

**Supplementary information S2 (Table).**  
**Expression patterns of VEGF ligands, receptors and co-receptors**

<i>GENE</i>	<i>PREDOMINANT EXPRESSION</i>
<i>VEGFA</i>	Ubiquitous <sup>1,2</sup>
<i>VEGFB</i>	Muscle, heart <sup>3</sup>
<i>VEGFC</i>	Heart, lung, mesenchymal cells <sup>4</sup>
<i>VEGFD</i>	Heart, lung, skeletal muscle, colon, small intestine, mesenchymal cells <sup>5</sup>
<i>PLGF</i>	Placenta, tumour cells <sup>6,7</sup>
<i>VEGFR1</i>	Endothelial cells, monocytes, macrophages, megakaryocytes, smooth muscle cells, osteoclasts <sup>8,9</sup>
<i>sVEGFR1</i>	Placenta <sup>10</sup>
<i>VEGFR2</i>	Endothelial cells (highest expression during development), CD34/AC133-positive bone-marrow precursors, neuronal cells <sup>11</sup>
<i>VEGFR3</i>	Embryonic expression in vascular endothelial cells, fenestrated capillaries, tumour endothelial cells, lymphatic endothelial cells, monocytes, macrophages <sup>12,13</sup>
<i>Neuropilin-1</i>	Developing nervous and vascular system, arterial endothelial cells, many non-neuronal adult tissues, bone marrow stromal cells, tumour cells <sup>14-17</sup>
<i>Neuropilin-2</i>	Developing nervous and vascular system, venous and lymphatic endothelial cells, tumour cells <sup>16,18-20</sup>

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