Experimental data
- cell source and treatments
- protein activities and gene expression

Scaling and normalization
- background subtraction and normalization
- scaling of protein data to ANIMO model
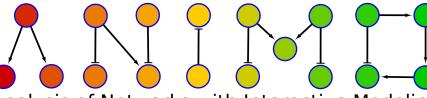
Model construction
- simplified prior knowledge network
- scenario choice and parameter fitting
- run model

Model use
- optimization of topology
- explanation of output
- interpretation of data
- new hypotheses

suggest new experiments

Analysis of Networks with Interactive Modeling