

| Environment features | Surroundings nutrient concentration (M) | Nutrient diffusivity (dm ² s ⁻¹) | Volume of environment | | | |
|----------------------------|--|---|--|---|---|---|
| | 10 ⁻² | 10 ⁻⁷ | 1mm ³ | | | |
| Gene network features | Transcription speed (s ⁻¹) | Translation speed (s ⁻¹) | Basal polymerase binding probability | Polymerase binding probability with activator | Ploymerase binding probability with repressor | Sizer degradation rate (s ⁻¹) |
| | 10 ⁻¹ | 10 ⁻¹ | 10 ⁻² | 0.5 | 0 | |
| | Timestep for TF dynamics (s.) | k _{on} BSTF (M ⁻¹ s ⁻¹) | k _{off} BSTF (s ⁻¹) | mRNA degradation rate (s ⁻¹) | Protein degradation rate (s ⁻¹) | |
| | 30 | 10 ² | 10 ⁵ | 10 ⁻³ | 10 ⁻³ | |
| Energy processing features | Nutrient to energy rate | ATP basal cost per V=1m3 (ATP units/μm ³) | Basal metabolism allometry coefficient | Growth cost of one cascade | Nutrient cost of one cascade | 0.2 |
| | 30 | 10 ⁶ | 5:6 | 10 ³ | 10 ² | |
| Cell features | Mutation rate per base (per gen.) | Daughter cells ratio | Lethal cell volume | Basal death rate | | |
| | 10 ⁻⁴ | 1:1 | 1 | 0.25 | | |
| Initial conditions | Initial sizer threshold | Initial cell volume (μm ³) | Initial number of transcripts | Initial number of cells | Theoretical simulation duration (s.) | |
| | 100 | 30 | 5 | 500 | 10 ⁷ | |