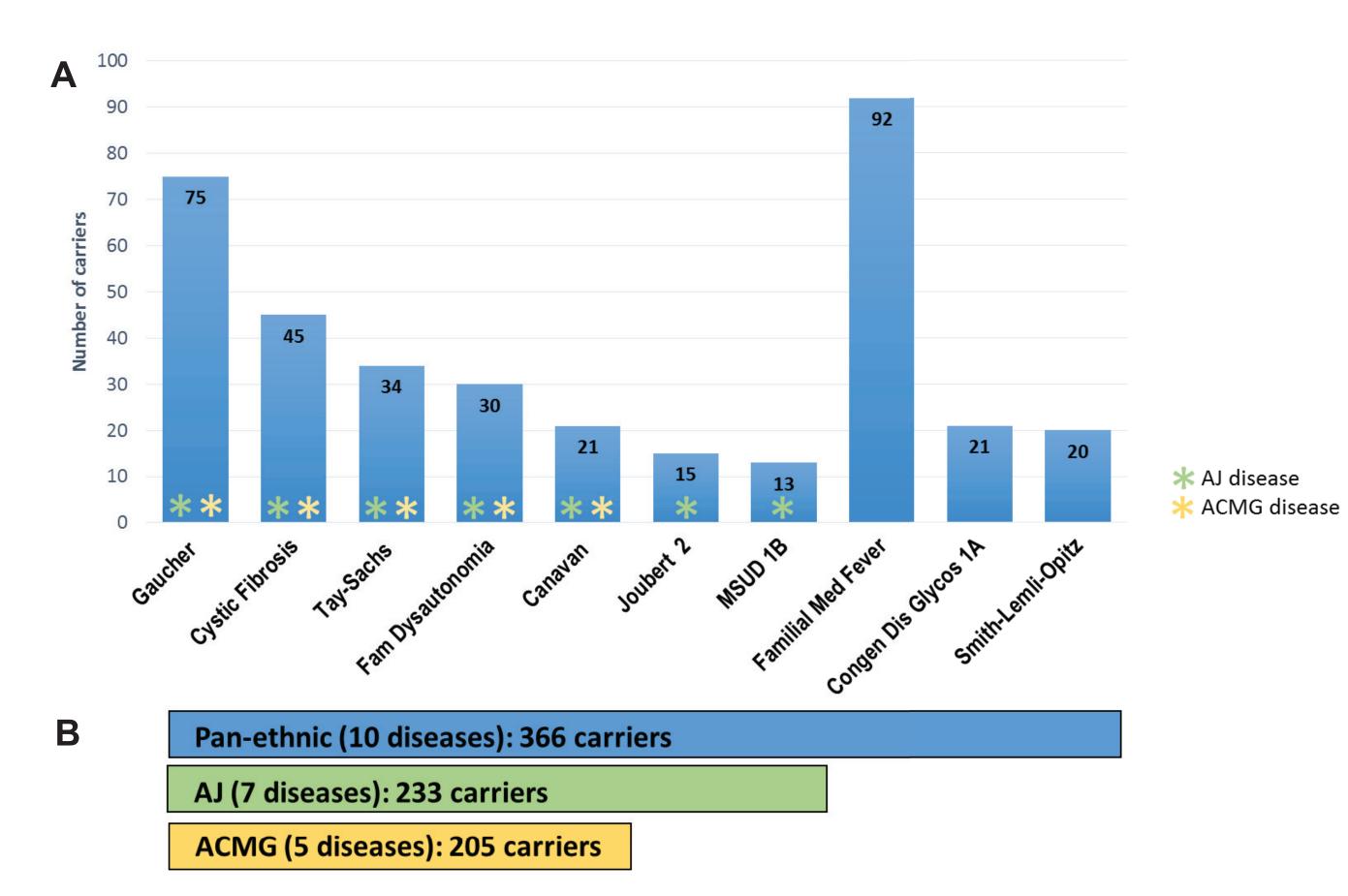
Disease	Gene	Disease	Gene
Beta Hemoglobinopathy, including Sickle Cell Disease	HBB	Neuronal Ceroid-Lipofuscinosis, TTP1-Related	TTP1
Cobalamin C Disease (Methylmalonic Aciduria and Homocystinuria Type cblC)	MMACHC	Niemann-Pick Disease Type C, NPC1-Related	NPC1
Congenital Disorder of Glycosylation Type 1a	PMM2	Niemann-Pick Disease Type C, NPC2-Related	NPC2
Cystinosis	CTNS	Pompe Disease (Glycogen Storage Disease Type II)	GAA
Dihydropyrimidine Dehydrogenase Deficiency	DPYD	Primary Hyperoxaluria Type 1	AGXT
Familial Mediterranean Fever	MEFV	Smith-Lemli-Opitz Syndrome	DHCR7
Galactosemia, GALT-Related	GALT	Sulfate Transporter-Related Osteochondrodysplasias	SLC26A2
Glycogen Storage Disease Type Ib	SLC37A4	Tyrosinemia Type 1	FAH
Hereditary Fructose Intolerance	ALDOB	Wilson Disease	ATP7B
Homocystinuria, CBS-Related	CBS	Neuronal Ceroid-Lipofuscinosis, TTP1-Related	TTP1
Mucopolysaccharidosis Type I (Hurler Syndrome)	IDUA		

DISCUSSION: The pan-ethnic panel identified carriers for 38 diseases among Ashkenazi Jews; however, carrier status for only 17 of these diseases could be assessed using the AJ panel. Therefore, 21 "non-AJ" diseases accounted for the difference in carrier rates between the pan-ethnic and AJ panels (Table 3)

Figure 2.

A. The ten diseases with the largest number of carriers among Ashkenazi Jewish individuals screened in the pan-ethnic panel.

B. Comparison showing the subset of these ten diseases and carriers that would be identified if only the 18 AJ panel genes or 9 ACMG recommended genes were assessed.



DISCUSSION: Of the ten diseases with the largest number of carriers among Ashkenazi Jewish individuals screened in the pan-ethnic panel, three diseases would have been missed if the AJ panel had been used: familial Mediterranean fever (92 carriers), congenital disease of glycosylation type 1a (21 carriers), and Smith-Lemli-Opitz syndrome (20 carriers). With the ACMGrecommended panel, two more of the most commonly identified diseases would have been missed: Joubert syndrome type 2 (15 carriers) and maple syrup urine disease type 1B (13 carriers). (Figure 2)

IV. Conclusions

- > A pan-ethnic expanded carrier screening panel of 87 genes increased the carrier detection rate in Ashkenazi Jewish individuals by approximately 50%, compared with a panel of 18 genes considered to be relevant to the Ashkenazi Jewish population.
- > The detection rate would have increased by approximately 100% if the pan-ethnic panel were compared to just the ACMG recommended genes in this data set.
- > These data show that a pan-ethnic panel is more effective than targeted AJ panels in carrier detection among individuals of Ashkenazi Jewish descent.

V. References

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