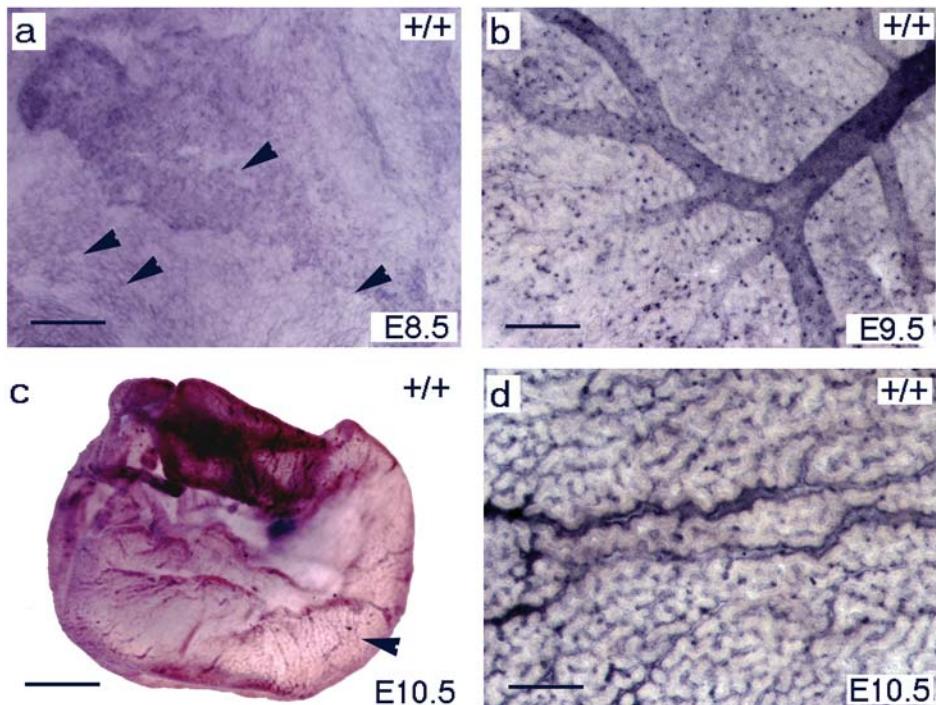


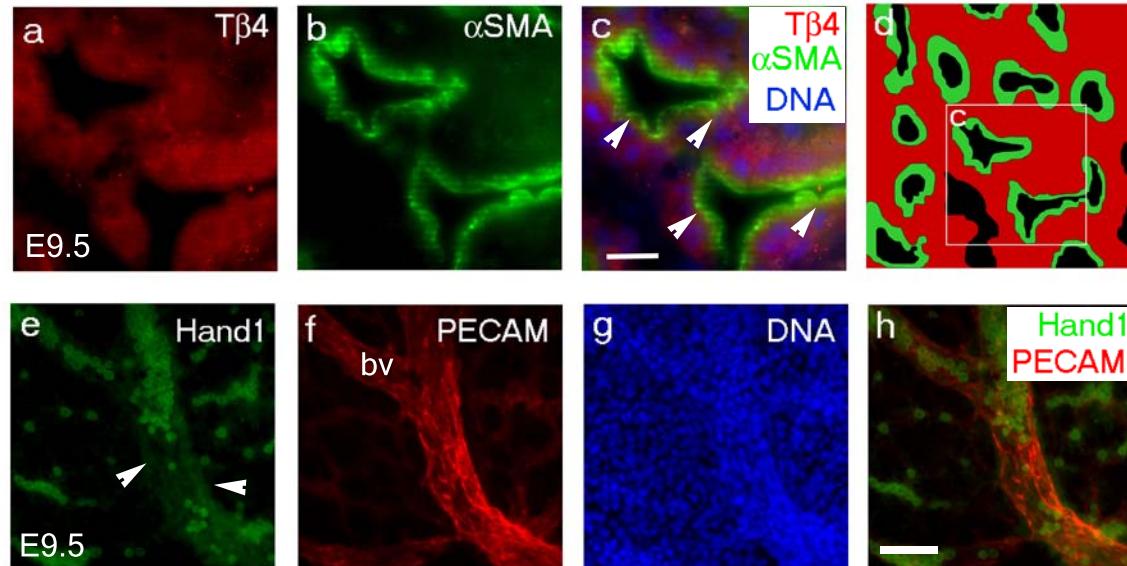
Supplementary Figure S1



***Tβ4* is expressed in the developing yolk sac vasculature**

In situ hybridisation revealed *Tβ4* expression in the blood islands of the yolk sac at E8.5 (**a**) and in the developing vessels at E9.5 (**b**) and E10.5 (**c, d**). Scale bars, **a, b**, 100 μm ; **c**, 250 μm ; **d**, 50 μm .

Supplementary Figure S2



TB4 is not expressed in smooth muscle cells within the developing yolk sac vasculature and Hand1 is down-regulated during vessel maturation

TB4 is not expressed in smooth muscle cells in the yolk sac vasculature at E9.5 (**a-c**) consistent with Hand1 expression at the same stage (see Fig. 4f, g); white arrowheads in **c** highlight α -SMA+ cells in green, lining TB4+ endothelium in red. Schematic of the representative region encompassing panel **c**, at lower magnification, to highlight plexus and network of vessels in red lined with green smooth muscle cells; black regions represents gaps between neighboring vessels and not individual vessel lumens (**d**). As the yolk sac endothelium matures with ongoing yolk sac angiogenesis Hand1 is down-regulated (**e-h**). Weak Hand1 immunostaining is highlighted in the vessel wall by white arrowheads (**e**). PECAM staining reveals elongated mature endothelial cells within the vessel wall (**f, h**). bv, blood vessel. Scale bars; **c**, 20 μ m; **h**, 10 μ m.

Supplementary Table S1. Primer sequences

Gene	Primer Sequence (5'-3')	
qRT-PCR PRIMERS		
Tβ4	Forward Reverse	ATGTCTGACAAACCGATATGGC CCAGCTTGCCTCTCTGTTCA
Hand1	Forward Reverse	TGAGTGCATCCCCAATGTG GCCAGCACGTCCATCAAGTA
Pecam	Forward Reverse	CTGCCAGTCCGAAAATGGAAC CTTCATCCACCAGGGGCTATC
Vegfa	Forward Reverse	GCACATAGAGAGAACATGAGCTTCC CTCCGCTCTGAACAAGGCT
Flt1	Forward Reverse	CTCAGACAAGTCAAACCTGGAG GGGAACATTCACTGGGTCCATAA
Tgfβ1	Forward Reverse	CTCCC GTGGCTTCTAGTGC GCCTTAGTTGGACAGGATCTG
TgfβR2	Forward Reverse	CCGCTGCATATCGCCTGTG AGTGGATGGATGGCCTATTACA
Smad2	Forward Reverse	ATGTCGTCCATCTGCCATT AACCGTCCTGTTCTTAGCTT
Smad5	Forward Reverse	TTGTT CAGAGTAGGAACTGCAAC GAAGCTGAGCAAACCTTGAT
Endoglin	Forward Reverse	CCCTCTGCCATTACCCCTG GTAAACGTCACCTCACCCCTT
Notch1	Forward Reverse	CCCTTGCTCTGCCTAACGC GGAGTCCTGGCATCGTTGG
Notch2	Forward Reverse	ATGTGGACGAGTGTCTGTTGC GGAAGCATAGGCACAGTCATC
Notch4	Forward Reverse	CTCTGCCACTCAATTCCCT TTGCAGAGTTGGGTATCCCTG
Hand1 GENOTYPING PRIMERS		
WT Forward (proximal exon 2)		CCACTAGGATCGCACGTGCA
Mutant Forward (neo ^r cassette)		GCAAAGCTGCTATTGGCCGC
Common Reverse (distal exon 2)		CCAGCAACGATTGGGAACGC