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Paracrine vs Autocrine Regulation of Early Vascular Patterning

During embryonic vasculogenesis, the earliest mechanism of blood vessel morphogenesis, isolated vascular cell progenitors called angioblasts assemble into a characteristic network-like pattern. So far, however, the mechanisms underlying the coalescence and patterning of angioblasts remain unclear.

We consider a hybrid cell-based approach similar to that used for a similar *in vitro* process [1,2]. However, contrary to previous mathematical models that assume chemotaxis towards an autocrine signal [1,2,3,4], we favour an alternative mechanism based on matrix-binding of paracrine signals. Detailed morphometric analysis of simulated vascular networks and confocal microscopy images obtained from *in vivo* quail embryos reveals our model can reproduce the vascular patterns with high accuracy.

The work to be reported has been made in collaboration with W. de Back, J. Starruf and A. Deutsch (Center for High Performance Computing, Technische Universität Dresden), M. A. Herrero (Department of Applied Mathematics and IMI, Universidad Complutense de Madrid) and A. Mattiotti and J. M. Pérez-Pomares (Laboratory of Cardiovascular Development and Angiogenesis, Universidad de Málaga).

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