Lyons_Supplementary Data Fig 2

A

Sheep IgG
@Sheep-Dy647
PECAM1-Dy550

Rabbit IgG
@Rabbit-Dy647
PECAM1-Dy550

Streptavidin-Dy647
PECAM1-Dy550

B

Efnb2:GFP/WT
Efnb2:GFP
Prox1
Ephb4

C

PECAM1
Ephb4
Efnb2:GFP
Prox1

E18

FA

Superior region

Inferior region
<table>
<thead>
<tr>
<th>Family</th>
<th>Group</th>
<th>Sex, Age</th>
<th>Clinical History/examination</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Unrelated controls</td>
<td></td>
<td>No varicose veins</td>
</tr>
<tr>
<td>GLD &lt;sub&gt;UK&lt;/sub&gt; I.2</td>
<td>Mosaic, 37% in blood</td>
<td>M 56</td>
<td>Varicose veins, bilateral persistent peripheral lymphedema in the lower limbs since age 15.</td>
</tr>
<tr>
<td>GLD &lt;sub&gt;UK&lt;/sub&gt; II.2</td>
<td>Constitutive</td>
<td>F 36</td>
<td>Varicose veins, no clinical signs of persistent peripheral lymphedema, but lower limb lymphoscintigraphy showed bilateral impaired lymphatic drainage.</td>
</tr>
<tr>
<td>GLD &lt;sub&gt;NOR&lt;/sub&gt; II.3</td>
<td>Mosaic, 50%</td>
<td>F 39</td>
<td>Varicose veins since late teens (operated). No clinical signs of persistent peripheral lymphedema, but lower limb lymphoscintigraphy showed bilateral impaired lymphatic drainage.</td>
</tr>
<tr>
<td>GLD &lt;sub&gt;NOR&lt;/sub&gt; III.9</td>
<td>Constitutive</td>
<td>M 7</td>
<td>Mildly prominent (but not varicose) veins on posterior leg. No clinical signs of peripheral lymphedema.</td>
</tr>
<tr>
<td>GLD &lt;sub&gt;NOR&lt;/sub&gt; II.2</td>
<td>Mosaic, 13-31%</td>
<td>F 39</td>
<td>Varicose veins since age 23 (operated). No clinical signs of persistent peripheral lymphedema, but lower limb lymphoscintigraphy showed multiple tortuous lymphatic tracts.</td>
</tr>
<tr>
<td>GLD &lt;sub&gt;NOR&lt;/sub&gt; III.6</td>
<td>Unaffected relative (control)</td>
<td>M 9</td>
<td>Normal</td>
</tr>
</tbody>
</table>
Supplementary Table 1 Legend

Family identifiers refer to Refs (1, 2). The EPHB4 (NM_004444.4) mutation in GLD\textsubscript{UK} is c.2216G>A, p.Arg739Glu, and in GLD\textsubscript{NOR} it is c.2345T>G, p.Ile782Ser. GLD\textsubscript{NOR} II.2 and II.3 are monozygotic twins and mosaic carriers of the EPHB4 variant; GLD\textsubscript{NOR} II.3 was almost 50:50 wildtype to mutant variant (i.e. similar to a heterozygous constitutive carrier) in most tissues, but GLD\textsubscript{NOR} II.2 had a lower mutation load (13-31\% in the tissues measured) and an associated milder VV phenotype (i.e. higher mean number of valves than GLD\textsubscript{NOR} II.3, Fig. 1B).(1) GLD\textsubscript{UK} I.2 was previously identified as a constitutive mutation carrier, but has since been confirmed to be mosaic, with approximately 37\% mutation load in his blood.(2) Clinical history related to the venous and lymphatic phenotype is shown here, for a detailed clinical history the reader is referred to references.(1, 2)

Supplementary Methods

Human VV connexin histology:

For localization of connexins in human VVs (obtained from patients undergoing coronary artery bypass grafting) 10\(\mu\)m frozen sections were thawed and fixed in -20\ce{^\circ}c acetone prior to quenching of endogenous peroxidase using 3\%\ce{H2O2}, blocking (X0909, DAKO), incubation with primary antibodies, and amplification (MP-XCP, Menarini) according to the manufacturer’s instructions. Signal detection was with alkaline phosphatase (DAKO), and the counterstain was Nuclear Fast Red (Vector). Sections were photographed using a Micropublisher 3.3RTV camera mounted on a Leitz DMRB microscope. Primary antibodies were raised in rabbit to CX43 (Invitrogen 71-0700), CX47 (Sigma SAB2100924). Controls were incubated with non-immune rabbit IgG (R&D)
**Supplementary References**


**Supplementary Figure Legends**

**Supplementary Figure 1**

A) The mean number of VVs per vein is shown for the unrelated controls, and for each participant from the affected families. Each data point indicates a single vein (8 veins per participant). Mosaic carriers are arranged in order of approximate *EPHB4* mutation load, see Supplementary Table 1. (P<0.0001, ANOVA)

B) The mean duration of reflux is shown for each participant from the affected families. Each data point represents the left or right popliteal vein in that individual. A reflux duration >1s indicates severe deep venous reflux. A single functioning valve near the analysed venous segment may prevent reflux, which may explain why some veins in affected individuals did not exhibit reflux, despite significant reductions in the overall number of valves. Deep venous reflux was not assessed in all of the unrelated control population because it is rare and reference values are well established. (P=0.024, ANOVA)
C) The mean VV leaflet lengths are shown, for those VVs detected, for each participant and also, in (D), by genotype. No VVs were detected in GLDNORIII.9. P=ns for C and D. Data points represent individual valves. Green = control, Orange = mosaic for EPHB4 mutations, Red = heterozygous for EPHB4 mutation. Error bars indicate sem.

Supplementary Figure 2

A) Isotype immunofluorescence staining controls, using the indicated fluorophore-conjugated secondary antibodies, following incubation of samples with the appropriate non-immune IgG (for sheep and rabbit), or without primary antibody (for Streptavidin). The valve is outlined in the combined image (dotted white line). Arrowheads indicate residual autofluorescent erythrocytes in the vein lumen.

B) This figure relates to Fig.2D, and shows uncropped and unrotated images of 6µm z-projections to visualise an approximate single cell layer of the upper and lower regions of a valve. The regions of the valve reproduced in Fig.2D are indicated by dotted boxes.

C) A lower magnification image of Figure 2A (outlined by dotted box). FA = femoral artery. The site of a tributary is circled. Bars = 20µm

Supplementary Figure 3

A) Two adjacent representative TEM micrographs of part of an adult murine control valve leaflet, indicating the distribution of interstitial cells embedded within the matrix core. Endothelial cells are indicated by ‘ec’, extracellular matrix by ‘m’, and interstitial cells embedded in matrix by white arrowheads.
B) A further example of an interstitial cell nucleus embedded in matrix, surrounded by a layer of endothelial cells on each leaflet surface.

C) The mean number of interstitial cells identified per 10µm leaflet length is shown (N=100 TEM sections, from three levels of a VV at P6). Error bars indicate sem. (A-C are wildtype BALB/C mice)

D) Two adjacent TEM micrographs of part of an adult human great saphenous VV leaflet are shown, with the endothelial cell layer to bottom right, and the extensive matrix of the leaflet core to the top left of the images. Multiple morphologies of interstitial cells were identifiable, frequently extending laterally under the endothelial cell layer.

E) A further example of an adult human great saphenous VV leaflet. Images in D-E are representative of multiple sections obtained from four valves. Abbreviations as in A-B.

F) Light micrographs showing Connexin43 and Connexin47 immunolocalised to interstitial cells (arrowheads), as well as endothelial cells (ec) in adult human VV leaflet, but absent immunostain in isotype controls. NFR = Nuclear Fast Red counterstain.

Bars = 2µm in A-B, 1µm in D, 10µm in E, 20µm in F