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Author for Correspondence:

A. Pierick, Emory University School of Medicine, Department of Pediatrics, 49 Jesse Hill Jr Dr SE, Atlanta, GA, USA 30303. Tel: +(319)541-8325. E-mail: Alysonpierick8@gmail.com

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Comparison of growth and feeding method in infants with and without genetic abnormalities after neonatal cardiac surgery

Alyson R. Pierick¹, Trudy A. Pierick² and Benjamin E. Reinking²

¹Emory University School of Medicine, Department of Pediatrics, Atlanta, GA, USA and ²University of Iowa Stead Family Children's Hospital, Department of Pediatrics, Iowa City, IA, USA

Abstract

Introduction: Congenital heart disease (CHD) is multifactorial in origin, resulting from an interaction between environmental and genetic factors. Multifactorial growth delay is common in infants with CHD. The impact of a genetic abnormality and CHD on the growth of an infant is lacking in the literature. The aim of this study is to compare the growth and method of feeding following neonatal cardiac surgery in infants with normal versus abnormal genetic testing. Methods: A retrospective chart review of neonates who underwent a Risk Adjustment in Congenital Heart Surgery IV-VI procedure between 1 January, 2006 and 22 September, 2016 was performed at our institution. Weight, length, head circumference measurements, and feeding method were collected at birth, time of neonatal surgery, and monthly up to 6 months of age. Results: A total of 53 infants met inclusion criteria, of which 22 had abnormal genetic testing. Approximately 90% of infants were discharged following neonatal cardiac surgery with supplemental tube feeds. At each monthly follow-up visit, more infants were exclusively fed orally: 80% of infants with normal genetics at 5 months post-operative follow-up versus 60% of infants with abnormal genetic testing, although statistically insignificant. Growth was not different among the two groups. Conclusions: Infants with critical CHD with or without genetic abnormalities are at risk for growth delays and many need supplemental tube feeds post-operatively and throughout follow-up. Infants with genetic abnormalities are slower to achieve oral feeds and more likely to require tube feedings. It is important to have a systematic protocol for managing these high-risk infants.

Introduction

Congenital heart disease (CHD) is the most common congenital malformation, occurring in nearly 1% of newborns. While many infants with CHD have normal life expectancies and require minimal medical care, those with critical CHD (defined as a lesion requiring a Risk Adjustment in Congenital Heart Surgery (RACHS) IV–VI neonatal surgery) frequently require surgery in the neonatal period. Over the last few decades, advances in procedural techniques and post-operative care have improved survival. As a result, attention has shifted to decreasing morbidities and improving long-term quality of life.

Optimising nutrition and growth, specifically in the peri- and post-operative period, improves outcomes. ⁴⁻⁹ Poor nutrition and growth in the pre-operative period are associated with poor neurodevelopmental outcomes, impaired wound healing, higher infection rates, and longer hospital length of stay. ^{4,5,10} Growth delay is multifactorial and can be secondary to cardiac failure, non-cardiac anomalies, tachypnea, increased energy expenditures, inadequate caloric intake, gastroesophageal reflux, dysphagia, oral aversion, and vocal cord dysfunction. ^{4,11,12} Extracardiac malformations, often associated with genetic abnormalities, may also contribute to growth delay. ¹³

Determining genetic abnormalities in infants with CHD is important when considering non-cardiac morbidities. Neurodevelopmental disability, respiratory disease, renal disease, and growth delay are all common in patients with genetic defects. The growth of patients with trisomy 21 and Turner syndrome, for example, is monitored using syndrome specific growth charts. The presence of non-cardiac or genetic abnormalities in patients undergoing neonatal cardiac surgery is associated with increased mortality, longer intensive care unit stay, and prolonged hospital stay. The presence of non-cardiac mortality is associated with increased mortality.

The aim of this study was to determine the impact of genetic abnormalities on the growth and method of feeding from birth to 6 months of age in infants with critical CHD following neonatal cardiac surgery.

Materials and methods

Internal Institutional Review Board approval was obtained for a retrospective chart review of all patients who underwent a RACHS IV–VI procedure in the neonatal period at the University of Iowa Stead Family Children's Hospital between 1 January, 2006 and 22 September, 2016.

Table 1. Demographic data

	Normal genetics N (%)	Abnormal genetics N (%)	
Total	31 (58.5)	22 (41.5)	
Male	23 (74)*	6 (27.2)	Z = 3.38, p < 0.001
Gestational age (days)	271 +/- 11	272 +/- 8	p = 0.59
Prenatal diagnosis	17	11	Z = 0.476 p = 0.634
Main cardiac diagnosis			
Hypoplastic Left Heart Syndrome variant (14)	10	4	p = 0.252
NON HLHS single ventricle (9)	6	3	p = 0.304
Unbalanced AV Canal (2)	1	1	
DORV variants (4)	2	2	
Ebstein's anomaly (3)	3	0	
Arch anomalies (15)	4	11*	Z = 2.954 p = 0.003
Coarctation (7)	4	3	
Coarctation + VSD (7)	0	7	
Interrupted aortic arch (1)	0	1	
D-TGA variants (7)	5	2	p = 0.803
D-TGA, VSD, coarctation (2)	2	0	
D-TGA VSD (5)	3	2	
TAPVR variant (6)	4	2	p = 0.66
Truncus arteriosus	1	0	p = 0.395
Critical aortic stenosis	1	0	p = 0.395

Abbreviations: D-TGA = dextro-transposition of the great arteries; DORV = double outlet right ventricle; HLHS = hypoplastic left heart syndrome; TAPVR = total anomalous pulmonary venous return; unbalanced AV canal = unbalanced atrioventricular canal; VSD = ventricular septal defect.

All neonates without genetic testing, those who were lost to follow up and those who died before 6 months of age, were excluded from analysis.

Growth parameters including weight, length, and head circumference were extracted from the electronic medical record at birth, the time of surgery, and at monthly intervals until 6 months of age. The route of feeding (enteral tube, oral, or combination of oral and enteral tube) was collected at birth, at the time of hospital discharge, and monthly from the time of surgery for the first 5 months. The number of subjects receiving feedings by each method was determined at each time point. Individuals were separated into groups based on genetic testing results (normal or abnormal). Study data were collected and managed using REDCap electronic data tools hosted at the University of Iowa.

Statistical analysis

Demographic and clinical variables were compared among groups with normal and abnormal genetic testing using Pearson Chi-square test for the categorical variables. Linear regression models were used to assess the association between the growth end points (weight, length, and head circumference) and explanatory

Table 2. Genetic defects and associated cardiac diagnoses. Arch anomalies were defined as coarctation, hypoplastic aortic arch, or interrupted aortic arch. D-TGA variants included isolated D-TGA +/- ventricular septal defect and coarctation

Cardiac diagnosis	Genetic defect		
Hypoplastic Left Heart Syndrome	94 kb deletion of 13q13.1		
	1.4 Mb deletion at 3q13.12q13.13		
	translocation at 5p14.1 and 18q11.2		
	505 kb duplication of 15q11.2		
Non-HLHS single ventricle	914 kb duplication at 11p14.1		
	520 kb duplication of 15q11.2		
	319 kb deletion at 15q11.2		
Arch anomalies	350 kb duplication at 15q22.33		
	22q11.2 deletion		
	22q11.2 deletion		
	142 kb duplication of 1q44		
	45 XO		
	47 XY + 21		
	171 kb deletion of 10p15.2, 268 kb duplication of 21q22.3		
	917 kb deletion of 16q24.1; 1.6 Mb duplication of 2p12		
	308 kb deletion of 13q12.11, Xo		
	512 kb deletion of 15q11.2		
	47 XY + 21		
D-TGA variants	951 kb deletion at 20p12.1		
TAPVR	191 kb duplication at 6q26		
	7.9 Mb deletion of 18p11.32p11.23 (18p deletion syndrome)		

 $Abbreviations: \ HLHS = hypoplastic \ left heart syndrome; TAVPR = total \ anomalous \ pulmonary \ venous \ return; \ D-TGA = D-transposition \ of \ the \ great \ vessels.$

variables (single ventricle versus two ventricle CHD, gender, gestational age at birth, and genetic testing result). Multivariable logistic regression models were performed to identify independent variables associated with feeding method outcomes. Independent variables for regression models were defined as single ventricle versus two ventricle CHD, gender, gestational age at birth, and genetic testing result. Significance was assessed at the 0.05 level.

Results

Demographics

During the study period, 97 patients underwent RACHS IV–VI procedures. Of those neonates, 7 were excluded because they were lost to follow up and 18 were excluded due to death before 6 months of age. Of the remaining 72 newborns, 53 had genetic testing performed and were included for further analysis. In total, 29 infants (54.7%) were male. Genetic defects were identified in 22 infants (41.5%). There were significantly more males (p < 0.001) in the group with normal genetic testing. Gestational age and rate of prenatal diagnosis were similar in both groups. Demographic data including cardiac diagnosis can be found in Table 1. Information on the identified genetic defects and the associated cardiac lesions can be found in Table 2.

^{* =} significant difference on Chi-square analysis.

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Anthropomorphic measurements

Weight, length, and head circumference measurements were collected for each infant at birth, time of surgery, and each month following surgery until 6 months of age. Although infants with genetic defects weighed less than those with normal genetic testing at every time point, the difference was not statistically significant. Similarly, length and head circumference measurements were less in those with abnormal genetic testing than in those with normal genetic testing at every time point. Those with genetic defects were noted to have significantly smaller head circumference at 5 months of age (p < 0.01) and shorter lengths at 3 months of age (p = 0.02) and 6 months of age (p < 0.01). Length and head circumference measurements were not significantly different between the two groups at any other time point. Growth data is summarised in the Figure 1. Data was not available for each subject at all time points. The number of subjects included at each time point can be found in the figures. Significant differences between the normal and abnormal genetic groups are designated with a star.

Linear regression models assessing anthropometric measurements with potential confounders showed that increasing gestational age was significantly associated with increasing weight, length, and head circumference at each time point. Although statistically insignificant, abnormal genetic test results, female gender, and single ventricle physiology were negatively associated. Results of the linear regression can be found in Table 3.

Method of feeding

Feeding method from the post-operative period to 6 months of age is summarised in Table 4. In both groups, most infants received feeds via an enteral tube in the immediate post-operative period. At each subsequent follow up visit, the percentage of infants exclusively orally fed increased. Those with normal genetic testing transitioned to all oral feeds more quickly than those with genetic defects, with significantly more infants taking all oral feeds at 3 months and 4 months of age. By 5 months of age, those with abnormal genetic testing taking feeds orally increased and there was no significant difference between the two groups. Multinomial logistic regression models found that no variable significantly impacted the method of feeding between the immediate post-operative period and at 5 months post operation. Results of the regression model can be found in Table 5.

Discussion

In this retrospective chart review, we evaluated newborns with critical CHD who underwent a neonatal cardiac surgery and compared growth and method of feeding of those with normal and abnormal genetic testing. We found that at our centre, there was no difference in growth parameters from birth to 6 months of age in these infants regardless of genetic testing. Infants with genetic abnormalities were slower to reach full oral feeds even at 5 months after surgery.

The rate of genetic abnormalities in our study population was similar to published values. ¹⁶ There was, however, an important difference in the demographics of the two groups. The percentage of females was significantly greater in the abnormal genetic testing group. While the reason for the female predominance in this study could be related to gender differences in specific congenital heart defects, ¹⁷⁻¹⁹ it could also be related to the small sample size of our single center study.

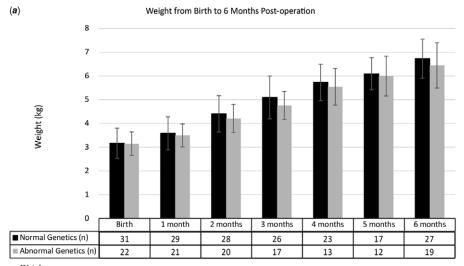
Growth following neonatal cardiac surgeries, specifically following the Norwood procedure, has been well studied. 4,5,8,20-22,20,23 With the recognition that infants post Norwood procedure have increased caloric needs, most institutions have instituted specific home-monitoring programs focussing on weight gain and increased surveillance during the inter-stage period. Poor weight gain has been associated with more post-operative complications, longer hospital stays, and worse neurodevelopmental outcomes. 4,10,21,24 Length has also been studied, with these infants on average being shorter at birth and not recovering to a normal height even with proper nutrition, 4,21,25 indicating there may be some other component contributing to growth in height. Most of these studies have excluded infants with recognised chromosomal abnormalities.

While the infants with abnormal genetic testing were smaller at almost every time point than those with normal genetic testing, few statistically significant differences were found among the groups in this study. We hypothesise that this is due to a programmatic emphasis on post-operative growth and nutrition in all infants who undergo complex neonatal cardiac surgeries. During the study period, our institution joined the National Paediatric Cardiology Quality Improvement Collaborative (NPC-QIC) in 2012. Membership in this collaborative changed discharge and feeding protocols during the study period.

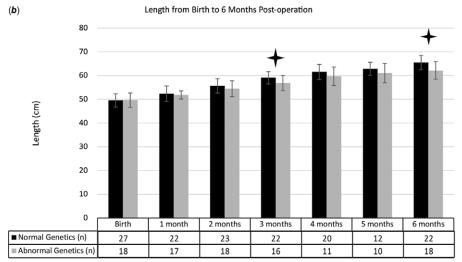
Prior to joining the collaborative, we discharged all neonatal cardiac patients in the same manner, without respect to diagnosis or genetic defects. Parents received extensive teaching from nursing staff regarding their infant's cares, including mixing formula or breast milk correctly, inserting nasogastric tubes, and giving medications and feedings per nasogastric tube. Once competence was demonstrated, parents completed an apartment experience for 24–48 hours in which they provided all cares for the infant using home equipment. The patient was then discharged with a daily home worksheet documenting feedings, output, medications, vitals from home nursing visits, and oxygen saturations if the patient was a single ventricle. Patients were followed by the primary cardiologist at routine intervals with outpatient feeding management by the cardiology nurse practitioner and dietitian, adjusting to maintain an average weight gain of 15–20 gm/day.

After joining the collaborative, many of these practices regarding feeding, discharge preparation, and outpatient management remained the same. An official home-monitoring program for inter-stage care of single ventricle and shunt dependent infants was developed in 2012. Families were now given a baby scale to document daily weights and learned how to calculate weight changes with a goal of 60 gm over a 4-day period. Outpatient management of this group was restructured to a team of two cardiologists, a cardiology nurse practitioner, and a dietician. Interval growth goals remained the same. Those infants followed outside of the inter-stage clinic had growth managed by the primary cardiologist, cardiology nurse practitioner, and dietitian, with less frequent weight collection (at clinical visits) but similar growth goals to the single ventricle patients. The consistency of feeding management by these same care providers, regardless of cardiac or genetic diagnosis, validates our findings over this 10-year period, even with programmatic changes in care.

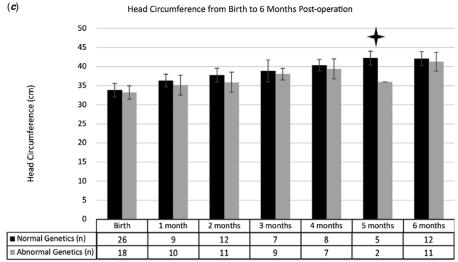
At the time of discharge, $\sim 90\%$ of infants in this study required nutrition via an enteral tube, regardless of genetic testing results. Those with normal genetic testing, however, progressed more quickly in their oral feeding skills, with a significantly greater number receiving at least a portion of their nutrition orally at 3 and 4 months of age. This difference could be due to the increased risk for neurodevelopmental delay, abnormal muscle tone, and



Weight



Length



Head circumference

Figure 1. Growth parameters from birth through 6-months post-op. Black bars denote patients with normal genetics, light grey bars denote patients with abnormal genetic testing. Small lines in the bars represent standard deviation. Tables under the bars correspond to the number of patients with a measurement at that time frame for each group. P-value <0.05 denoted with stars above the corresponding time frame. P values for the length differences at 3 and 6 months were 0.025 and 0.003 respectively, 0.006 for the head circumference difference at 5 months.

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Table 3. Results of linear regression model evaluating the impact of single ventricle (SV*) versus two ventricle congenital heart disease, gender (male*), gestational age, and chromosomal microarray results (normal*) on weight, length, and head circumference between birth and 6 months post-operation

Variable Coefficient 95% CI p-value Birth weight SV versus 2V -0.111-0.423, 0.2 0.476 Gender -0.204-0.548, 0.1390.237 GA 0.013 0.011, 0.015 <0.001 CMA results -0.426, 0.233 0.559 -0.096SV versus 2V 1-month wt. -0.016-0.347, 0.3150.922 Gender -0.415-0.795, -0.036 0.033 0.012, 0.016 0.014 <0.001 CMA results -0.497, 0.24 0.485 -0.1292-month wt. SV versus 2V 0.088 -0.32, 0.41970.666 Gender -0.191 -0.641, 0.2580.396 GA 0.013, 0.018 0.015 < 0.001 CMA results -0.3, 0.563-0.1310.544 3-month wt. SV versus 2V 0.013 -0.481, 0.506 0.959 Gender -0.22-0.772, 0.3270.418 GA 0.015, 0.021 0.019 < 0.001 CMA results 0.282 -0.247, 0.8110.287 4-month wt. SV versus 2V 0.202 -0.313, 0.717 0.43 Gender -0.492-1.106, 0.1220.112 0.02 0.017, 0.024 <0.001 CMA results -0.006-0.602, 0.5890.983 5-month wt. SV versus 2V -0.316, 0.944 0.315 0.314 Gender -0.462-1.091, 0.167 0.143 GΑ 0.0147, 0.025 <0.001 0.021 -0.497, 0.649CMA results 0.076 0.786 6-month wt. SV versus 2V -0.536, 0.533-0.0010.996 Gender -0.226-0.827, 0.376 0.453 GΑ 0.021, 0.028 <0.001 0.024 CMA results 0.211 -0.369, 0.7910.466 Birth length SV versus 2V 0.255 -1.412, 1.923 0.759 Gender -2.545, 1.324 0.527 -0.611GΑ 0.171, 0.194 0.182 < 0.001 CMA results 0.058 -1.824, 1.94 0.951 1-month lt. SV versus 2V -0.592 -2.148, 0.963 0.445 Gender -1.018-2.862, 0.8260.270 GΑ 0.196 0.186, 0.206 <0.001 CMA results 0.146 -1.573. 1.865 0.864 2-month lt. SV versus 2V 0.554 -1.272, 2.381 0.542 Gender -0.443-2.434, 1.5480.655 0.198 0.186, 0.209 <0.001 CMA results 1.569 -0.33, 3.468 0.103 3-month lt. SV versus 2V -0.674-2.837, 1.48960.531 Gender -0.177 -2.482, 2.128 0.877 GΑ 0.212 0.198, 0.226 < 0.001 -0.041, 4.411 0.54 CMA results 2.185

(Continued)

Table 3. (Continued)

	Variable	Coefficient	95% CI	p-value
4-month lt.	SV versus 2V	1.148	-1.728, 4.024	0.42
	Gender	-0.922	-4.3221, 2.477	0.583
	GA	0.216	0.197, 0.234	<0.001
	CMA results	1.556	-1.765, 4.878	0.345
5-month lt.	SV versus 2V	0.51	-3.779, 4.779	0.806
	Gender	-3.17	-7.466, 1.125	0.138
	GA	0.23	0.205, 0.255	<0.00
	CMA results	1.454	-2.221, 5.128	0.417
6-month lt.	SV versus 2V		-1.086, 2.991	0.350
	Gender	-0.719	-3.186, 1.748	0.558
	GA	0.223	0.21, 0.237	<0.00
	CMA results	3.904	1.585, 6.223	0.002
Birth head	SV versus 2V	0.414	-0.519, 1.347	0.375
circumference	Gender	-0.489	-1.551, 0.573	0.358
	GA	0.121	0.115, 0.127	<0.00
	CMA results	0.727	-0.281, 1.734	0.153
1-month HC	SV versus 2V	1.987	-2.223, 6.196	0.332
	Gender	-0.176	-4.693, 4.341	0.935
	GA	0.12	0.094, 0.146	<0.00
	CMA results	0.189	-4.416, 4.525	0.927
2-month HC	SV versus 2V	0.301	-1.566, 2.167	0.740
	Gender	-1.183	-3.507, 1.141	0.300
	GA	0.134	0.12, 0.148	<0.00
	CMA results	1.134	-1.189, 3.457	0.320
3-month HC	SV versus 2V	1.776		0.129
	Gender	-1.558		0.155
	GA	0.13	0.115, 0.146	
	CMA results	1.589	-0.704, 3.882	
4-month HC		0.654	-2.426, 3.734	
		-1.22		
	GA	0.145	0.129, 2.077	<0.00
	CMA results		-2.374, 3.201	
5-month HC	SV versus 2V	-0.279	-6.997, 6.439	
			-7.297, 8.313	
	GA		0.094, 0.174	0.00
			-0.052, 13.101	
6-month HC			-1.73, 1.562	
			-2.978, 0.899	
			0.144, 0.164	
			-0.981, 2.678	

Abbreviations: SV = single ventricle; 2V = two ventricle; CHD = congenital heart disease; GA = gestational age; CMA = chromosomal microarray; HC = head circumference; mo = months; $month{w}$ = weight; $month{h}$ t = length.

p-value < 0.05 in bold.

^{*}indicated reference category for regression model.

Table 4. Comparison of the feeding methods seen in infants with normal and abnormal genetic testing from the post-operative period to 5 months post-operation

	Post Operation	2 Months	3 Months	4 Months	5 Months
	Feeding route				
	Normal genetic testing n (%)				
Enteral tube	30 (96)	6 (19)	5 (17)	5 (17)	3 (10)
Oral	1 (4)	16 (52)	21 (70)	22 (76)	22 (80)
Combo	0 (0)	9 (29)	4 (13)	2 (7)	3 (10)
	Abnormal genetic testing n (%)				
Enteral tube	20 (95)	2 (10)	2 (10)	2 (12.5)	2 (13)
Oral	1 (5)	7 (37)	9 (45)	9 (56.25)	9 (60)
Combo	0 (0)	10 (53)	9 (45)	5 (31.25)	4 (27)
Z	z = 0.283	$X^2 = 3.513$	$X^2 = 6.259$	$X^2 = 6.016$	$X^2 = 3.11$
р	0.777	0.173	0.04	0.04	0.211

non-cardiac abnormalities in patients with genetic abnormalities. ¹⁶ All patients in the study, regardless of genetic testing result, received an increasing amount of feeds orally during the study, in line with previous studies. ⁸ Overall, it is our opinion that the number of infants in both groups who require enteral tubes to meet nutrition and growth goals is too high. It is our hope to decrease the need for enteral tube feeds in the future through standardisation of preoperative feedings, promoting early post-operative feeds, and early, ongoing involvement of therapists with expertise in infant feeding.

This study has several important limitations. First, the small sample size makes detection of small but significant impacts of the various clinical variables on growth and feeding impossible. Second, clinical practice variation between providers and over the study time period also limits the strength of study findings. As an example, feeding neonates prior to cardiac surgery was almost unheard of 10 years ago. Pre-operative oral feeding is now standard at our institution in infants deemed stable. Third, almost half of eligible patients during our study time frame were excluded due to various reasons, including death prior to the end of the study time frame (6 months post-operative). Fourth, there are patients with missing data especially length or head circumference, which are typically not measured at each cardiology clinic visit and would only be included in our electronic medical record if their pediatrician was in our medical system. Finally, other variables not evaluated such as feeding therapy, vocal cord dysfunction, or length of intubation may have important impacts on growth and ability to feed orally. 12,26

Our retrospective chart review of growth and feeding methods of infants following neonatal surgery for critical CHD emphasises some of the important factors that still need to be optimised in these children. Continued work on early oral feedings and ensuring optimal caloric intake for each infant is needed, regardless of genetics. Those with genetic abnormalities are more likely to have multifactorial difficulties with growth and oral feeding than their counterparts with normal genetics. It has been shown that a strong emphasis on growth and nutrition regardless of genetic testing can help mitigate growth differences. Our findings show that our centre has adequate protocols in place to alleviate any growth differences between those with and without genetic abnormalities, but children with genetic abnormalities have more difficulty achieving oral feedings and require longer tube feeds to maintain

Table 5. Results of multinomial logistic regression model evaluating the association between method of feeding (enteral tube feed as base outcome) and single ventricle (SV*) versus two ventricle congenital heart disease, gender (male*), gestational age, and chromosomal microarray results (normal*) between the post-operative period and 5 months post-operation

	Variable	Odd ratio (95% CI)	p-value
Post-Operation	SV versus 2V	4.079 (0.023, 736.58	0.596
	Gender	30.766 (0.02, 48,042)	0.361
	GA	0.968 (0.928, 1.01)	0.138
	CMA results	1.627, (0.092, 29.9)	0.740
	RRR	(95% CI)	
1-month	SV versus 2V	1.198 (0.219, 6.552)	0.835
	Gender	1.717 (0.225, 13.12)	0.602
	GA	0.997 (0.985, 1.01)	0.602
	CMA results	2.124 (0.263, 17.134)	0.480
2-month	SV versus 2V	-0.768 (0.128, 4.616)	0.773
	Gender	2.876 (0.338, 24.434)	0.333
	GA	1.005 (0.994, 1.017)	0.389
	CMA results	0.441 (0.06, 3.261)	0.422
3-month	SV versus 2V	1.655 (0.197, 13.92)	0.643
	Gender	1.297 (0.133, 12.608)	0.823
	GA	1.002 (0.989, 1.015)	0.738
	CMA results	0.22 (0.024, 2.016)	0.180
4-month	SV versus 2V	3.051 (0.472, 19.714)	0.241
	Gender	0.327 (0.04, 2.65)	0.295
	GA	0.999 (0.986, 1.012)	0.866
	CMA results	1.102 (0.118, 10.269)	0.932
5-month	SV versus 2V	1.123 (0.113, 11.127)	0.921
	Gender	1.821 (0.167, 19.909)	0.623
	GA	1.968E-4 (0.986, 1.016)	0.897
	CMA results	1.958 (0.218, 17.584)	0.549

Abbreviations: RRR = relative risk ratio; CI = confidence intervals; SV = single ventricle; $2V = two \ ventricle$; CHD = congenital heart disease; GA = gestational age; CMA = chromosomal microarray; HC = head circumference; mo = months; wt = weight; lt = length. p-value < 0.05 in bold.

growth. Multicentre studies concentrating on specific cardiac abnormalities with larger numbers of infants are required to further elucidate differences.

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Conflicts of Interest. None.

Ethical Standards. The authors assert that all procedures contributing to this work comply with the ethical standards of the relevant national guidelines on human experimentation and with the Helsinki Declaration of 1975, as revised in 2008, and has been approved by the institutional committees (Institutional Review Board at the University of Iowa).

^{*}indicated reference category for regression model.

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