



Differential Equation System “CELL DEATH”

1 Rate Laws

1.1 Reaction: re2

$$v_1 = [s_7] \cdot k_{B0034} \quad (1)$$

1.2 Reaction: re3

$$v_2 = \frac{k_1 \cdot K_m^{hill}}{K_m^{hill} + \left(\frac{[s_{13}]}{vol \cdot N}\right)^{hill}} \cdot [s_3] \quad (2)$$

1.3 Reaction: re4

$$v_3 = [s_{29}] \cdot k \quad (3)$$

1.4 Reaction: re5

$$v_4 = [s_{13}] \cdot [s_{23}] \cdot k - [s_{15}] \cdot k \quad (4)$$

1.5 Reaction: re7

$$v_5 = [s_{13}] \cdot d \quad (5)$$

1.6 Reaction: re8

$$v_6 = [s_{15}] \cdot d \quad (6)$$

1.7 Reaction: re9

$$v_7 = [s_{29}] \cdot d \quad (7)$$

1.8 Reaction: re10

$$v_8 = [s_{28}] \cdot d \quad (8)$$

1.9 Reaction: re17

$$v_9 = d \cdot [s_{23}] \quad (9)$$

1.10 Reaction: re1

$$v_{10} = k \cdot [s_2] \quad (10)$$

1.11 Reaction: re11

$$v_{11} = d \cdot [s_7] \quad (11)$$

1.12 Reaction: re21

$$v_{12} = [s_{23}] \cdot [s_{26}] \cdot k \quad (12)$$

1.13 Reaction: re22

$$v_{13} = [s_{30}] \cdot k \quad (13)$$

1.14 Reaction: re23

$$v_{14} = [s_{30}] \cdot k \quad (14)$$

2 Equations

2.1 Species: s1 (transcriptie LuxR)

$$\frac{d[s_1]}{dt} = -v_{10} \quad (15)$$

2.2 Species: s2 (gen LuxR)

$$\frac{d[s_2]}{dt} = 0 \quad (16)$$

2.3 Species: s3 (gen_ccdB)

$$\frac{d[s_3]}{dt} = 0 \quad (17)$$

2.4 Species: s4 (translatie LuxR)

$$\frac{d[s_4]}{dt} = -v_1 \quad (18)$$

2.5 Species: s6 (transcriptie_ccdB)

$$\frac{d[s_6]}{dt} = -v_2 \quad (19)$$

2.6 Species: s7 (mRNA LuxR)

$$\frac{d[s_7]}{dt} = -v_{11} + v_{10} \quad (20)$$

2.7 Species: s8 (sa37_degraded)

$$\frac{d[s_8]}{dt} = v_{11} \quad (21)$$

2.8 Species: s9 (sa44_degraded)

$$\frac{d[s_9]}{dt} = v_7 \quad (22)$$

2.9 Species: s10 (sa38_degraded)

$$\frac{d[s_{10}]}{dt} = v_5 \quad (23)$$

2.10 Species: s12 (translatie_ccdB)

$$\frac{d[s_{12}]}{dt} = -v_3 \quad (24)$$

2.11 Species: s13 (C0062_LuxR)

$$\frac{d[s_{13}]}{dt} = -v_5 - v_4 + v_1 \quad (25)$$

2.12 Species: s14 (sa45_degraded)

$$\frac{d[s_{14}]}{dt} = v_8 \quad (26)$$

2.13 Species: s15 (HSL_LuxR_complex)

$$\frac{d[s_{15}]}{dt} = -v_6 + v_4 \quad (27)$$

2.14 Species: s23 (HSL)

$$\frac{d[s_{23}]}{dt} = -v_{12} - v_9 - v_4 + v_{14} + v_6 \quad (28)$$

2.15 Species: s24 (sa23_degraded)

$$\frac{d[s_{24}]}{dt} = v_9 \quad (29)$$

2.16 Species: s26 (lactonase)

$$\frac{d[s_{26}]}{dt} = -v_{12} + v_{14} + v_{13} \quad (30)$$

2.17 Species: s25 (hydroxyacid)

$$\frac{d[s_{25}]}{dt} = v_{13} \quad (31)$$

2.18 Species: s28 (P1010_CcdB)

$$\frac{d[s_{28}]}{dt} = -v_8 + v_3 \quad (32)$$

2.19 Species: s29 (mRNA_CcdB)

$$\frac{d[s_{29}]}{dt} = -v_7 + v_2 \quad (33)$$

2.20 Species: **s30 (EnzymeComplex)**

$$\frac{d[s_{30}]}{dt} = -v_{14} - v_{13} + v_{12} \tag{34}$$

2.21 Species: **s16 (HSL)**

$$\frac{d[s_{16}]}{dt} = 0 \tag{35}$$

2.22 Species: **s17 (C0062_LuxR)**

$$\frac{d[s_{17}]}{dt} = 0 \tag{36}$$

3 Parameters

Parameter	Value
<i>vol</i>	7.0E-16
<i>N</i>	6.023E23

4 Species

Species	Initial concentration	compartment
<i>s</i> ₁	0.0	default
<i>s</i> ₂	1.0	default
<i>s</i> ₃	1.0	default
<i>s</i> ₄	0.0	default
<i>s</i> ₆	0.0	default
<i>s</i> ₇	0.0	default
<i>s</i> ₈	0.0	default
<i>s</i> ₉	0.0	default
<i>s</i> ₁₀	0.0	default
<i>s</i> ₁₂	0.0	default
<i>s</i> ₁₃	0.0	default
<i>s</i> ₁₄	0.0	default
<i>s</i> ₁₅	0.0	default
<i>s</i> ₂₃	0.0	default
<i>s</i> ₂₄	0.0	default
<i>s</i> ₂₆	0.0	default
<i>s</i> ₂₅	0.0	default
<i>s</i> ₂₈	0.0	default
<i>s</i> ₂₉	0.0	default
<i>s</i> ₃₀	0.0	default
<i>s</i> ₁₆	4.9E-324	
<i>s</i> ₁₇	4.9E-324	

5 Compartments

Compartment	Volume
<i>default</i>	1.0