

Supplementary Material Table S1. Shows the details of pathogenic copy number variations in 38 CHD fetuses.

case	Cardiac and Extra Cardiac Phenotype	Chr	pCNVs	Type	Size (Mb)	Syndromes	Major genes included	Pattern of Inheritance	Outcomes
1	VSD, endocardial fibroelastosis, MCDK	1	849,466-5,747,184	del	4.8	1p36 deletion syndrome	<i>PRDM16</i> , <i>SKI</i>	/	TP, Autopsy ^a
2	TOF, FGR, limb abnormality	5	36,069,681-40,557,872	del	4.4	Cornelia de Lange	<i>NIPBL</i>	/	TP
3	VSD, spina bifida	6	163,732,938- 170,914,297	del	7.1	6q terminal deletion syndrome	<i>DLL1</i>	/	LTF
4	CoA, FGR	7	72,653,992-74,154,209	del	1.5	WBS	<i>ELN</i> , <i>LIMK1</i>	/	LTF
5	CAT, VSD, horseshoe kidney	8	160,101-7,120,108	del	6.9		<i>CSMD1</i> , <i>ANGPT2</i>	/	TP, Autopsy ^a
6	HRHS, VSD, callosal dysplasia, SUA	10	98,561,770-105,586,041	dup	7.0		<i>LBX1</i> , <i>BTRC</i> , <i>POLL</i>	/	TP, Autopsy ^{b1}
7	HLHS, hemivertebra	11	12,674,297-13,486,847	del	8.8		<i>FLI1</i>	/	TP, Autopsy ^a
8	Single ventricle, DORV, PS	11	81,354,834-106,214,708	del	24.8		<i>FZD4</i>	/	TP
9	Aortic stenosis, VSD, hydrocephalus	12	103,403,037- 133,777,562	dup	30.3	Noonan syndrome 1	<i>PTPN11</i> , <i>TBX5</i>	/	TP
10	VSD, CCAM	16	15,438,164-18,172,468	dup	2.7	16p13.11 recurrent microduplication	<i>MYH11</i>	/	LTF
11	TOF, hyperechogenic bowel, ventriculomegaly	17	16,736,262-20,417,235	del	3.6	SMS	<i>RAI1</i>	/	TP, Autopsy ^a
12	VSD, osteogenic dysplasia	17	14,077,970-15,482,833	del	1.4	HNPP	<i>PMP22</i>	/	LTF
13	PA, VSD, RAA, choroid plexus cyst	21	33,623,086-35,352,874	del	1.7	Tokita-Kim syndrome	<i>RUNX1</i> , <i>SON</i> , <i>SYNJ1</i>	/	TP

14	VSD, PS, tricuspid valve dysplasia	17	87,009–1,184,534	del	1.1	MDLS	<i>LIS1</i>	/	TP, Autopsy ^c
		22	18,919,942–21,440,514	del	2.5	22q11.2 DS	<i>TBX1</i>		
15	d-TGA, VSD	22	18,901,424-21,790,211	del	2.8	22q11.2 DS		/	TP
16	VSD, cor triatriatum, mitral valve dysplasia, SUA	22	18,984,187-21,927,646	dup	2.9	22q11.2 microduplication syndrome		/	TP, Autopsy ^a
17	TOF(APVS), VSD	22	18,916,842-21,798,907	del	2.8	22q11.2 DS		denovo	TP
18	VSD, aortic stenosis, hydramnios	22	18,636,749-21,800,471	del	3.1	22q11.2 DS		denovo	TP
19	TOF	22	18,644,790-21,465,659	del	2.8	22q11.2 DS		denovo	TP, Autopsy ^c
20	RAA, APLPA	22	16,888,899-20,725,309	del	3.8	22q11.2 DS		/	TP, Autopsy ^a
21	VSD, PS	22	18,636,749-21,800,471	del	3.1	22q11.2 DS		denovo	TP
22	VSD, CoA	22	18,648,855-21,800,471	del	3.1	22q11.2 DS		/	TP
23	DORV, VSD, PA	22	18,939,748-21,721,712	del	2.7	22q11.2 DS		denovo	TP, Autopsy ^c
24	RAA, VSD, left superior vena cava	22	19,004,071-21,800,471	del	2.7	22q11.2 DS		/	TP, Autopsy ^a
25	RAA, VSD	22	18,901,201-21,480,190	del	2.5	22q11.2 DS		/	TP
26	TOF	22	18,906,341-21,460,123	del	2.5	22q11.2 DS		/	TP, Autopsy ^a
27	TOF	22	18,978,201-21,926,534	del	2.9	22q11.2 DS		/	TP, Autopsy ^a
28	TOF, RAA, absence of ductus arteriosus	22	18,636,750-21,800,471	del	3.1	22q11.2 DS		/	TP, Autopsy ^a
29	VSD, CoA	22	18,648,856-21,800,471	del	3.1	22q11.2 DS		/	TP, Autopsy ^{b2}
30	TOF, RAA, thymus aplasia	22	18,648,856-21,058,888	del	2.4	22q11.2 DS		/	TP, Autopsy ^a
31	IAA, type A, aortic stenosis, ARSA	22	18,648,856-21,800,471	del	3.1	22q11.2 DS		/	TP, Autopsy ^c
32	TOF	22	18,882,825-21,796,237	del	3.1	22q11.2 DS		/	TP
33	TOF (PA-VSD)	22	18,927,431-21,427,431	del	2.5	22q11.2 DS		/	TP
34	Single ventricle, single atrium, CAT	22	46933489-51219152	del	4.2	PHMDS	<i>SHANK3</i>	denovo	TP

35	TAPVC, accessory ear, micrognathia	22	16050230-18679614	dup	2.6	CES		/	TP, Autopsy ^a
36	RAA	X	2,703,661-6,609,379	del	3.9		<i>ARSL</i>	denovo	Birth
37	TOF (PA-VSD), RAA, left superior vena cava, SUA	X	168,552-55,508,714	Del	55.3	LWD, Xq28 duplication	<i>SHOX</i> , <i>DMD</i> , <i>USP9X</i> , <i>GATA1</i> , <i>PIGA</i> <i>MECP2</i>	/	TP, Autopsy ^a
			55,509,386-155,233,098	dup	99.7				
38	VSD, holoprosencephaly, facial abnormality	X	134,743,723- 155,043,723	dup	20.3	Xq28 duplication	<i>MECP2</i>	/	TP

MCDK, multicystic dysplastic kidney; SUA, single umbilical artery; CoA, coarctation of aorta; CAT, common arterial trunk; CCAM, congenital cystic adenomatoid malformation; APVS, absent pulmonary valve syndrome; APLPA, anomalous origin of left pulmonary from the ascending aorta; ARSA, aberrant right subclavian artery; DS: deletion syndrome; del: deletion; dup: duplication; WBS: Williams-Beuren Syndrome; SMS: Smith-Magenis syndrome; HNPP: Hereditary Liability to Pressure Palsies; MDLS: Miller-Dieker lissencephaly syndrome; PHMDS: Phelan-McDermid syndrome; CES: Cat eye syndrome; LWD: Leri-Weill dyschondroostosis; LTF, lost to follow-up; TP, terminal pregnancy.

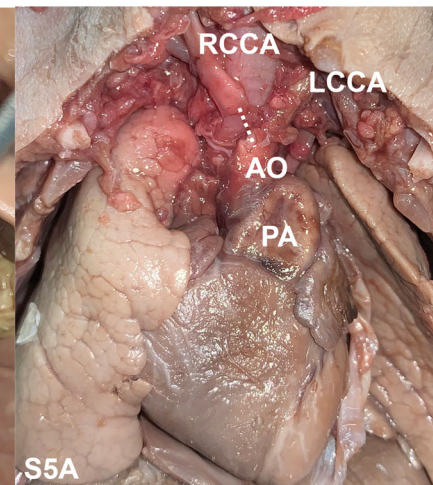
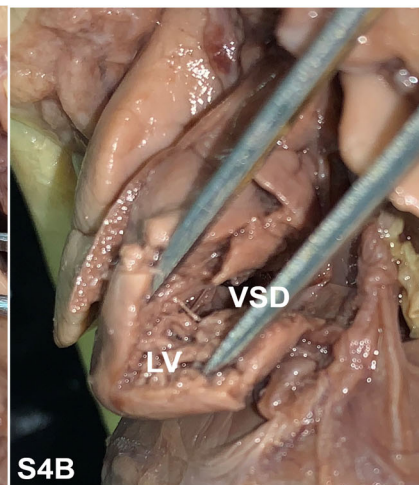
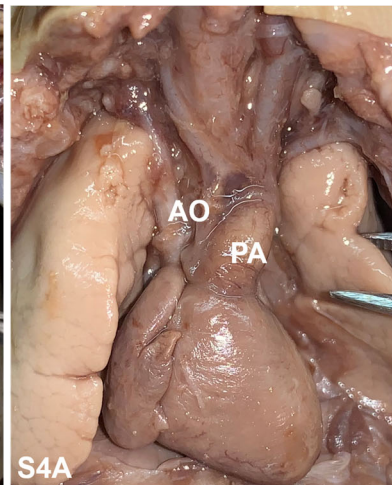
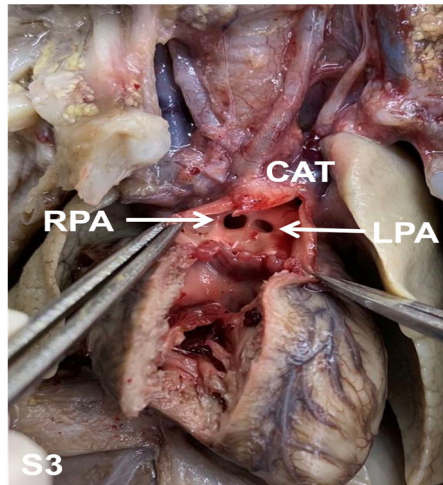
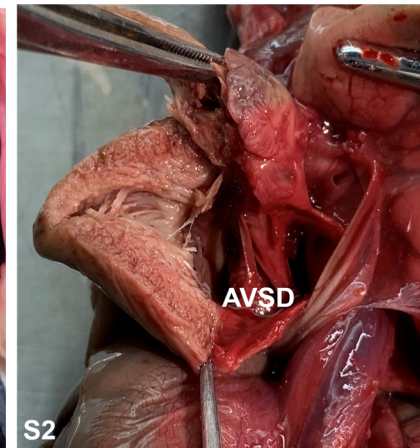
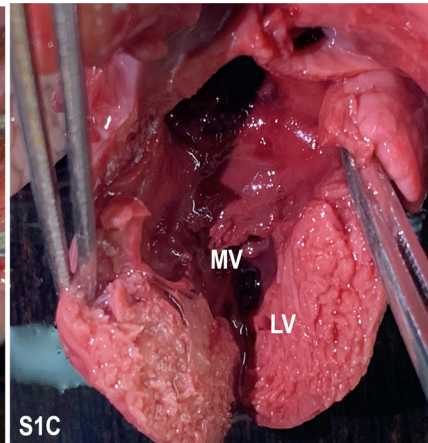
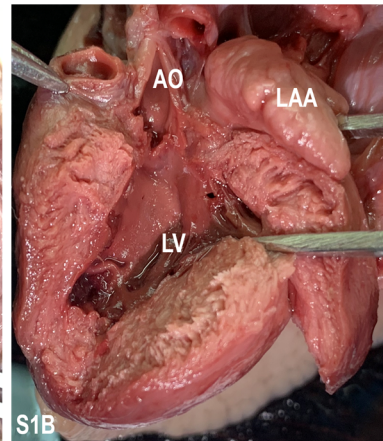
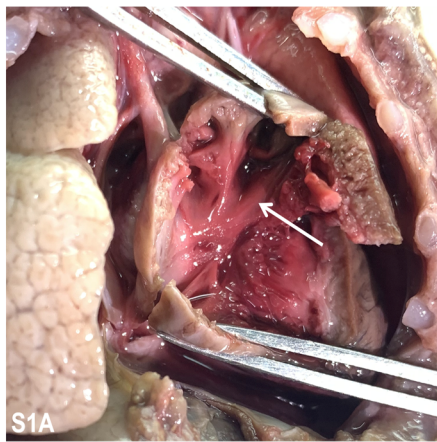
^a autopsy results are consistent with prenatal ultrasound.

^{b1} autopsy revealed repeated bladder and anal atresia, others were consistent with prenatal ultrasound.

^{b2} autopsy result is IAA, type B.

^c autopsy also found thymus hypoplasia/aplasia.

Bold font is the genes discussed in the article.



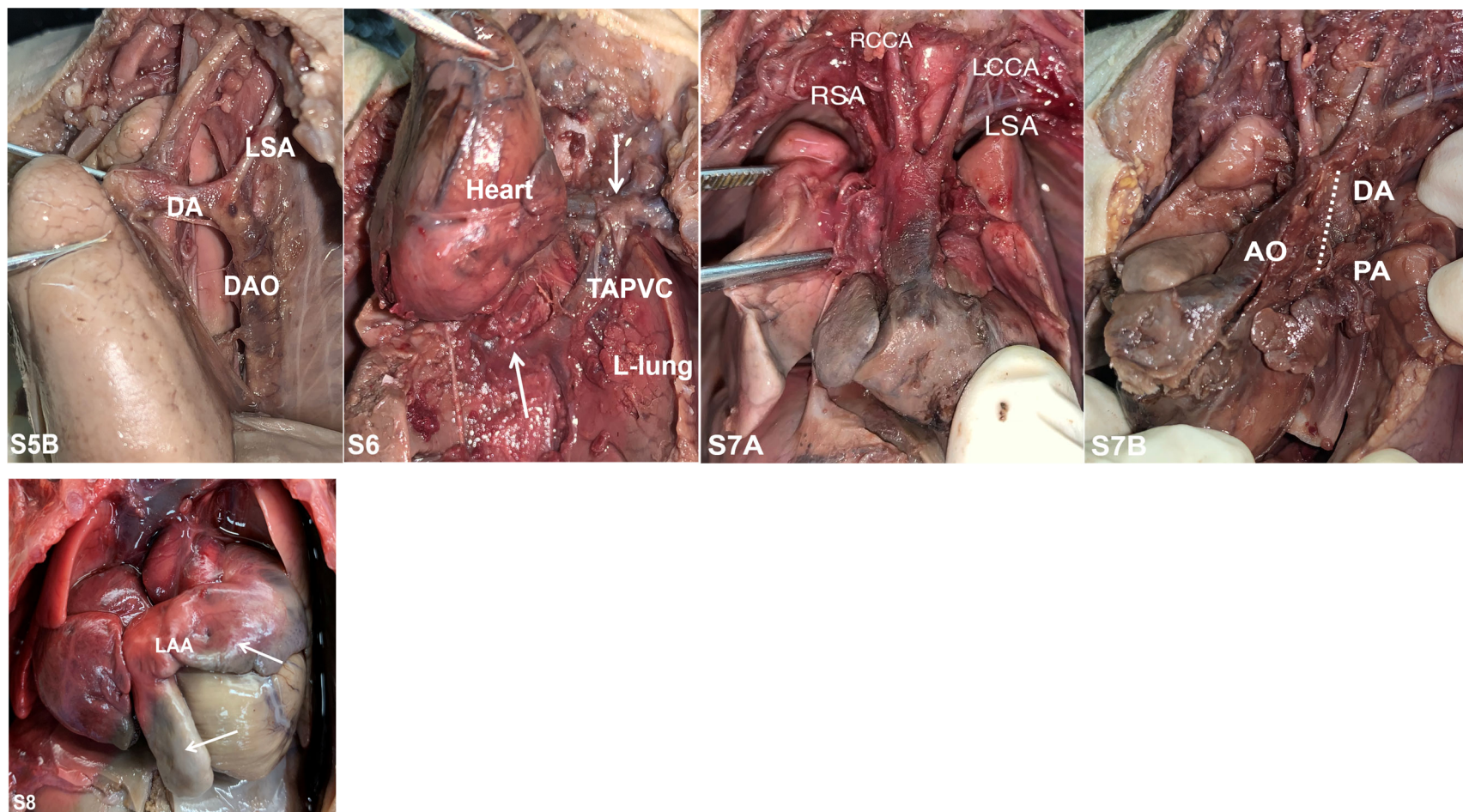


Figure S1-S8 shows the autopsy of CHD fetus.

Fig S1, mitral valve dysplasia; S1A, arrow shows no foramen ovale; S1B, LV and LAA thickening, stenosis aorta; S1C, mitral valve dysplasia, CNV result was VUS; Fig S2, complete type AVSD, trisomy 18; Fig S3 shows case 5 in table 2, autopsy results show CAT (type II); Fig S4 shows

case 7 in table 2; S4A, aorta dysplasia; S4B, LV is very small, VSD can be seen. Fig S5 shows case 29; S5A, only two blood vessels are sent out in ascending aorta; S5B, LSA is sent from descending Ao. Figure S6 shows case 35, and the arrows show common vena cava and vertical vein, respectively; Figure S7 shows case 37; S7A, RAA with mirror image branching; S7B, PA atresia, left and right pulmonary arteries supplied by DA. Dotted line, blood vessel accidentally broken; LAA, left atrial appendage; Ao, aorta; PA, pulmonary artery; LV, left ventricle; MV, mitral valve; CAT, common arterial trunk; LCCA, left common carotid artery; RCCA, right common carotid artery; LSA, subclavian artery; DA, ductus arteriosus. Fig S8, prenatal diagnosis of abnormal cardiac function, autopsy found a huge LAA, CNV result was VUS.