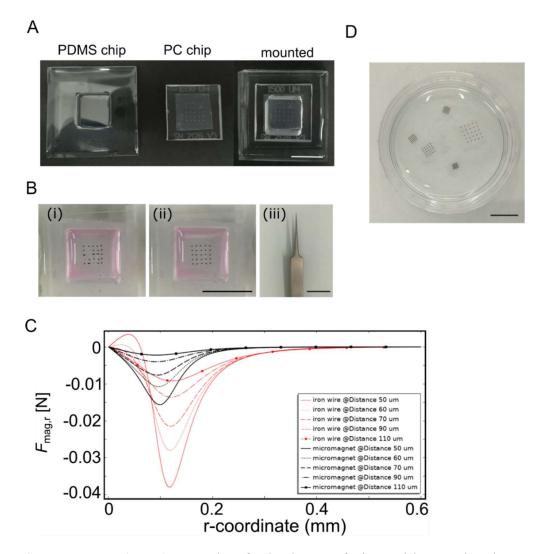
SUPPLEMENTARY FIGURES

A comprehensive toolkit for manipulation and analysis of sprouting capillary networks based on magnetic ordering of multiple EC-coated microcarriers and their use in tissue modelling and drug testing.

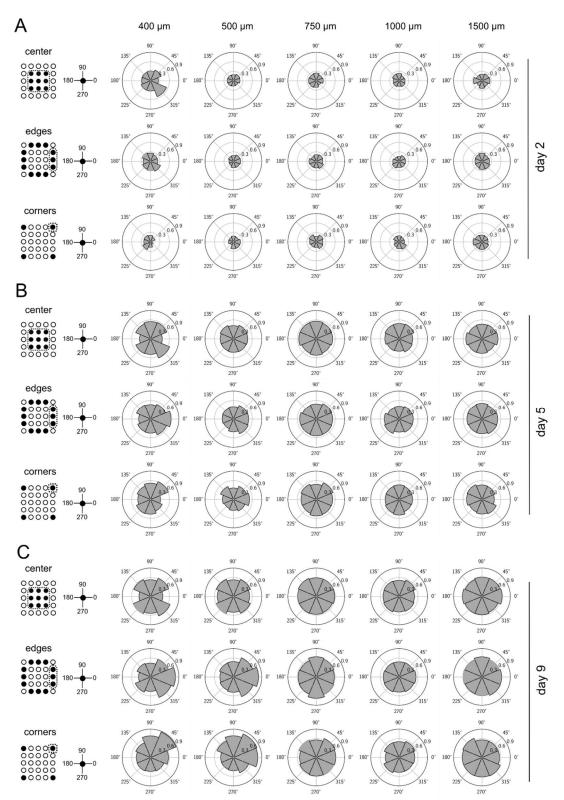
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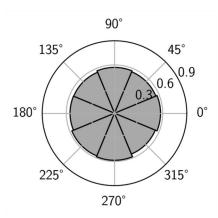
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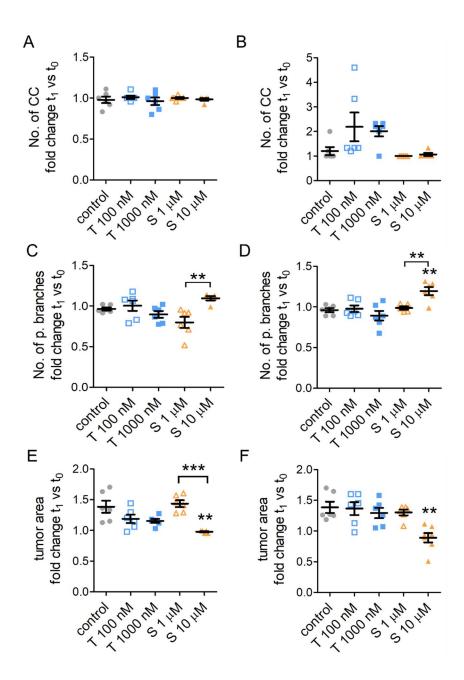
Supplementary Figure 1. Generation of ordered arrays of microparticles over the substrate using magnetic field patterning. A) Images of PDMS culture chamber, PC chip with embedded micromagnets and the assembled device. B) The magnetic microbeads that did not spontaneously migrate to the magnetic-field hotspots can be manually corrected using titanium fine-tip tweezers. C) Results of the numerical calculation of the magnetic force $F_{\rm mag,r}$ acting at a probe microparticle depending on its radial displacement r from the micromagnet axis for the experimentally relevant spacing $H = 90 \, \mu \text{m}$. Positive direction of the force is outward (i.e., negative values correspond to magnetic attraction). D) Isolated hydrogel sheets with embedded arrays of EC-coated microparticles in PBS in P35 Petri dish. Scale bars 1 cm.



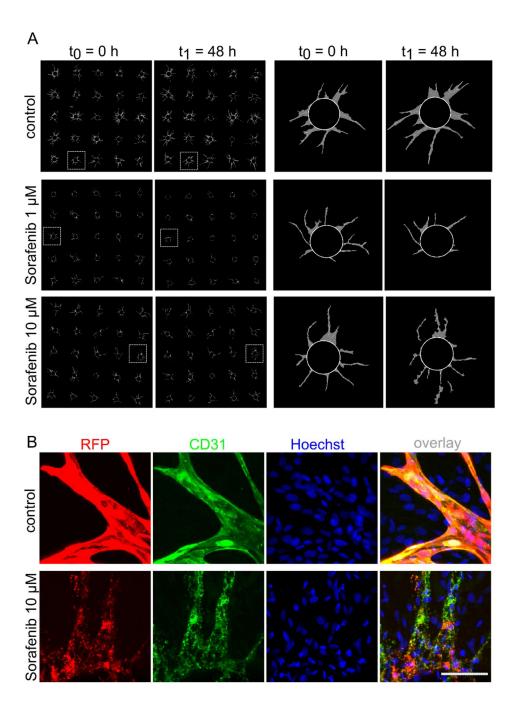
Supplementary Figure 2. Directionality analysis of the sprouting microcapillary networks. **A-C**) Radar charts illustrating the angular (azimuthal) distribution of the area occupied by capillaries at day (**A**) 2, (**B**) 5 and (C) 9 of culture. EC-coated microbeads were classified into 3 groups based on the number of nearest neighbors: center (8 neighbors), edge (5 neighbors) and corner (3 neighbors). Asymmetrical results from corner and edge regions were transformed to a common coordinate system, with all regions oriented to match the region indicated with a dotted frame (eg. top bar is transformed by -90 degrees, etc.). The numbers of the arrays in the assay (biological repetitions) were as follows: Spacing 400 μm, n = 3; 500 μm, n = 5; 750 μm, n = 7; 1000 μm, n = 6; 1500 μm, n = 7).



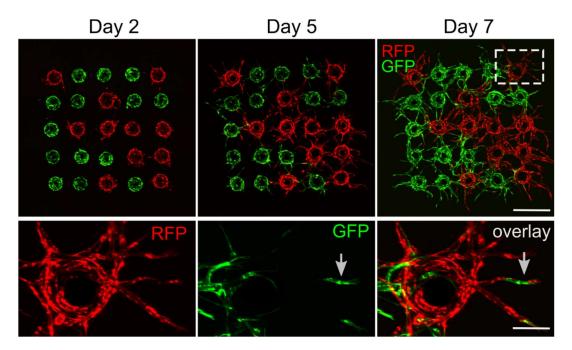
Supplementary Figure 3. Directionality analysis of the sprouting microcapillary networks formed around the central beads (3 x 3 subarray) seeded at $d = 500 \mu m$. Radar chart illustrating the angular (azimuthal) distribution of the area occupied by capillaries at day 7 of culture normalized by the area of the Voronoi-cell subsection in each octant. The number of the arrays in the assay (biological repetitions) n = 5.



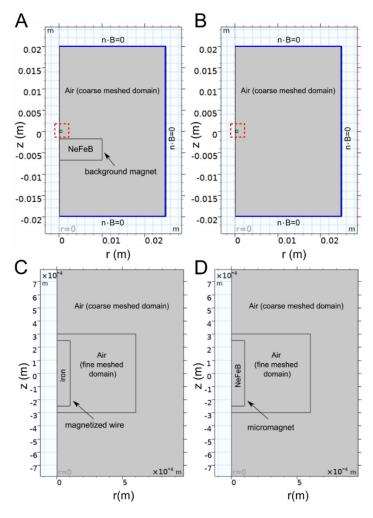
Supplementary Figure 4. Morphometric analysis of the fold-change in (**A,B**) the number of connected components (CC), (**C, D**) the number of primary branches and (**E,F**) the tumor area at time t_1 relative to t_0 for the arrays of HUVEC-coated microbeads with (**A, C, E**) d=1500 µm and (**B, D, F**) d=500 µm spacing cocultured with HeLa cells and exposed to anticancer drugs. Spacing 500 µm: Control, n=6; Taxol (T) 1000 nM, n=6; Sorafenib (S) 1 µM, n=5; Sorafenib (S) 10 µM, n=6. Spacing 500 µm: Control, n=6; Taxol (T) 100 nM, n=6; Taxol (T) 1000 nM, n=6; Sorafenib (S) 1 µM, n=6; Sorafenib (S) 10 µM, n=6; Sorafenib



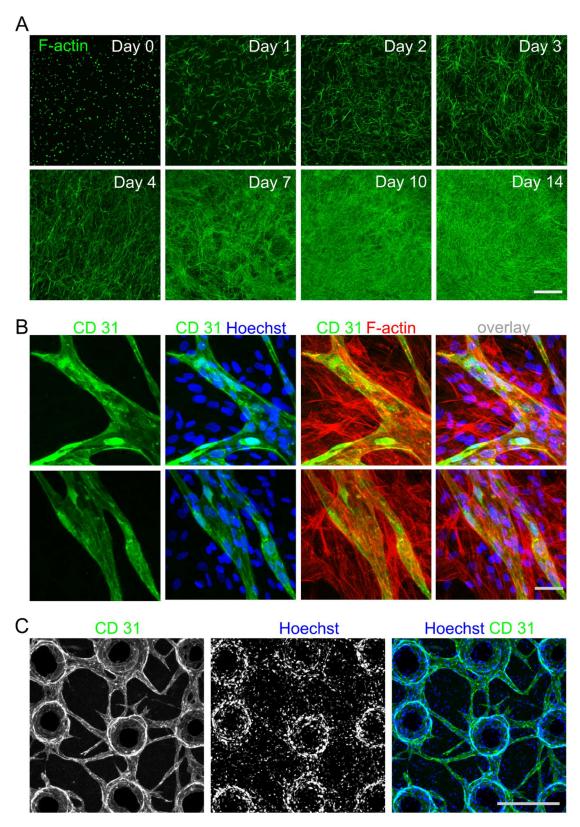
Supplementary Figure 5. Effect of Sorafenib treatment of vasculature integrity. **A)** Skeletons of the arrays of fluorescently tagged HUVEC- coated microbeads co-cultured with HeLa cells (not visualized) and seeded at $d = 1500 \,\mu\text{m}$. At day 5 Sorafenib (1 or 10 μM) was administered to the structures; t_0 – day 5 of culture, before drug exposure; t_1 – day 7 of culture, 48 h after drug application. High magnification images (right panels) show selected areas of the arrays just before drug application (t_0) and 48 h after drug exposure (t_1). Please not fragmentation of skeleton after 10μM Sorafenib treatment (bottom panel). **B)** High magnification confocal images of representative areas of control array and an array treated with 10 μM Sorafenib. RFP (red) was used to visualize HUVECs. Tight junctions were immunostained with anti-CD31 antibody (green). Hoechst was used to visualize nuclei. Scale bar 100 μm.



Supplementary Figure 6. An array of GFP and RFP-tagged HUVEC-coated microbeads seeded at $d=500~\mu m$ and imaged at days 2, 5 and 7 of culture. Bottom panel shows high magnification images of the selected area. Arrow indicates a single GFP tagged cell that migrated from the GFP-HUVEC coated microbead to the distal part of RFP-HUVEC coated microbead. Scale bar 500 μm (upper panel) and 100 μm bottom panel.



Supplementary Figure 7. Schematic representation of (**A**) Geometry 1 and (**B**) Geometry 2 showing the air (coarse meshed domain), and the region of interest (red dashed box). At the external boundaries of the system (marked with blue color), a 50 μm geometry layer was added to serve as an artificial domain with a physical width equal to 47 m. In (**A**) there is a background magnet placed beneath ROI. (**C**,**D**) Zoom-in of ROI (**A**) and (**B**) respectively. Air domain around the micromagnet/iron wire is meshed with smaller elements.



Supplementary Figure 8. Confocal images illustrating network of fibroblasts surrounding microcapillaries. **A)** NHDFs stained with F-actin (green) and visualized at the time of seeding the array (day 0) and after 1, 2, 3, 4, 7, 10 and 14 days of culture. **B)** High magnification images of microcapillaries and surrounding them NHDFs acquired at day 7 of culture. HUVECs were immunostained with anti CD31 antibody (green). All cells were stained with Hoechst dye (blue) to visualize nuclei and F-actin (red). **C)** Array of EC-coated microbeads co-cultured with NHDFs, 7 day of culture. HUVECs were immunostained with anti CD31 antibody, Hoechst dye was used to visualize nuclei. Scale bar 500 μm in A and C, 50 μm in B.