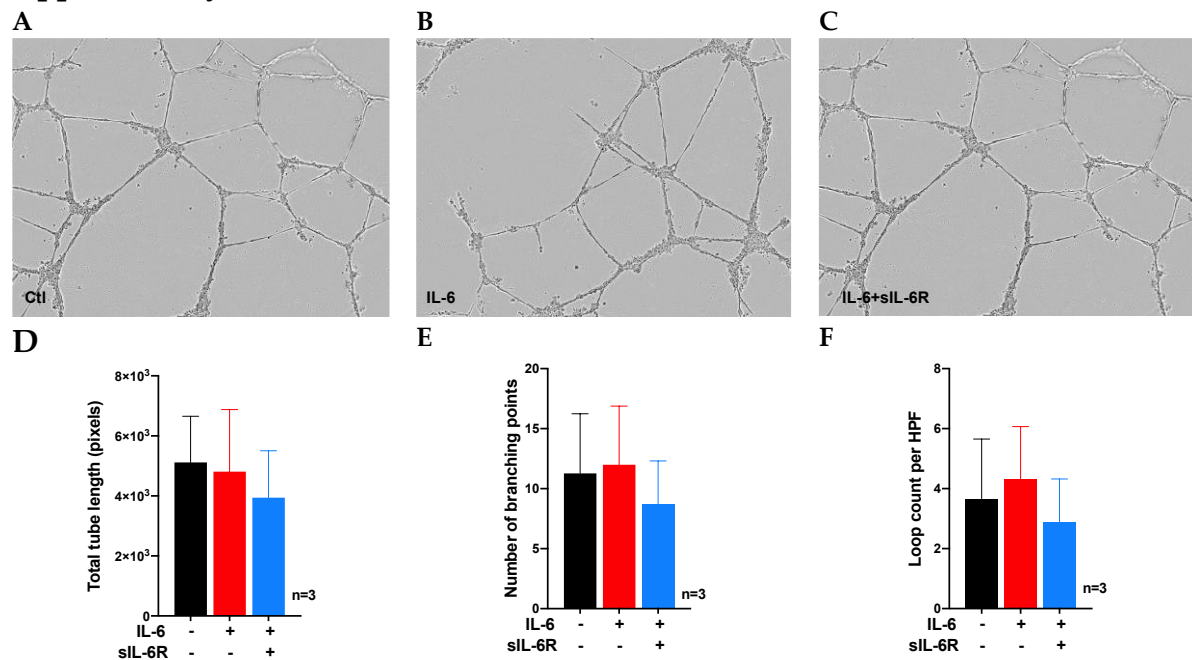
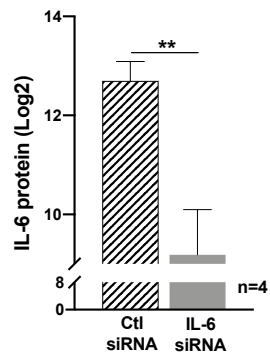


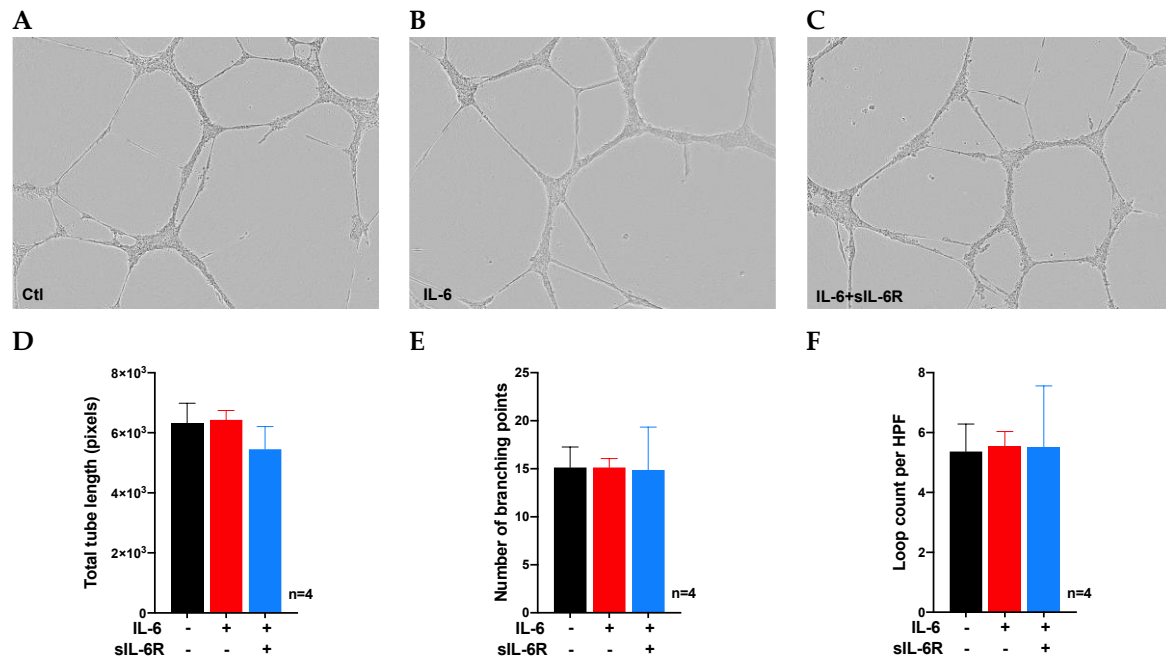
Supplementary data



Supplementary figure 1. The effect of IL-6 signaling in tube formation of vascular ECs when added to the Matrigel at the same time as the cells. Representative images from tube formation of (A) unstimulated ECs or (B) ECs treated with IL-6 alone or (C) in combination with sIL-6R. Quantification of the tube formation is presented as (D) total tube length, (E) number of branching points and loop count (F). Data is presented as mean $\pm$ SEM of 3 experiments each run-in duplicate.



Supplementary figure 2. Olink normalized protein expression (NPX) data on the expression of IL-6 after knockdown using stealth siRNAs. Data is presented as mean $\pm$ SEM of 4 experiments each run-in duplicate. **p < 0.01 compared to control.



Supplementary figure 3. Tube formation of vascular ECs which have been pre-treated with IL-6 alone or in combination with sIL-6R on growth-factor-reduced (GFR) Matrigel. Representative images from tube formation of (A) unstimulated ECs or (B) ECs treated with IL-6 alone or (C) in combination with sIL-6R. Quantification of the tube formation is presented as (D) total tube length, (E) number of branching points and loop count (F). Data is presented as mean $\pm$ SEM of 4 experiments each run-in duplicate.

Supplementary table 1. List of genes with Assay IDs in the angiogenesis array and respective $\Delta\Delta C_t$ values, p-values and BH-p-values.

Assay ID	Target Name	$\Delta\Delta C_t$ values	P-value	BH p-value
Hs00171042_m1	<i>CXCL10</i>	6,433	0,004	0,031
Hs00171467_m1	<i>SERPINF1</i>	5,929	0,020	0,065
Hs00300159_m1	<i>HGF</i>	5,002	0,003	0,031
Hs00236077_m1	<i>CEACAM1</i>	4,723	0,020	0,065
Hs00383235_m1	<i>PTN</i>	4,386	0,045	0,100
Hs00601975_m1	<i>CXCL2</i>	4,309	0,009	0,048
Hs00171022_m1	<i>CXCL12</i>	3,887	0,001	0,022
Hs00998026_m1	<i>PDGFRA</i>	3,401	0,012	0,052
Hs00427220_g1	<i>PF4</i>	2,885	0,022	0,065
Hs00900373_m1	<i>CHGA</i>	2,292	0,190	0,278
Hs00559786_m1	<i>ANGPTL1</i>	2,181	0,001	0,018
Hs01001469_m1	<i>ITGB3</i>	2,116	0,004	0,031
Hs00196470_m1	<i>ENPP2</i>	1,934	0,000	0,018
Hs00913333_m1	<i>TNNI1</i>	1,911	0,328	0,442
Hs00272659_m1	<i>LYVE1</i>	1,886	0,101	0,175
Hs00157317_m1	<i>TYMP</i>	1,140	0,060	0,126
Hs01105174_m1	<i>BAI1</i>	0,776	0,024	0,066
Hs01077958_s1	<i>IFNB1</i>	0,716	0,089	0,158
Hs00896294_m1	<i>PROX1</i>	0,685	0,018	0,063
Hs01922614_s1	<i>S1PR1</i>	0,651	0,162	0,253
Hs00199608_m1	<i>ADAMTS1</i>	0,582	0,133	0,218
Hs00176096_m1	<i>TEK</i>	0,544	0,017	0,063
Hs02379000_s1	<i>ANG</i>	0,431	0,102	0,175
Hs00187290_m1	<i>NRP2</i>	0,411	0,210	0,301
Hs00234042_m1	<i>PDGFB</i>	0,410	0,356	0,464
Hs00223332_m1	<i>TNMD</i>	0,392	0,735	0,793
Hs00765775_m1	<i>ANGPTL2</i>	0,388	0,138	0,222
Hs01568063_m1	<i>THBS2</i>	0,370	0,329	0,442
Hs00265254_m1	<i>FGF1</i>	0,263	0,068	0,132
Hs00170014_m1	<i>CTGF</i>	0,217	0,370	0,472
Hs00900054_m1	<i>VEGFA</i>	0,121	0,515	0,611
Hs00963711_g1	<i>GRN</i>	0,051	0,785	0,826
Hs00962914_m1	<i>THBS1</i>	0,049	0,922	0,945
Hs00171064_m1	<i>MDK</i>	0,040	0,627	0,714
Hs01011995_g1	<i>F2</i>	0,024	0,958	0,970
Hs01101127_m1	<i>ANGPTL4</i>	-0,015	0,970	0,970
Hs00826128_m1	<i>NRP1</i>	-0,065	0,732	0,793
Hs00176573_m1	<i>FLT1</i>	-0,118	0,163	0,253
Hs00611096_m1	<i>AMOT</i>	-0,130	0,381	0,473
Hs00176676_m1	<i>KDR</i>	-0,132	0,374	0,472
Hs00169777_m1	<i>PECAM1</i>	-0,133	0,646	0,725
Hs00181613_m1	<i>ANGPT1</i>	-0,143	0,750	0,798
Hs00166654_m1	<i>SERPINC1</i>	-0,199	0,280	0,389
Hs00234278_m1	<i>TIMP2</i>	-0,204	0,039	0,095
Hs00266645_m1	<i>FGF2</i>	-0,222	0,522	0,611
Hs00264877_m1	<i>PLG</i>	-0,245	0,895	0,929
Hs00233808_m1	<i>ITGAV</i>	-0,260	0,386	0,473
Hs00173634_m1	<i>VEGFB</i>	-0,282	0,044	0,100
Hs00993254_m1	<i>LECT1</i>	-0,356	0,663	0,735
Hs00153458_m1	<i>VEGFC</i>	-0,377	0,357	0,464
Hs00169867_m1	<i>ANGPT2</i>	-0,411	0,170	0,253
Hs00168433_m1	<i>ITGA4</i>	-0,423	0,083	0,154
Hs01549940_m1	<i>FN1</i>	-0,427	0,003	0,031
Hs00174344_m1	<i>CDH5</i>	-0,517	0,008	0,044
Hs00246256_m1	<i>FST</i>	-0,562	0,169	0,253
Hs00168405_m1	<i>IL12A</i>	-0,563	0,035	0,086
Hs00178500_m1	<i>TIE1</i>	-0,579	0,005	0,031
Hs01047677_m1	<i>FLT4</i>	-0,609	0,055	0,120

Hs00194179_m1	<i>HSPG2</i>	-0,648	0,015	0,060
Hs00165949_m1	<i>TIMP3</i>	-0,654	0,010	0,049
Hs00197064_m1	<i>FBLN5</i>	-0,667	0,022	0,065
Hs00189521_m1	<i>FIGF</i>	-0,676	0,123	0,206
Hs00208609_m1	<i>VASH1</i>	-0,703	0,021	0,065
Hs01098873_m1	<i>COL4A2</i>	-0,750	0,089	0,158
Hs00153304_m1	<i>CD44</i>	-0,868	0,066	0,132
Hs00266237_m1	<i>COL4A1</i>	-0,913	0,064	0,131
Hs00362096_m1	<i>EPHB2</i>	-0,936	0,016	0,063
Hs00181017_m1	<i>COL18A1</i>	-0,972	0,004	0,031
Hs00234422_m1	<i>MMP2</i>	-0,992	0,024	0,066
Hs99999918_m1	<i>TGFB1</i>	-1,016	0,001	0,022
Hs00188273_m1	<i>SEMA3F</i>	-1,151	0,030	0,081
Hs00270951_s1	<i>FOXC2</i>	-1,177	0,008	0,046
Hs00174781_m1	<i>EDIL3</i>	-1,248	0,003	0,031
Hs00266332_m1	<i>COL15A1</i>	-1,321	0,275	0,389
Hs00387364_m1	<i>PDGFRB</i>	-1,360	0,395	0,476
Hs00184728_m1	<i>SERPINB5</i>	-1,382	0,555	0,641
Hs99999083_m1	<i>CSF3</i>	-2,283	0,012	0,052
Hs00608187_m1	<i>TGFA</i>	-2,622	0,082	0,154
Hs00232618_m1	<i>HEY1</i>	-3,120	0,001	0,018
Hs00270802_s1	<i>TNFSF15</i>	-3,353	0,033	0,085
Hs00174103_m1	<i>IL8</i>	-3,457	0,042	0,099
Hs00174029_m1	<i>KIT</i>	-5,242	0,007	0,044
Hs99999901_s1	<i>18s</i>	HKG		
Hs99999905_m1	<i>GAPDH</i>	HKG_used for analyses		
Hs99999909_m1	<i>HPRT1</i>	HKG		
Hs99999908_m1	<i>GUSB</i>	HKG		
Hs00241027_m1	<i>FGA</i>	Not expressed		
Hs00168730_m1	<i>PRL</i>	Not expressed		
Hs00173564_m1	<i>FGF4</i>	Not expressed		
Hs00174877_m1	<i>LEP</i>	Not expressed		
Hs00260905_m1	<i>PROK1</i>	Not expressed		
Hs00174128_m1	<i>TNF</i>	Not expressed		
Hs00174143_m1	<i>IFNG</i>	Not expressed		
Hs00205581_m1	<i>ANGPTL3</i>	Not expressed		
Hs01022527_m1	<i>COL4A3</i>	Not expressed		
Hs00211115_m1	<i>ANGPT4</i>	Not expressed		