







1 **Supplementary Table 1: Study participants**

Family	Group	Sex, Age	Clinical History/examination
Unrelated controls		M 25	No varicose veins
		F 30	
		F 24	
		M 24	
		M 37	
		F 42	
		F 42	
		F 53	
		F 58	
		M 69	
		M 23	
		M 66	
GLD _{UK} I.2	Mosaic, 37% in blood	M 56	Varicose veins, bilateral persistent peripheral lymphedema in the lower limbs since age 15.
GLD _{UK} II.2	Constitutive	F 36	Varicose veins, no clinical signs of persistent peripheral lymphedema, but lower limb lymphoscintigraphy showed bilateral impaired lymphatic drainage.
GLD _{NOR} II.3	Mosaic, 50%	F 39	Varicose veins since late teens (operated). No clinical signs of persistent peripheral lymphedema, but lower limb lymphoscintigraphy showed bilateral impaired lymphatic drainage.
GLD _{NOR} III.9	Constitutive	M 7	Mildly prominent (but not varicose) veins on posterior leg. No clinical signs of peripheral lymphedema.
GLD _{NOR} II.2	Mosaic, 13-31%	F 39	Varicose veins since age 23 (operated). No clinical signs of persistent peripheral lymphedema, but lower limb lymphoscintigraphy showed multiple tortuous lymphatic tracts.
GLD _{NOR} III.6	Unaffected relative (control)	M 9	Normal

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4 **Supplementary Table 1 Legend**

5 Family identifiers refer to Refs (1, 2). The *EPHB4* (NM_004444.4) mutation in GLD_{UK}
6 is c.2216G>A, p.Arg739Glu, and in GLD_{NOR} it is c.2345T>G, p.Ile782Ser. GLD_{NOR} II.2
7 and II.3 are monozygotic twins and mosaic carriers of the *EPHB4* variant; GLD_{NOR}
8 II.3 was almost 50:50 wildtype to mutant variant (i.e. similar to a heterozygous
9 constitutive carrier) in most tissues, but GLD_{NOR} II.2 had a lower mutation load (13-
10 31% in the tissues measured) and an associated milder VV phenotype (i.e. higher
11 mean number of valves than GLD_{NOR} II.3, Fig. 1B).(1) GLD_{UK} I.2 was previously
12 identified as a constitutive mutation carrier, but has since been confirmed to be
13 mosaic, with approximately 37% mutation load in his blood.(2) Clinical history related
14 to the venous and lymphatic phenotype is shown here, for a detailed clinical history
15 the reader is referred to references.(1, 2)

16

17 **Supplementary Methods**

18 *Human VV connexin histology:*

19 For localization of connexins in human VVs (obtained from patients undergoing
20 coronary artery bypass grafting) 10µm frozen sections were thawed and fixed in -
21 20°C acetone prior to quenching of endogenous peroxidase using 3% H₂O₂, blocking
22 (X0909, DAKO), incubation with primary antibodies, and amplification (MP-XCP,
23 Menarini) according to the manufacturer's instructions. Signal detection was with
24 alkaline phosphatase (DAKO), and the counterstain was Nuclear Fast Red (Vector).
25 Sections were photographed using a Micropublisher 3.3RTV camera mounted on a
26 Leitz DMRB microscope. Primary antibodies were raised in rabbit to CX43 (Invitrogen
27 71-0700), CX47 (Sigma SAB2100924). Controls were incubated with non-immune
28 rabbit IgG (R&D)

29

30 **Supplementary References**

- 31 1. Martin-Almedina S, et al. EPHB4 kinase-inactivating mutations cause
32 autosomal dominant lymphatic-related hydrops fetalis. *J Clin Invest.*
33 2016;126(8):3080-8.
- 34 2. Martin-Almedina S, et al. Janus-faced EPHB4-associated disorders: novel
35 pathogenic variants and unreported intrafamilial overlapping phenotypes.
36 *Genetics in medicine : official journal of the American College of Medical*
37 *Genetics.* 2021.

38

39 **Supplementary Figure Legends**

40 **Supplementary Figure 1**

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

45 B) The mean duration of reflux is shown for each participant from the affected
46 families. Each data point represents the left or right popliteal vein in that individual. A
47 reflux duration >1s indicates severe deep venous reflux. A single functioning valve
48 near the analysed venous segment may prevent reflux, which may explain why some
49 veins in affected individuals did not exhibit reflux, despite significant reductions in the
50 overall number of valves. Deep venous reflux was not assessed in all of the
51 unrelated control population because it is rare and reference values are well
52 established. (P=0.024, ANOVA)

53 C) The mean VV leaflet lengths are shown, for those VVs detected, for each
54 participant and also, in (D), by genotype. No VVs were detected in GLDNORIII.9.
55 P=ns for C and D. Data points represent individual valves.
56 Green = control, Orange = mosaic for *EPHB4* mutations, Red = heterozygous for
57 *EPHB4* mutation. Error bars indicate sem.

58

59 **Supplementary Figure 2**

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

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

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

71 Bars = 20µm

72

73 **Supplementary Figure 3**

74 A) Two adjacent representative TEM micrographs of part of an adult murine control
75 valve leaflet, indicating the distribution of interstitial cells embedded within the matrix
76 core. Endothelial cells are indicated by 'ec', extracellular matrix by 'm', and interstitial
77 cells embedded in matrix by white arrowheads.

78 B) A further example of an interstitial cell nucleus embedded in matrix, surrounded by
79 a layer of endothelial cells on each leaflet surface.

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

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

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

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

95 Bars = 2 μ m in A-B, 1 μ m in D, 10 μ m in E, 20 μ m in F