

Prox1 dosage controls the number of lymphatic endothelial cell progenitors and the formation of the lymphovenous valves

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Arteries, veins, and lymphatic vessels are functionally linked, and their physical interaction is tightly regulated. The lymphatic vessels communicate with the blood vessels only at the junction of the jugular and subclavian veins. Here, we characterize the embryonic lymphovenous valves controlling this vital communication and show that they are formed by the intercalation of lymphatic endothelial cells (LECs) with a subpopulation of venous endothelial cells (ECs) at the junction of the jugular and subclavian veins. We found that unlike LEC progenitors, which move out from the veins and differentiate into mature LECs, these Prox1-expressing ECs remain in the veins and do not acquire LEC features. We demonstrate that the development of this Prox1-expressing venous EC population, and therefore of lymphovenous valves, requires two functional copies of *Prox1*, as the valves are absent in *Prox1* heterozygous mice. We show that this is due to a defect in the maintenance of Prox1 expression in venous ECs and LEC progenitors promoted by a reduction in Coup-TFII/Prox1 complex formation. This is the first report describing the molecular mechanism controlling lymphovenous communication.

[*Keywords:* LEC progenitors; Prox1; lymphatics; lymphovenous valves]

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In mammals, the two vascular systems—the blood vasculature, which transports oxygen and nutrients, and the lymphatic vasculature, which transports fluid and macromolecules from tissues back to the blood circulation—are tightly interconnected. Because of differences in oncotic pressure, the plasma fluids that continuously extravasate from the blood vasculature are collected from the tissue space by the lymphatic vasculature that then returns it back to the blood circulation so that fluid homeostasis is properly maintained (Oliver and Srinivasan 2008).

The stepwise process leading to the formation of the lymphatic vascular network starts when differentiating Prox1-expressing lymphatic endothelial cell (LEC) progenitors leave the embryo's cardinal vein (CV) to form intermediate structures called lymph sacs. The first lymph sacs to develop are the paired jugular lymph sacs. Each jugular lymph sac retains a connection to the adjacent vein (Sabin 1902; Kampmeier 1969; van der Putte 1975) controlled by lymphovenous valves (van der Putte 1975). Because this connection is where reabsorbed tissue fluids collected by the lymphatics are released back into the blood circulation, any alteration in the genesis of the lympho-

venous valves is supposed to severely compromise the maintenance of normal fluid homeostasis.

The anatomical identification of these lymphovenous valves has been hampered by the lack of specific markers (Lewis 1905; Gittenberger-De Groot et al. 2004) and no information is yet available about the molecular and cellular mechanisms leading to their formation. However, in the last decade, genes controlling different aspects of developmental and postnatal lymphangiogenesis have been identified, and useful animal models have been generated (Oliver and Srinivasan 2008). Taking advantage of these tools, we have now performed a detailed molecular characterization of these lymphovenous valves during developmental lymphangiogenesis. We conclusively show that the lymphovenous connection is mediated by two nearby valves located at the junction of the jugular and subclavian veins. We report for the first time that these valves are formed by the intercalation of lymph sac-derived Prox1⁺ LECs with a previously unidentified population of Prox1⁺ endothelial cells (ECs) present in the adjacent veins. As LECs originate from the embryonic veins (Srinivasan et al. 2007), we argue that these Prox1⁺ ECs arise from a subpopulation of the Prox1-expressing venous ECs; some will become LEC progenitors and leave the vein, while others will remain on the vein and participate in the formation of the lymphovenous valves. We also show that in *Prox1* heterozygous embryos, the number of Prox1-

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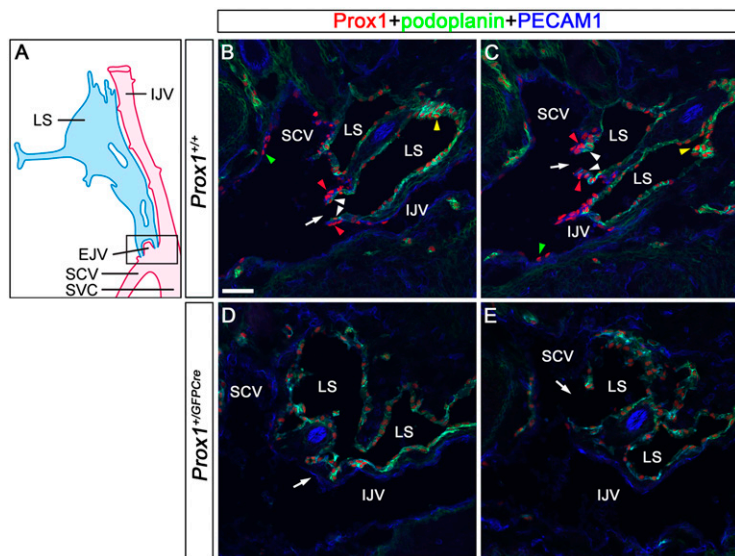


Figure 1. Prox1 is expressed on a previously unidentified population of ECs in the lymphovenous valves. (A) Schematic representation (adapted with modifications from van der Putte, 1975) of the area of an E13.5 embryo where the jugular and subclavian veins join to form the lymphovenous valves. The head (anterior) is oriented toward the top and the heart (posterior) is oriented toward the bottom of the figure. These veins merge to form the SVC. The posterior portion of the jugular lymph sac (LS) opens into the SCV where the veins merge. To visualize the boxed region of the diagram, E13.5 wild-type (B,C) or *Prox1*^{+/GFP-Cre} (D,E) embryos were frontally sectioned from the dorsal to the ventral side and immunostained for the LEC markers Prox1 and podoplanin and the pan-endothelial PECAM1. (B) The first valve (arrow) is adjacent to the IJV on one side and the SCV on the other side. (C) The second valve (arrow) in this orientation is on the SCV. Each of the valve's two leaflets consists of two layers of Prox1⁺ ECs: an inner Prox1⁺ podoplanin⁺ layer that is continuous with the lymph sac (white arrowheads) and an outer Prox1⁺ podoplanin⁻ layer that is continuous with the veins (red arrowheads).

Note the relatively high levels of Prox1 in the valves (red arrowheads) and in some cells of the lymph sac (yellow arrowheads). (B,C) Few Prox1⁺ cells are also seen on the veins away from the lymphovenous valves (green arrowheads). (D) In *Prox1*^{+/GFP-Cre} embryos, Prox1 is not expressed on the walls of the veins (arrow). (E) Occasionally, the lymph sacs abruptly fuse with the IJV (arrow). The brain is oriented toward the right, the heart is oriented toward the left, and the thymus is oriented toward the bottom of B–E. Bar, 50 μ m.

expressing ECs on the vein is reduced, and, consequently, the number of LECs is reduced and the lymphovenous valves fail to form. We demonstrate that this defect is because of a reduction in the amount of Coup-TFII/Prox1 complex formation necessary for the maintenance of *Prox1* expression. We thereby provide a mechanistic explanation for the haploinsufficiency observed in *Prox1* heterozygous mice (Harvey et al. 2005).

Results

Molecular identification and characterization of the lymphovenous valves

Previous detailed anatomical studies in mouse embryos identified the junction of the jugular and subclavian veins as the location where the communication between the lymphatic and blood vasculatures is established (Fig. 1A; van der Putte 1975). Internal and external jugular veins return the deoxygenated blood from the brain and other anterior organs to the heart, and the subclavian vein (SCV) returns deoxygenated blood from the limb buds (van der Putte 1975). The internal jugular vein (IJV) runs in an anterior to posterior orientation, and the SCV runs more or less perpendicular to the IJV (Fig. 1A). The external jugular vein (EJV) takes a lateral route after leaving the skull, and because of its orientation only the site of its fusion with the SCV is represented in the scheme shown in Figure 1A. These three veins merge at approximately the level of the thymus to form the superior vena cava (SVC), which in turn opens into the atrium of the heart (van der Putte 1975). All three veins are symmetrically paired structures with one vein of each pair present on each side of the body (van der Putte 1975).

Taking this anatomical information into account, we performed a detailed anatomical and molecular characterization of the formation of the lymphovenous valves in mouse embryos. First, embryonic day 13.5 (E13.5) embryos were frontally sectioned (dorsal to ventral) to facilitate the visualization of the lymph sacs along with the IJV, the SCV, and the valves through which they interact (Fig. 1A). Then, sections were coimmunostained using the LEC markers Prox1 and podoplanin and the pan-endothelial marker PECAM1. As expected, two valves connecting the lymph sacs with the surrounding veins were identified at the junction between the jugular and subclavian veins (Fig. 1B,C, arrows). One of these valves lies more dorsomedially than does the other one (Fig. 1B, arrow). Veins and lymph sacs are nearly symmetrically located on the left and right sides of the embryo. Similarly, the lymphovenous valves are also present on both sides, and their locations and overall structures are nearly identical (data not shown).

We found that each valve is composed of two leaflets, each containing two layers of Prox1⁺ cells. The inner layer is contributed by the lymph sacs; therefore, it is composed of Prox1⁺ podoplanin⁺ LECs (Fig. 1B,C, white arrowheads). However, the outer layer is composed of a previously unidentified population of PECAM1⁺/Prox1⁺/podoplanin⁻ ECs extending from the walls of the surrounding veins (Fig. 1B,C, red arrowheads).

To better understand the structural organization of these valves, we performed immunostainings for Prox1 and PECAM1 on transverse sections along the anterior–posterior axis. As expected, Prox1 expression is detected lining the entire lymph sac at the most anterior level but is not observed on the IJV (Supplemental Fig. S1A). However, as we move posterior to the region shown within the box in Figure 1A, Prox1 expression is detected on the IJV's wall

but only within the region that is in closest contact with the lymph sac (Fig. 2A, arrow). Prox1 is detected on the walls of the IJV and the SCV as the SCV approaches the jugular vein (Supplemental Fig. S1B). Around this level, where the lymph sac is split into two portions by the vertebral artery (Fig. 2B, arrowhead), the more medial segment intercalates with Prox1⁺ cells originating from the IJV on one side and from the SCV's wall on the other side (Fig. 2B, arrows). This structure corresponds to the first valve shown on the frontal sections of Figure 1A. Posterior sections show that the IJV and the SCV are fused together, forming a single blood vessel containing a cluster of Prox1⁺ cells representing the tip of the valve (Fig. 2C, arrow). The

more lateral portion of the lymph sac lies close to the SCV (Fig. 2D). The EJV, which was not apparent in the frontal sections, becomes obvious in adjacent sections where it branches off from the SCV and moves laterally (Fig. 2E, arrow). High levels of Prox1 expression are detected on the EC located on the wall of the EJV that is in close contact with the wall of the Prox1-expressing lymph sac (Fig. 2E). Because of the close proximity between these two structures (i.e., lymph sac and EJV), the two cell layers (one of each structure) appear to merge together, forming a single cell layer (Fig. 2E, arrowhead). The most posterior sections show a valve-like opening through which the lymph sac communicates with the veins (Fig. 2F, arrow). There-

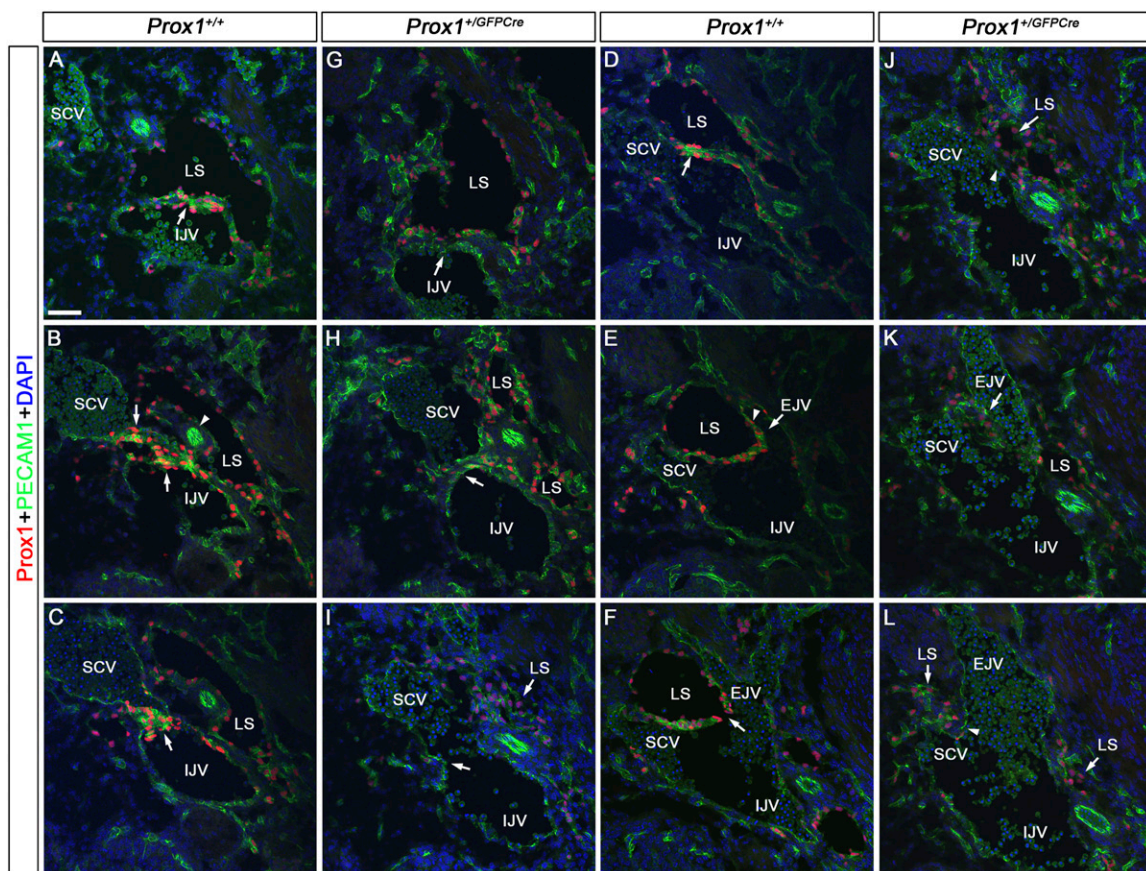


Figure 2. Lymphovenous valves are formed by the fusion of lymph sacs with two adjacent veins. E13.5 wild-type (A–F) or *Prox1*^{+/GFPcre} (G–L) embryos were transversely sectioned in an anterior to posterior orientation in the region where the jugular and subclavian veins interact (box in Fig. 1A) and were immunostained for the LEC marker Prox1 and the pan-endothelial marker PECAM1. (A) In wild-type embryos, Prox1 is expressed uniformly in LECs forming the lymph sac (LS) and in a polarized manner on the IJV (arrow). Note the relatively high levels of Prox1 on the ECs in the vein and in the lymph sacs' LECs that are facing the vein. (B) The lymph sac is split into two portions by the vertebral artery (white arrowhead). Both walls of the medial portion of the lymph sac intercalate with the wall of the IJV medially and the SCV laterally (arrows). (C) The IJV and the SCV have completely merged together, and the valve rudiment is seen in the middle (arrow). (D) The lateral portion of the lymph sac (LS) runs adjacent to the SCV. Note the relatively higher levels of Prox1 in the venous ECs and in the LECs in the lymph sac facing the vein (arrow). (E) The EJV is branching off from the SCV (arrow), and Prox1 is expressed on the walls of this vein in a polarized manner (arrowhead). This wall is also adjacent to the lymph sac. (F) The opening of the valve (arrow) is now seen and is formed by the fusion of the two layers of Prox1⁺ ECs. (G,H) In *Prox1*^{+/GFPcre} embryos, Prox1 is expressed in LECs forming the lymph sac, but very few Prox1⁺ cells are seen on the walls of the IJV and SCV (arrows). (I) At the point where the IJV and the SCV merge, no Prox1⁺ cells or valve-like structures exist (arrow). (J) No Prox1 expression was observed on the wall of the SCV that lies close to the lymph sac (arrowhead). (K) Posterior to that, the EJV branches off from the SCV. No Prox1⁺ cells are seen on the walls of the EJV (arrow). (L) No communication is observed between the lymph sac and the veins (arrowhead). Also, note an overall reduction in the number of LECs in G–L. The neural tube is oriented toward the right, the heart is oriented toward the left, and the thymus is oriented toward the bottom in each panel. Bar, 50 μ m.

fore, the lymphovenous valves are formed where the junction of two veins (subclavian and internal or external jugular) comes in close contact with the nearby jugular lymph sacs, and these valves are formed by two different types of Prox1-expressing EC: LECs from the lymph sac and a specialized EC type provided by the veins' walls.

Prox1 heterozygous embryos lack lymphovenous valves

The ECs in the leaflets of the lymphovenous valves expressed high levels of Prox1 (Fig. 1B,C, red arrowheads). However, in the lymph sacs, the level of Prox1 expression among cells was variable, being particularly high in some regions (Fig. 1B,C, yellow arrowheads). To evaluate whether these variable levels of Prox1 expression were functionally relevant, we took advantage of two available *Prox1* heterozygous strains (Wigle et al. 1999; Srinivasan et al. 2010). Using antibodies against Prox1, PECAM1, and podoplanin, we first evaluated the junction of the jugular and subclavian veins in frontal sections of E13.5 *Prox1*^{+/*GFP*Cre} embryos. The differences in the levels of Prox1 expression among LECs in the lymph sacs of these mutant embryos (Fig. 1D,E) were not as obvious or widespread as in normal controls (Fig. 1B,C). Furthermore, no obvious lymphovenous valve-like structures were evident in *Prox1* heterozygous embryos, and venous ECs down-regulated the expression of Prox1 (Fig. 1D, arrow). Instead, some *Prox1* heterozygous embryos' lymph sacs abruptly fused with veins (Fig. 1E, arrow) and were occasionally blood-filled (data not shown).

Analysis of transverse sections of *Prox1* heterozygous embryos at this same region confirmed that Prox1 is down-regulated in the venous walls and that lymphovenous valves are absent (Fig. 2G–L). Additionally, compared with their wild-type littermates, the total number of LECs in *Prox1*^{+/*GFP*Cre} embryos at this stage appears to be reduced; consequently, the lymph sacs were consistently smaller than those of control littermates (Fig. 2).

To confirm whether *Prox1*^{+/*GFP*Cre} embryos were missing valves, we performed additional immunostainings using the LEC markers Vegfr3 (Dumont et al. 1998) and Lyve-1 (Banerji et al. 1999) and the blood EC marker von Willebrand factor (vWF) (Ginsburg et al. 1985). As expected, wild-type control animals had Vegfr3 on lymph sac LECs but not on the venous side (Supplemental Fig. S2A). Lyve-1 was detected on most LECs in the lymph sacs and occasionally on ECs on the venous walls (Supplemental Fig. S2 arrows). In contrast, vWF was detected on venous ECs but not on LECs (Supplemental Fig. S2A). The expression of these markers was similar in *Prox1*^{+/*GFP*Cre} embryos (Supplemental Fig. S2B).

In collecting lymphatics, Foxc2 labels the lymphatic valves, and its activity is crucial for their normal development (Petrova et al. 2004). In the lymphovenous valves of control embryos, Foxc2 expression is restricted to the venous Prox1⁺ cells (Fig. 3A, arrow) (occasionally some localized expression was seen also on the LEC side). Integrin- α 9 is an additional lymphatic valve marker required for valve development in the collecting lymphatics (Bazigou et al. 2009). This gene was also strongly expressed

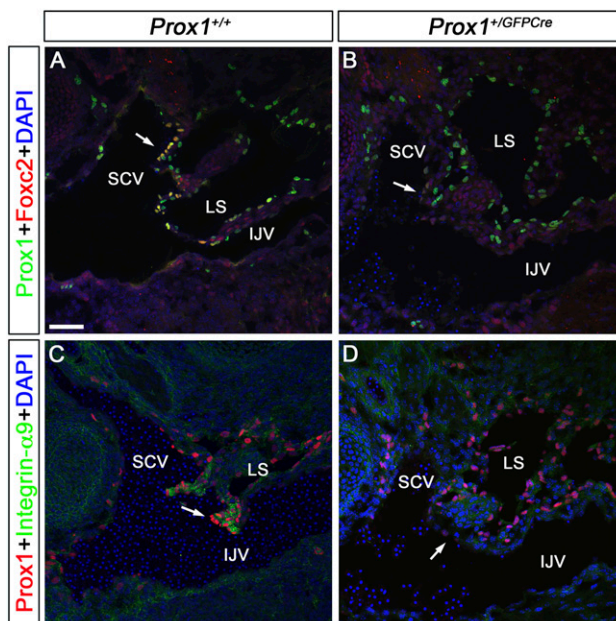


Figure 3. Foxc2 and Integrin- α 9 are expressed in the lymphovenous valves but are absent in *Prox1* heterozygous embryos. E13.5 wild-type (A,C) and *Prox1*^{+/*GFP*Cre} (B,D) embryos were frontally sectioned and immunostained for either Prox1 and Foxc2 (A,B) or Prox1 and Integrin- α 9 (C,D). (A) In wild-type embryos, Foxc2 is specifically expressed on the outer layer of the valve's leaflets, which are continuous with the walls of the veins (arrow). (B) In sections from an identical region in *Prox1*^{+/*GFP*Cre} embryos, Foxc2 is not expressed on the walls of the veins (arrow). Integrin- α 9 is strongly expressed in the valves of control embryos (C, arrow) but is absent in *Prox1*^{+/*GFP*Cre} littermates (D, arrow). The head is oriented toward the right, the heart is oriented toward the left, and the thymus is oriented toward the bottom in both panels. (IJV) Internal jugular vein; (SCV) subclavian vein; (LS) lymph sac. Bar, 50 μ m.

in the lymphovenous valves of wild-type controls (Fig. 3C, arrow). A similar analysis of *Prox1*^{+/*GFP*Cre} embryos failed to detect Foxc2 or Integrin- α 9 expression in this region, confirming that lymphovenous valves are missing in these embryos (Fig. 3B,D).

The pool of Prox1-expressing venous ECs and LEC progenitors is reduced in Prox1 heterozygous embryos

To better characterize the development of these lymphovenous valves, we compared the expression of Prox1 and Foxc2 in frontal sections of E11.5 and E12.5 wild-type control and *Prox1*^{+/*GFP*Cre} littermates. At E11.5, only weak Foxc2 expression is observed in Prox1⁺ venous ECs (Fig. 4A, arrowhead) and, occasionally, a single valve rudiment starts to be detected at this stage in the CV (future IJV) of control embryos (Fig. 4A, arrow). In contrast, no valves or valve rudiments were present in *Prox1*^{+/*GFP*Cre} embryos (Fig. 4B). We also noticed that the number of Prox1⁺ ECs on and outside the CV was reduced in *Prox1* heterozygous embryos (Fig. 4C). At E12.5, Foxc2⁺/Prox1⁺ lymphovenous valve rudiments were evident on the wall of the CV of control embryos, juxtaposed to Prox1⁺/Foxc2⁻ lymph sacs (Fig. 4D, arrows). These Foxc2⁺ cells were the only Prox1⁺

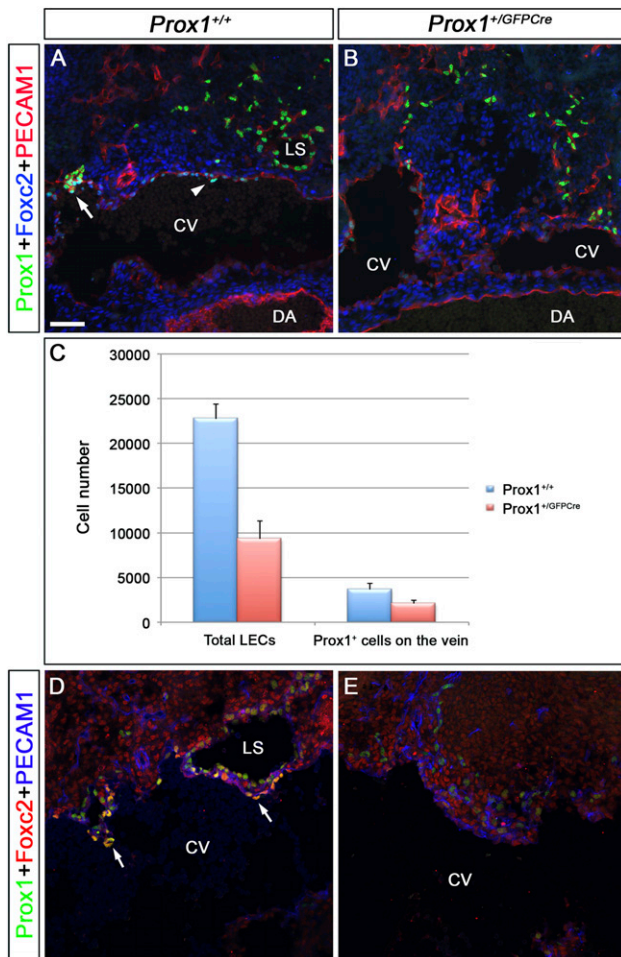


Figure 4. The number of *Prox1*-expressing LEC progenitors and *Prox1*-expressing venous ECs that will form the lymphovenous valves is reduced in *Prox1* heterozygous embryos. (A,B) E11.5 control (A) and *Prox1*^{+/GFP-Cre} (B) embryos were frontally sectioned and immunostained for *Prox1*, *Foxc2*, and *PECAM1*. (A) In the control embryos, *Foxc2* is weakly expressed in the *Prox1*⁺ cells on the vein (arrowhead). In addition, lymph sacs and occasionally one valve rudiment (arrow) could be observed. (B) In *Prox1*^{+/GFP-Cre} embryos, the total number of LECs appears to be reduced, and no lymph sacs or valve rudiments are seen. (C) Compared with wild-type littermates, the total number of *Prox1*⁺ *PECAM1*⁺ ECs on the vein and outside the vein were reduced in E11.5 *Prox1*^{+/GFP-Cre} embryos ($n = 3$ for each genotype, $P < 0.05$). (D,E) E12.5 control (D) and *Prox1*^{+/GFP-Cre} (E) embryos were frontally sectioned and immunostained for *Prox1*, *Foxc2*, and *PECAM1*. (D) In control embryos, *Foxc2* is expressed on the valve rudiments (arrows), which lie adjacent to the primitive lymph sacs. (E) In *Prox1*^{+/GFP-Cre} embryos, the total number of LECs appears to be reduced, and no valve rudiments are seen. Lymph sacs are also not observed in this location. The head is oriented toward the right, the heart is oriented toward the left, and the thymus is oriented toward the bottom in all panels. (CV) Cardinal vein; (DA) dorsal aorta; (LS) lymph sac. Bar, 50 μ m.

cells on the CV at this stage. In contrast to that finding but similar to what we found at later stages, no *Prox1*⁺/*Foxc2*⁺ cells were present on the veins of E12.5 *Prox1*^{+/GFP-Cre} littermates (Fig. 4E), and the total number of LECs outside

the veins was also reduced at this stage (Fig. 4D,E). It could be possible that the observed progressive reduction in the number of *Prox1*-expressing venous ECs in *Prox1*^{+/GFP-Cre} embryos might be partially responsible for the lack of lymphovenous valves.

To investigate whether the lack of lymphovenous valves could also be due to a developmental delay, we analyzed the expression of *Prox1*, podoplanin, and *PECAM1* in E16.5 *Prox1*^{+/LacZ} embryos. At this stage, the edematous phenotype seen at earlier stages is rescued (Supplemental Fig. S3). Also at this stage, the general appearance of the lymphovenous valves of control embryos is similar to that of those analyzed at E13.5 (Supplemental Fig. S4A, arrow), although the outer *Prox1*⁺ podoplanin⁻ cell layer in the leaflets appeared longer (Supplemental Fig. S4A, dotted line); *Prox1* expression in venous valves could also be seen at this stage (Supplemental Fig. S4A, arrow; see below). In contrast, in *Prox1*^{+/LacZ} embryos, *Prox1*⁺ cells were rarely seen on venous ECs, *Prox1* expression remains significantly down-regulated in the walls of the IJV, and no valves are present (Supplemental Fig. S4B). In some areas, the lymph sac was abruptly fused with the IJV (Supplemental Fig. S4C); however, only a small number (<10%) of *Prox1* heterozygous embryos had blood-filled superficial lymphatics (Supplemental Fig. S3). This finding suggests that additional alterations are required for blood to breach the lymphatic vasculature. Occasionally, we also saw lymphatic vessels from the para-tracheal lymph plexus opening into the pleural cavity of *Prox1* heterozygous embryos (Supplemental Fig. S4D, arrow), a defect that could be partially responsible for the chylothorax (i.e., accumulation of chyle in the thoracic cavity) seen in newborn *Prox1* heterozygous pups (Harvey et al. 2005). Taken together, our findings show that *Prox1* activity is required for the proper formation of the lymphovenous valves in a dose-dependent manner; therefore, these valves are defective in *Prox1* heterozygous embryos with abnormal lymphovenous connections. These abnormal connections are likely used to relieve the excessive pressure due to fluid accumulation in the lymphatic vessels.

The lack of lymphovenous valves in Prox1 heterozygous embryos is caused by defective maintenance of venous Prox1 expression

It was previously suggested that two steps are necessary during the initial LEC differentiation process: an early *Prox1* initiation step dependent on Coup-TFII and Sox18, and a later *Prox1* maintenance step dependent on *Prox1* itself (Francois et al. 2008; Srinivasan et al. 2010). Our results showing that losing a copy of *Prox1* triggers a reduction in the number of *Prox1*⁺ cells in the veins without affecting their rate of proliferation or cell death (data not shown) suggested that the *Prox1*-dependent maintenance step could be affected in these heterozygous embryos. To evaluate this possibility, we performed lineage-tracing analysis by crossing *Prox1*^{+/CreERT2}; *R26R*^{+/YFP} mice (Srinivasan et al. 2001; Srinivasan et al. 2007) with *Prox1*^{+/LacZ} mice (Wigle et al. 1999) and injecting the pregnant females with 2 mg of tamoxifen per 40 g of body weight at E10.5. *Prox1*^{+/CreERT2}; *R26R*^{+/YFP} and *Prox1*^{LacZ/CreERT2}; *R26R*^{+/YFP}

embryos isolated from those crosses were analyzed at E13.5. Using this approach, LEC progenitors should be labeled as soon as *Prox1* expression is initiated (around E10.0). In this case, if *Prox1* maintenance is normal, all labeled YFP⁺ ECs also will be *Prox1*⁺; however, if *Prox1* maintenance is defective, those ECs will be *Prox1*⁻ YFP⁺.

As described before, numerous *Prox1*⁺ cells were detected in the lymphovenous valve region of control *Prox1*^{+/CreERT2};R26R^{+/YFP} embryos, and some of these coexpressed YFP, a result indicating that these cells are descendants of LEC progenitors that were present at E10.5 (Supplemental Fig. S5A,B). Importantly, *Prox1*⁻ YFP⁺ cells were not detected in these embryos (Supplemental Fig. S5A,B). Instead, *Prox1* expression was down-regulated in the putative lymphovenous valve territory of *Prox1* heterozygous (*Prox1*^{LacZ/CreERT2};R26R^{+/YFP}) littermates (Supplemental Fig. S5C), although few YFP⁺ cells were detected in this region (Supplemental Fig. S5D, arrow). Therefore, these YFP⁺ cells correspond to the original LEC progenitors that failed to maintain *Prox1* expression and thus were unable to differentiate into valve cells.

Prox1^{+/CreERT2}-mediated labeling is mosaic and occurs only during the short developmental time frame when tamoxifen is active; therefore, we performed a similar analysis in E13.5 *Prox1*^{+/GFPcre};R26R^{+/YFP} embryos. Likewise, *Prox1* expression was down-regulated; however, numerous YFP⁺ cells were present in the putative lymphovenous valve region (Fig. 5A, arrows). As indicated before, some *Prox1* heterozygous embryos have abnormal lymphovenous connections and blood-filled lymphatics at this stage (Fig. 5B, arrow). We also detected several *Prox1*⁻/YFP⁺ cells in the IJV in this area (Fig. 5B, arrowheads). These observations suggest that in *Prox1* heterozygous embryos there is a defect in the maintenance of *Prox1* expression in LECs and lymphovenous valve progenitors.

CoupTFII/Prox1 complex formation is reduced in *Prox1* heterozygous LEC progenitors

Next, we aimed to identify the mechanisms leading to the silencing of *Prox1* expression in *Prox1* heterozygous embryos. As mentioned above, two steps are likely involved

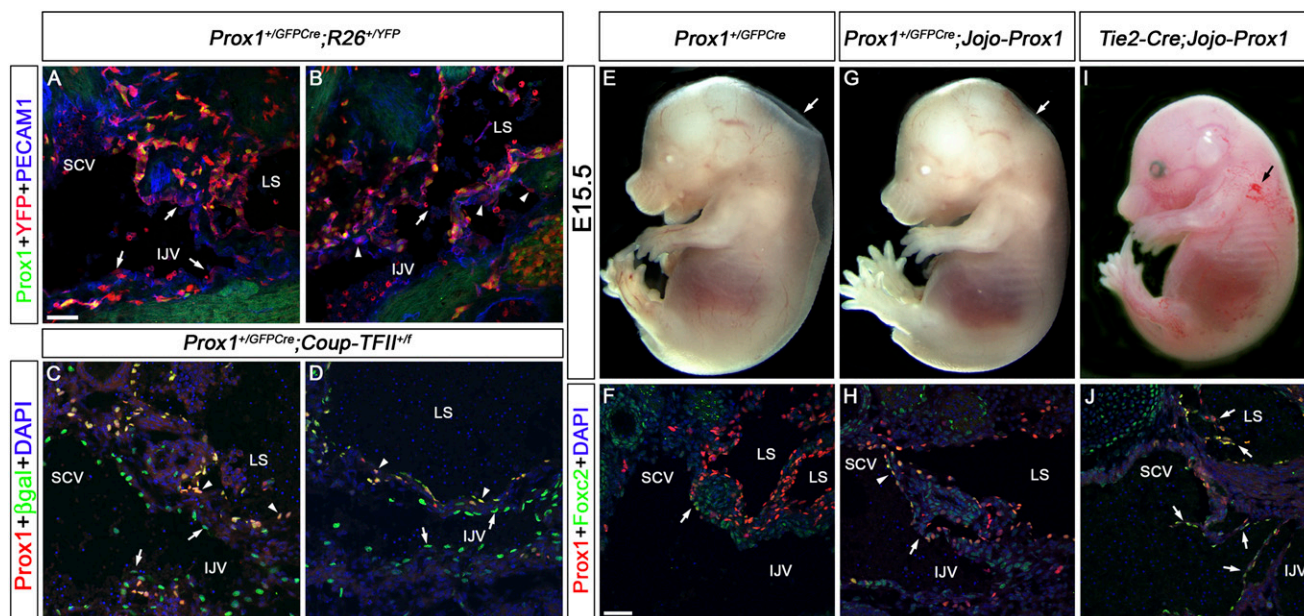


Figure 5. *Prox1* heterozygous embryos do not maintain *Prox1* expression in a subpopulation of LEC and lymphovenous valve progenitors. (A,B) E13.5 *Prox1*^{+/GFPcre};R26R^{+/YFP} embryos were sectioned and immunostained for *Prox1*, YFP, and PECAM1. (A) No *Prox1*⁺ ECs are seen in the area that should have developed into the lymphovenous valve; however, numerous YFP⁺ cells can be detected (arrows). (B) In some *Prox1* heterozygous embryos that develop blood-filled lymphatics at this stage, abnormal connections between the lymph sacs and the IJV are observed (arrow); several *Prox1*⁻ YFP⁺ cells are seen in this area (arrowheads). The YFP⁺ PECAM1⁻ cells in the lumen of the blood vessel and the lymph sacs are hematopoietic cells. (C,D) E15.5 *Prox1*^{+/GFPcre};Coup-TFII^{+/fl} embryos were immunostained for *Prox1* and β-galactosidase. (C,D) The blood-filled lymph sac is clearly seen and is lined by *Prox1*⁺β-gal⁺ LECs (arrowheads). (C) No *Prox1* expression is seen on veins in the region where the valves are normally formed (arrows), but β-gal⁺ cells are detected in this region. (D) *Prox1*⁻β-gal⁺ cells line the IJV (arrows). (E–J) *Jojo-Prox1* mice were bred to *Prox1*^{+/GFPcre} or *Tie2-Cre* mice and the resulting embryos were analyzed at E15.5 by whole mount or by immunohistochemistry on frontal sections with antibodies against *Prox1* and *Foxc2*. *Prox1*^{+/GFPcre} embryos display edema (E, arrow), and the expression of *Prox1* and *Foxc2* is reduced in the region where the valves are normally formed (E, arrow). (G) Expression of *Prox1* in *Prox1*^{+/GFPcre};Jojo-*Prox1* embryos rescues the lymphatic vascular phenotype, as these embryos display no obvious edema (arrow). (H) Expression of *Prox1* and *Foxc2* in the venous ECs (arrowhead) and the lymphovenous valves (arrow) is observed in *Prox1*^{+/GFPcre};Jojo-*Prox1* embryos. (I) Misexpression of *Prox1* in blood ECs in *Tie2-Cre*;Jojo-*Prox1* embryos results in a blood-filled lymphatic phenotype (arrow). (J) Numerous abnormal valve-like structures composed of *Prox1*⁺*Foxc2*⁺ cells were observed in the IJV and lymph sac (LS) of *Tie2-Cre*;Jojo-*Prox1* embryos (arrows). In the sections, the head is oriented toward the right, the heart is oriented toward the left, and the thymus is oriented toward the bottom in all panels. (IJV) Internal jugular vein; (SCV) subclavian vein; (LS) lymph sac. Bar, 50 μm.

in regulating *Prox1* expression (Srinivasan et al. 2010). The *Prox1*-dependent maintenance step, which occurs when *Prox1* expression is induced and LEC differentiation is progressing, can be further subdivided into an early Coup-TFII/*Prox1*-dependent step that occurs on the vein and a later Coup-TFII-independent step that occurs when LECs move out from the vein (Srinivasan et al. 2010). The observed reduction in *Prox1* expression in the veins of *Prox1* heterozygous embryos could be caused by defects in the maintenance step (i.e., the Coup-TFII/*Prox1*-dependent step). Directly quantifying the amount of Coup-TFII/*Prox1* complex is technically difficult because there are few LEC progenitors in *Prox1* heterozygous embryos, there are no available surface markers to sort these cells, and there are two different *Prox1*⁺ cell populations on the vein at any given time (those that will maintain and those that will not maintain *Prox1* expression). Therefore, as an alternative method to study the Coup-TFII/*Prox1* complex, we used an algorithmic approach. According to Langmuir isotherm (Copeland 2000), Coup-TFII/*Prox1* interaction could be represented as a bimolecular reaction as follows: [Coup-TFII] + [*Prox1*] → [Coup-TFII/*Prox1*] (brackets represent the equilibrium concentrations). The amount of Coup-TFII/*Prox1* complex could be represented as [Coup-TFII/*Prox1*] = k_a [Coup-TFII][*Prox1*], where k_a is the equilibrium association constant and is dependent on the intrinsic characteristics of the interacting proteins (i.e., the strength of electrostatic and hydrophobic interactions and extrinsic properties such as temperature, osmolarity, and pH).

By considering k_a and [Coup-TFII] to be the same, it is easy to appreciate that the amount of Coup-TFII/*Prox1* complex formed in *Prox1* heterozygous LECs will be half that formed in wild-type LECs. We hypothesized that this reduction in the amount of Coup-TFII/*Prox1* complex formed could be the reason for the reduction in *Prox1* expression. Then, if Coup-TFII levels are reduced in addition to those of *Prox1*, the amount of Coup-TFII/*Prox1* complex formed should be further reduced, and the resulting phenotype should be more severe. We used *Prox1*^{+/GFP^{Cre} mice that were bred with *Coup-TFII*^{+/f} mice to generate double-heterozygous animals; most of those pups die at birth in mixed backgrounds or have a significantly low survival (2.5%) in the NMRI background (Srinivasan et al. 2010). Accordingly, we collected double-heterozygous embryos at either E11.5 or E15.5. At E11.5, significantly fewer *Prox1*⁺ cells were on or outside the veins of *Prox1*^{+/GFP^{Cre}; *Coup-TFII*^{+/f} embryos compared with their *Prox1*^{+/GFP^{Cre} littermates (Supplemental Fig. S6A). At E15.5, all of the double-heterozygous embryos had severe edema and variable blood-filled lymphatics (Supplemental Fig. S6B–E). Lineage-tracing analysis using the *LacZ* reporter that is activated when the *Coup-TFII* allele is deleted (Takamoto et al. 2005) showed that the lymphovenous valves were absent in the double-heterozygous embryos and were replaced by *Prox1*[−]βgal⁺ cells as expected (Fig. 5C, arrows). Additionally, we observed numerous *Prox1*[−]βgal⁺ cells lining the IJV, indicating a dramatic reduction in the number of differentiating LECs (Fig. 5D, arrows). In contrast, in E13.5 *Prox1*^{+/CreERT2}; *Coup-TFII*^{+/f} embryos exposed to tamox-}}}

ifen at E10.5, no *Prox1*[−]βgal⁺ ECs were seen in the vein (Supplemental Fig. S7A,B).

To confirm that LECs is the original fate of the labeled cells (*Prox1*[−]/YFP⁺ cells in *Prox1*^{+/GFP^{Cre}; *R26R*^{+/YFP} embryos and *Prox1*[−]/βgal⁺ cells in *Prox1*^{+/GFP^{Cre}; *Coup-TFII*^{+/f} embryos) and that they do not arise because of aberrant or leaky expression of the *Prox1*^{+/GFP^{Cre} line, we performed a rescue experiment using a previously reported *Prox1* conditional transgenic line (Lavado et al. 2010). This conditional gain-of-function strain (*Jojo-Prox1*) will express *Prox1* only in cells exposed to Cre recombinase. We then crossed this transgenic strain with the *Prox1*^{+/GFP^{Cre} strain and observed that in contrast to their *Prox1*^{+/GFP^{Cre} littermates (Fig. 5E, arrow), E15.5 double-heterozygous embryos were completely devoid of edema (Fig. 5G, arrow). Immunostainings of frontal sections of *Prox1*^{+/GFP^{Cre}; *Jojo-Prox1* embryos against *Prox1* and *Foxc2* confirmed that the expression of these two markers as well as the lack of the lymphovenous valves were rescued in these embryos (Fig. 5F,H). In contrast, double-heterozygous embryos generated by crossing the *Prox1* transgenic strain with *Tie2-Cre* (Kisanuki et al. 2001) that express *Prox1* in blood ECs display blood filled lymphatics (Fig. 5I) and ectopic abnormally shaped valves within the vein and the lymph sac (Fig. 5J, arrows). This result confirmed that the original fate of the labeled *Prox1*[−] cells is to be LECs and demonstrated that *Prox1* heterozygous embryos have a defect in the proper maintenance of *Prox1* expression, resulting in the loss of LEC progenitors and lymphovenous valves.}}}}}}

Next, we studied the contribution of k_a and [Coup-TFII] to the reaction kinetics. To determine the role of k_a , we used previously generated *Prox1*^{+/NR} mice in which the nuclear hormone interaction motif (and therefore the Coup-TFII interaction motif) is mutated (Srinivasan et al. 2010). Although mutating the nuclear hormone interaction motif affects the k_a between Coup-TFII and *Prox1* (Srinivasan et al. 2010), *Prox1*^{+/NR} embryos are normal (Srinivasan et al. 2010), and the development of the valves is not affected in these mutant embryos (Fig. 6A,B). Therefore, the reduction in k_a caused by the mutation in the nuclear hormone interaction motif is not sufficient to cause a valve defect.

We then investigated the role of [Coup-TFII] in the formation of the valves in *Tie2-Cre*; *Coup-TFII*^{+/f} embryos. A small number of *Tie2-Cre*; *Coup-TFII*^{+/f} embryos had mild edema (data not shown); otherwise, these EC-specific *Coup-TFII* heterozygous embryos were normal (Fig. 6C) and their lymphovenous valves were indistinguishable from those of wild-type controls (Fig. 6D). However, a blood-filled lymphatics phenotype and lack of lymphovenous valves were observed in E13.5 *Prox1*^{+/CreERT2}; *Coup-TFII*^{+/f} embryos exposed to tamoxifen at E10.5 (Supplemental Fig. S7C).

Next we tested the additive contribution of k_a and [Coup-TFII] in *Tie2-Cre*; *Coup-TFII*^{+/f}; *Prox1*^{+/NR} embryos and pups. In contrast to single-heterozygous animals, these compound-mutant animals were indistinguishable from *Prox1* heterozygous mice, as they had severe edema at E13.5 (data not shown) and at E15.5 (Fig. 6E) and lacked lymphovenous valves (Fig. 6F, arrows). Furthermore, sim-

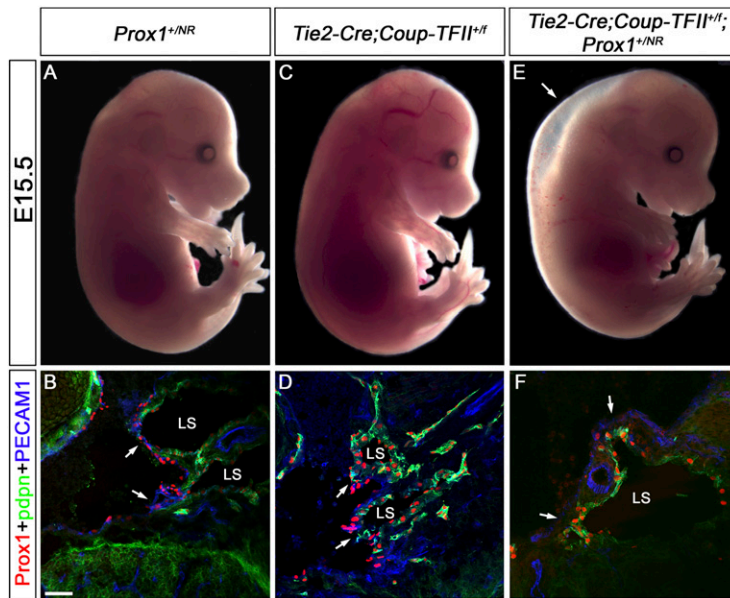


Figure 6. *Prox1* haploinsufficiency is the result of a reduction in Coup-TFII/*Prox1* interaction. *Prox1*^{+/*NR*} and *Tie2-Cre;Coup-TFII*^{+/*+*} mice were bred and the resulting embryos were analyzed at E15.5 by whole mount or by immunohistochemistry on frontal sections with antibodies against *Prox1*, podoplanin (pdpn), and PECAM1. (A–D) *Prox1*^{+/*NR*} and *Tie2-Cre;Coup-TFII*^{+/*+*} embryos are phenotypically normal with unremarkable valves. Instead, *Tie2-Cre;Coup-TFII*^{+/*+*}; *Prox1*^{+/*NR*} embryos are edematous (E, arrow), devoid of *Prox1*⁺ cells in the veins (F, arrows), and lack lymphovenous valves. In the sections, the head is oriented toward the right, the heart is oriented toward the left, and the thymus is oriented toward the bottom in all panels. (LS) Lymph sac. Bar, 50 μ m.

ilar to *Prox1* heterozygotes, compound mutants also had chylothorax, a reduced survival rate after birth (data not shown), and a reduced number of *Prox1*⁺ ECs on or outside the vein at E11.5 (Supplemental Fig. S8). A less severe reduction in the number of *Prox1*⁺ cells was also observed on or outside the veins of E11.5 *Prox1*^{+/*NR*} and *Tie2-Cre;Coup-TFII*^{+/*+*} embryos (Supplemental Fig. S8), a result that could explain the lack of any obvious lymphatic defects in these mutant embryos. These data further suggests that the lack of lymphovenous valves is likely due to a reduction in the amount of Coup-TFII/*Prox1* complex that forms during LEC specification.

Prox1 dosage is also important for the formation of venous valves

Valves are important for the normal physiology of the cardiovascular system and are also present in collecting lymphatic vessels, heart, and veins. *Prox1* expression was previously reported in cardiac valves (Risebro et al. 2009) and in collecting lymphatic vessels (Bazigou et al. 2009; Norrmen et al. 2009), but we did not observe any obvious defect in the development of these valves in *Prox1* heterozygous embryos (Fig. 7A–C; Supplemental Fig. S9). We decided to analyze whether *Prox1* is expressed in venous valves and whether the development of these valves is defective in *Prox1* heterozygous embryos. Venous valves develop later than the lymphovenous valves and can be seen at the junction of jugular and subclavian veins guarding their entry point (Kampmeier 1928). Accordingly, we found that at E16.5, the venous valves are also *Prox1*⁺ (Fig. 7D, arrowheads). In contrast, similar to the lymphovenous valves, no venous valves were detected in *Prox1* heterozygous embryos (Fig. 7E) at this stage, although few *Prox1*⁺ cells were present on the veins.

A possible explanation for the lack of venous and lymphovenous valves in *Prox1* heterozygous embryos could be that they have a common origin. To test this possibil-

ity, we generated E16.5 *Prox1*^{+/*CreERT2*}; *R26R*^{+/*LacZ*} embryos and exposed them to tamoxifen (5 mg per 40 g of body weight of pregnant dams) at E10.5, a stage when LECs and lymphovenous valve progenitors arise. As expected, we observed several labeled cells in the lymphovenous valves (Fig. 7F, arrows); however, few, if any, labeled cells were observed in the venous valves (Fig. 7F, arrowheads), a result indicating that the venous valves originate later than LECs and lymphovenous valves. The few *Prox1*⁺ cells observed in the veins of E13.5 embryos away from the lymphovenous valves (Fig. 1B,C, green arrowheads) are the likely source of these valves.

Discussion

“The duct lies for some distance against the vein, the two being separated only by a double layer of endothelium, one for the vein and one for the lymph duct. Finally, in each series, one can see that, just at the edge of the lymph duct, these two layers are continuous” (Sabin 1902).

Almost a century after the existence of lymphovenous valves at the junction of the jugular lymph sacs with the jugular and subclavian veins was first reported, our work provides a detailed molecular framework for Sabin’s accurate anatomical description (Sabin 1902), showing that, indeed, there is a gradual merger of the ECs of the lymph sacs with the walls of the two adjacent veins (Supplemental Fig. S10). Thus, the lymph sac merges at a site where two veins branch off, effectively converting the branch point into a valve.

We previously suggested that the formation of the lymphatic vasculature is a stepwise process that starts with the expression of *Prox1* in a subpopulation of venous ECs (Oliver and Harvey 2002). The new data presented in this study indicate that although the majority of those *Prox1*-expressing ECs (LEC progenitors) will eventually move out from the veins, acquire the expression of additional

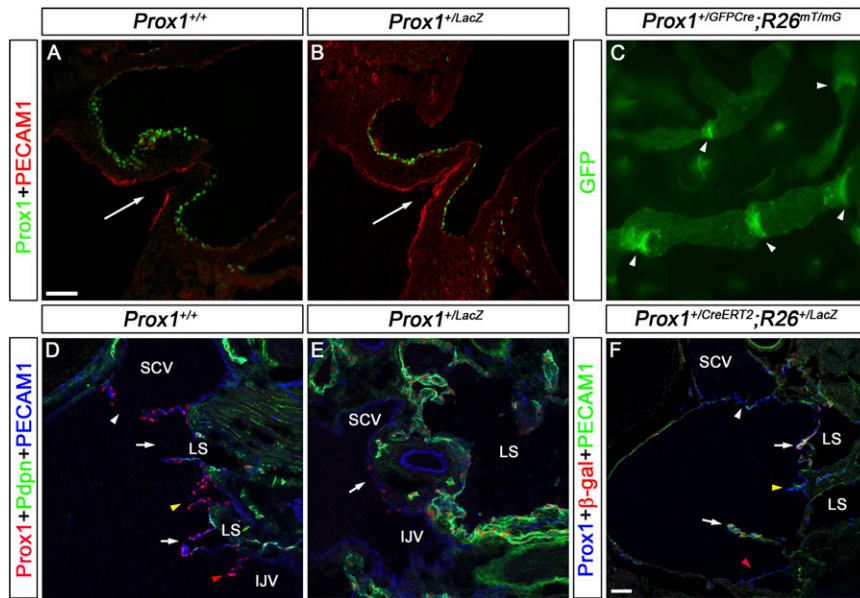


Figure 7. *Prox1* heterozygous embryos have defective venous valves. (A,B) The semilunar valve in the outflow tract from the right ventricle was analyzed in E16.5 wild-type (A) and *Prox1* heterozygous (B) embryos using *Prox1* and PECAM1 antibodies. Arrows indicate the direction of blood flow. In both embryos, *Prox1* is expressed on the downstream side of the valves and no obvious defect was observed in the morphology of the valves. (C) The lymphatic vessels in the ears of adult *Prox1*^{+/GFP-Cre;R26^{mT/mG} mice were analyzed by the autofluorescence of the membrane tagged GFP activated by Cre expression. The valves in the lymphatic vessels could be seen clearly and appear normal. (D,E) On frontal sections, the venous valves were analyzed by immunostaining for *Prox1*, PECAM1, and podoplanin (Pdpn) in E16.5 wild-type (D) and *Prox1* heterozygous (E) embryos at the junction of the jugular and subclavian veins. (D) In control embryos, the venous valves could}

be seen at the outlets of the SCV (white arrowhead), EJV (yellow arrowhead), and IJV (red arrowhead). The lymphovenous valves are also seen (arrows). (E) In contrast, in *Prox1* heterozygous embryos, few *Prox1*⁺ cells are seen on the veins (arrow) but no valve is detected. (F) *Prox1*^{+/CreERT2} and *R26^{+/LacZ}* mice were bred and the pregnant mice were exposed to tamoxifen at E10.5, a stage when LECs are specified. The lymphovenous valves were subsequently analyzed at E16.5 on frontal sections by immunostaining for *Prox1*, PECAM1, and β-gal. While β-gal⁺ cells could be observed on the lymphovenous valves (arrows), hardly any labeled cells were observed in the venous valves guarding the subclavian (white arrowhead), external jugular (yellow arrowhead), and internal jugular (red arrowhead) veins, indicating a later origin of these cells. The head is oriented toward the right of A, B, and D–F. (IJV) Internal jugular vein; (SCV) subclavian vein; (LS) lymph sac. A–E have the same image magnification. Bar, 50 μm.

LEC markers (LEC specification), and further differentiate, a smaller subpopulation will remain in the veins and help form the lymphovenous valves. This finding also indicates that *Prox1*-expressing venous ECs represent the pool of common progenitors that give rise to both LECs and lymphovenous valves. In the case of *Prox1* heterozygous embryos, the original number of *Prox1*-expressing venous ECs is reduced. This alteration leads to a reduction in the number of LEC progenitors and the lack of lymphovenous valves. Most likely, whether a *Prox1*-expressing EC becomes an LEC or a lymphovenous valve EC is determined by the process that regulates the exit of LEC progenitors from the veins; i.e., only those ECs that move out from the veins will become fully differentiated LECs.

One interesting question is why lymphovenous and venous valves, but not cardiac and collecting lymphatic valves, are dependent on *Prox1* dosage. One possible explanation could be that, as we previously showed, Coup-TFII is dispensable in fully differentiated LECs after E13.5 (Srinivasan et al. 2010), and, as collecting lymphatic valves develop after E16.5 (Norrmen et al. 2009), they are probably under the control of a different genetic program. Consistent with this proposal, we were unable to observe any significant Coup-TFII expression in the cardiac valve, which also develops normally in *Prox1* heterozygous embryos (Supplemental Fig. S11). Alternatively, blood flow is known to regulate endothelial gene expression (Dekker et al. 2002), and the valves in collecting lymphatics are not exposed to blood flow. Interestingly, in the cardiac valve, *Prox1* is ex-

pressed opposite to the flow side in a perfectly complementary pattern to that of *Klf2*, a flow-regulated gene (Lee et al. 2006). The above observation suggests that *Prox1* could be negatively regulated by shear force of blood flow. Consistent with this proposal, the lymphovenous and venous valves are directly exposed to blood flow and might require two copies of *Prox1* for their normal development and maintenance.

Our finding that the lymphovenous valve needed to connect the lymphatic and blood vasculatures is missing in *Prox1* heterozygous embryos, allowing lymph sacs to abruptly merge with nearby blood vessels or the pleural cavity, might partially explain the abnormal lymphangiography patterns observed in adult *Prox1* heterozygous mice (Harvey et al. 2005). Additionally, the importance of this valve in efficiently regulating the return of lymph fluid to the blood circulation and in preventing blood from entering the lymphatic vasculature is evident from our characterization of *Prox1* heterozygous embryos, a percentage of which had a blood-filled lymphatics phenotype. The total number of LECs could be the crucial determinant of whether blood can breach the lymph sacs, as we observed a significant reduction in the number of these cells in *Prox1*^{+/GFP-Cre;Coup-TFII^{+/f} embryos that displayed blood-filled lymphatics. This phenotype resembles the one previously described in *Syk*⁻, *Slp-76*⁻, and podoplanin-null embryos (Abtahian et al. 2003; Uhrin et al. 2010). These molecules have been shown to play an important role in regulating LEC–platelet interaction}

during lymphatic–blood separation. It could be possible that somehow these molecules also regulate the number of Prox1-expressing venous EC and LEC progenitors. In this case, we will assume that lymphovenous valves should also be absent in these mutant embryos.

We previously reported that there are two phases of *Prox1* regulation in LECs: an early initiation step dependent on Coup-TFII and Sox18, and a later maintenance step dependent on *Prox1* autoregulation (Srinivasan et al. 2010). Here we show that this autoregulation of *Prox1* expression is dose-dependent and that the total number of Prox1-expressing progenitors in the vein of *Prox1* heterozygous mice is reduced because the amount of available Coup-TFII/*Prox1* protein complex during the early stages of *Prox1* regulation is reduced. This may be because, although the concentration of Coup-TFII is at steady-state levels during LEC specification, *Prox1* is just beginning to be expressed during this stage and is therefore the only true variable involved in this protein–protein interaction kinetics. This, in combination with a weak residual interaction between the mutated *Prox1* and Coup-TFII, could be the reason for the milder phenotype observed in *Prox1*^{+/*NR*} and *Tie2-Cre;Coup-TFII*^{+/*f*} compared with that of *Tie2-Cre;Coup-TFII*^{+/*f*}; *Prox1*^{+/*NR*} embryos. It is possible that some transcription factor may compete with Coup-TFII/*Prox1* complexes in a dose-dependent manner to negatively regulate *Prox1* expression and thereby control the number of LEC progenitors that are generated in the veins. This scenario would be similar to that which occurs during angiogenesis when the correct number of tip versus stalk cells is determined by the Notch signaling activator VEGF-A and its inhibitor, Dll4 (Hellstrom et al. 2007). In the case of *VEGF-A* and *D114*, this balance goes awry in embryos heterozygous for either of these genes, indicating the importance of a delicate balance between two opposing pathways in this process (Hellstrom et al. 2007). Our results highlight a similar role for *Prox1* in determining lymphatic versus blood EC fates in a concentration-dependent manner during developmental lymphangiogenesis. Identifying and characterizing *Prox1*'s blood EC counterpart will provide further insight into the mechanisms that precisely control the blood EC versus LEC differentiation process.

In conclusion, we now provide a detailed molecular framework to understand the formation of the lymphovenous valves. Characterizing these valves in patients with primary lymphedema or other lymphatic alterations should offer further insight into the function of these structures in health and disease.

Materials and methods

Mice

The generation of *R26R*^{+/*LacZ*}, *Prox1*^{+/*LacZ*}, *Tie2-Cre*, *R26R*^{+/*YFP*}, *Coup-TFII*^{+/*f*}, *R26R*^{*mT/mG*}, *Prox1*^{+/*CreERT2*}, *Jojo-Prox1*, *Prox1*^{+/*NR*}, and *Prox1*^{+/*GFP**Cre*} mutant mice has been described (Soriano 1999; Wigle et al. 1999; Kisanuki et al. 2001; Srinivas et al. 2001; Takamoto et al. 2005; Muzumdar et al. 2007; Srinivasan et al. 2007, 2010; Lavado et al. 2010). All described mouse experiments

were approved by the St. Jude Children's Research Hospital Animal Care and Use Committee.

Antibodies

The primary antibodies used were rabbit anti-*Prox1* (AngioBio), goat anti-*Prox1* (R&D Systems), rat anti-PECAM1 (BD Pharmingen), rabbit anti-GFP (Molecular Probes), guinea pig anti-Lyve-1 (in-house), hamster anti-podoplanin (Hybridoma Bank), goat anti-Vegfr3 (R&D Systems), rabbit anti-vWF (DAKO), goat anti-Integrin- α 9 (R&D Systems), and rabbit anti-Foxc2 (in-house). The secondary antibodies used were Cy3-conjugated donkey anti-rabbit (Jackson ImmunoResearch Laboratories), Cy3-conjugated donkey anti-rat (Jackson ImmunoResearch Laboratories), Alexa 488-conjugated donkey anti-rat (Molecular Probes), Alexa 488-conjugated goat anti-hamster (Molecular Probes), Alexa 488-conjugated donkey anti-goat (Molecular Probes), Dylight 649-conjugated donkey anti-rabbit (Jackson ImmunoResearch Laboratories), DyLight 649-conjugated donkey anti-guinea pig (Jackson ImmunoResearch Laboratories), and Cy5-conjugated anti-rat (Jackson ImmunoResearch Laboratories).

Tamoxifen administration

Tamoxifen preparation and intraperitoneal injections of pregnant dams were performed as previously described (Srinivasan et al. 2007).

Immunohistochemistry

Fluorescent immunohistochemistry, X-gal staining, and horse-radish peroxidase staining using 3,3'-diaminobenzidine (DAB) as a substrate were performed as previously described (Srinivasan et al. 2007). The sections were mounted by using mounting medium containing DAPI (Vectashield), and confocal microscopy was performed as described previously (Srinivasan et al. 2010). To image the valves in the lymphatic vessels of the ear, adult wild-type and *Prox1*^{+/*GFP**Cre*}; *R26R*^{*mT/mG*} mice were perfused and the ears were collected and processed for whole mount as described previously (Harvey et al. 2005). Subsequently, wild-type ears were immunostained with an antibody specific for podoplanin, whereas the *Prox1*^{+/*GFP**Cre*}; *R26R*^{*mT/mG*} ears were visualized directly using the autofluorescence of the activated GFP reporter (Muzumdar et al. 2007). To image the valves in the collecting lymphatic vessels of the mesentery, guts of newborn *R26R*^{*mT/mG*} and *Prox1*^{+/*GFP**Cre*}; *R26R*^{*mT/mG*} pups were harvested and whole-mount immunohistochemistry was performed for Foxc2 or *Prox1*.

Statistical analysis

Microsoft Excel was used to evaluate the statistical significance by unpaired Student's *t*-test.

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