

SUPPLEMENTARY FIG. S3. Cell orientation and spatial correlation length. (a) A large mosaic image of the cell colony was divided into smaller subimages. Scale bar: 500 μ m. In each subimage, the principal orientation (*red arrows*) of the cells was found using an orientation algorithm described in the Methods section. Then, the directional autocorrelation between any two subimages was calculated. If the orientation is parallel, DA(R)=1, while if it is orthogonal, DA(R)=-1. (b) The directional autocorrelation was then plotted in terms of the distance between the two subimages, R. If the average value of DA(R) is 0, and the standard deviation of DA(R) is large, then at a distance R, cell orientation is the most uncorrelated since DA(R) is randomly distributed.