Functional analysis of biochemical signaling pathways mediating the acute inflammatory response

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Outline

- Dose-response, time-course, and risk assessment
- Nonlinearities in cell signaling
- Analysis of IL-1α-mediated signaling in acute inflammation
- What's needed to develop a computational model

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Predicting health risks: Dose-response and exposure assessments



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Typical high dose rodent data – what do they tell us?



Not much!















Response

Possibilities



Biological mechanisms determine dose-response















How might signaling behaviors modulate dose-response? MAPK





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Apical scheme





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Molecular-level scheme



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Expected characteristics of cellular signaling

- Signal transduction
- Amplification (ultrasensitivity)
- Switch-like response to input
- Regulatory crosstalk
- Self-limiting after signal cutoff
- Redundancy
- Timeliness

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Molecular-level scheme



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What do we know?

- IL-1 α activates IL-1R1
- Small GTPases are switches
- MAPKs, amplifier and switch
- PIP, switch
- NFκB, oscillatory switch
- IL-6 mediates inflammatory response



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The IB–NF-B Signaling Module: Temporal Control and Selective Gene Activation Alexander Hoffmann, Andre Levchenko, Martin L. Scott, David Baltimore Science 298:1241 – 1245, 2002



6 hr



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Qualitative insights

- Signal is amplified but also terminated
- Parallel but convergent signaling
 - Redundancy?
 - Regulatory crosstalk?
- Positive and negative signals in the cross-talk
 - What determines who wins?
 - Or does this just sharpen the signal?



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Actually, we know a lot...

- We know that activation of IL -1R1 leads to IL-6 upregulation.
- For a given duration of IL-1R1 activation, small GTPases, PIP, MyD-88 and MAPK determine the time-course and intensity (?) of IL-6 upregulation
- Overall signaling is self-limiting

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How to move on to a quantitative model?

- Need dose-response and timecourse data for each module
 - (Eungdamrong & Iyengar, Modeling Cell Signaling Networks, Biol. of the Cell, 355-362, 2004)

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Types of models

- Qualitative
- Quantitative-1
 - Functional relationships not tested against data and parameter values not known
 - Hypothesis generation
- Quantitative-2
 - Data used to check functional relationships and parameter values
 - Prediction

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Iteration of experiments in the laboratory and the computer



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Questions

- Why should PIP be a negative regulator of the signaling?
 - Conditions under which the cell does not want to generate an inflammatory response?
 - Help to generate more switch-like transitions?

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- Generic aspects of signal transduction pathways
- Qualitative analysis of existing literature suggests initial topology for computational model
- Targeted experimentation needed in support of computational model



Best time to think about signal transduction...





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"And that's why we need a computer."

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