Human Heterotaxy Syndrome

– From Molecular Genetics to Clinical Features, Management, and Prognosis –

Isao Shiraishi, MD, PhD; Hajime Ichikawa, MD, PhD

Human heterotaxy syndrome is characterized by a wide variety of cardiac and extracardiac congenital malformations that are primarily induced by disorders of the left-right axis determination during early embryonic development. The cellular and molecular mechanisms of the left-right asymmetry have been extensively investigated in the past decade and the developmental mechanisms of the syndrome have been considerably elucidated. Medical and surgical management and treatment of heterotaxy syndrome have advanced as well. However, prognosis of the disease still remains unsatisfactory because the syndrome is often associated with a combination of complicated congenital heart diseases. Management of heterotaxy patients, particularly those who have undergone the Fontan procedure, is now one of the most important issues in pediatric and adult congenital heart disease clinics. In this review, we focus on the recent advances in knowledge of the genetic and molecular pathogenesis of heterotaxy syndrome, as well as its clinical features, management, and prognosis. (*Circ J* 2012; **76**: 2066–2075)

Key Words: Congenital heart disease; Genetics; Molecular biology; Surgery

Congenital heart diseases (CHD) affect approximately 0.75–0.9% of newborns and are the leading cause of death in neonates and infants.^{1,2} The clinical issues of CHD were once limited to pediatric cardiologists and pediatric cardiac surgeons, because the survival rate of patients was, in general, low. Because medical and surgical treatments of CHD have become extremely advanced, over 90% of CHD patients survive until adulthood.^{3,4} Consequently, the total number of adult patients with CHD currently exceeds that of children^{4,5} and CHD is now becoming an essential part of healthcare not only for pediatric but also for adult cardiologists.

Among the various kinds of CHD, heterotaxy syndrome is one of the most serious. This syndrome occurs in approximately 1 to 5,000–7,000 of live births with CHD.^{6,7} Heterotaxy syndrome is primarily induced by disorders of left-right axis determination during early embryonic development. Recently, the molecular mechanisms of the left-right axis determination have been extensively investigated in animal models including genetic engineering of mice.^{8–11} Briefly, an initial break in symmetry occurs at the primitive node as a leftward “nodal flow”, which is created by unidirectional rotation of nodal cell cilia. This flow provokes asymmetry signals that are transmitted toward the left lateral plate mesoderm (LPM), where downstream left-side specific growth and transcription factors, Nodal, Lefty2, and Pitx2c, are activated. As a result, Pitx2c and other undetermined factors regulate genetic programs in the left side of the body and create asymmetric organ morphogenesis. Human

genetics have also revealed several genes that are responsible for left-right laterality and heterotaxy syndrome, including ZIC3, NODAL, LEFTY2.^{11,12}

In the clinical field, precise diagnosis and surgical treatment of patients with heterotaxy syndrome have recently advanced as well. However, outcomes and the survival rate of patients are not satisfactory because this syndrome is often associated with combinations of serious CHDs.^{13,14} The mid- to long-term prognosis of heterotaxy patients who undergo the Fontan procedure for single ventricular physiology is now an important issue in pediatric and adult CHD clinics.^{15–18}

Morphological Characteristics of Heterotaxy Syndrome

Patients with heterotaxy syndrome are subdivided into “bilateral right-sided” (right isomerism) or “bilateral left-sided” (left isomerism) according to the characteristic morphology of the atrial appendages of the heart.^{13,14} However, there is a wide spectrum of pathology, with considerable overlap of the anatomical features. The morphological characteristics of heterotaxy syndrome are summarized in [Table](#).

Molecular and Cellular Mechanisms of Left-Right Determination

To elucidate the etiology of heterotaxy syndrome, it is neces-

Received July 22, 2012; accepted July 23, 2012; released online August 3, 2012

Department of Pediatric Cardiology (I.S.), Department of Pediatric Cardiac Surgery (H.I.), National Cerebral and Cardiovascular Center, Suita, Japan

Mailing address: Isao Shiraishi, MD, PhD, Department of Pediatric Cardiology, National Cerebral and Cardiovascular Center, 5-7-1 Fujishirodai, Suita 565-8565, Japan. E-mail: isao@hsp.ncvc.go.jp

ISSN-1346-9843 doi:10.1253/circj.CJ-12-0957

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Table. Morphological Characteristics of Heterotaxy Syndrome

Right isomerism	Left isomerism
Cardiovascular malformations	Cardiovascular malformations
Common atrium with bilateral right atrial appendages	Bilateral left atrial appendages
Mesocardia/dextrocardia	Complete/incomplete AV septal defect
Atrioventricular discordance	Unbalanced ventricles
Single right ventricle	Persistent left superior vena cava sometimes draining into the left atrium
AV septal defect	Interrupted hepatic portion of the inferior vena cava
Common AV connection associated with AV valve regurgitation	Partial anomalous pulmonary venous drainage
Double-outlet right ventricle	Hypoplastic sinus node (sick sinus syndrome)
Malposition of the great arteries	Single/paired AV nodes
Pulmonary stenosis or atresia	Interruption between AV node and His bundles (congenital AV block)
Total anomalous pulmonary venous drainage (with/without PVO)	Extracardiac malformations
Right aortic arch	Bilateral left-sided lungs and bronchi
Bilateral superior vena cava	Bilateral hyparterial bronchi
Bilateral sinus node	Polysplenia
Paired (anterior/posterior) AV nodes with sling formation	Midline liver
Extracardiac malformations/dysfunctions	Extrahepatic biliary atresia/hypoplasia
Bilateral right-sided lungs and bronchi	Extrahepatic portal vein atresia
Asplenia (susceptibility to <i>Streptococcus pneumoniae</i>)	
Symmetrical liver	
Right-sided stomach	
Malrotation of the intestine	
Bronchial cilia dysfunction	

AV, atrioventricular; PVO, pulmonary venous obstruction.

sary to understand the underlying molecular and cellular mechanisms of determination of the left-right body axis during early embryogenesis.^{8–12} This process initiates in the primitive node at E7.5 in mice and develops through the following pathways:^{19,20} (1) breaking of symmetry as a result of the leftward nodal flow generated by rotational movement of primary cilia in the node; (2) transmission of asymmetric signals to the LPM; (3) asymmetric expression of Nodal and Lefty2, which encodes a feedback inhibitor of Nodal signaling, in the LPM; and (4) situs-specific morphogenesis mediated by asymmetric expression of Pitx2, which encodes a transcription factor and is regulated by Nodal signaling.

Node Cell Monocilia Create Leftward “Nodal Flow” and Activate Asymmetry Signaling Around the Node

The determination of left-right asymmetry starts as leftward nodal flow generated by rotational movement of monocilia in the primitive node^{21,22} (Figures 1A–D). Monocilia consist of 2 different, highly sensitive subtypes: motile and immotile monocilia (Figure 1A).^{23–27} Motile cilia are predominantly located in the central region of the node and contain left-right dynein (*lrd*) in mice,²⁸ and the gene responsible for Kartagener's syndrome, *DNAH5*, in human²⁹ as a motor protein on microtubules (Figure 1B). In contrast, the immotile mechanosensory cilia expressing the calcium ion channel polycystin-2 (*Pkd2*) are located in the peripheral region of the node.^{23,30} Clockwise rotation of the motile cilia creates a unidirectional leftward flow at the surface of the node because the rotational axes of the cilia tilt in the caudal direction of the embryo (Figure 1C).^{31,32}

Although the nodal flow is the initial break in body sym-

metry, the precise mechanism of how this flow is perceived by nodal and perinodal cells remains uncertain. One hypothesis (chemosensory model) is that the nodal flow produces a gradient of left determinant particles (node vesicular parcels, NVPs) containing hedgehog proteins and retinoic acid secreted by node pit cells (Figure 1D).^{27,33} The secretion of NVPs is regulated by fibroblast growth factor 8.³³ These morphogens, in association with Nodal secreted in the node, activate downstream signaling of Nodal in the left-side perinodal cells. For instance, a transcription enhancer, ANE and Smad2/3, is activated by Nodal signaling and exhibits left dominant asymmetric expression in perinodal cells.²⁰ Consequently, the left-side dominant active Nodal in the node is translated into asymmetry in the left LPM.

An alternative hypothesis (mechanosensory model) is that the leftward nodal flow provokes an asymmetrical increase influx of Ca²⁺ ion in the sensory cilia cells through *PKD2*, a causative gene for human polycystic kidney disease.^{23,32} This Ca²⁺ influx is linked to the activation of Nodal in the left-side perinodal cells, which is consequently transferred to the left LPM (Figure 1D).

Experimental creation of the reverse nodal flow results in a reverse pattern of asymmetric gene expression.³⁴ Mutations in the left-right dynein (*Dnahc11*)³⁵ or kinesin (*KIF13*),^{36,37} which are the motor proteins of microtubules, give rise to randomization of the left-right asymmetry. Mutations in dynein assembly factor, *DNAAF3*, induce primary cilia dyskinesia and disturbance of left-right asymmetry in mice.³⁸ In embryos with mutations of *Noto*, a transcription factor regulating cilia formation, left-right asymmetry of internal organs and expression of laterality markers is randomized.³⁹ The Notch ligand *Dll1*-

Figure 1. Breaking of symmetry as a result of the leftward nodal flow generated by rotational movement of primary cilia in the node. Nodal cell cilia comprise motile and mechanosensory ones (**A**). Cilia consist of microtubules and dynein arms, which express left-right dynein (LRD) in mice (**B**). The rotational axes of the cilia tilt caudally in embryos, which creates unidirectional leftward “nodal flow” (**C**). Signaling mechanism of Nodal and related molecules in the nodal pit cells and surrounding node crown cells (**D**). NVP, nodal vesicular parcels.

mediated Notch signaling pathway plays a primary role in the establishment of left-right asymmetry by directly regulating expression of *Nodal* around the node.⁴⁰ A t-box transcription factor, *Tbx6*, is also involved in left-right determination through Notch-mediated *Nodal* signaling and formation of node cell monocilia.⁴¹ A member of the GLI superfamily zinc finger transcription factor, *Zic3*, is also involved in left-right patterning.^{42,43} The precise mechanism of the laterality defect remains unknown, but *Zic3* deficiency is associated with decreased expression of cardiac-specific genes, including *Nkx2.5*, *Tbx5* and *ANF*.⁴⁴ Epigenetic regulation with a chromatin modifier, *Baf60c*, has also been shown to control left-right asymmetry via Notch-mediated induction of *Nodal*.⁴⁵ In addition to activation of *Nodal* in the left side of the node, an endogenous nodal antagonist, *Cerl2*, is expressed around the right side of the node and is responsible for downregulation of *Nodal* in the right side of the embryo.^{20,46,47}

An experimental perturbation of the left-right determination process induced by maternal administration of retinoic acid at E6.5 to E7.0 is shown in **Figure 2**. **Figures 2A–D** and **Figures 2E–I** demonstrate dissected late fetal mice with right and left atrial isomerism, respectively. *Nodal* mRNA expression is limited in the left-side in a control mouse embryo (**Figure 2J**), and *Nodal* expression is randomly distributed in a RA-treated one (**Figure 2K**).

Asymmetry Signaling Transmits to the Left LPM and Upregulates Left Determinants Such as *Nodal*, *Lefty2*, and *Pitx2*

Transmission of *Nodal* to the left LPM followed by *Pitx2* ac-

tivation and the consequent heart morphogenesis in normal subjects is summarized in **Figure 3A**. Possible mechanisms of right/left isomerism and situs inversus from the viewpoint of nodal flow and the distribution of left determinants are shown in **Figures 3B,C**.

Activated *Nodal* in the left side of the primary node is transmitted toward the left LPM and midline. A member of the transforming growth factor (TGF)- β superfamily growth/differentiation factor 1 (GDF1) plays a role in the transportation of *Nodal*.^{48,49} *Nodal* activates other members of the TGF- β superfamily: *lefty1* in the left LPM and *lefty2* in the midline.⁵⁰ *Lefty1* and *lefty2* compete and antagonize nodal activity to restrict the extent and duration of nodal signaling.⁵¹ Bone morphogenic protein (BMP) also plays a role in negatively regulating *Nodal* expression. Chordin and noggin, which are BMP signaling antagonists, promote *Nodal* signaling around the node, while BMP signaling represses *Nodal* expression in the left LPM.^{52,53} The *Nodal* signaling pathway is mediated by an activin receptor complex composed of a dimer of the type I activin receptor, ActRIB, and a dimeric type II receptor, ActRII or ActRIIB.⁵⁴ Different from conventional Activin signaling, the *Nodal* pathway requires activities of membrane-bound EGF-CFC co-receptors, Cripto and Cryptic.⁵⁵ Loss-of-function mutations in the EGF-CFC genes are associated with left-right laterality defects.⁵⁶

Consequently, *Nodal* in the left LPM upregulates a transcription factor, *Pitx2*. *Pitx2* is a major laterality gene and plays a direct and pivotal role in asymmetric organogenesis of the heart and other visceral organs.⁵⁷ *Pitx2c*, a left-right asymmetric isoform, is expressed in the left LPM, outflow tract myocar-

Maternal RA administration at E6.5

Right atrial isomerism, dextrocardia, common AV valve, double outlet right ventricle, doubly committed VSD, right Ao arch

Maternal RA administration at E7.0

Left atrial isomerism, dextrocardia, common AV valve, double outlet right ventricle, subaortic VSD, pulmonary stenosis, IVC interruption, left Ao arch

Nodal
Control RA-treated **Left isomerism**
Polysplenia

Figure 2. Experimental perturbation of the left-right determination process induced by maternal administration of retinoic acid at E6.5 to E7.0. A late fetal mouse treated with retinoic acid at E6.5 shows right atrial isomerism (A–D). A late fetal mouse treated with retinoic acid at E7.0 shows left atrial isomerism with multiple spleen (E–I). Nodal mRNA expression is limited in the left side of a control mouse embryo (J), while Nodal is randomly distributed in a RA-treated one (K).

dium, left side of the brachial arch, artery, and the left atrium.^{58,59} The loss-of-function of *Pitx2c* induces severe cardiac malformations including right isomeric heart.⁵⁹ Recently, *Pitx2* was identified as a suspect for human atrial electric and structural remodeling arrhythmias, because, for example, *PITX2C* is significantly decreased in human patients with sustained atrial fibrillation.⁶⁰ *Cited2*, a transcriptional co-activator and is a negative regulator of HIF-1 α , is also involved in left-right patterning.^{61,62} In association with AP2, *Cited2* activates *Nodal*-mediated gene transcription such as *Nodal*, *Lefty2*, and *Pitx2* in the LPM.^{61,63}

Genes Associated With Human Heterotaxy Syndrome

Recent human and animal model studies have provided insights into the genetic and developmental etiology of the heterotaxy syndrome. In humans, genes that are associated with heterotaxy syndrome are *ZIC3*, *NODAL*, *CFC1*, *ACVR2B*, *LEFTY2*, *CITED2*, and *GDF1*.^{44,56,64–67}

Clinical Manifestation of the Heterotaxy Syndrome

Because the overall short- to mid-term prognosis of heterotaxy patients is not satisfactory, treatment is determined by the nature and severity of the cardiac and extracardiac abnormali-

ties.^{68,69} Factors that deteriorate the prognosis of heterotaxy patients have been described as complications with pulmonary venous obstruction, pulmonary arterial distortion, regurgitation of the atrioventricular valve, elevated pulmonary vascular resistance, and impaired ventricular function.⁷⁰

Right Isomerism

Neonates with right isomerism typically show a single atrium, single right ventricle, and a univentricular atrioventricular connection often associated with atrioventricular valve regurgitation. First stage palliation of such patients is control of pulmonary blood flow. If the pulmonary artery is atresic or severely stenotic, intravenous administration of prostaglandin E1 is necessary to open the left ductus, followed by the surgical operation of systemic pulmonary shunt (modified Blalock-Taussig shunt). If the pulmonary artery is not stenotic, pulmonary artery banding is necessary to protect the pulmonary vasculature until the Glenn and Fontan procedures can be performed. Both surgical interventions are generally performed approximately 2–4 weeks after birth, depending on the patient's body weight and complications. Pulmonary venous obstruction because of total anomalous pulmonary venous drainage should be precisely diagnosed and immediately repaired by surgical operation. However, the complication of pulmonary venous obstruction in patients with right isomerism is prob-

Figure 4. Surgical procedures of right isomerism (A-1-A-3) and left isomerism (B-1-B-3). Representative right atrial isomerism (A-1) after bilateral bidirectional Glenn shunt (A-2), and completion of the Fontan procedure (total cavopulmonary connection, TCPC) (A-3). Representative left atrial isomerism (B-1) after bilateral bidirectional Glenn anastomosis (total cavopulmonary shunt, TCPS) (B-2), and completion of the Fontan procedure (TCPC) (B-3).

long-term prognosis of the conventional procedure (atriopulmonary connection, APC) has proved to be unsatisfactory, characterized by enlargement of the atrium, intractable atrial tachyarrhythmias, and thromboembolism.

After successful completion of the TCPC, cyanosis disappears and the general condition of the patient improves. However, the number of right isomerism patients who have undergone successful Fontan procedure is approximately 50%, because right isomerism is often accompanied by a combination of severe and complicated CHDs.⁷⁷ Biventricular repair can be achieved in only a few patients with right isomerism.⁷⁸ The surgical procedures for right isomerism are illustrated in Figure 4A.

Left Isomerism

In contrast to right isomerism, the combination of the cardiac malformations in left isomerism is not so complicated. Left isomerism is typically associated with atrioventricular septal defect (AVSD), persistent left superior vena cava, interrupted hepatic portion of the inferior vena cava, and atrioventricular conduction disturbance.¹³ Patients with the complete type of

AVSD may undergo pulmonary artery banding in the neonatal period to protect the pulmonary vasculature. In left isomerism, the sinus node and atrioventricular node are usually hypoplastic and sinus bradycardia or complete atrioventricular block is a frequently accompaniment. In patients with severe bradycardia because of complete atrioventricular block from the fetal period, implantation of a pacemaker should be considered immediately after birth. Several months after pulmonary arterial banding, cardiac catheterization should be performed to ensure decreased pulmonary arterial pressure and resistance. Biventricular repair with a 2-patch method is usually performed for the complete type of AVSD with balanced ventricles.⁷⁹

AVSD with unbalanced ventricles is sometimes seen in patients with left isomerism, where either of the ventricles and the inflow tract is significantly hypoplastic, for instance in less than 30%.⁸⁰ In these cases, biventricular repair is not feasible. Bidirectional Glenn anastomosis followed by extracardiac shunt between the hepatic veins and pulmonary artery should be considered according to the criteria of the operation. If the patient's systemic outflow tract exhibits signs of potential stenosis, the double-barrel Damus-Kaye-Stansel operation is also

Figure 5. Possible mid- to long-term complications after completion of the Fontan procedure for heterotaxy patients.

performed in order to prevent ventricular outflow obstruction after the bidirectional Glenn shunt.

In either case of right or left isomerism, a hepatic to pulmonary connection is mandatory to avoid the development of pulmonary atriovenous fistula.⁸¹ Congenital absence of the portal vein is a rare but noteworthy complication of left isomerism.⁸² Pulmonary arterial hypertension may develop because of massive portosystemic shunt via the ductus venosus or other venous collaterals. Surgical procedures for left isomerism are illustrated in **Figure 4B**.

Long-Term Prognosis of Heterotaxy Patients

Although the medical and surgical treatments of heterotaxy syndrome are remarkably advanced, the long-term prognosis of the patients remains unsatisfactory. Right isomerism has been recognized as one of the worst forms of CHD, with the overall 5-year survival ranging from 30% to 74%. The results are better for left isomerism, with 5-year survival rates ranging between 65% and 84%, which is still considerably lower than survival for most other forms of CHD.⁸³ The main reason is that the nature of the Fontan single ventricle physiology is fundamentally imperfect and the Fontan circulation is likely to fail earlier than the normal biventricular counterpart. Management of the complications of the Fontan circulation and improvement of the long-term prognosis are becoming important issues in the field of pediatric and adult CHD.^{17,18} Cardiac complications include intractable atrial or ventricular arrhythmia, atrioventricular valve regurgitation, progressive heart failure, thromboembolism, pericardial effusion, and infectious endocarditis.^{70,84–86} Extracardiac complications include pleural effusion, plastic bronchitis, progressive cyanosis because of pulmonary arteriovenous shunts or systemic venous shunt to the left heart, liver dysfunction, liver cirrhosis, hepatocellular

carcinoma,⁸⁷ glucose intolerance and diabetes,⁸⁸ renal dysfunction, protein-losing enteropathy (PLE),^{85,86} ascites, menstrual disorders and subfertility in female patients,⁸⁹ systemic edema, and systemic thromboembolism including cerebral infarction.^{70,84,85} High risks of pregnancy and delivery, including deterioration of arrhythmias, progression of heart failure, coagulopathy, and miscarriage, have also been reported in women with Fontan circulation.⁹⁰ Representative long-term complications of the Fontan operation are illustrated in **Figure 5**.

PLE

PLE, one of the most severe manifestation of the failing Fontan circulation, occurs in 5–10% of the total postoperative cases.^{70,85,86,91} Chronic loss of serum proteins into the gastrointestinal tract results in systemic edema, ascites, pleural effusion, diarrhea, gastrointestinal bleeding, susceptibility to infections, and ultimately cachexia. The underlying mechanism of PLE remains uncertain. Elevated concentrations of inflammatory factors, such as tumor necrosis factor (TNF)- α or IFN- α , dilatation of intestinal lymphatic vessels, and widening of gap junctions between intestinal epithelial cells may be involved in the protein and fluid losses.⁹² Liver cirrhosis is often associated with the postoperative state of the Fontan circuits, in which elevated portal resistance and mesenteric pressure may be involved in the pathogenesis of PLE. Steroids, high-molecular-weight heparin,⁹³ sildenafil,⁹⁴ surgical interventions (eg, creation of a systemic-to-pulmonary venous atrial-level communication (fenestration)⁹⁵ or conversion of the Fontan circuit)⁸⁶ have been described as effective. Recently, the efficacy of oral controlled-released steroid, budesonide, for PLE has been reported.^{96,97} To date, however, cardiac transplantation is considered to be the only and complete resolution of PLE pathophysiology.^{98,99}

Arrhythmias

Reentrant atrial tachyarrhythmias are the most common in patients after the Fontan operation and are often associated with deterioration of hemodynamics, either causally or as a result.⁷⁰ Hemodynamic abnormalities such as valve regurgitation or outflow obstruction, if present, should be aggressively treated by surgery. Catheter ablation is generally challenging because the anatomical structure of the patient is complex and catheter approach is difficult after TCPC, even with the development of the contemporary 3-dimensional mapping system.^{100,101} The Maze procedure may be considered, particularly where conversion of the circuit is necessary.¹⁰²

Heart Failure

The initial feature of heart failure long after the Fontan procedure is generally worsening of ventricular relaxation and compliance.^{70,103,104} These abnormalities may be caused by exposure to hypoxia and volume/pressure overload preceding the Fontan procedure, repetitive surgical operations, and the hemodynamic disadvantages of the Fontan circuit. These changes are primarily progressive, potentially reducing ventricular filling volume, elevating pulmonary arterial pressure, and consequently leading to failure of the Fontan circuit. Late after the Fontan procedure, systolic dysfunction becomes apparent in a group of patients. Administration of angiotensin-converting enzyme inhibitors or β -blockade may be beneficial for particular patients, although the clinical evidence and cellular mechanisms remain to be elucidated.^{105,106}

Hepatic Dysfunction

Hepatic dysfunction, liver fibrosis, and cirrhosis are common

complications in patients long after the Fontan operation.^{70,107,108} Sinusoidal dilatation and sinusoidal fibrosis are marked in Fontan patients.^{107,108} The cause of the hepatic changes may be chronic elevation of central venous and portal venous pressures.⁸⁷ Significant portal fibrosis had occurred in most of those who died soon after the Fontan procedure, suggesting that hepatic fibrosis begins before the Fontan operation.¹⁰⁹ Recently, cases of hepatocellular carcinoma after the Fontan operation have been reported.¹¹⁰ Careful observation is necessary to detect the hepatic changes long after the Fontan operation.

Management of the Failing Fontan Patient

The patients who undergo APC or the lateral tunnel procedure are likely to be complicated with thromboembolism or intractable arrhythmias because of enlargement of the right atrium.^{70,86} Surgical intervention with conversion to TCPC is required before such complications become irreversible.^{102,111} A right atrial Maze procedure is simultaneously considered in patients with intractable atrial tachyarrhythmia.^{102,112}

Cardiac transplantation may be the only option for patients with severe heart failure, intractable arrhythmias, or recurrent PLE.^{98,99,113}

Future Directions

In the basic science field, embryonic development of left-right asymmetry has been uncovered by means of genetic engineering of mice. In addition, advanced human genetics has uncovered many of the genes responsible for human heterotaxy syndrome. By means of innovative technologies, such as next-generation sequencer or patient-based human inducible pluripotent stem cells, novel genes will be clarified and analyzed. In the clinical field, anatomical and physiological diagnosis from the fetal period, better clinical management after birth, tailor-made surgical operations, and systematic follow-up of the patients will improve their prognosis. Cell or tissue-based regeneration therapies could recover cardiac function of the failing Fontan patients. Multiple approaches including basic and clinical science are necessary to improve the prognosis and quality of life of heterotaxy patients.

Acknowledgments

The authors gratefully acknowledge Dr Masae Morishima for her excellent dissection, precise diagnosis, and in situ hybridization of the retinoic acid-treated mouse fetus. The authors also acknowledge members of pediatric cardiology and pediatric cardiac surgery at the National Cerebral and Cardiovascular Center for their clinical management and treatment of patients.

Disclosures

None.

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