

# Left Atrial Ligation Alters Intracardiac Flow Patterns and the Biomechanical Landscape in the Chick Embryo

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**Background:** Hypoplastic left heart syndrome (HLHS) is a major human congenital heart defect that results in single ventricle physiology and high mortality. Clinical data indicate that intracardiac blood flow patterns during cardiac morphogenesis are a significant etiology. We used the left atrial ligation (LAL) model in the chick embryo to test the hypothesis that LAL immediately alters intracardiac flow streams and the biomechanical environment, *preceding* morphologic and structural defects observed in HLHS. **Results:** Using fluorescent dye injections, we found that intracardiac flow patterns from the right common cardinal vein, right vitelline vein, and left vitelline vein were altered immediately following LAL. Furthermore, we quantified a significant ventral shift of the right common cardinal and right vitelline vein flow streams. We developed an *in silico* model of LAL, which revealed that wall shear stress was reduced at the left atrioventricular canal and left side of the common ventricle. **Conclusions:** Our results demonstrate that intracardiac flow patterns change immediately following LAL, supporting the role of hemodynamics in the progression of HLHS. Sites of reduced WSS revealed by computational modeling are commonly affected in HLHS, suggesting that changes in the biomechanical environment may lead to abnormal growth and remodeling of left heart structures. *Developmental Dynamics* 243:652–662, 2014. © 2013 Wiley Periodicals, Inc.

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## Introduction

Hypoplastic left heart syndrome (HLHS), is a rare but serious congenital heart defect, occurring in 1 of every 5,000 births (Go et al., 2013). The hallmarks of HLHS are an underdeveloped and nonfunctioning left ventricle and hypoplastic ascending and transverse aorta in association with stenosis or atresia of the mitral and/or aortic valves, and intra-uterine compensatory enlargement of right sided cardiac structures (Friedman et al., 1951; Noonan and Nadas, 1958). A genetic component for HLHS is supported by studies that examined heritability, which show that HLHS is linked to chromosomes 10q and 6q and genetically related to bicuspid aortic valve (Hinton et al., 2007, 2009), although the strength of this relationship is unknown (McBride et al., 2009). The genetic basis of HLHS is still largely undetermined and no transgenic animal models have recapitulated the human HLHS phenotype (Sedmera et al., 2005). Clinical innovations and scientific research has significantly improved the outlook for infants born with HLHS from a fatality rate of over 95% in 1980 to our current projections that 70% of infants born with HLHS are expected to survive to adulthood (Feinstein et al., 2012). These advances in diagnostic and treatment strategies are

remarkable; however, the pathogenesis of HLHS during embryonic and fetal life remains poorly understood. Fetal interventions have become available with the goal of positively impacting fetal and post-natal cardiac growth and remodeling.

For most of its history, HLHS has been classified as a “flow defect,” attributed to altered hemodynamic loading of the left heart structures, and fetal echocardiography has demonstrated that blood flow patterns have an important role in the development of HLHS (Grossfeld et al., 2009). An abnormally small or absent foramen ovale may be one key component, reducing flow to the left heart and impairing normal growth of left heart structures (Chin et al., 1990; Feit et al., 1991; Rychik et al., 1999), and one study has shown a correlation between diameter of the foramen ovale and relative right heart and/or left heart flow (Atkins et al., 1982). Obstructed inflow or outflow of the left ventricle due to valvular defects is more likely, however, as there is a strong correlation between the diameter of the left atrioventricular junction and left ventricle or aortic root (Sedmera et al., 2005). While the initial insult causing HLHS, genetic or structural, is unknown, the resulting hemodynamic alterations are significant and progressive. A typical diagnostic scenario in the clinic is detection of normal left heart dimensions with reduced function at mid-gestation, which is later followed by progressive involution of the left ventricle in the third trimester of pregnancy (McElhinney et al., 2010). One unifying hypothesis is that altered intracardiac flow patterns (ICFP) and altered mechanical loading

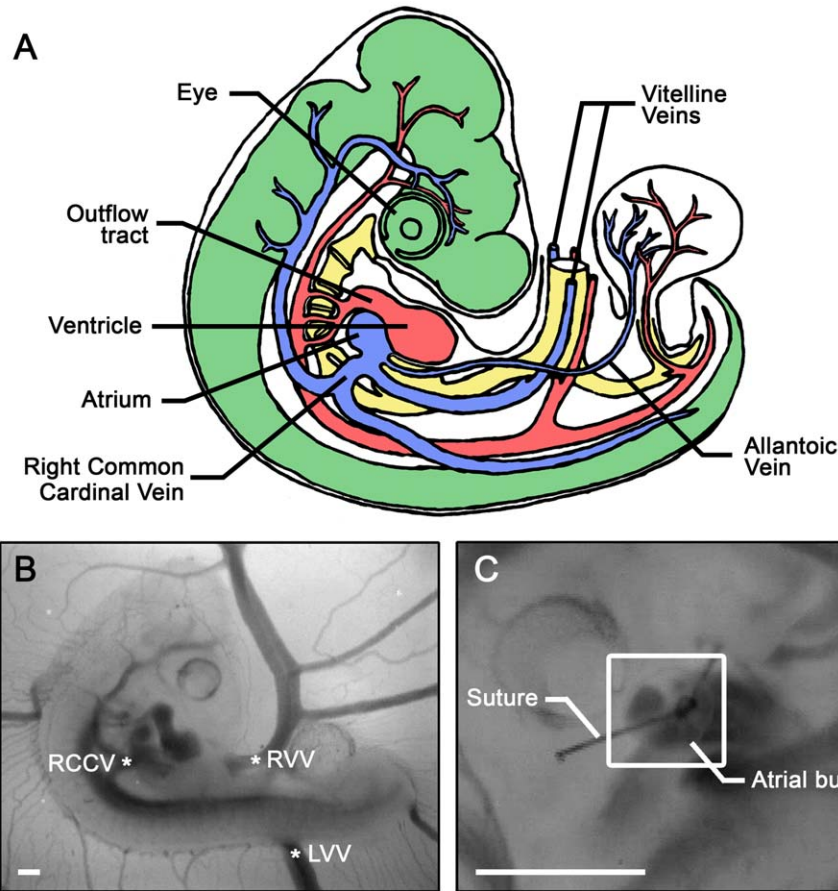
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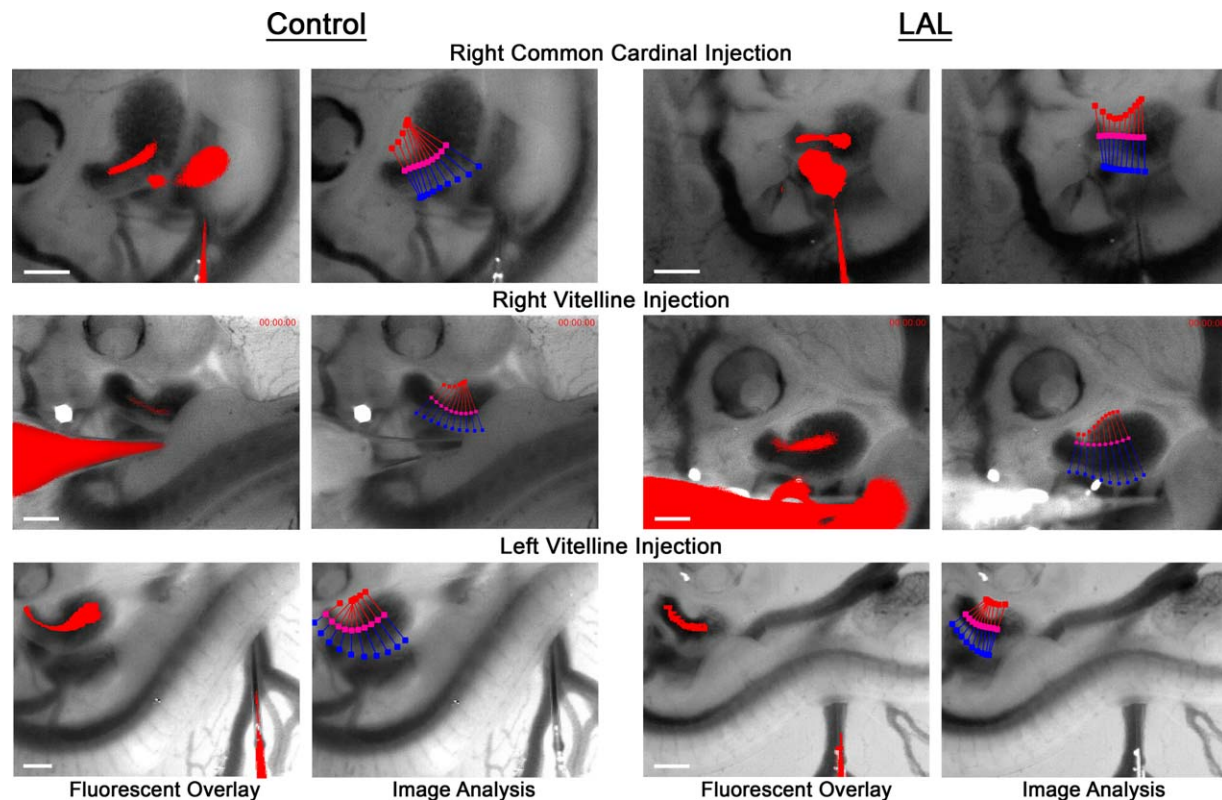
**Fig. 1.** The stage 21 chick embryo experimental model. **A:** Schematic of a stage 21 chick embryo, right lateral perspective. Vascular structures are labeled, where blue represents venous vessels and red arterial vessels. Green and yellow show neuronal and digestive structures, respectively. **B:** Stereomicroscope image of a normal stage 21 embryo, viewed from the right lateral perspective. Venous injection sites used in this study are labeled. The right common cardinal vein (RCCV) is the confluence of the right anterior and posterior cardinal veins, which drain blood from the embryo proper. The left and right vitelline veins (LVV, RVV) return blood from the extraembryonic vitelline bed. **C:** Stereomicroscope image of an LAL embryo, viewed from the left lateral perspective. The primitive left atrium is tied off with a 10-0 nylon suture. All scale bars = 500  $\mu\text{m}$ .

conditions result in left ventricular hypoplasia due to the lack of sufficient mechanical loading to stimulate cardiac growth and remodeling. This hypothesis has been applied as a rationale for fetal interventions, in which fetal balloon aortic valvuloplasty is performed to restore normal antegrade aortic flow and left ventricular loading conditions (McElhinney et al., 2010).

A large number of transgenic animal models have revealed key roles for signaling pathways and transcription factors in many of the events required for normal cardiovascular development, including outflow tract septation (Franz, 1989; Tallquist and Soriano, 2003), valve morphogenesis and remodeling (de la Pompa et al., 1998; Ranger et al., 1998; Dunker and Kriegelstein, 2002; Hurlstone et al., 2003), and myocardial contraction (Bartman et al., 2004). However, embryonic models that alter the mechanical environment in the setting of a normal genotype are limited. These studies are usually performed in avian embryos, mainly represented by vitelline vein ligation (Rychter and Lemez, 1965; Hogers et al., 1997), conotruncal banding (Clark et al., 1989), and left atrial ligation (Rychter and Lemez, 1958; Sedmera et al., 1999). Acquiring reliable, spatially resolved velocity measurements in the embryonic heart remains challenging, and many studies lack a quantitative analysis of the biomechanical environment after these perturbations. Newborn and juvenile models of

univentricular circulation have been developed in sheep to monitor cardiac function after repair of HLHS, although their long-term success and application is limited (Rodefeld et al., 2003; Myers et al., 2006). Further research using these existing mechanical models, and the development of novel models, is necessary to understand the role of hemodynamics and optimize fetal intervention strategies (Pekkan and Keller, 2013).

The chick embryo has been widely used as a model for vertebrate cardiovascular development (Pexieder, 1986; Martinsen, 2005). Left atrial ligation (LAL) in the chick embryo remains the only long-term prenatal animal model of HLHS (Rychter, 1962). In the LAL model, tying off the presumptive left atrium as a recoverable procedure disrupts inflow to the left side of the embryonic ventricle. Measured as early as one hour after ligation, cardiac output is transiently reduced, returning to normal levels by 32 hours post-LAL, while downstream cardiac pressures remain normal throughout (Lucitti et al., 2005). Signs of remodeling of the ventricular myocardium can be observed 2 days after ligation, including decreased myocardial volume, accelerated trabecular compaction, and delayed changes in transmural myofiber angle (Sedmera et al., 1999; Tobita et al., 2005). Circumferential and longitudinal strain increases in both ventricles after LAL, while the onset of preferential circumferential strain patterns in



**Fig. 2.** Representative images from the experimental fluorescent dye injections. Control embryos are on the left and LAL embryos are on the right. For each injection site, the overlay of the fluorescent and brightfield image is shown alongside the output of the image analysis to measure the distance between the flow stream (pink) and the ventral (red lines) and dorsal (blue lines) walls of the heart. All scale bars = 250  $\mu$ m.

the right ventricle are accelerated and the preferential longitudinal strain patterns in the left ventricle are abolished (Tobita and Keller, 2000). Cellular changes after LAL include reduced proliferation in the left ventricular compact layer and trabeculae, decreased expression of FGF-2 and PDGF-B throughout the heart, and an increased number of apoptotic cells in the right atrioventricular cushions (Sedmera et al., 2002), as well as increased microtubule density in the left ventricular compact layer (Schroder et al., 2002). Despite the limited number of studies focusing on vascular development compared with the ventricle, downstream vessels are also affected in LAL; flow distribution within the aortic arches is disrupted, and defects including aortic arch hypoplasia and interrupted aortic arch are observed as early as 32 hr after ligation (Hu et al., 2009).

These previous studies clearly demonstrate the morphologic, structural remodeling, and cellular consequences of LAL. However, to date there has not been a *quantitative* analysis of changes in ICFP that occur following LAL. While hemodynamic anomalies such as reduced left ventricular filling and mitral, tricuspid, and aortic valve regurgitation have been recorded, these were observed 4 days post-LAL (deAlmeida et al., 2007). The immediate effects of LAL (<1 hr after ligation) on flow patterns are therefore unknown. Furthermore, quantitative analysis of hemodynamic loading after LAL has not been performed. In this study, we test the hypothesis that ICFP are altered immediately following LAL. Validation of this hypothesis would demonstrate that changes in ICFP precede changes in ventricular morphology and structure, supporting the idea that flow patterns are a primary factor in the production of the HLHS phenotype after LAL.

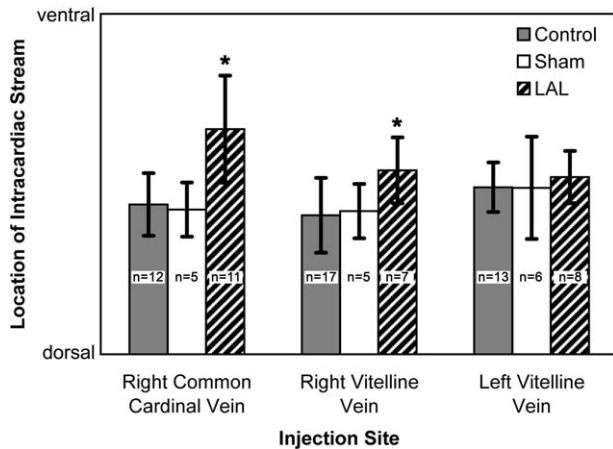
We applied a quantitative analysis of fluorescent dye microinjections to map the location of the intracardiac flow stream within the ventricle and outflow tract. We further developed a computational fluid dynamics (CFD) model of the normal and LAL heart loop to estimate changes in wall shear stress (WSS) that may occur due to the occlusion of the left atrium. Our results show that the position of the intracardiac flow stream is shifted significantly in two of the three venous injection sites and that WSS is reduced at the left atrioventricular canal and left ventricle. Thus, LAL immediately re-routes flow through the heart, likely causing reduced biomechanical loading and growth of left heart structures and leading to impaired remodeling and eventual HLHS.

## Results

### Intracardiac Flow Streams in Normal and LAL Embryos

We performed LAL at Hamburger-Hamilton stage 21 (Hamburger and Hamilton, 1951; Al Naieb et al., 2013). At stage 21, cardiac looping is largely complete, with only minor migration and rotation of the outflow tract progressing through stage 24 (Manner, 2000). The left and right atrial cavities have formed and ventricular trabeculation is present, although neither septum has fully formed (Martinsen, 2005). The dorsal and ventral atrioventricular canal cushions have started to form (De la Cruz et al., 1983), and four of the five primary outflow tract cushions are present (Qayyum et al., 2001). Blood enters the common atrium through several venous sites including the vitelline veins (VV), which drain the vitelline bed, the right and left common cardinal veins





**Fig. 3.** Results of the intracardiac stream measurements. The average location of the stream in control, sham, and LAL embryos is shown. Error bars represent the standard deviation. Numbers within each bar indicate the sample size. \*Indicates a statistically significant ( $P < 0.05$ ) difference vs. the control group.

(CCV), which return blood from the embryo proper, and the allantoic veins, which drain the allantois (Fig. 1). Blood is ejected through the common outflow tract and symmetric, bilaterally paired aortic arches. Most often three aortic arch pairs are present at stage 21, although a variety of patterns have been documented associated with flow and vessel caliber mismatch (Kowalski et al., 2013). The stage 21 embryonic heart rate is 155 beats per minute and cardiac output is  $1.28 \text{ mm}^3/\text{s}$  (Hu and Clark, 1989).

To visualize flow streams, we injected embryos with fluorescent dye at three different venous sites: (1) the right CCV (RCCV), (2) the right VV (RVV), and (3) the left VV (LVV) (Fig. 1). Each embryo was injected once; sham and LAL embryos were injected after a reincubation period lasting less than 1 hr. We computed a normalized intracardiac stream position from the recorded injections. Blood flow was laminar in all embryos. The heart continued to beat at a constant rate for a minimum of one minute after dye injection, allowing us to record multiple cardiac cycles. Low Reynolds number flow in the embryonic ventricle does not allow flow stream mixing. After removing the needle, blood hemorrhaged and the heart beat ceased rapidly afterward. Our fluorescent dye injections and image analysis allowed us to quantitatively compute the position of the intracardiac flow stream within the common ventricle and outflow tract. Results of our experimental dye injections are summarized in Figures 2 and 3, and fluorescent overlay movies of representative injections are provided in the Supplemental Material, which are available online. Our normalized stream position represents the distance from the dorsal wall of the embryonic heart, where 0 is exactly the dorsal wall and 1 is exactly the ventral wall. For control embryos, we found that dye injected at the RCCV took a slightly dorsal route, with an average stream location of  $0.44 \pm 0.09$  ( $n = 12$ ). The stream from the RVV was the most dorsal, with a position of  $0.41 \pm 0.11$  ( $n = 17$ ). The location of the LVV stream was  $0.49 \pm 0.07$  ( $n = 13$ ), directly through the midline of the heart. Sham embryos displayed similar ICFP, with no significant differences vs. the control group.

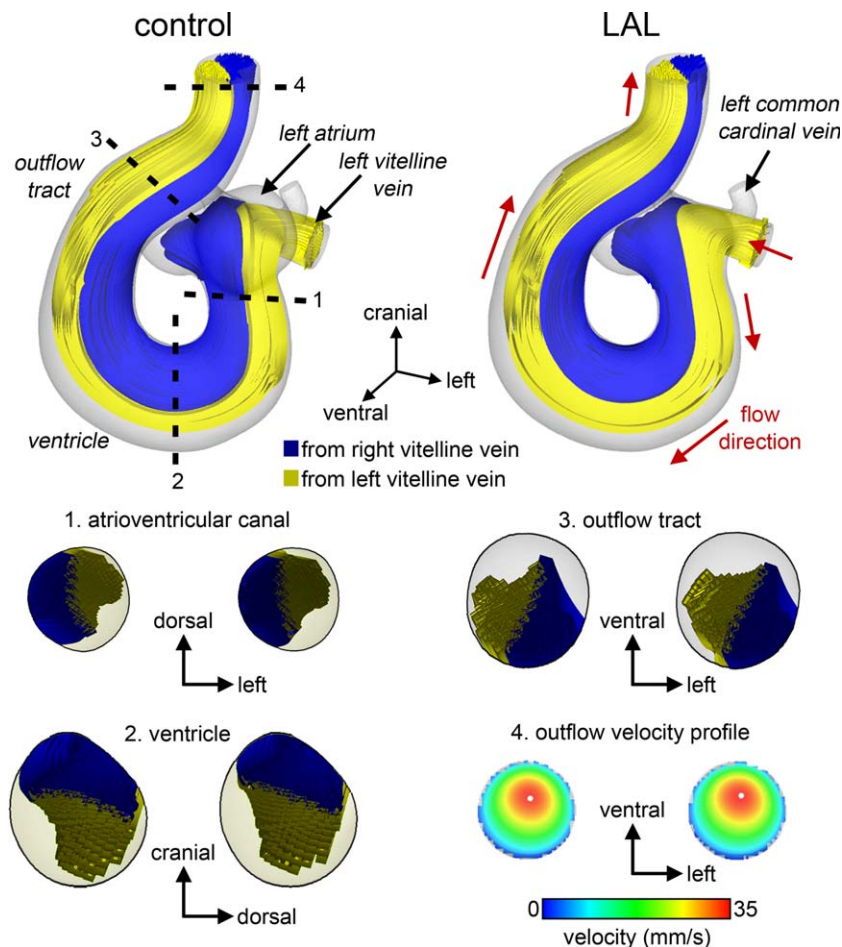
The response of LAL embryos to insertion of the micro needle was similar to control embryos with no observable effect on hemodynamics or heart rate. For LAL embryos, our quantitative

analysis revealed that the RCCV stream changed significantly ( $P < 0.05$ ), now taking a ventral course through the ventricle and outflow tract, with an average location of  $0.66 \pm 0.16$  ( $n = 11$ ). The RVV stream also shifted toward the ventral side of the heart ( $P < 0.05$ ), positioned at  $0.54 \pm 0.10$  ( $n = 7$ ). The intracardiac path of the LVV remained similar at  $0.52 \pm 0.08$  ( $n = 8$ ,  $P > 0.05$ ). In the LAL embryos, the RCCV and RVV streams were ventral to the LVV stream, the reverse of the control ICFP. Of interest, all flow patterns shifted ventrally, although only the RCCV and RVV were significantly changed compared with the controls. We also observed some retrograde flow in the LAL embryos, where dye injected at the RVV and LVV would flow into the cardinal veins. These results demonstrate that LAL significantly alters ICFP in the embryo, and that this change occurs immediately following the intervention.

### In Silico LAL Model

We performed three-dimensional (3D) CFD simulations of steady flow through control and LAL models of the stage 21 heart loop. Both models were created in the computer using our in-house surgical planning suite (Pekkan et al., 2008; Dur et al., 2011). Blood flow was modeled using our in-house immersed boundary cardiovascular flow solver, which has previously been validated for models of the stage 11, 13, and 18 heart loop, comparing accurately with the commercial second-order unstructured grid flow solver Fluent (version 6.3.26, ANSYS Inc., Canonsburg, PA) and experimental microfluidic models of the embryonic heart tube (Supp. Fig. S1; Pekkan et al., 2009). Mean total cardiac output was  $1.28 \text{ mm}^3/\text{s}$ , equivalent to stage 21 Doppler flow measurements (Yoshigi et al., 2000), and rigid, no-slip walls were assumed for simplicity, simulating mean quasi-steady flow conditions in the heart. We computed color-coded flow streamlines originating from the right and left vitelline veins to observe ICFP in our CFD models (Fig. 4). There were some subtle differences, including an overall ventral shift of the flow streams in the LAL model (Fig. 4). This movement of ICFP toward the ventral margin of the heart is similar to our experimental observation. Flow from the RVV seemed to follow the inner curvature of the heart more tightly in the LAL model as well. However, the variation between the control and LAL CFD models is far less pronounced than the experimental results, and we did not observe the reversal of RVV and LVV stream positions (Fig. 4), prompting future improvements in computational model. We computed the velocity profile at the outflow tract and found that it was slightly more skewed in the ventral and left directions in the LAL vs. control model.

To examine changes in biomechanical loading that occur due to removal of the left atrium, we calculated WSS from our CFD results (Fig. 5). The maximum WSS levels in the heart were 3.5 Pa, found at the CCV and VV inlets and the ventral side of the outflow tract outlet, consistent with WSS values derived from particle image velocimetry in similar stage chick embryos (Poelma et al., 2010). Compared with the control model, WSS in the LAL model was reduced at the left side of the atrioventricular canal and left side of the common ventricle (Fig. 5). WSS was also slightly increased at the ventral surface of the distal outflow tract. These simplified, static CFD models show that redistribution of the left atrial volume is sufficient to alter ICFP and WSS loading, while the disparity with the experimental results supports the



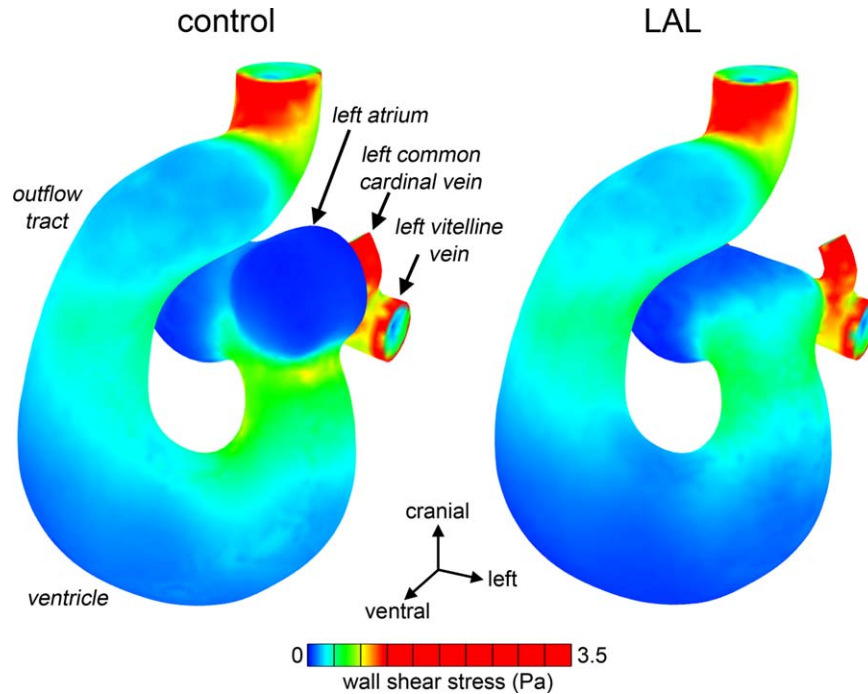
**Fig. 4.** Intracardiac flow patterns in CFD models of control and in silico LAL embryonic cardiac geometries. The control model is on the left and the LAL model is on the right. In the upper figures, color coded streamlines are shown, with blue representing flow emanating from the RVV and yellow depicting flow from the LVV. Red arrows give the forward flow direction. The bottom panels show cross-sections at each of the sites labeled in the control flow stream figure. Control cross-sections are on the left and LAL on the right. The cross-sections reveal a slight overall ventral shift of flow paths in the LAL model. The velocity profile at the outflow tract is shown for cross-section 4, where a slight ventral shift of the peak flow location is observed in the LAL model.

additional roles of the endocardial cushions and dynamic ventricular suction during relaxation in determining ICFP.

## Discussion

Qualitative description of normal ICFP in the chick embryo has been taken up by several authors, giving rise to some contradictions (Bremer, 1932; Rychter and Lemez, 1965; Jaffee, 1967; Yoshida et al., 1983; Hogers et al., 1995). Early studies attempted to follow red blood cells as they coursed through the heart, and led to the widely perceived notion that two separate streams spiraled around each other through the outflow tract (Bremer, 1932; Jaffee, 1967). The presence of these streams was thought to shape the spiral aortopulmonary septum, with anomalous spiraling proposed as a major cause of septation defects (Jaffee, 1965). This notion was challenged when researchers began injecting dyes or India ink and did not observe spiraling outflow streams (Yoshida et al., 1983; Hogers et al., 1995). In their comprehensive study of chick embryos between stages 14 and 22, Yoshida et al. observed two types of intracardiac streams which depended on both injection site and developmen-

tal age (Yoshida et al., 1983). These streams did not spiral around one another, but rather followed a dorsal or ventral path through the ventricle and outflow tract. These patterns were later confirmed by Hogers et al. using India ink injections of embryos between stages 12 and 17 (Hogers et al., 1995). Their study revealed distinct ICFP for blood emanating from different yolk sac regions, further demonstrating the significance of venous return sites and the presence of multiple simultaneous ICFP. Our ICFP observed in control stage 21 embryos matched those of Yoshida et al. (1983), with blood from the RVV taking a more dorsal path compared with blood from the LVV. No study has examined either CCV. In no cases of these embryonic did we observe evidence of spiraling streams through the outflow tract. The work of Yoshida et al. also suggested that blood from the RVV and LVV perfused only the ipsilateral aortic arches at stage 21 (Yoshida et al., 1983). This pattern was contradicted by other studies, which showed that the distribution through the aortic arches was not a simple left and right current (Rychter and Lemez, 1965; Hogers et al., 1995; Hu et al., 2009). While we did not examine ICFP beyond the outflow tract in this study, our previous investigations dedicated to the aortic arches



**Fig. 5.** WSS computed from CFD models of control and *in silico* LAL embryonic cardiac geometries. The control model is on the left and the LAL model is on the right. Reduced WSS can be seen at the left side of the atrioventricular canal and left region of the common ventricle. Increased WSS is located at the ventral side of the distal outflow tract. To display a broader range of WSS, a peak WSS of 1.0 Pa is shown.

corresponds with a complex perfusion pattern (Wang et al., 2009; Kowalski et al., 2013).

Yoshida et al. observed nearly the exact reverse ICFP in embryos stage 17–18: blood from the RVV took a more ventral route compared with blood from the LVV, and the two streams perfused the contralateral aortic arches (Yoshida et al., 1983). These patterns transitioned at around stage 19, and the authors attributed the dramatic reversal to the changes in the vitelline venous system. Our data show that in LAL stage 21 embryos, blood from the RVV courses more ventrally compared with blood from the LVV, similar to the stage 17–18 patterns described by Yoshida et al. (Fig. 2). It may be that obstruction of the left atrium prevents the change in RVV and LVV patterns or returns ICFP to their earlier state. Before stage 18, the atrial cavities are small and hardly distinguishable from the heart tube. Between stages 18 and 21, however, the atria significantly increase in size, particularly the left atrium, which becomes larger than the right (Manner, 2000; Kim et al., 2011; van den Berg and Moorman, 2011; Yalcin et al., 2011). This expansion of atrial volume and the atrial contribution to ventricular filling may relate to the change in ICFP, which occurs during the same interval. As the expansion of the left atrium is more pronounced than the right, its ligation may return the cardiac geometry to resemble stage 17, causing ICFP to revert. Ventricular trabeculation may also have a role in the changing flow patterns at stage 19. Beginning at stage 16/17, trabeculation progresses rapidly during this period, and is largely responsible for increases in myocardial mass (Sedmera et al., 2000). Another key morphologic event occurring during the period between stages 17 and 25 involves substantial growth and remodeling of the endocardial cushions lining the atrioventricular canal and outflow tract, brought on by the invasion of cells that begin to undergo epithelial to mesenchymal transfor-

mations at stage 17 (Noden, 1991; Butcher et al., 2007). LAL may interfere with these events by preventing growth of the left atrium, limiting the expansion of the atrioventricular canal, or displacing the endocardial cushions.

Our numerical control vs. LAL CFD models confirmed the variation in the location of the flow streams originating from the RVV and LVV, but these are less pronounced than the *in vivo* injections (Fig. 4). A major difference is that we do not see the dramatic ventral shift of the RVV flow stream after LAL. Presently, our CFD model did not incorporate the endocardial cushions or ventricular trabeculae and applied rigid, no-slip walls and steady, unidirectional flow, ignoring the AV synchronous contractions and pulsatile flow environment. Furthermore, the external boundary conditions specified at the vessel inlet and outlets, as well as the neglected wall movement, may suppress the *in vivo* observed internal flow differences. These limitations influence the numerical model of ICFP and mask key elements of the biomechanical environment such as oscillatory WSS. Effects of flow pulsatility may be less important, illustrated in our previous pulsatile CFD models of the stage 18 heart loop, which revealed that Womersley numbers were low (0.44), suggesting that the effects of pulsatile flow are small (Pekkan et al., 2009). The assumptions described above were necessary to reduce computational time and produce a solvable model. Differences between the computational model and experiment may indicate that those aspects absent from the model, such as cardiac contractions, endocardial cushions, and ventricular trabeculae, are vital to establishing ICFP. Even without these features, however, our models show that simply removing the left atrium can alter the course of blood and WSS levels in the heart realistically. In particular, WSS was reduced at the left side of the common ventricle and left wall of the atrioventricular canal and slightly increased on the ventral



surface of the outflow tract (Fig. 5). WSS differed by 0.5 Pa at these sites, which is above the known threshold to initiate an endothelial response (Egorova et al., 2011). These areas are commonly affected in HLHS, suggesting that LAL may redirect flow to unload the presumptive left ventricle and its inflow. The velocity profile change in the *in silico* LAL model may indicate a reason for altered aortic arch perfusion and defects in LAL embryos (Hu et al., 2009). However, our recent simulations of aortic arch flow using various velocity profile shapes at the outflow tract do not show significant changes in flow distribution or WSS, again demonstrating the importance of the dynamic contractions (Kowalski et al., 2013).

Morphogenesis of the veno-atrial region is ongoing at stage 21, and the veno-atrial connections may differ between embryos (Manner and Merkel, 2007; van den Berg and Moorman, 2011). Our model geometry includes proximal separation of the right and left vitelline veins, a centrally located venous sinus, and symmetric cardinal veins, all of which are characteristic of the early state of veno-atrial morphogenesis (Fig. 4). By stage 24, however, the proximal vitelline veins have fused, the sinus connection is limited to the right atrium, and the left cardinal vein is elongated. As this transformation likely influences ICFP, we modified our model geometry to correspond with this later state venous anatomy and repeated our control and LAL flow models (Supp. Fig. S2). As with our original geometry, flow patterns in the LAL model were not substantially different from the control. This later state model, therefore, reinforces our previous reasoning that those features not incorporated, *i.e.* cardiac contractions, endocardial cushions, and ventricular trabeculae, have a role in directing intracardiac blood flow. We did find, however, that the outflow velocity profile was more skewed in the ventral direction after changing the venous morphology (Fig. 4 and Supp. Fig. S2). These two models suggest that morphogenesis of the sinuatrial region can influence ICFP and may provide an explanation for the changes in flow patterns that occur in normal embryos from stages 18 to 24 (Yoshida et al., 1983).

Static CFD models of flow in the human embryonic heart show the presence of multiple simultaneous intracardiac streams, with patterns similar to those observed in the chick embryo, demonstrating the relevance of these animal models to human development (DeGroff et al., 2003). While success has been made in reconstructing static embryonic cardiac geometries from polymeric casting (Wang et al., 2009; Yalcin et al., 2011), there is no existing model of the dynamic embryonic heart. Recent techniques have simulated blood flow in the contractile outflow tract (Liu et al., 2011), although this work is in nascent stages and has yet to incorporate the upstream heart and downstream vessels. Validation of such models will require information of 3D velocity profile shapes *in vivo*. We have recently explored a combined particle image velocimetry and optical coherence tomography technique to acquire velocity measurements noninvasively, which can be applied to future studies (Chen et al., 2012).

Numerous studies using chick (Gessner, 1966; Hogers et al., 1997, 1999; Reckova et al., 2003; Lucitti et al., 2006) and zebrafish (Hove et al., 2003; Chen et al., 2011; Corti et al., 2011) embryos support the role of hemodynamics in modifying cardiovascular growth, morphogenesis, and remodeling. It is well established that WSS is a key environmental factor affecting cardiovascular development in the embryo and adaptation in the adult (Rodbard, 1975; Langille and O'Donnell, 1986; Girerd et al., 1996; Bayer et al., 1999; Culver and Dickinson, 2010). Using an

RVV ligation model, Hogers et al. found that ICFP emanating from four yolk sac regions followed altered routes through the heart and outflow tract, leading to defects such as small to large ventricular septal defects, bicuspid aortic valve, and aortic arch anomalies (Hogers et al., 1997). This ligation model was further shown to cause spatial and temporal changes in the transcriptional profile of genes encoding Krüppel-like factor (increased), endothelin-1 (decreased), and endothelial nitric oxide synthase (decreased), suggesting increased WSS (Groenendijk et al., 2005). The increased WSS in the outflow tract was also associated with higher Tgf $\beta$ /Alk5 signaling, a phenomenon which was sensitive to the level of WSS (Egorova et al., 2011). This mechanosensitivity of endocardial cells, coupled with the changed ICFP after LAL, likely results in abnormal morphogenesis and remodeling of the atrioventricular and outflow tract cushions, leading to valve defects normally present in HLHS. Although our technique does not capture total flow through the atrioventricular canal, the quantitative measurements through the outflow tract show a significant ventral shift of the RVV and RCCV streams (Fig. 3). In the RVV ligation model of Hogers et al., flow streams from two of the four yolk sac regions saw a similar dorsal to ventral shift (Hogers et al., 1997). It is interesting that the outlet of the left ventricle is dorsal to that of the right ventricle in the mature chick. The lack of a dominant dorsal stream after LAL may cause reduced WSS on the dorsal cushions present at stage 21, impairing the formation of the left ventricle outflow tract and aortic valve.

Our experiments support the hypothesis that LAL immediately alters ICFP in the chick embryo. We quantitatively showed a significant ventral shift in both the RVV and RCCV flow streams. While the LVV flow stream did not change significantly, the LVV and RVV streams switched relative positions in LAL compared with control hearts, resembling earlier stage 18 patterns (Yoshida et al., 1983). Our static CFD model of the embryonic heart loop further revealed that removing the left atrium volume was sufficient to alter WSS at critical locations within the left heart. These results suggest that changes in ICFP following LAL precede structural defects such as bicuspid aortic valve and the characteristic undersized left ventricle. The concordance between control and sham embryos suggests that the manipulations required to perform LAL do not cause a change in flow patterns. However, the ligation itself may cause structural damage to the heart, which can affect its further development. While we take great care and reject any embryos that show bleeding after LAL, we cannot rule out this trauma as a cause for HLHS defects. Therefore, we conclude that the change in ICFP occurs early in development of HLHS after LAL, but may not be the only or initial insult.

## Experimental Procedures

### Experimental Measurement of Intracardiac Stream Position

#### *Embryo preparation and injection*

Fertile white Leghorn chicken eggs were incubated blunt end up in a forced draft incubator at 37°C and 50% humidity for 84 hr to Hamburger-Hamilton stage 21 (Hamburger and Hamilton, 1951; Al Naieb et al., 2013). To access the developing embryo, we gently cracked the eggshell above the air sac and peeled back the shell and membrane to form a 1-cm<sup>2</sup> window. We then removed

the overlying extraembryonic membrane using microforceps (Dumont M5, WPI, Sarasota, FL). Limb size, spinal curvature, and the presence of body tissues were used to stage embryos, and any that were dysmorphic, such as lacking retina, displaying axial twisting, or left side up were excluded. To perform LAL, we positioned embryos *in ovo* under a stereomicroscope (Leica M165F, Leica Microsystems GmbH, Germany), dissected the amniotic membrane, and flipped the embryo to reveal the presumptive left atrium. We used microforceps to make a slit-like opening in the thoracic wall above the atrium and tightened an overhand knot formed from 10-0 nylon suture around the primitive left atrium, decreasing its effective volume (Fig. 1) (Tobita et al., 2005). We repositioned the embryo to its original right-side up orientation, sealed the window with Parafilm (Pechiney Plastic Packaging, Menasha, WI), and reincubated for 1 hr to restore normal heart rhythms. Control embryos had only the overlying membrane removed and were injected immediately with no reincubation. Sham embryos had a slit made in the thoracic wall above the left atrium and were reincubated for one hour.

To inject embryos, we pulled needles from 1.0 mm inner diameter glass capillary pipettes to 15  $\mu\text{m}$  inner diameter and a 30° bevel (PC-10 Puller, EG-44 Microgrinder, Narishige Inc., Japan). Needles were connected by means of polyethylene tubing (PE100, Braintree Scientific, Braintree, MA) to a 10  $\mu\text{l}$  glass Hamilton gastight syringe (1701LT, Fisher Scientific, Waltham, MA). The syringe was mounted on a three-axis mechanical micromanipulator (Leica Microsystems GmbH, Germany). We primed the syringe, tubing, and needle with chick ringer's solution and then drew up 1  $\mu\text{l}$  of Cy5-diethyl fluorescent dye diluted 1:1 in phosphate buffered saline (excitation wavelength 626 nm, emission wavelength 666 nm). Cy5 dyes were prepared at the Molecular Biosensor Imaging Center at CMU (Mujumdar et al., 1993). We injected embryos at three different venous sites: (1) RCCV, (2) RVV, and (3) LVV, with all embryos viewed in the right side up orientation (Fig. 1). Each embryo was placed under the stereomicroscope and the yolk sac membrane (RVV, LVV sites) or amnion (RCCV site) was removed above the injection site. The needle was inserted into the selected vein and a brightfield movie was recorded using a low light monochrome digital camera (Leica DFC 350FX, Leica Microsystems, GmbH, Germany). We then injected 0.3  $\mu\text{l}$  of dye, visualized using an external fluorescent light source (Leica EL6000, Leica Microsystems, GmbH, Germany) and recorded the flow streams. This volume of injected dye has been shown to minimally affect normal hemodynamics, resulting in less than a 10% change in stroke volume and mean arterial pressure (Wagman et al., 1990). A total of 84 embryos were injected: 28 RCCV (12 control, 5 sham, 11 LAL), 29 RVV (17 control, 5 sham, 7 LAL), and 27 LVV (13 control, 6 sham, 8 LAL). Movies of each injection site for representative control and LAL embryos are provided in the . All injections were performed *in ovo* and only one site was injected per embryo.

#### Quantitative measurement of the intracardiac flow stream

We applied a quantitative approach to determine the location of the intracardiac streams. End-diastolic still images ( $696 \times 520$  pixels) were extracted from the recorded brightfield movies and used to manually identify the ventral and dorsal boundaries of the ventricle and outflow tract. The manually selected points were used to form polynomial expressions defining the shape of the ventral and dorsal margins, giving two continuous functions.

We used the horizontal position as the independent variable, and rotated images, if necessary, so that the long axis of the heart was approximately horizontal. We then extracted still frames from the fluorescent recordings of the injections, which showed the flow stream as it passed through the ventricle and outflow tract. The centerline of the flow stream was manually identified and used to form a third polynomial expression. Any rotations applied to the brightfield image were applied exactly to the fluorescent image. For each point along the flow stream, we were then able to mathematically calculate the two points of intersection between the line normal to the stream and the ventral and dorsal walls of the heart. Using these intersection points, we then calculated the Euclidean distance between the centerline of the stream and the ventral ( $\varepsilon$ ) and dorsal ( $\delta$ ) walls. The normalized intracardiac stream position was defined as  $\delta/(\varepsilon+\delta)$ . We computed the position for 10 evenly distributed points along the stream centerline and took the mean value as the final intracardiac stream position (Fig. 2). The entire process was performed using a semi-automated ad hoc program developed in MATLAB (R2011a, Mathworks, Natick, MA). We performed the analysis once per embryo.

The brightfield movie was recorded after insertion of the needle into the vessel, but before injection of the dye. The injection movie was then recorded without moving the embryo or changing the zoom level to facilitate direct overlapping. As a check, we can clearly observe that the dye within the needle seen in the fluorescent image directly overlaps with the bore of the needle visible in the brightfield image (Fig. 2). When performing our analysis, we rejected any overlays that did not meet this registration test. In addition, as the aortic arches, dorsal aorta, and extra-embryonic vessels were filled with dye, their patterns directly overlapped with these vessels in the brightfield image, further demonstrating that no movement occurred between the recording of the brightfield and fluorescent data. When manually selecting the ventral and dorsal boundaries, we chose points along the outer surface of the heart, as it was easier to identify. Although these boundaries are not the inner lumen, we were consistent in our selection, allowing for a one to one comparison between control and LAL embryos.

We performed two-tailed, unpaired *t*-tests to determine differences in ICFP between groups. A *p*-value of less than 0.05 was considered significant.

#### Computational Fluid Dynamics Model of an *In Silico* LAL Stage 21 Heart

##### Creation of geometric heart loop models

We based the geometry and proportions of our control stage 21 heart loop model on a variety of anatomical imaging studies including serial histology (van den Berg and Moorman, 2011), scanning electron microscopy (Manner, 2000), micro computed tomography (Kim et al., 2011), magnetic resonance imaging (Yelbuz et al., 2003), and optical coherence tomography (Davis et al., 2008). The general "looped tube" structure was designed in Pro-Engineer (Parametric Technology Corp., Needham, MA) based on previous sketches and models (Patten, 1920; Manner, 2004). Atrial and ventricular bulbs, outflow tract tapering, and the venous inlets (RVV, LVV, and both CCV) were added using our in-house sketch-based anatomical editing tool, SketchCAD. The heart geometry was modeled at end diastole. The model was



completed and made CFD-ready using Geomagic Studio 10 (Geomagic Inc., Durham, NC). An in silico LAL surgery was performed by removing the presumptive left atrium volume and creating a smoothed surface to connect the left wall of the atrioventricular canal to the remaining right atrium, using our in-house surgical planning suite (Pekkan et al., 2008; Dur et al., 2011). The finished control and LAL models were rendered in Geomagic Studio 10 (Geomagic Inc., Durham, NC) and exported into GAMBIT (ANSYS Inc., Cannonsburg, PA), to create a surface triangular mesh. The triangular mesh was imported into our in-house CFD preprocessor, which generates an unstructured 3D Cartesian immersed boundary mesh. The immersed boundary grid had a spatial resolution of 14  $\mu\text{m}$  for both the control and LAL models.

### Computational fluid dynamics

Blood flow through the embryonic heart was modeled using our in-house immersed boundary cardiovascular flow solver incorporating a validated second-order accurate multi-grid artificial compressibility numerical method, described in our previous studies (Payli et al., 2007; Pekkan et al., 2009; Menon et al., 2012). Flow was simulated in terms of inlet normalized spatio-temporal units, on a high-resolution unstructured Cartesian immersed boundary grid with finite-difference numerical treatment. The objective of the CFD model was to determine if the geometric change of removing the left atrial volume was sufficient to alter ICFP and WSS. Therefore, we performed a steady-state simulation with fixed walls and no slip boundary conditions. The mean cardiac output of the stage 21 chick embryo, 1.28  $\text{mm}^3/\text{s}$  (Yoshigi et al., 2000), was used to compute inflow boundary conditions at each of the four venous inlets. As reliable data for the venous flow rates was not available, we prescribed an inflow distribution of 25/75 between the blood returned from the embryo by the CCV and blood returned from the vitelline bed by the VV (Hu et al., 1996). Furthermore, a 50/50 split was specified between the cumulative inflows from the laterals. The Reynolds number (Re) of each of the CCV inlets was therefore 0.25, the Re of the VV inlets was 0.76, and the Re at the outflow tract was 2. A plug flow profile was assigned to all inlets. An instantaneous mass-flow preserving outflow boundary condition was adopted. Blood was treated as a Newtonian fluid with constant hemodynamic properties ( $\rho = 1,060 \text{ kg/m}^3$ ,  $\mu = 3.71 \times 10^{-3} \text{ Pa}\cdot\text{s}$ ) (Al-Roubaie et al., 2011). In the case of the later stage model, which has only three inlets, the cardiac output remained the same, with 75% from the vitelline vein and 12.5% from each of the CCV. All other boundary conditions and assumptions were unchanged.

CFD simulations were conducted at Pittsburgh Supercomputing Center's Blacklight supercomputing cyber-infrastructure. Each simulation cost an average of 22 hr at 32 core parallelism. Convergence of the steady inflow CFD solution was monitored using the running average velocity field, which converged to steady state relatively quickly.

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