



# Prevalence and Findings of Genetic Testing in Scimitar Syndrome

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## BACKGROUND

- Scimitar syndrome (SS) is a rare congenital heart defect where the right-sided pulmonary veins drain aberrantly to the inferior vena cava instead of the left atrium.
- Several studies have highlighted the genetic implications in total anomalous pulmonary venous return (TAPVR), however the genetics of SS are less well-elucidated.
- Until recently, SS was believed to occur largely in isolation, but recent reports there is an increased prevalence of SS in 22q11 and VACTERL syndrome.

## OBJECTIVE

We sought to identify the prevalence of genetic testing, either with chromosomal microarray (CMA) or whole exome sequencing (WES), in a cohort of patients with SS at our institution. We also sought to characterize the results of the studies.

## METHODS

- Data Source: Texas Children's Hospital electronic database
- Inclusion criteria: SS, defined as partial anomalous pulmonary venous return of the right pulmonary veins to the inferior vena cava – right atrial junction forming a “Scimitar” vein
- Exclusion criteria: TAPVR
- CMA and WES results were reviewed. Variants noted as pathogenic or likely pathogenic were considered “positive”. Additionally, all reported genetic results were reviewed and variants of unknown significance also reported.

## RESULTS

Patient Characteristics (N=77)		
	N	Percentage
<b>Sex</b>		
Female	52	68%
Male	25	32%
<b>Testing</b>		
CMA performed	17	22%
CMA positive	2	12%
WES performed	5	6%
WES positive	3	60%

CMA	WES	CMA info	Genes	Interpretation	Other CHD	Extracardiac defects
Y		GAIN 22q11.21	>20 genes	<b>22q11.2 duplication syndrome</b>	Dextroposition, 2° ASD, RPA narrowing, arch elongation	R lung hypoplasia, AP collateral, absent RUL, CDH, post. rib fusions, hemivertebra,
Y		GAIN 10q21.3q23.1	>20 genes	<b>10q22.3-q23.2 deletion syndrome</b>	2° ASD, coarctation, pulmonary valve stenosis	CDH, rib fracture, R lung hypoplasia
Y	Y	-	<b>NAA15</b>	<b>Aut. dom. mental retardation, with CHD</b>	Dextroposition, TOF, mild pulmonary valve stenosis, HCM, 2° ASD	R lung hypoplasia
Y	Y	LOSS 3q13.2; LOSS 8q21.3	<b>MYRF</b>	<b>Cardiac urogenital syndrome</b>	HLHS (MA/AS), severe coarctation, pulm HTN	R lung hypoplasia, global delay paraesophageal hernia,
Y	Y	GAIN 2q.37.3	<b>EP300</b>	<b>Rubinstein-Taybi syndrome</b>	Aberrant R subclavian, L SVC	FTT, global delay, laryngeal cleft, CDH
Y	Y	LOSS 16q22.1	-	Unknown significance	TOF, ASD	R lung hypoplasia, R hand ectrodactyly, thoracic vert hypoplasia, scoliosis
Y		GAIN 16p13.11	-	Unknown significance	Dextroposition, ASD, pulm HTN, SVT	FTT, Pyloric stenosis, OSA, bronchomalacia, CDH
Y		LOSS 8p11.21	-	Unknown significance	ASD, VSD	Scoliosis, torticollis, leg-length discrepancy, asthma
Y		DER (X) T (X; 2) (2-6.1 q31.1)	-	Unknown significance	Dextroposition, PDA, VSD	Global delay, thrombocytopenia, scoliosis

CHD = congenital heart disease; HLHS = hypoplastic left heart syndrome; 2° ASD = secundum ASD; PDA = patent ductus arteriosus; VSD = ventricular septal defect; TOF = tetralogy of Fallot; HCM = hypertrophic cardiomyopathy; CDH = congenital diaphragmatic hernia

- An additional patient clinically has Marfan syndrome, but *FBN1* sequencing has not been performed

## LIMITATIONS

This was a retrospective study, with non-universal genetic testing, and WES performed only in a fraction of patients. The sample may be biased if testing was conducted in patients with greater degrees of dysmorphism or extracardiac birth defects.

## CONCLUSIONS

- 22% of patients had genetic testing performed, and this was predominantly composed of CMA testing.
- Pathogenic variants were found in 5/17 (30%) patients, accounting for at least 6% of our entire cohort.
- Three of the 5 pathogenic variants involved changes in genes related to different phases of cell growth/proliferation, including TGF-beta, transcription coactivators and post-translational acetylation processing. One involved oligodendrocyte cell proliferation. One involved microduplication of the 22q11.21 region.
- Genetic testing – CMA with reflex to WES if CMA is negative – should be considered in patients with SS.

CMA/WES Results in Patients with Genetic Testing

