

Supplemental figures for case (ii-a)

This SI shows results from simulations where a continuous drug infusion during 24 h is assumed. Therefore the blood borne drug concentration is given by $s^v = 1$ for $t < 24 h$ and 0 else.

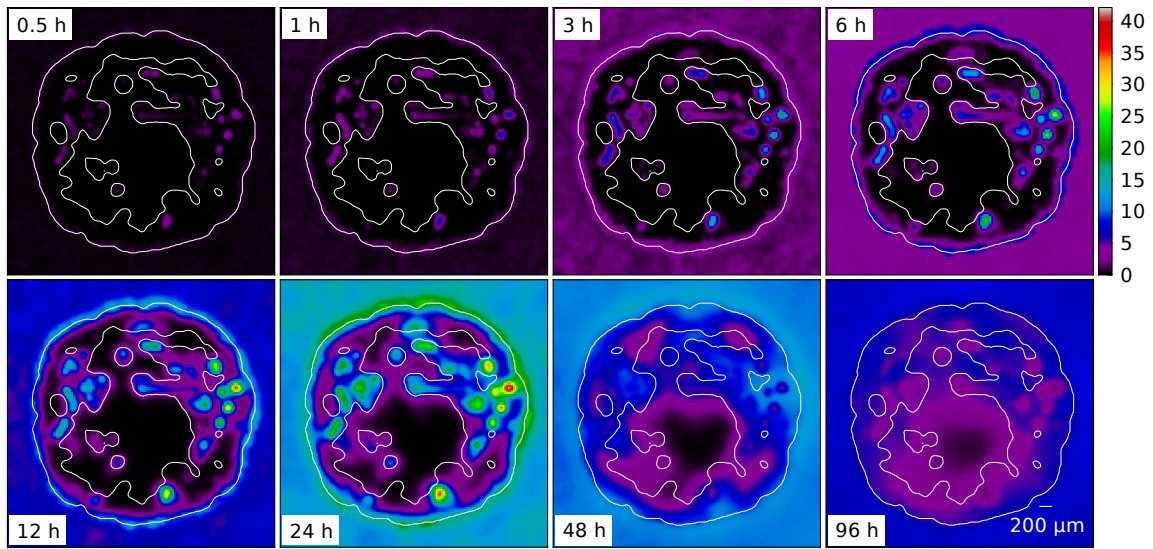


Figure 1: Drug distribution s in a series of snapshots. Corresponds to Figure 7 in the paper.

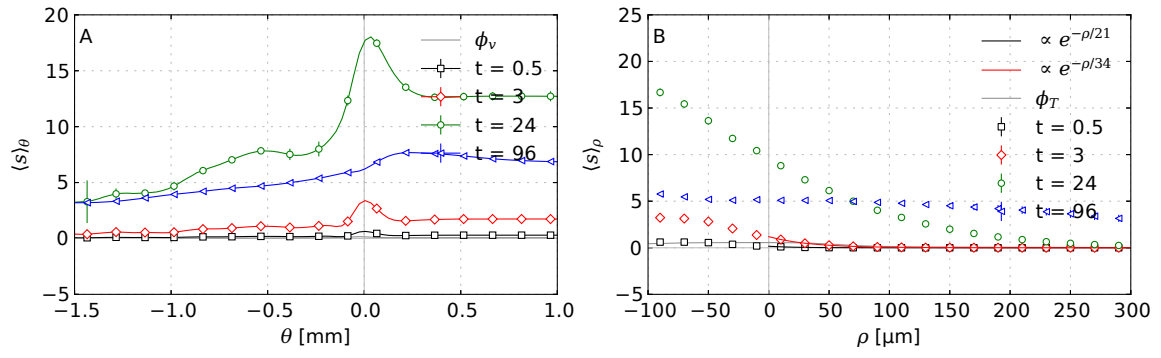


Figure 2: **Drug concentration profiles at different times.** (A) plotted vs. θ , and (B) vs. distance from vessels ρ . Corresponds to Figure 8 in the paper.

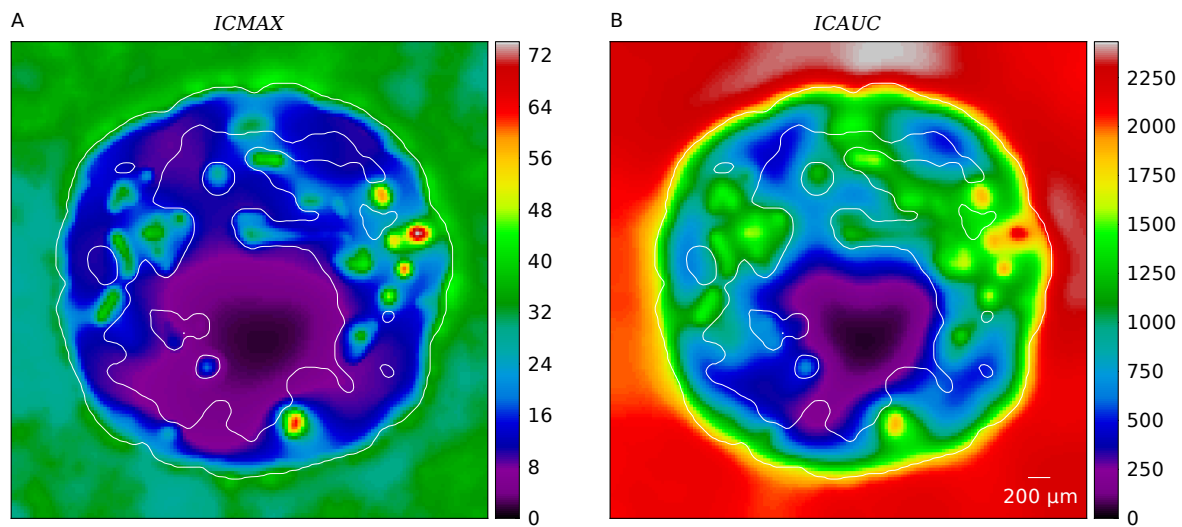


Figure 3: **Spatial distribution of drug exposure metrics.** (A) maximum concentration ICMAX and (B) the AUC ICAUC, taken from a slice through the origin of the system. Corresponds to Figure 9 in the paper.

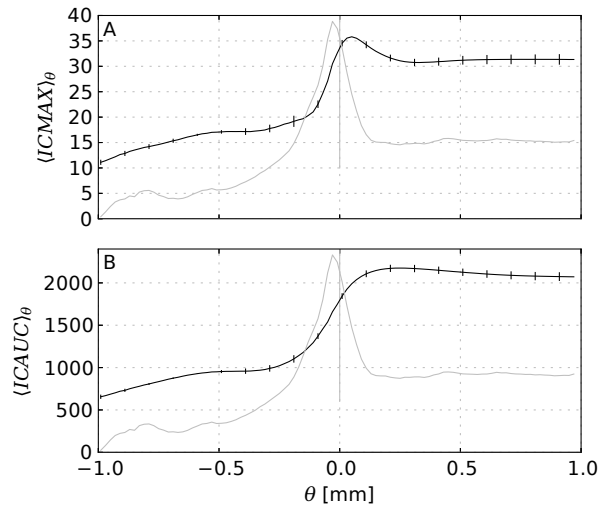


Figure 4: **Drug exposure metrics profiles.** Maximal concentration IC_{MAX} (A) and area under curve IC_{AUC} (B) plotted vs. θ . Corresponds to Figure 10 in the paper.

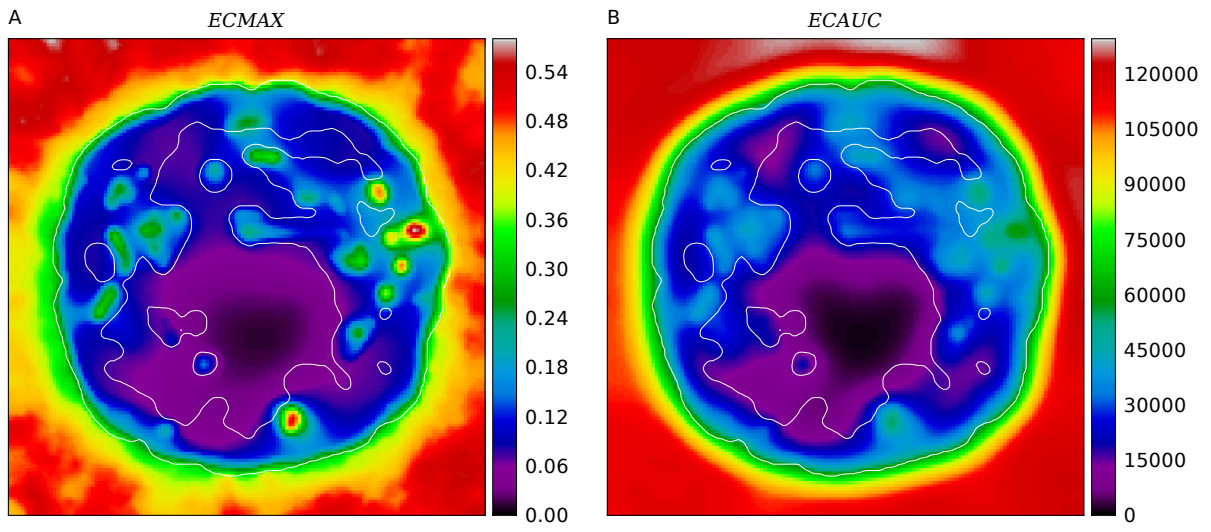


Figure 5: Maximal concentration (A) and area under curve (B) for the concentration in the interstitial compartment.

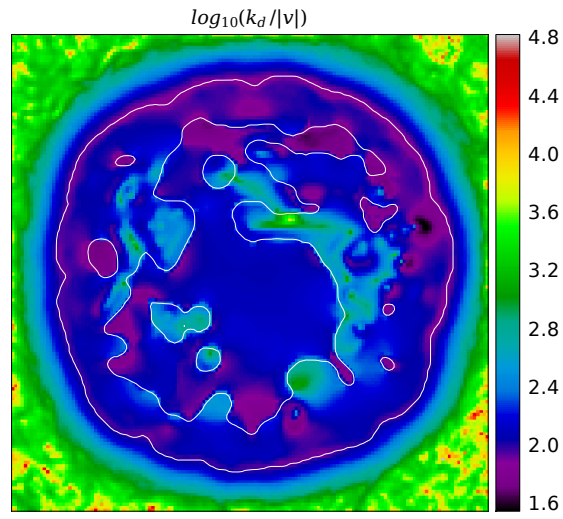


Figure 6: **Logarithmic plot of the length scale L_{dc} .** It is defined by $L_{dc} = k_d/|v|$ following the requirement that the Peclet number equals one, i.e. $1 = Pe = L_{dc}|v|/k_d$. The data is scaled logarithmically.