

Two branches

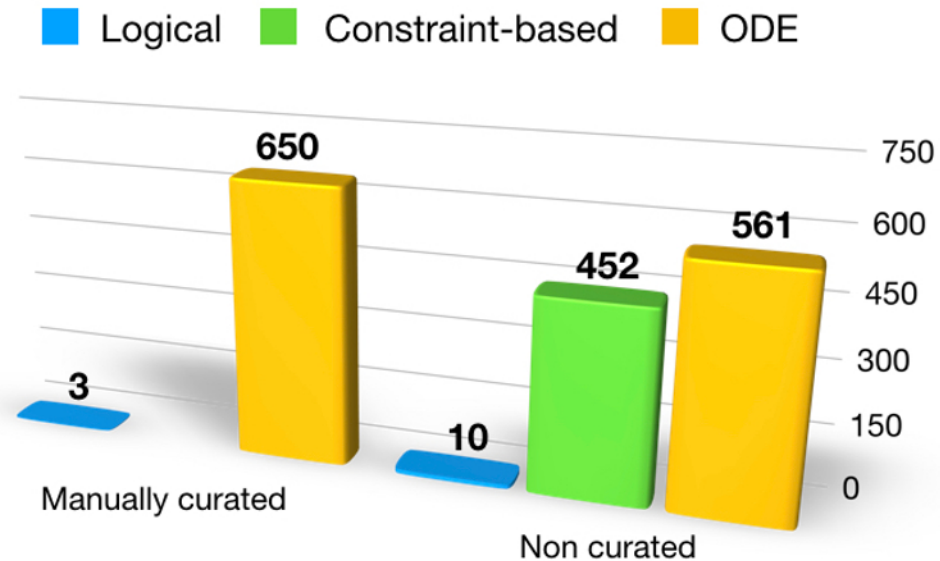
- ☒ Manually curated
- ☒ Non curated

Model formats

- ☒ SBML
- ☒ CellML
- ☒ Matlab
- ☐ ...

Modelling approaches

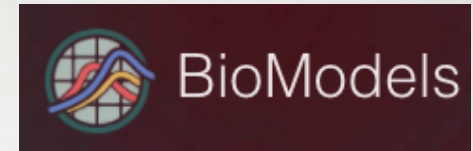
- ☒ Ordinary Differential Equation
- ☒ Logical
- ☒ Constraint-based
- ☐ ...



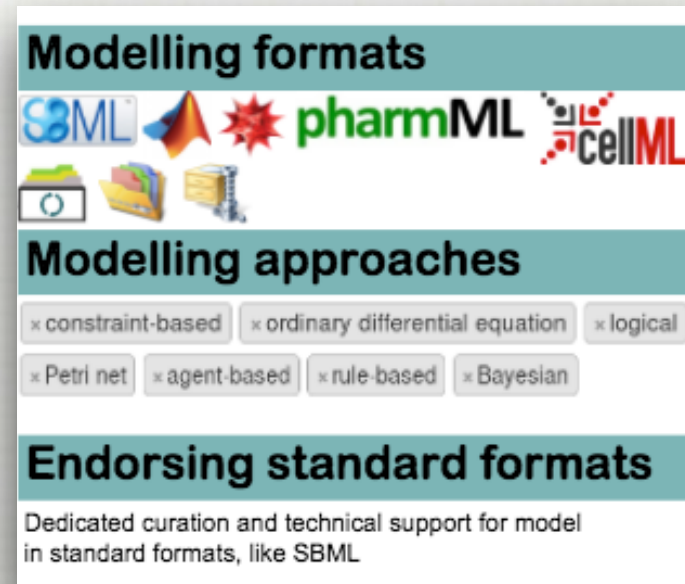
BioModels



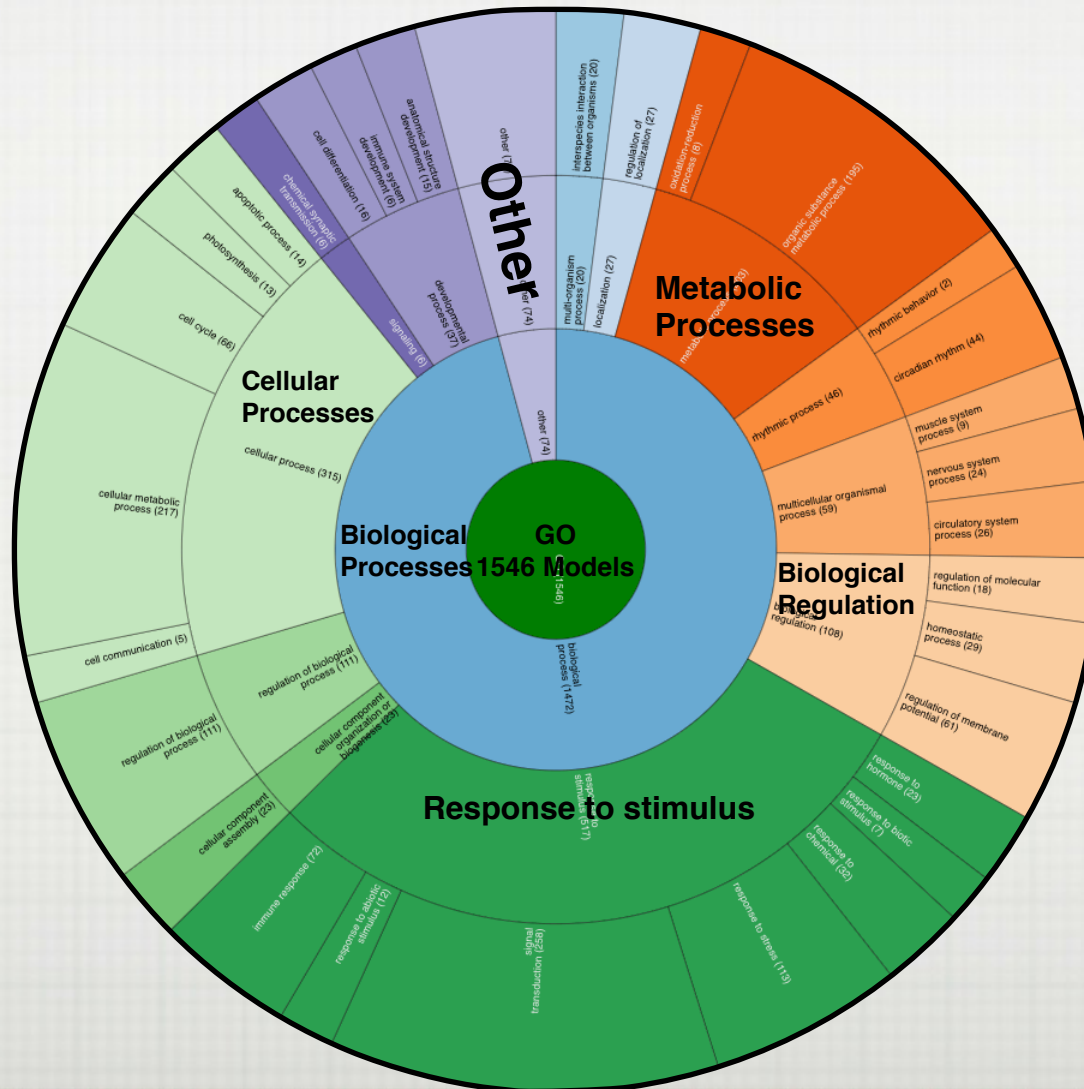
Biomodels Database



- Repository of published models
 - MIRIAM compliant, SBML standard format
- Provides reproducible, high-quality, freely-accessible models
- Some useful features:
 - Pathway Map Generation
 - Online converters
 - Advanced filtering
 - Modeling approach
 - Organism
 - Disease
 - Gene Ontology



Available models based on GO



Examples of Models

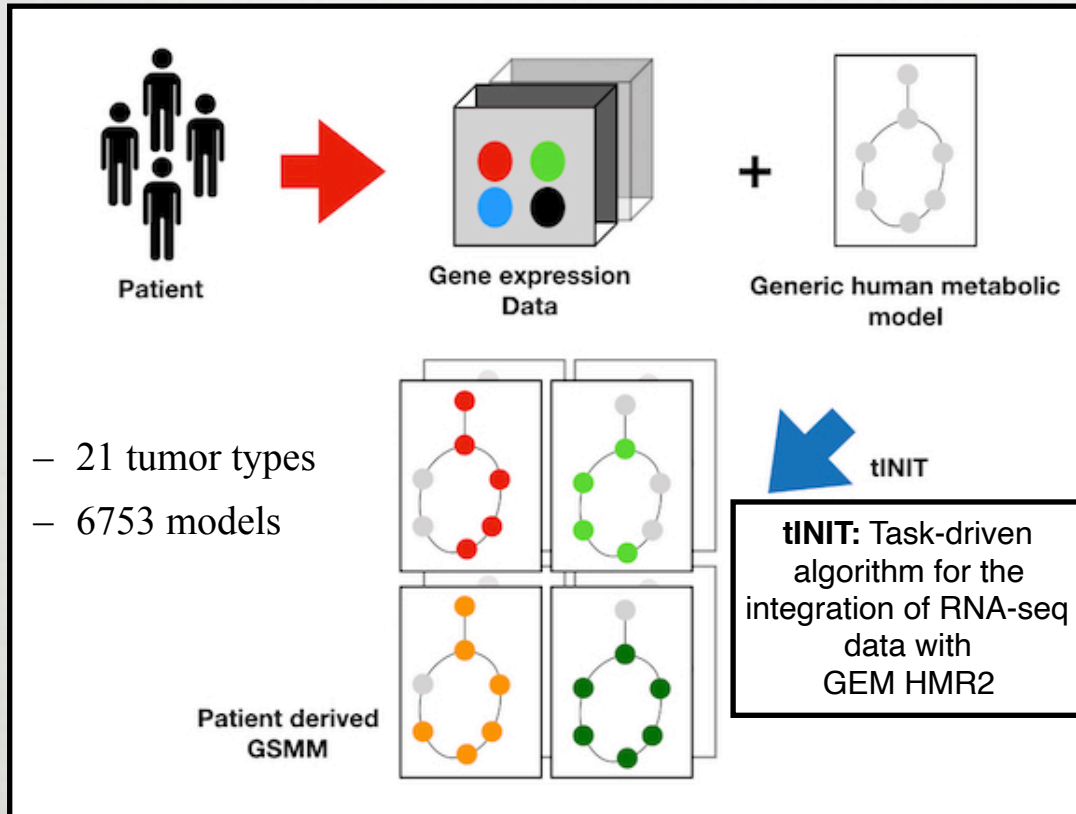
- *Homo sapiens* (7225+ models)
 - iAB-RBC-283 (RBC Metabolic Network)
 - Blood stem cell regulatory network (LMPP Network)
 - ~455 Patient specific Kidney Renal Clear Cell Carcinoma models
 - ERK, P13K/Akt and Wut Signaling network models (Normal and 6 different mutations)
 - Stochastic model of Alzheimer's disease
 - Integrated calcium homeostasis and bone remodeling

Examples of Models

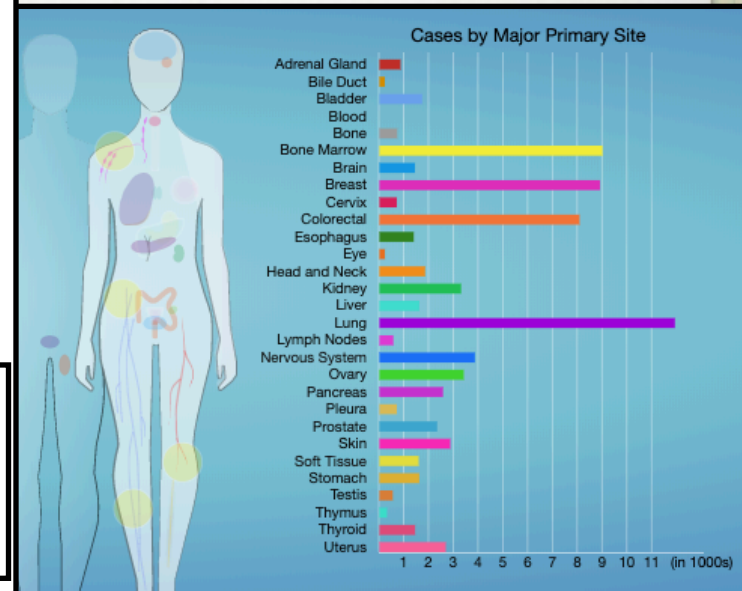
- *Patient-derived GEMs*
- *Neurodegenerative disease processes*
- *Complex Plant Circadian Clock*
- *Meiotic initiation in *S. cerevisiae**
- *$\alpha\beta$ Immunization in Alzheimer's disease*
- *Zombie Apocalypse*

PDGSMM

Patient Derived Genome Scale Metabolic Models

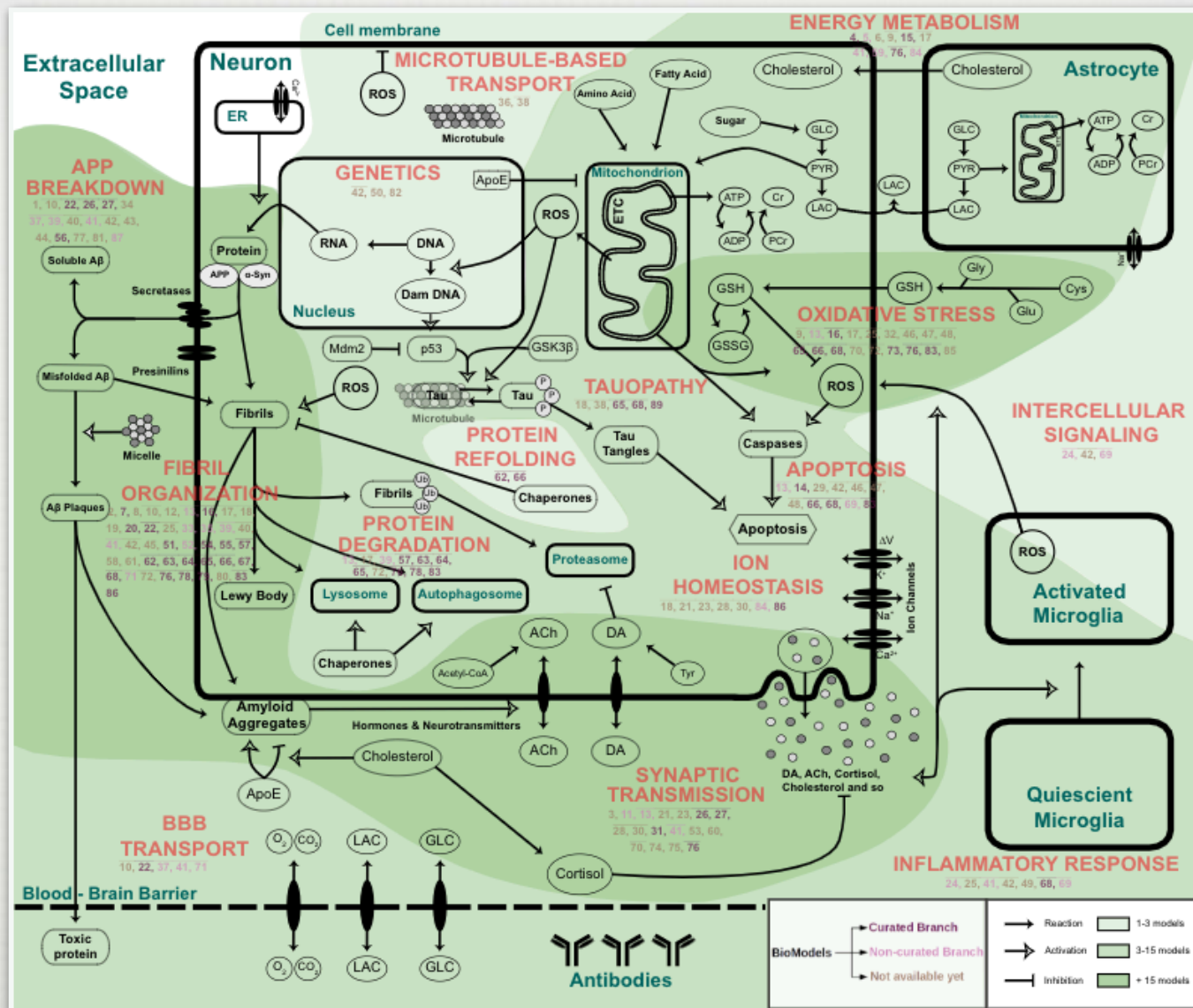


The Cancer Genome Atlas (TCGA)

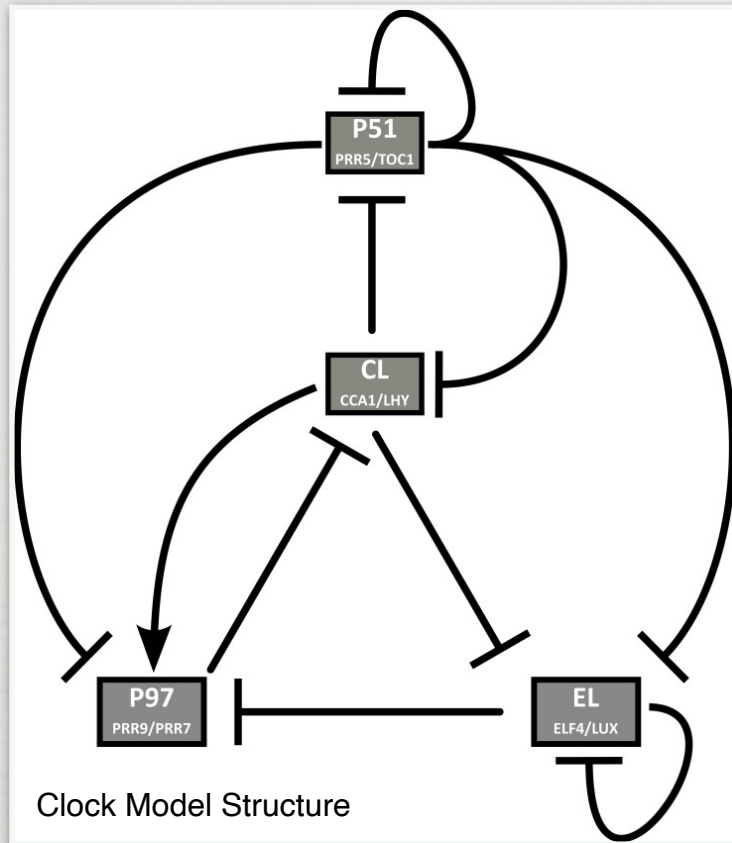


<https://portal.gdc.cancer.gov/>

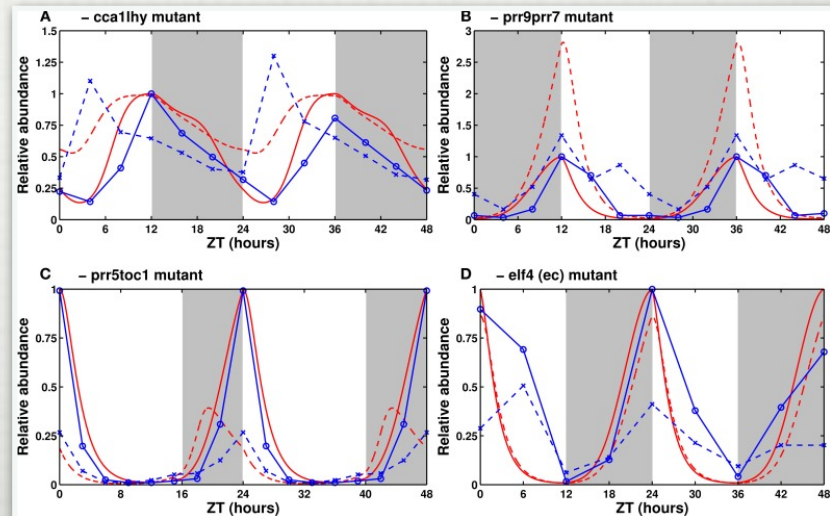
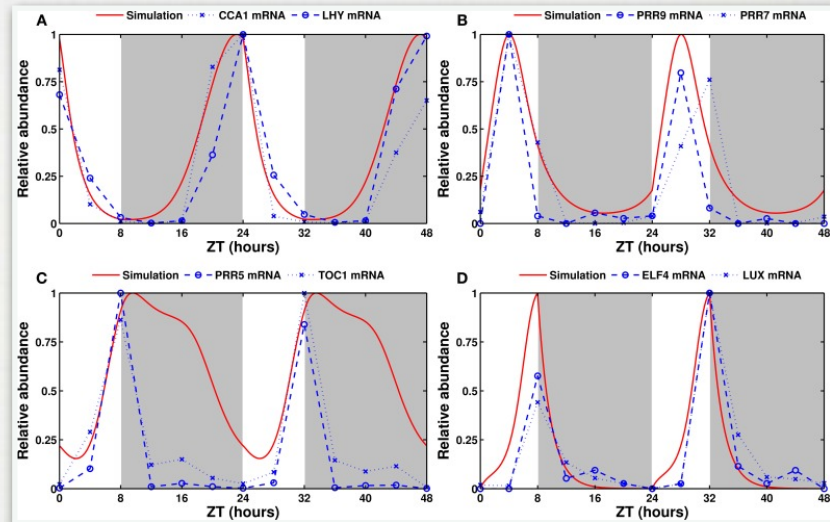
Neurodegenerative disease processes



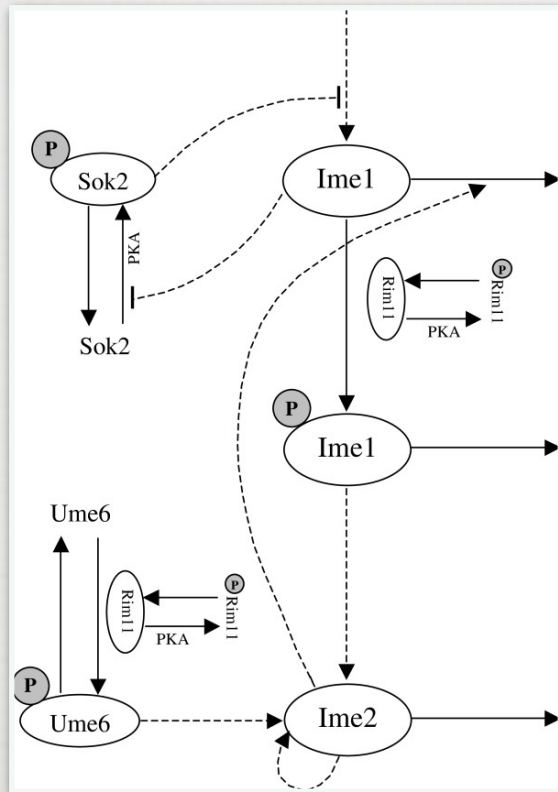
Complex Plant Circadian Clock



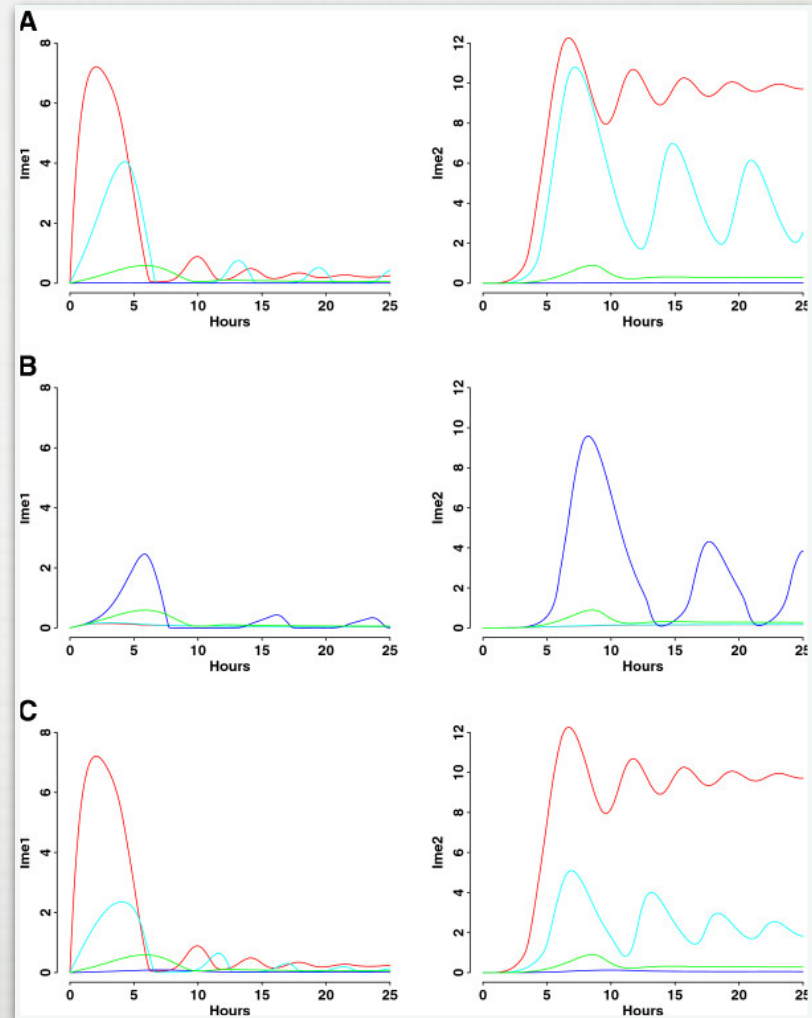
De Caluwé J, Xiao Q, Hermans C, Verbruggen N, Leloup J-C, Gonze D. A Compact Model for the Complex Plant Circadian Clock. *Frontiers in Plant Science*. 2016;7:74. doi:10.3389/fpls.2016.00074.



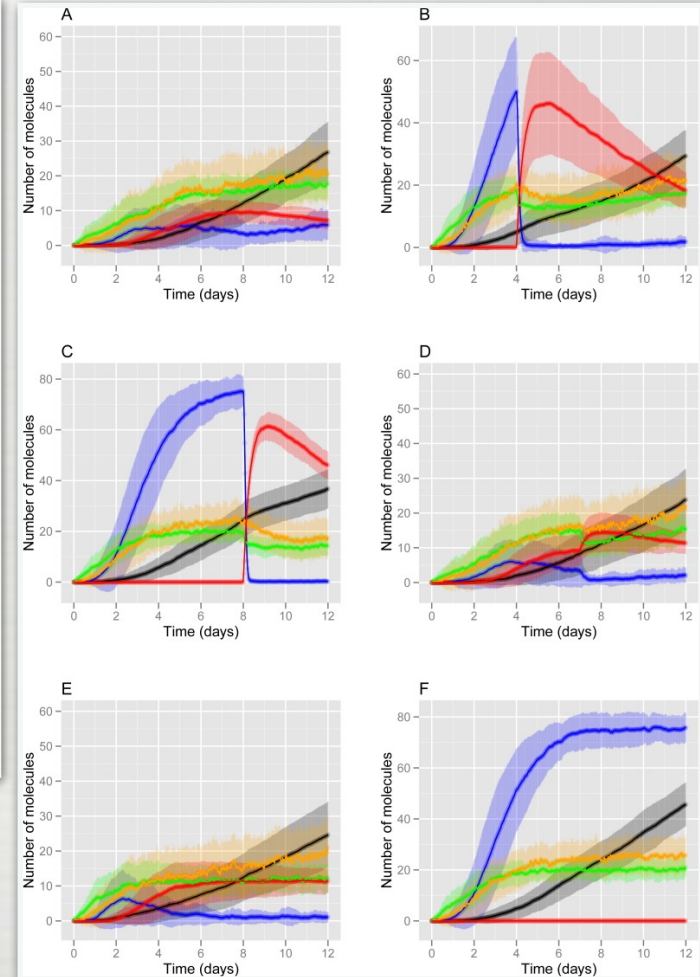
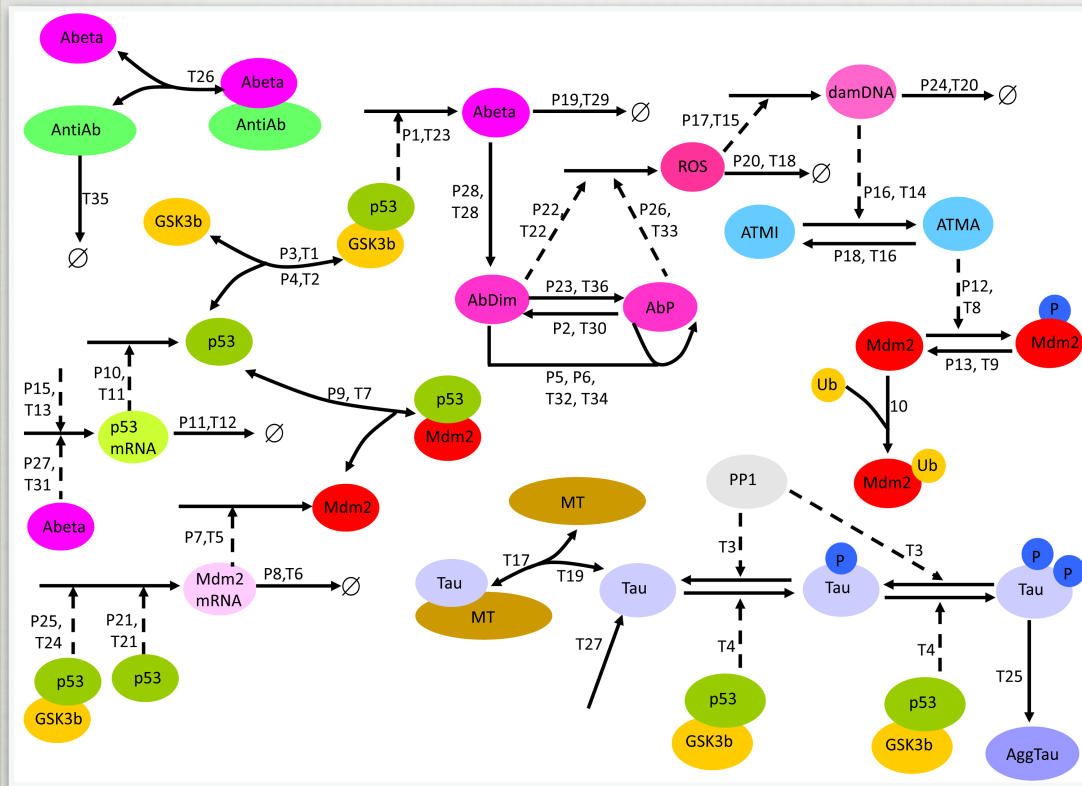
Meiotic initiation in *S. cerevisiae*



Ray D, Su Y, Ye P. Dynamic modeling of yeast meiotic initiation. *BMC Systems Biology*. 2013;7:37. doi: 10.1186/1752-0509-7-37.

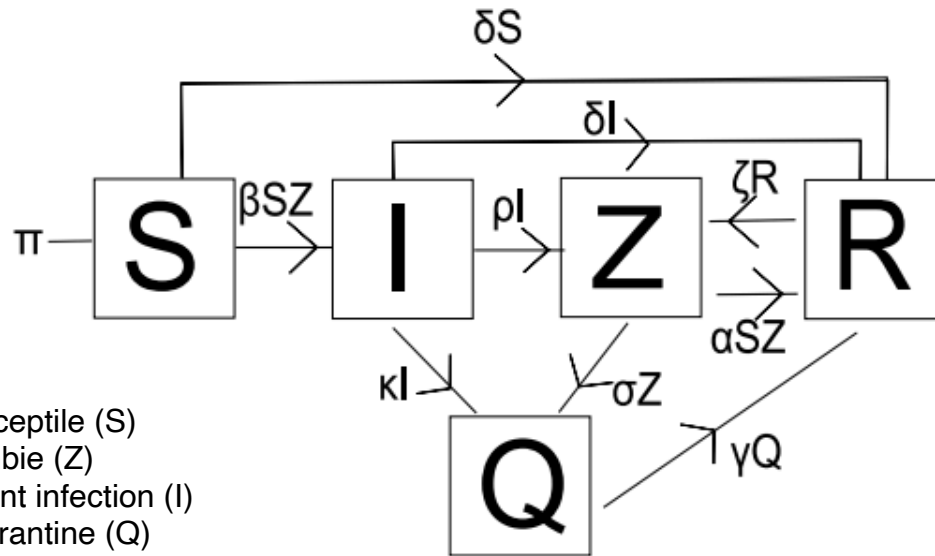


$\alpha\beta$ Immunization in Alzheimer's disease

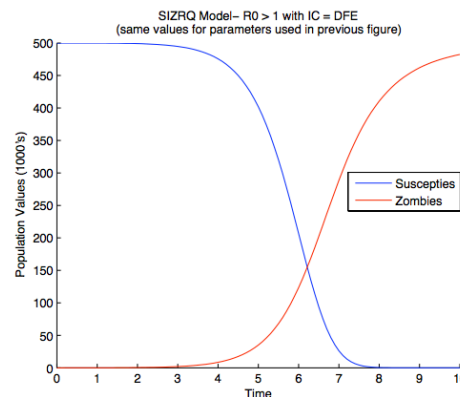
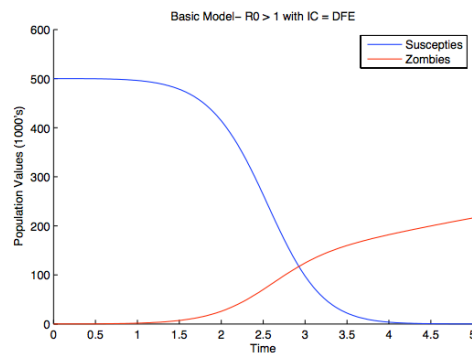
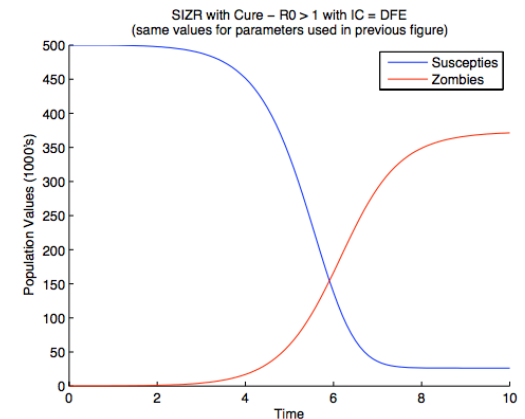
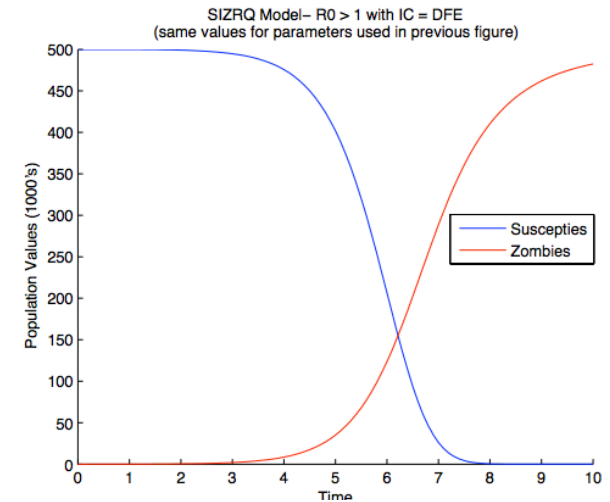


Proctor CJ, Boche D, Gray DA, Nicoll JAR. Investigating Interventions in Alzheimer's Disease with Computer Simulation Models. Padmanabhan J, ed. *PLoS ONE*. 2013;8(9):e73631. doi: 10.1371/journal.pone.0073631.

Zombie Apocalypse



Susceptible (S)
Zombie (Z)
Latent infection (I)
Quarantine (Q)
Removed/Dead with Zombie potential (R)



Usefulness

- Provides a framework of what can be considered a high-quality, reproducible model.
 - Syntactic correctness, semantic soundness, and correspondence with reference publication
- Annotation of models removes ambiguity.
- Unlike other pathway databases, contains quantitative information representing quantities or kinetics.
- Having over 150,000+ quality models is a significant resource for the scientific community.
- <https://www.ebi.ac.uk/biomodels-main/>