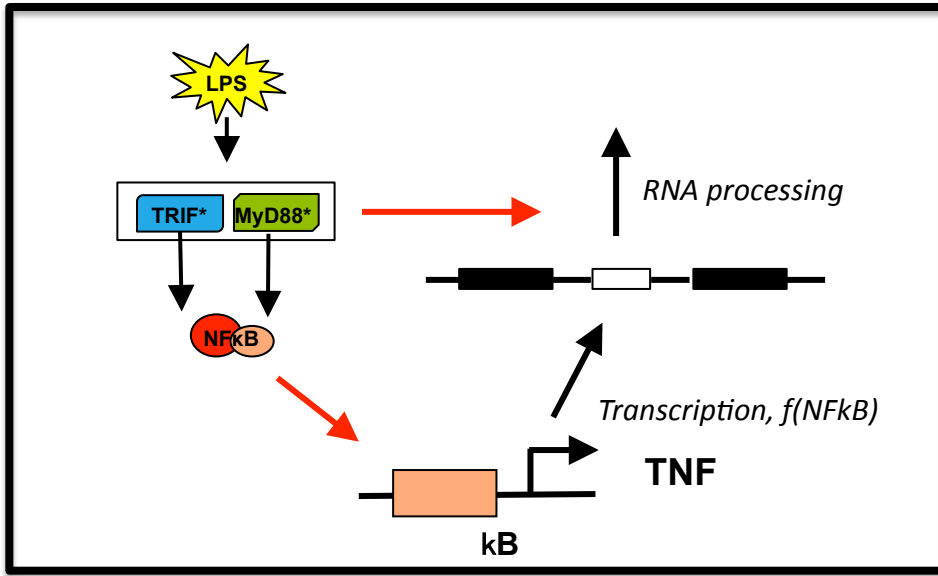
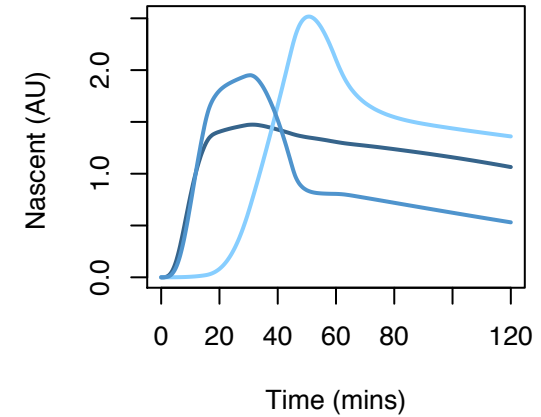


Module1 : Transcription + RNA processing



Simulation



Equation

$$\frac{d[Nascent]}{dt} = V_{tr} \frac{[NF\kappa B_n]^{n_H}}{[NF\kappa B_n]^{n_H} + K_{dir}^{n_H}} - k_{pr}[Nascent]$$

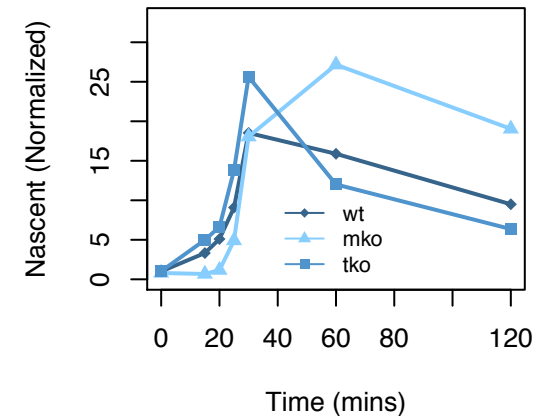
Initial condition

$$[Nascent]_{t=0} = \frac{V_{tr}}{k_{pr}} \frac{[NF\kappa B_n]_{t=0}^{n_H}}{[NF\kappa B_n]_{t=0}^{n_H} + K_{dir}^{n_H}}$$

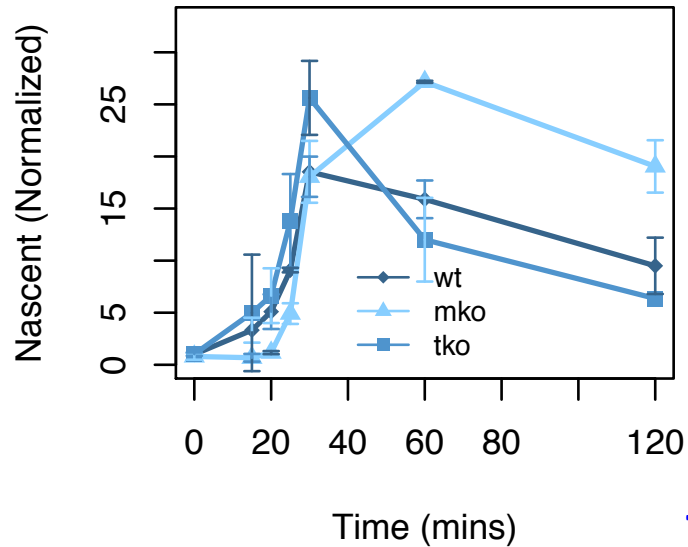
Parameters ($n_p=5$)

$$V_{tr} = 1(\text{fixed}), K_{dir} = 0.65, n_H = 2, k_{pr}^{wt} = 0.4 \text{ min}^{-1}, k_{pr}^{mko} = \frac{k_{pr}^{wt}}{4.2}, k_{pr}^{tko} = \frac{k_{pr}^{wt}}{1.5}$$

Experiment



Experiment



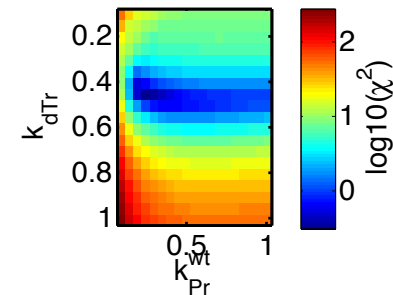
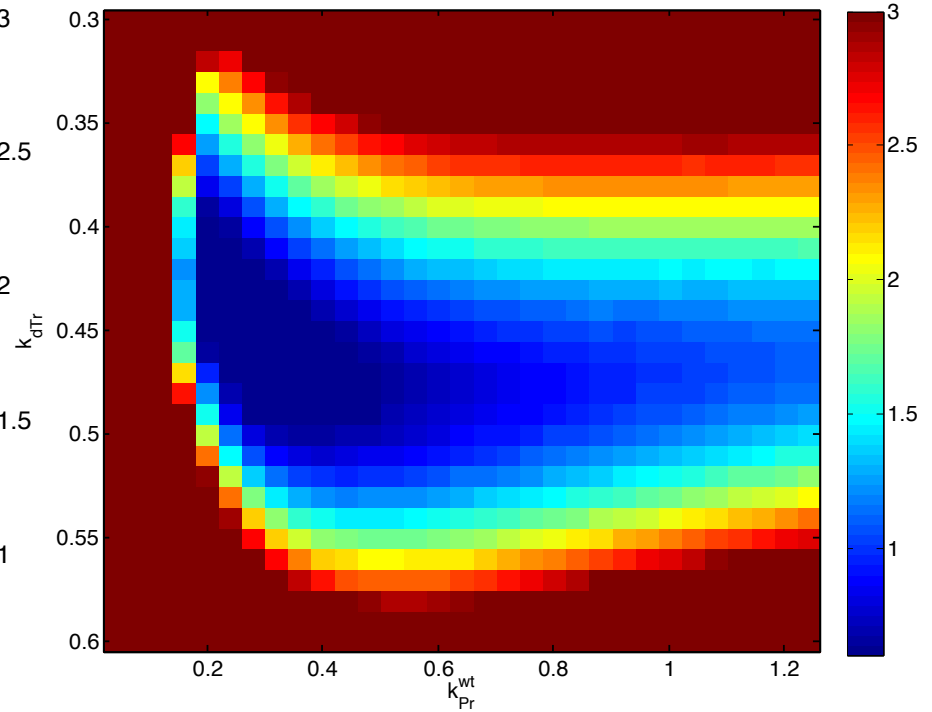
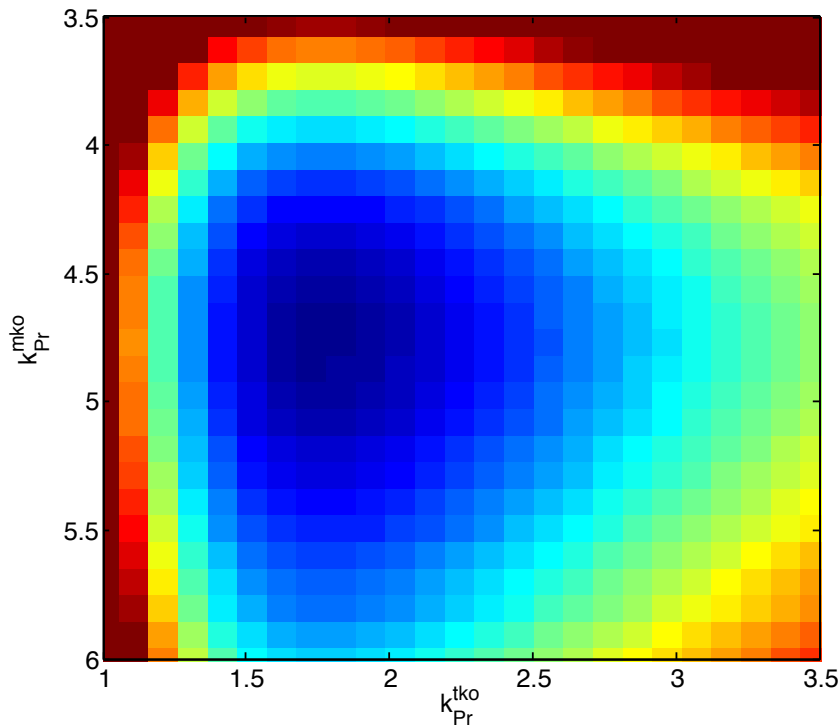
Score function (modified chi square)

$$\chi_m^2 = \frac{1}{n_F - n_p - 1} \sum_{i=1}^{n_F} \frac{(F_i^s - F_i^e)^2}{\sigma_i^2}$$

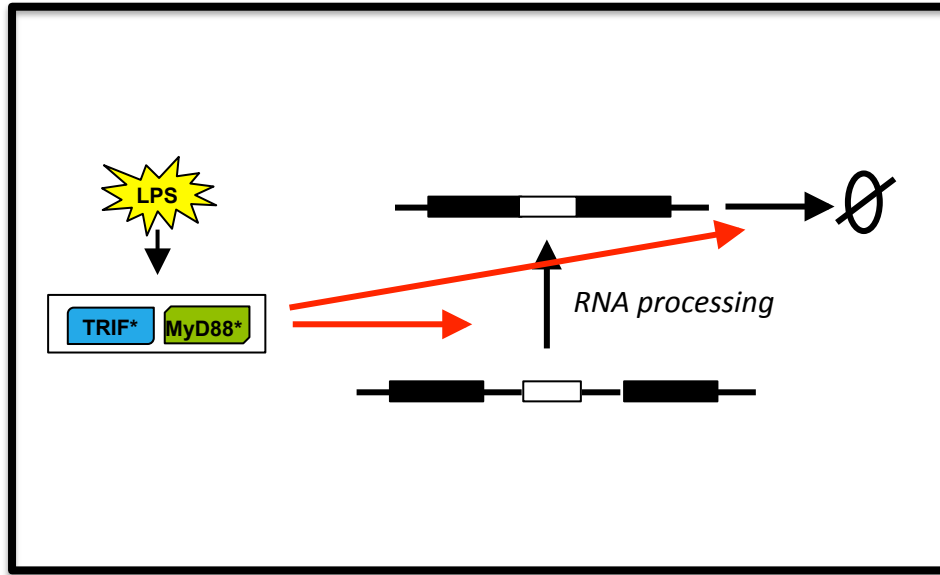
Table of features ($n_F=9$)

| Index(i) | Feature | Value(F_i^e) | Error(σ_i) |
|----------|------------------|------------------|---------------------|
| 1 | Peak time (wt) | 30 min | 5 min |
| 2 | Peak time (mko) | 60 min | 10 min |
| 3 | Peak time (tko) | 30 min | 5 min |
| 4 | Peak_wt/Peak_mko | 0.68 | 0.11 |
| 5 | Peak_wt/peak_tko | 0.75 | 0.22 |
| 6 | Wt(60)/tko(60) | 1.55 | 0.67 |
| 7 | Wt(60)/mko(60) | 0.59 | 0.07 |
| 8 | Wt(120)/mko(120) | 0.52 | 0.21 |
| 9 | Wt(120)/tko(120) | 1.49 | 0.43 |

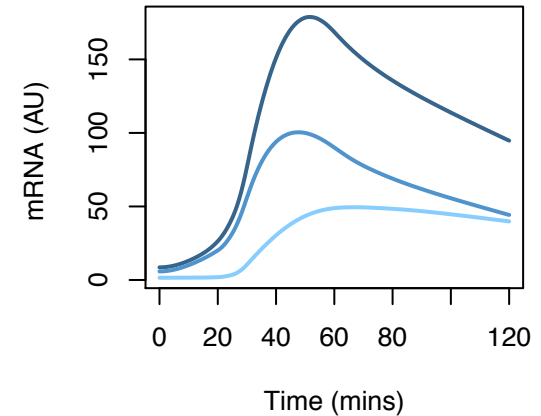
How well the model capture the features.
Score heat map for different parameters.



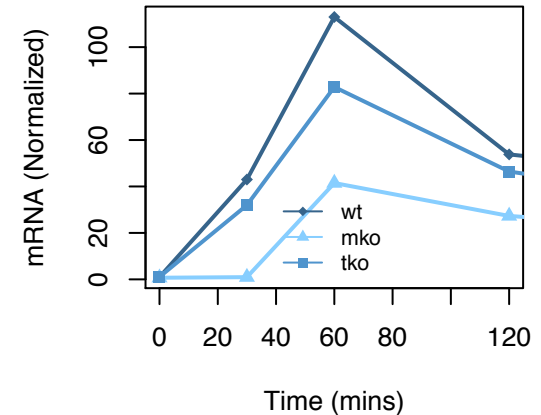
Module 2: RNA processing + stabilization



Simulation



Experiment



Equation

$$\frac{d[mRNA]}{dt} = k_{pr}[Nascent] - k_{deg mRNA}[mRNA]$$

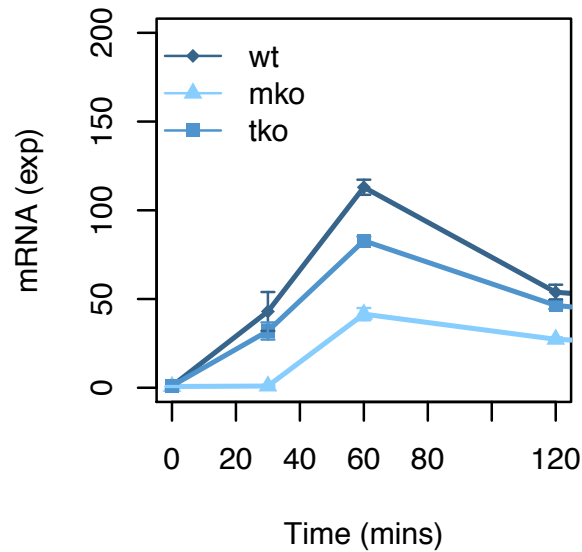
Initial condition

$$[mRNA]_{t=0} = \frac{k_{pr}}{k_{deg mRNA}}[nascent]_{t=0}$$

Parameters ($n_p = 5$)

$$k_{pr}^{wt} = 0.6 \text{ min}^{-1}, k_{pr}^{mko} = \frac{k_{pr}^{wt}}{4.5}, k_{pr}^{tko} = \frac{k_{pr}^{wt}}{1.5}, k_{deg mRNA}^{tko} = 0.07 \text{ min}^{-1}, k_{deg mRNA}^{wt/mko} = \begin{cases} -\frac{0.05}{30}t + 0.07 \text{ min}^{-1} (0 \leq t < 30 \text{ min}) \\ \frac{0.05}{30}(t - 30) + 0.02 \text{ min}^{-1} (30 \text{ min} \leq t < 60 \text{ min}) \\ 0.07 \text{ min}^{-1} (60 \text{ min} \leq t) \end{cases}$$

Experimental data



Score function (modified chi square)

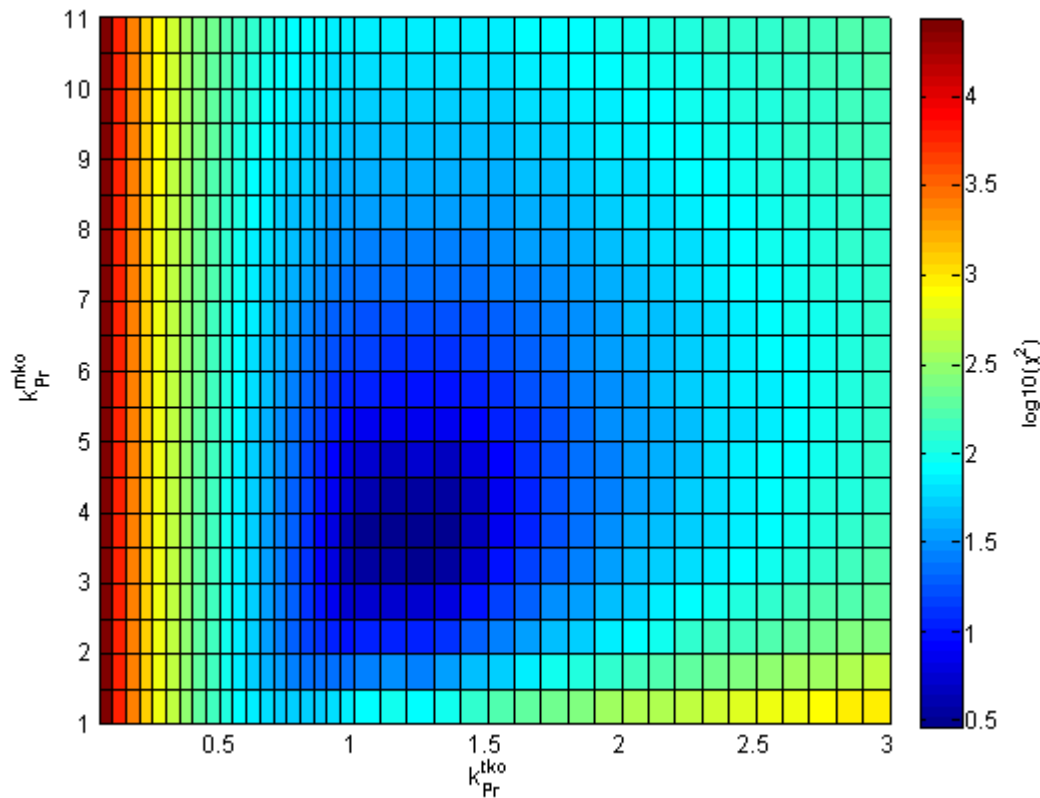
$$\chi_m^2 = \frac{1}{n_F - n_p - 1} \sum_{i=1}^{n_F} \frac{(F_i^s - F_i^e)^2}{\sigma_i^2}$$

Table of features ($n_F=7$)

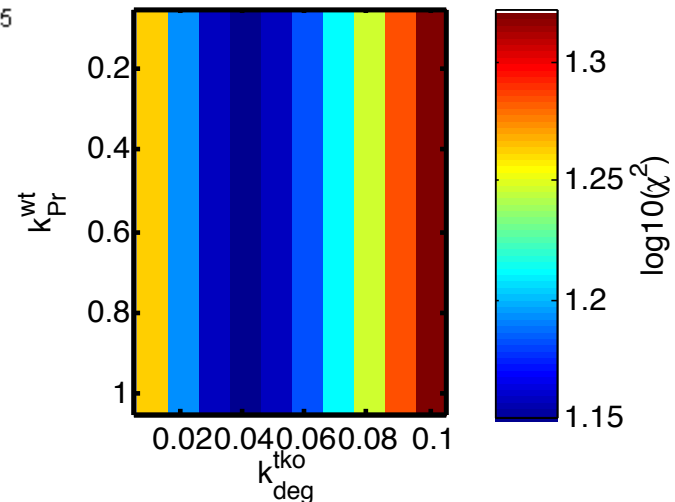
| Index(i) | Feature | Value(F_i^s) | Error(σ_i) |
|----------|-------------------|------------------|---------------------|
| 1 | Peak time (wt) | 60 min | 10 min |
| 2 | Peak time (mko) | 60 min | 10 min |
| 3 | Peak time (tko) | 60 min | 10 min |
| 4 | Peak_tko/Peak_wt | 0.73 | 0.05 |
| 5 | Peak_mko/peak_tko | 0.50 | 0.06 |
| 6 | wt_120/mko_120 | 1.98 | 0.23 |
| 7 | Wt_120/tko_120 | 1.17 | 0.14 |

How well the model capture the features.

Score heat map for different parameters.



As long as keep the same ratios of processing rate between wt, mko and tko, the fit doesn't depend on the processing rate.



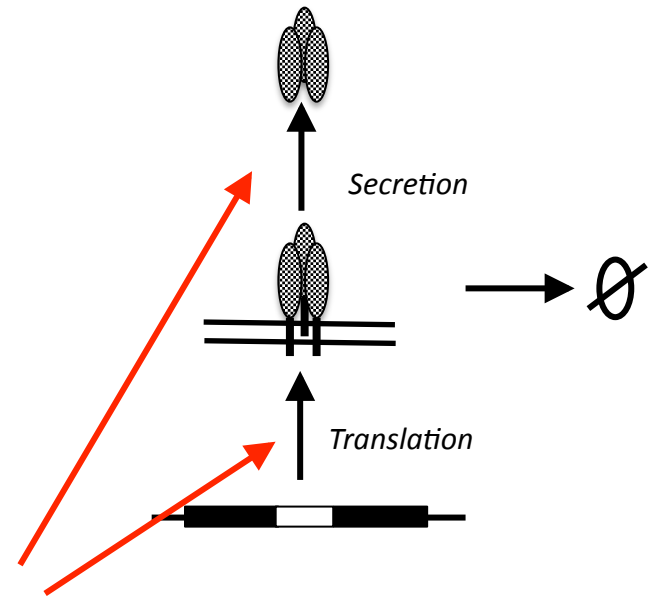
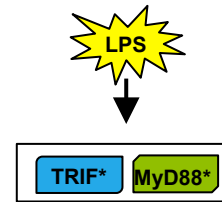
Module 3: Translation + Secretion

$$\frac{d[proTNF]}{dt} = k_{tl}[mRNA] - k_{degP}[proTNF] - k_{sec}[proTNF]$$

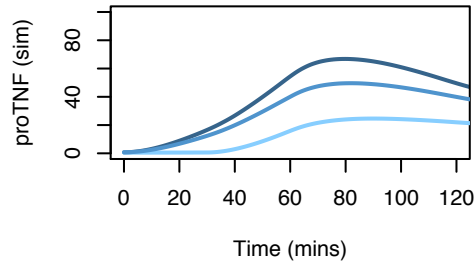
$$[secTNF](t) = \int_0^t k_{sec}[proTNF] dt$$

$$k_{tl}^{wt/mko} = 0.05 \text{ min}^{-1}, k_{tl}^{tko} = \frac{k_{tl}^{wt}}{1.5}, k_{degP} = 0.07 \text{ min}^{-1},$$

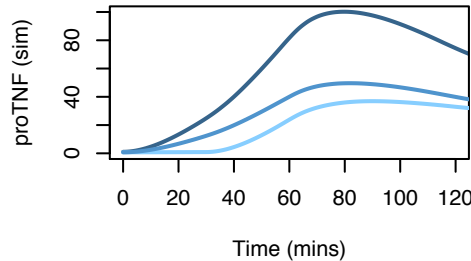
$$k_{sec}^{wt/mko} = 0.07 \text{ min}^{-1}, k_{sec}^{tko} = \frac{k_{tl}^{wt}}{5}$$



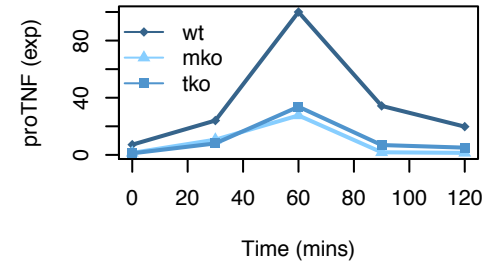
No tl regulation



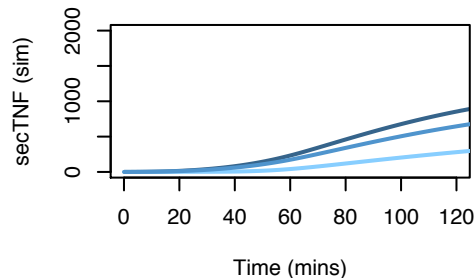
tl regulation



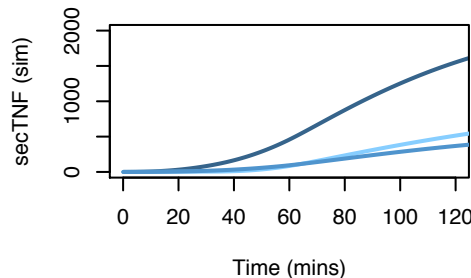
Experimental data



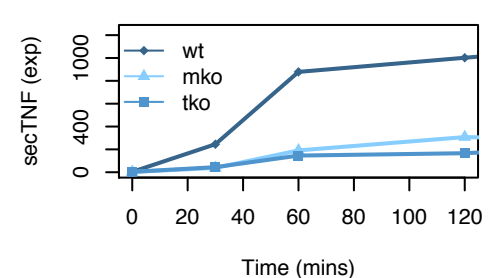
No sec regulation



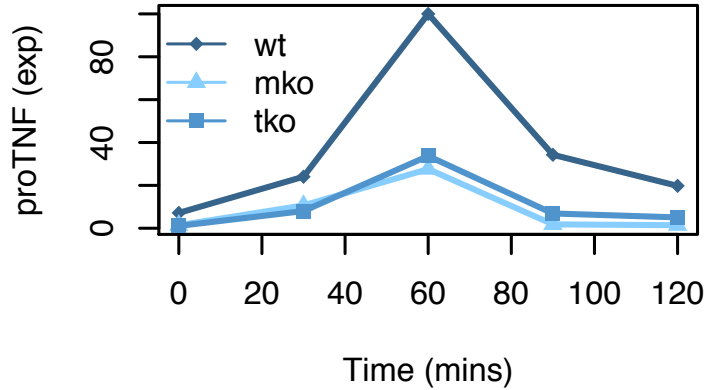
Sec regulation only



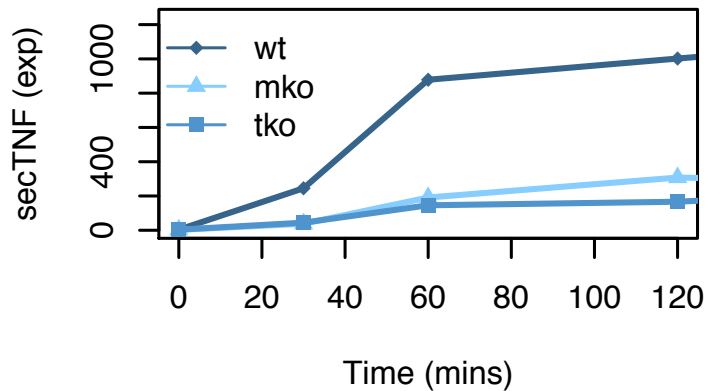
Experimental data



Experimental data



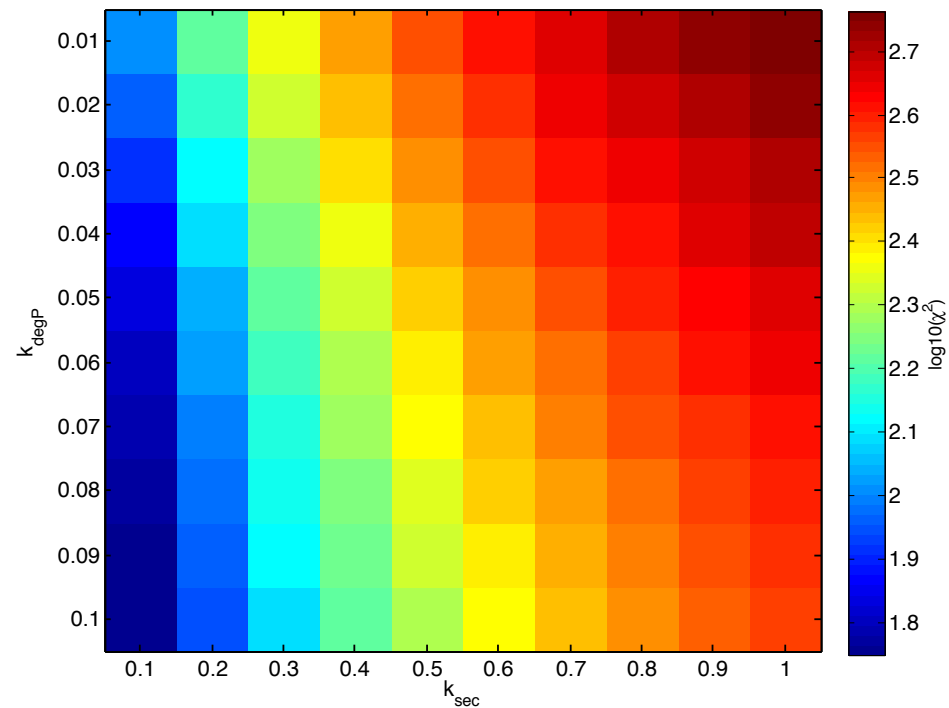
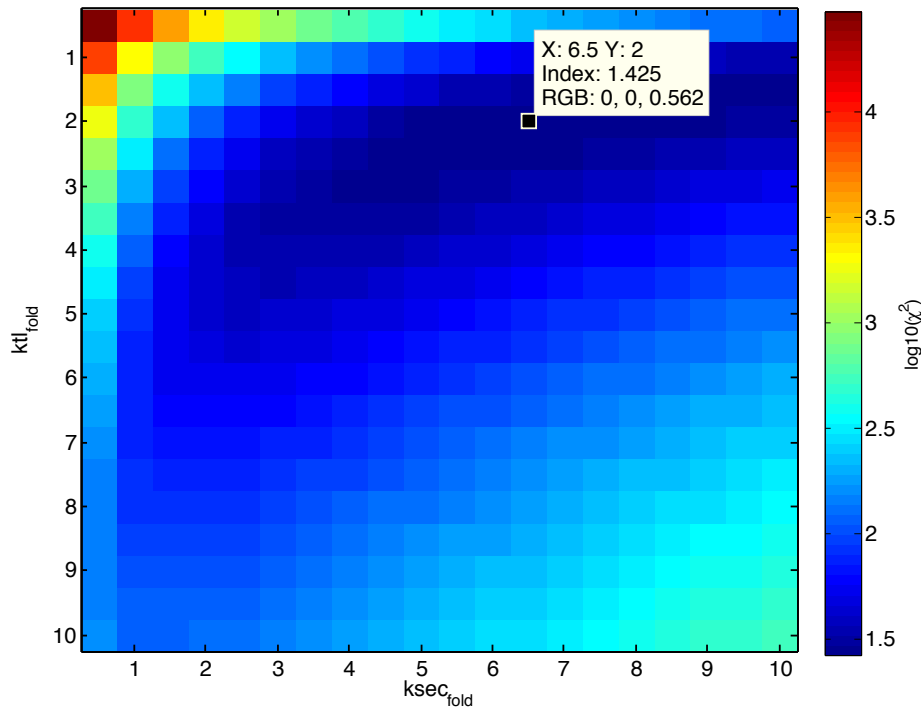
Experimental data



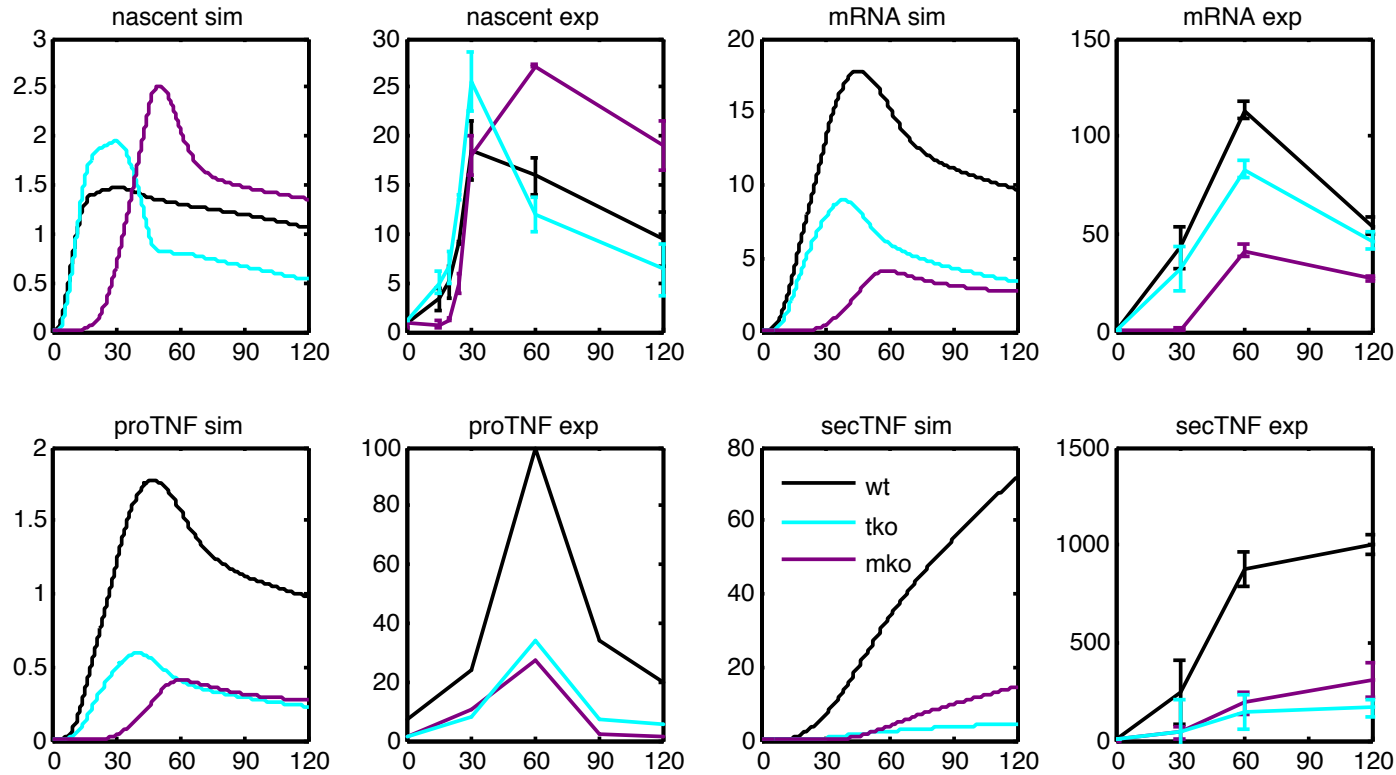
| Index(i) | Feature | Value(F^s_i) | Error(σ_i) |
|----------|-------------------|------------------|---------------------|
| 1 | Peak time (wt) | 60 min | 10 min |
| 2 | Peak time (mko) | 60 min | 10 min |
| 3 | Peak time (tko) | 60 min | 10 min |
| 4 | Peak_tko/Peak_wt | 0.34 | 0.06 |
| 5 | Peak_mko/peak_tko | 0.82 | 0.14 |
| 6 | wt_120/Peak_wt | 0.20 | 0.04 |
| 7 | Wt_120/tko_120 | 3.9 | 0.65 |
| 8 | wt_120/mko_120 | 14 | 2.3 |
| 9 | Tko_30/mko_30 | 0.75 | 0.12 |

| Index(i) | Feature | Value(F^s_i) | Error(σ_i) |
|----------|-----------------|------------------|---------------------|
| 1 | tko_60/wt_60 | 0.17 | 0.03 |
| 2 | Mko_60/tko_60 | 1.4 | 0.54 |
| 3 | Tko_120/wt_120 | 0.17 | 0.03 |
| 4 | Mko_120/tko_120 | 1.9 | 0.76 |
| 5 | wt_120/wt_60 | 1.2 | 0.16 |

How well the model capture the features.
Score heat map for different parameters.



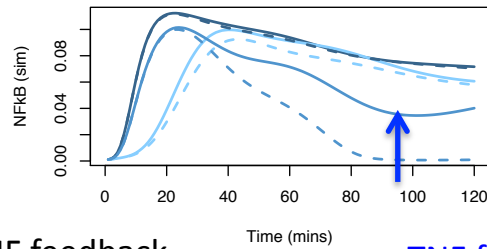
Module 1 to 3, together (NFkB as input)



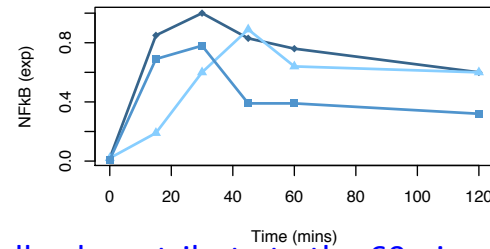
| ID | Name | Values |
|----|-------------|--------|
| 1 | Km_tr | 0.65 |
| 2 | Km_tr_fold | 2 |
| 3 | V_tr | 1 |
| 4 | k_pr | 0.4 |
| 5 | k_sec | 0.07 |
| 6 | k_tl | 0.05 |
| 7 | kdeg_m | 0.02 |
| 8 | kdeg_p | 0.07 |
| 9 | n | 2 |
| 10 | pr_fold_mko | 4.2 |
| 11 | pr_fold_tko | 1.5 |
| 12 | sec_fold | 5 |
| 13 | tl_fold | 1.5 |

Linking with TLR4 model

Simulation

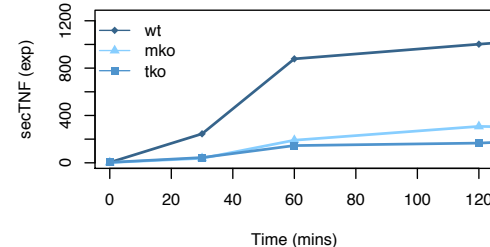
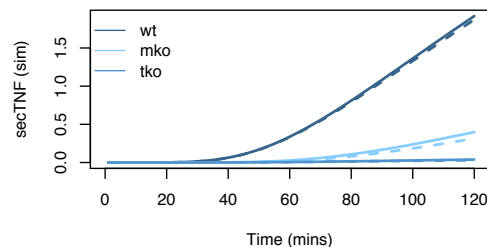
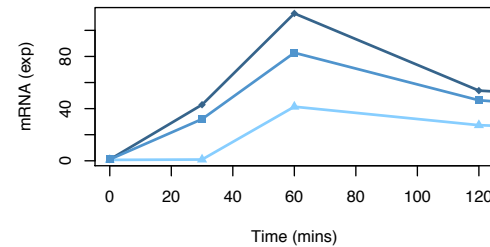
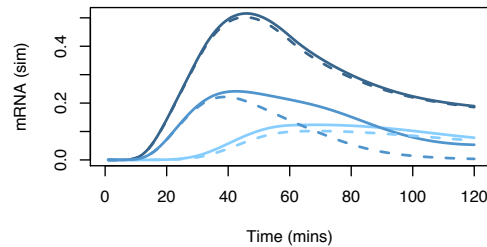
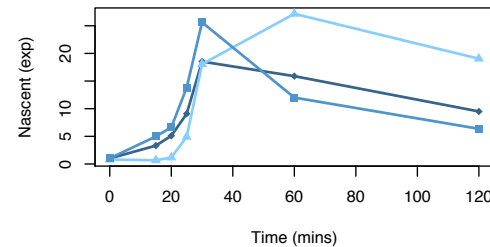
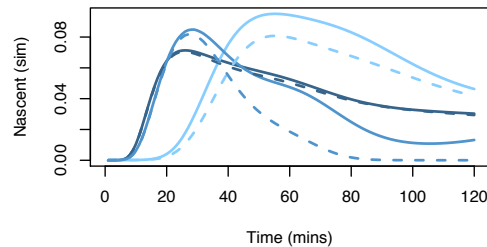


Experimental data



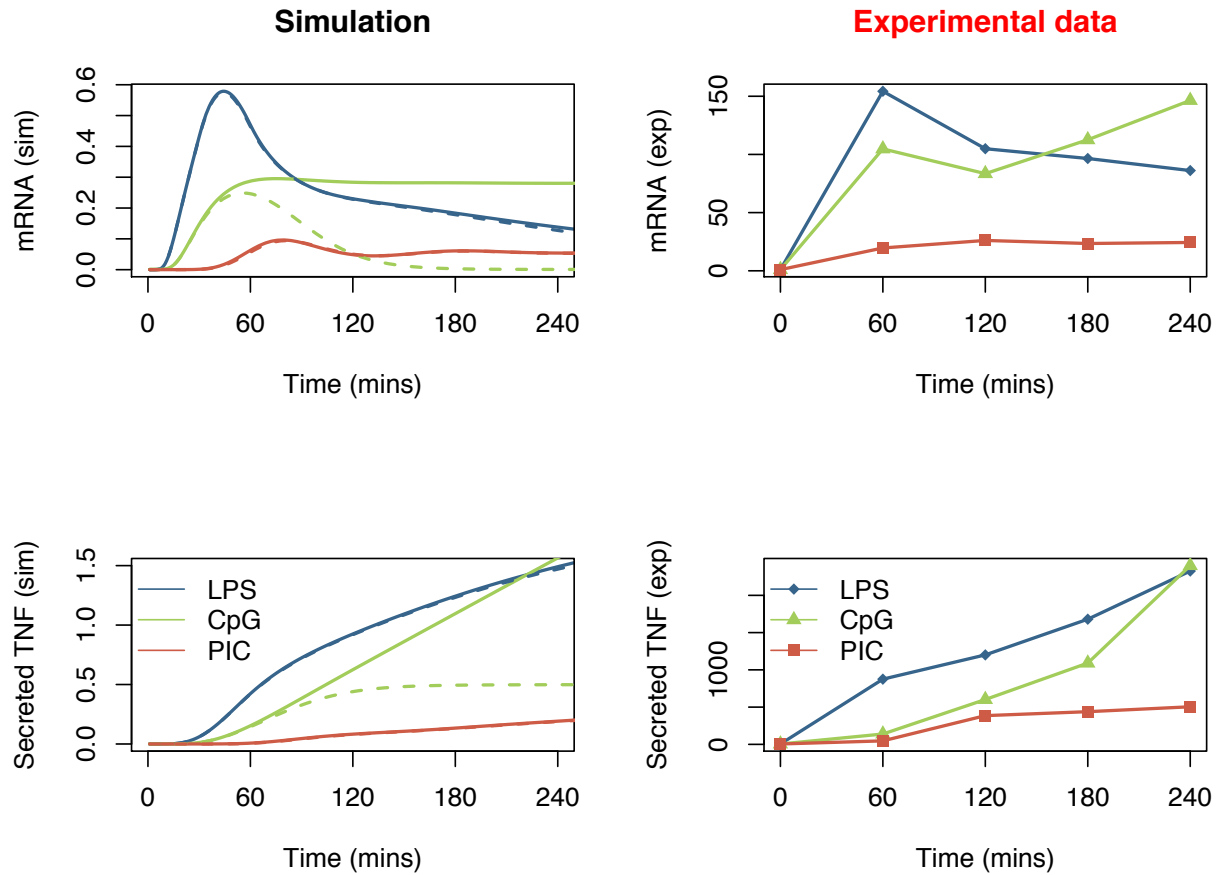
Dashed lines: without TNF feedback
Solid lines: with TNF feedback

TNF feedback contribute to the 60mins -120mins activity in trif knockout



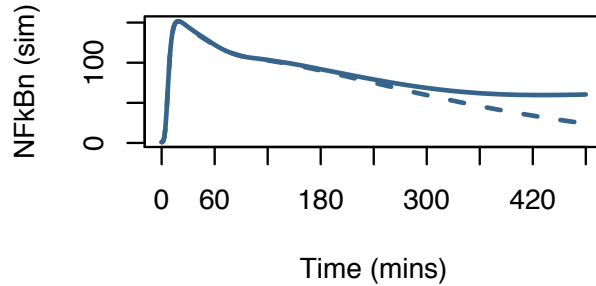
CpG and PIC stimulation (with / without TNF feedback)

Feedback toggled by TNFR synthesis

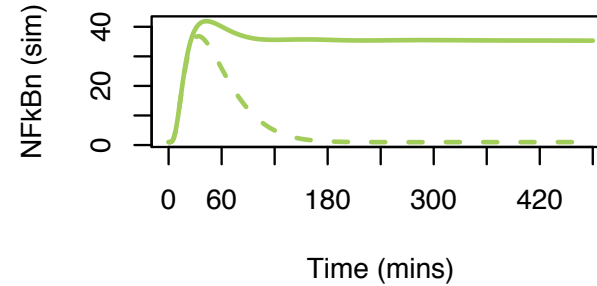


CpG and PIC stimulation (with / without TNF feedback)

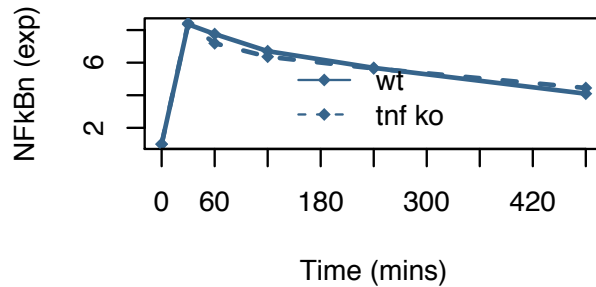
LPS stimulation (sim)



CpG stimulation (sim)



LPS stimulation (exp)



CpG stimulation (exp)

