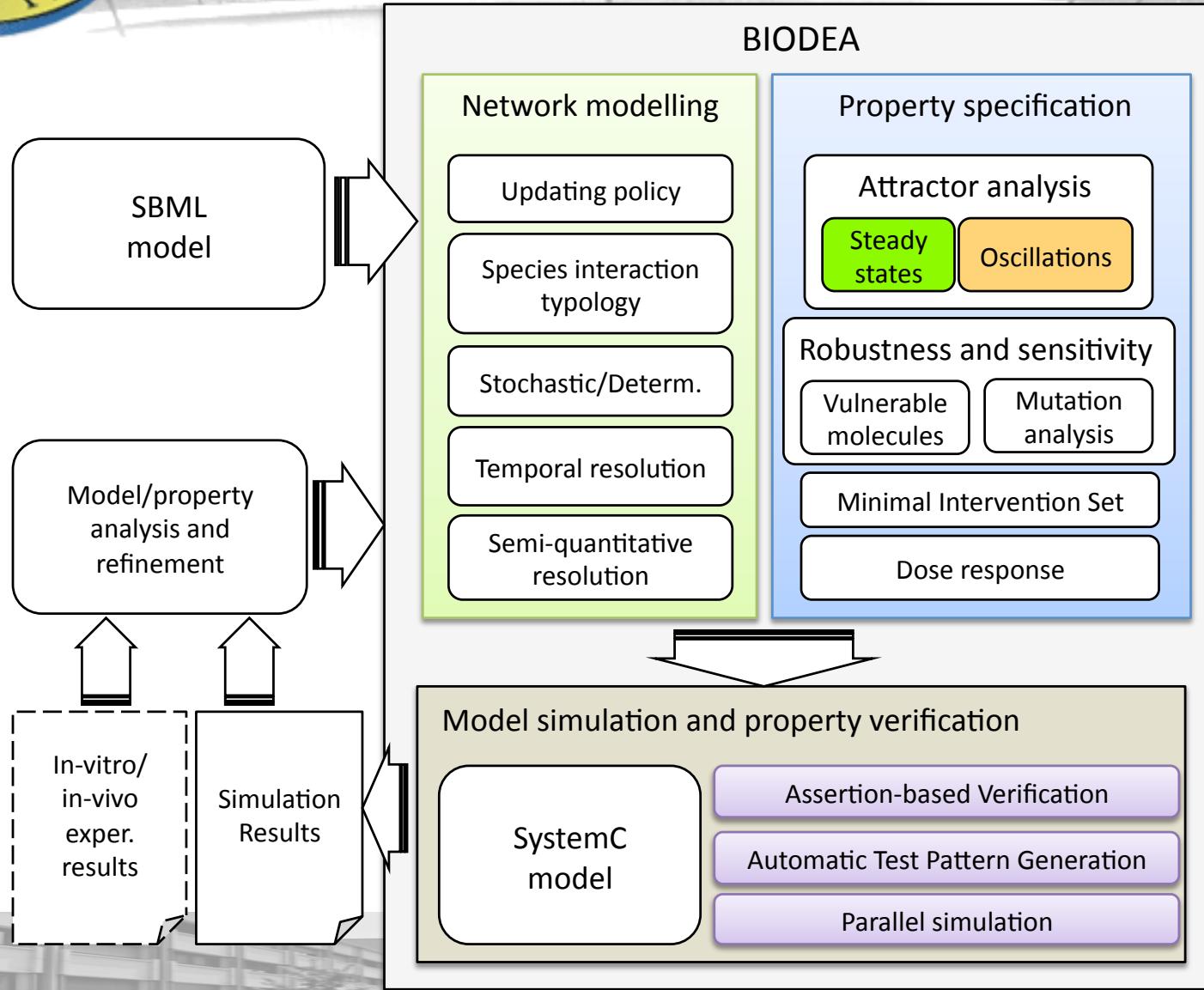




**Modeling,  
verification,  
parallel simulation:  
from Electronic Design Automation (EDA)  
to Systems Biology**



# Modeling and simulation framework

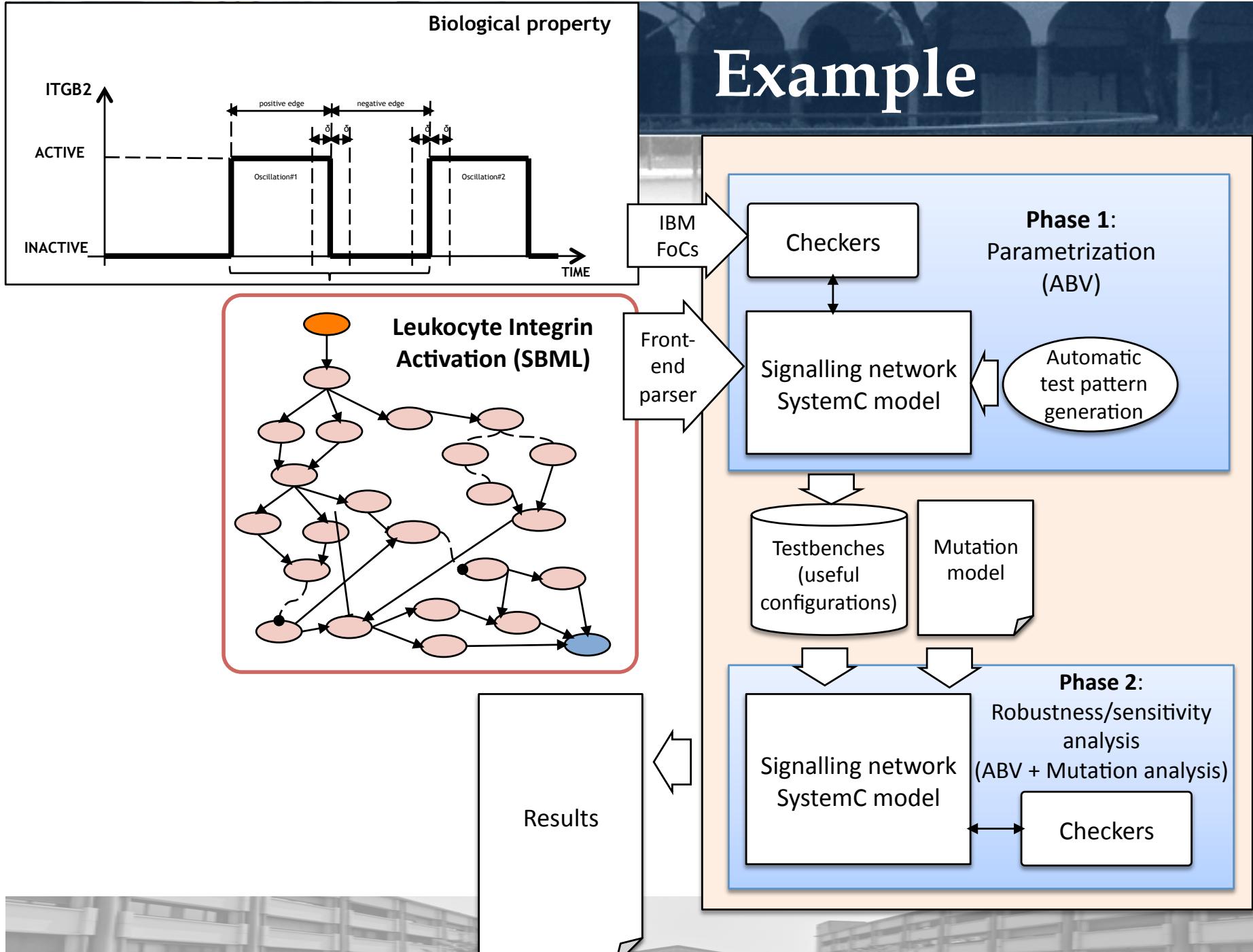




# Modeling Overview

Model	Type	Feat.	Properties
Interaction graphs		<ul style="list-style-type: none"><li>• No time</li><li>• No concentrations</li></ul>	Dependency matrix Source-target pathways Activators (strong/weak) Inhibitors (strong/weak) Feedback loops (pos, neg) Degree (scale free, etc.) Monotony Minimal intervention set (MIS)
Boolean models	Qualitative modeling	<ul style="list-style-type: none"><li>• Time (instants)</li><li>• No concentrations</li></ul>	+ <ul style="list-style-type: none"><li>• Simple attractors<ul style="list-style-type: none"><li>– Steady/stable states</li></ul></li><li>• Complex attractors<ul style="list-style-type: none"><li>– Oscillations</li></ul></li><li>• Robustness &amp; sensitivity</li></ul>
Extended Boolean models	Semi-quantitative modeling	<ul style="list-style-type: none"><li>• Time (instants)</li><li>• Concentrations (%)</li></ul>	+ <ul style="list-style-type: none"><li>• Accuracy...</li></ul>
Ordinary differential equations (ODEs)	Quantitative modeling	<ul style="list-style-type: none"><li>• Time (actual)</li><li>• Concentrations (actual)</li></ul>	<ul style="list-style-type: none"><li>• Parametrization<ul style="list-style-type: none"><li>– Using experimental data</li></ul></li></ul>

# Example

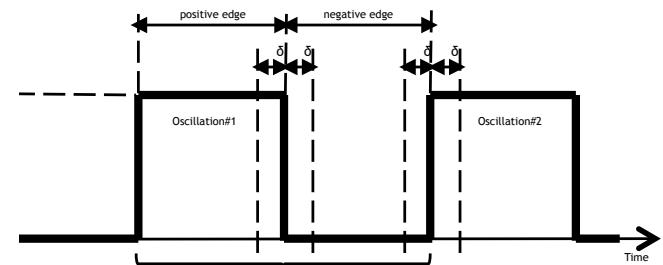




# Qualitative modeling: results

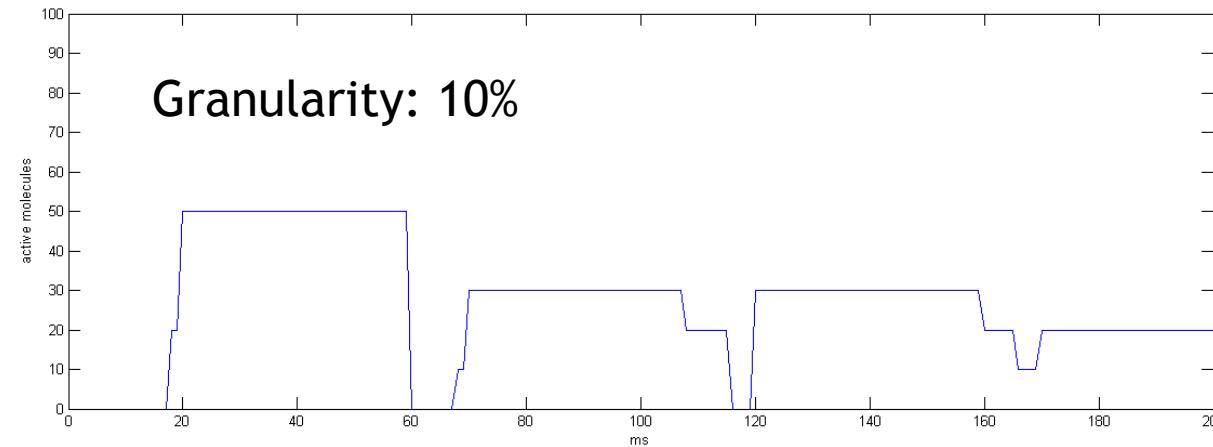
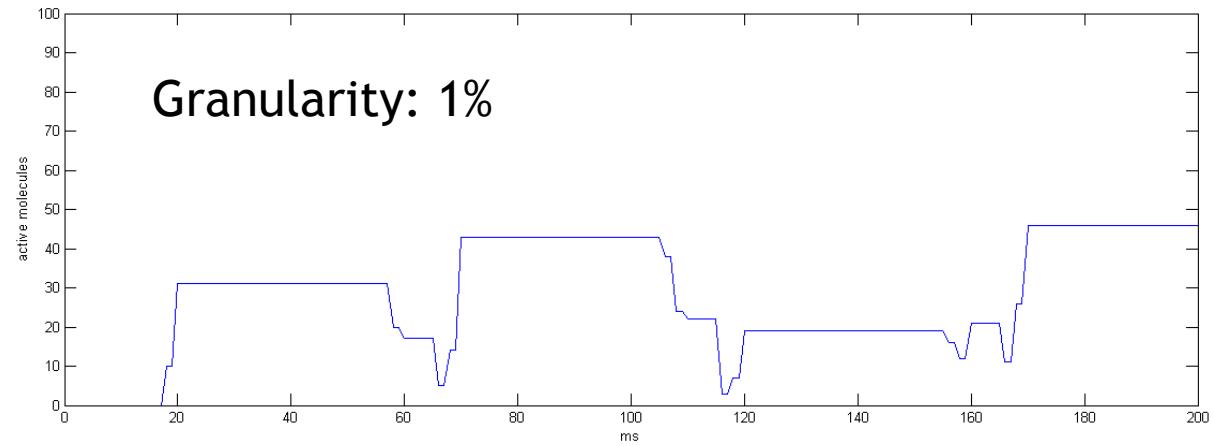
Mutated protein	Aperiodic config.	Useful conf. (periodic)	Oscill. (#)	Period ms
Golden model	-	2,008,188	2	50( $\pm 20\%$ )
CXCR4	0	0	0	0
JAK3	2,008,188	0	2-4	-
JAK2	2,008,188	0	2-4	-
ABG	0	2,008,188	2	50( $\pm 20\%$ )
VAV1	0	0	0	0
PLC	0	2,008,188	2	50( $\pm 20\%$ )
RAC1	860,652	1,147,536	1-3	50( $\pm 20\%$ )
RHOA	860,652	1,147,536	1-3	50( $\pm 20\%$ )
CDC42	2,008,188	0	2-4	-
IP3	0	2,008,188	2	50( $\pm 20\%$ )
DAG	17,640	1,990,548	1-2	50( $\pm 20\%$ )
PLDI	0	0	0	0
PIP5K1C	0	0	0	0
CA	0	2,008,188	2	50( $\pm 20\%$ )
RASGRP1	0	2,008,188	2	50( $\pm 20\%$ )
PA	0	0	0	0
RAP1A	0	2,008,188	2	50( $\pm 20\%$ )
PIP2	0	0	0	0
RIAM	0	2,008,188	2	50( $\pm 20\%$ )
RASSF5	0	2,008,188	2	50( $\pm 20\%$ )
TLN1	792,036	1,216,152	2-3	50( $\pm 20\%$ )
FERMT3	952,140	1,056,048	2-4	50( $\pm 20\%$ )

Parametrization over  
≈  $2 \times 10^9$  configurations  
Total simulation time:  
≈ 4h





# Semi-quantitative modeling: results

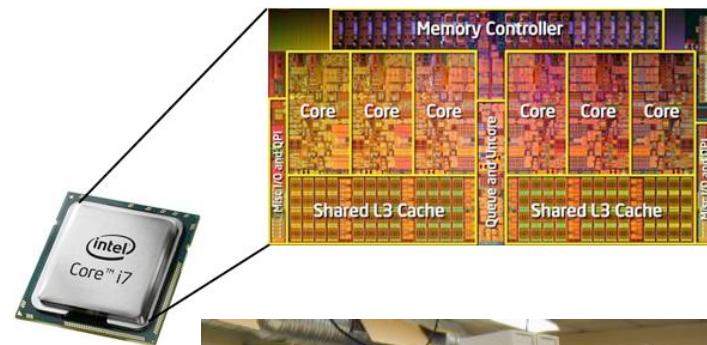
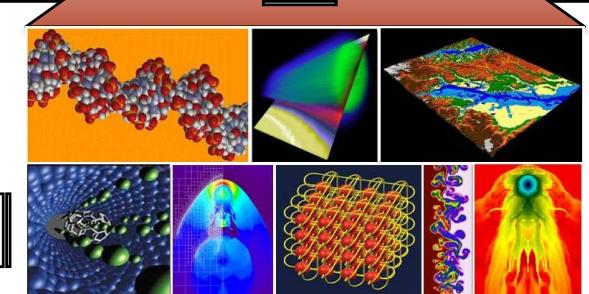


# Parallel simulation



## Parallelism and parallel architectures:

- GPUs
  - NVIDIA CUDA  
(Fermi, Kepler, Maxwell technology)
  - OpenCL/OpenACC  
(AMD APU technology)



- Multi-core CPUs
  - OpenMP
- Cluster architectures:
  - MPI

