

A multiscale model of complex endothelial cell dynamics in early angiogenesis

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Parameter	Units	Description	Value used in simulations	Ref.
R_s	μm	Interaction radius.	15	estim., [1, 2]
b_N	$\text{molec} \cdot \text{time}^{-1}$	Baseline Notch receptor expression.	500	[3, 4]
b_D	$\text{molec} \cdot \text{time}^{-1}$	Baseline Delta ligand expression.	800	[3, 4]
b_{R2}	$\text{molec} \cdot \text{time}^{-1}$	Baseline VEGFR2 expression.	800	[4]
I_0	molec	Activation threshold for NICD.	100	[4]
$R2^*_0$	molec	Activation threshold for activated VEGFR2.	200	[4]
$\lambda_{I,N}$	<i>dimensionless</i>	Weight factor characterising fold change of the production rate of Notch receptor depending on the NICD concentration.	4.0	[3, 4]
$\lambda_{R2^*,D}$	<i>dimensionless</i>	Weight factor characterising fold change of the production rate of Delta ligand depending on the activated VEGFR2 concentration.	2.0	[4]
$\lambda_{I,R2}$	<i>dimensionless</i>	Weight factor characterising fold change of the production rate of VEGFR2 depending on the NICD concentration.	0.0	[4]
n_N	<i>dimensionless</i>	Cooperativity parameter for Hill function for NICD-dependent Notch up-regulation.	2	[5]
n_D	<i>dimensionless</i>	Cooperativity parameter for Hill function for activated VEGF-dependent Delta up-regulation.	1	[5]
n_{R2}	<i>dimensionless</i>	Cooperativity parameter for Hill function for NICD-dependent VEGFR2 repression.	1	[5]
V	molec	External VEGF.	2500 (Fig 3E); 0 – 2500 (Fig 3F); {0, 2500, 25000} (in the rest of the simulations)	[3, 4]
D_{ext}	molec	External Delta ligand.	0 – 3000 (Fig 3E and 3F); calculated from adjacent cells (in the rest of the simulations)	[3, 4]
N_{ext}	molec	External Notch receptor.	1000 (Fig 3E and 3F); calculated from adjacent cells (in the rest of the simulations)	[3, 4]
k_t	$\text{molec}^{-1} \cdot \text{time}^{-1}$	Trans-binding rate for Notch receptor and Delta ligand.	$5.0e - 5$	[5]
k_c	$\text{molec}^{-1} \cdot \text{time}^{-1}$	Cis-interaction rate for Notch receptor and Delta ligand.	$6.0e - 4$	[5]
k_v	$\text{molec}^{-1} \cdot \text{time}^{-1}$	Binding rate for VEGFR2 and external VEGF.	$5.0e - 5$	[4]
η	<i>dimensionless</i>	Endocytic regulation of Notch signalling.	0.5	estim., [6]
γ	time^{-1}	Degradation rate of proteins.	0.1	[4]
γ_e	time^{-1}	Degradation rate of activated receptors.	0.5	[4]

S1 Table. Baseline parameter values for the VEGF-Delta-Notch subcellular model. Description and reference values used in simulations of the subcellular VEGF-Delta-Notch signalling.

References

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