

# 2nd E-Cell Sprint

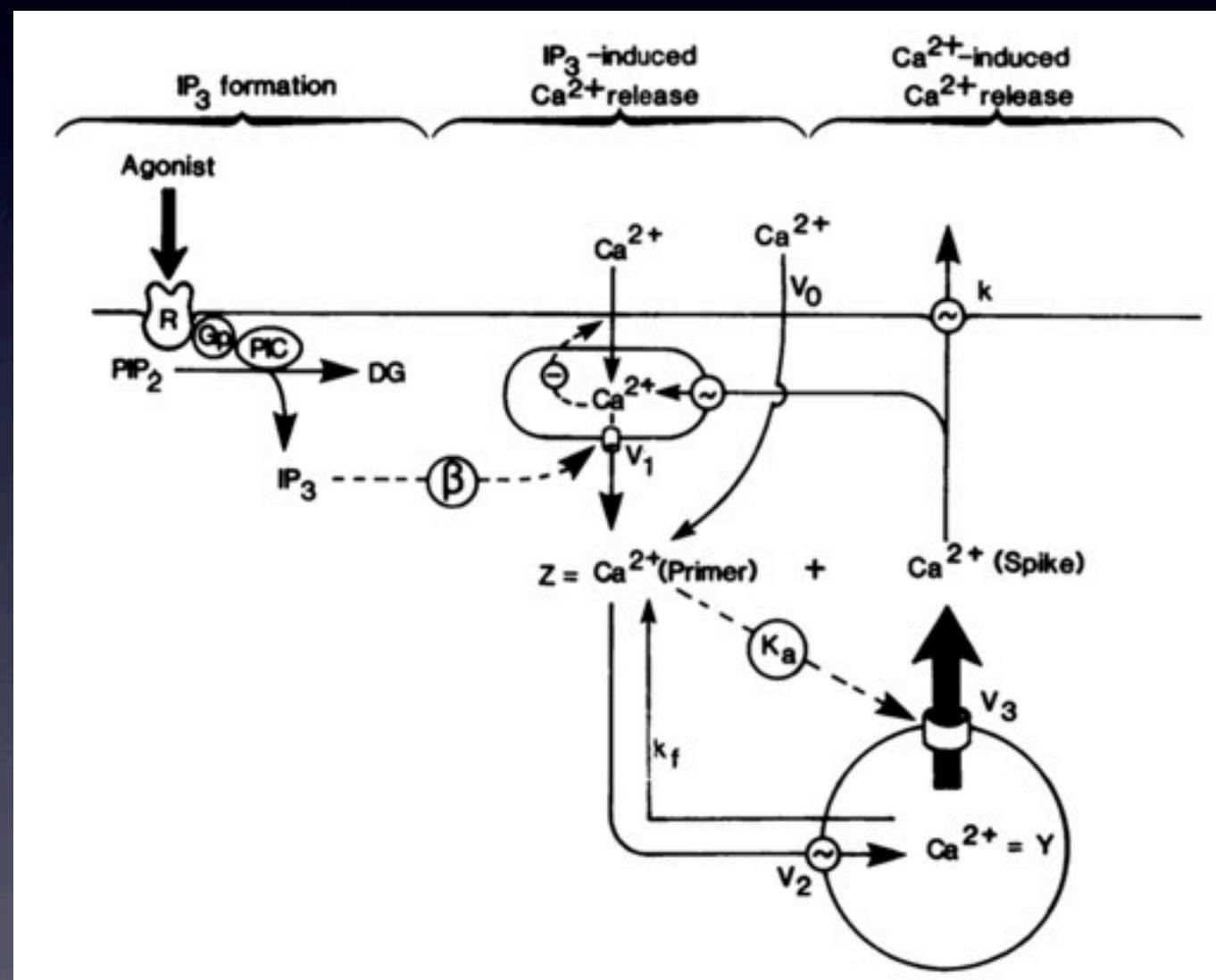
Keita Ashida

# Result

- Two Modelings by E-Cell 3
  - Simple  $\text{Ca}^{2+}$  Oscillation Model
  - Simple Cyclin Oscillation Model
- Pairwise Alignment Program by Python

# Modelings

## ● Simple $\text{Ca}^{2+}$ Oscillation Model

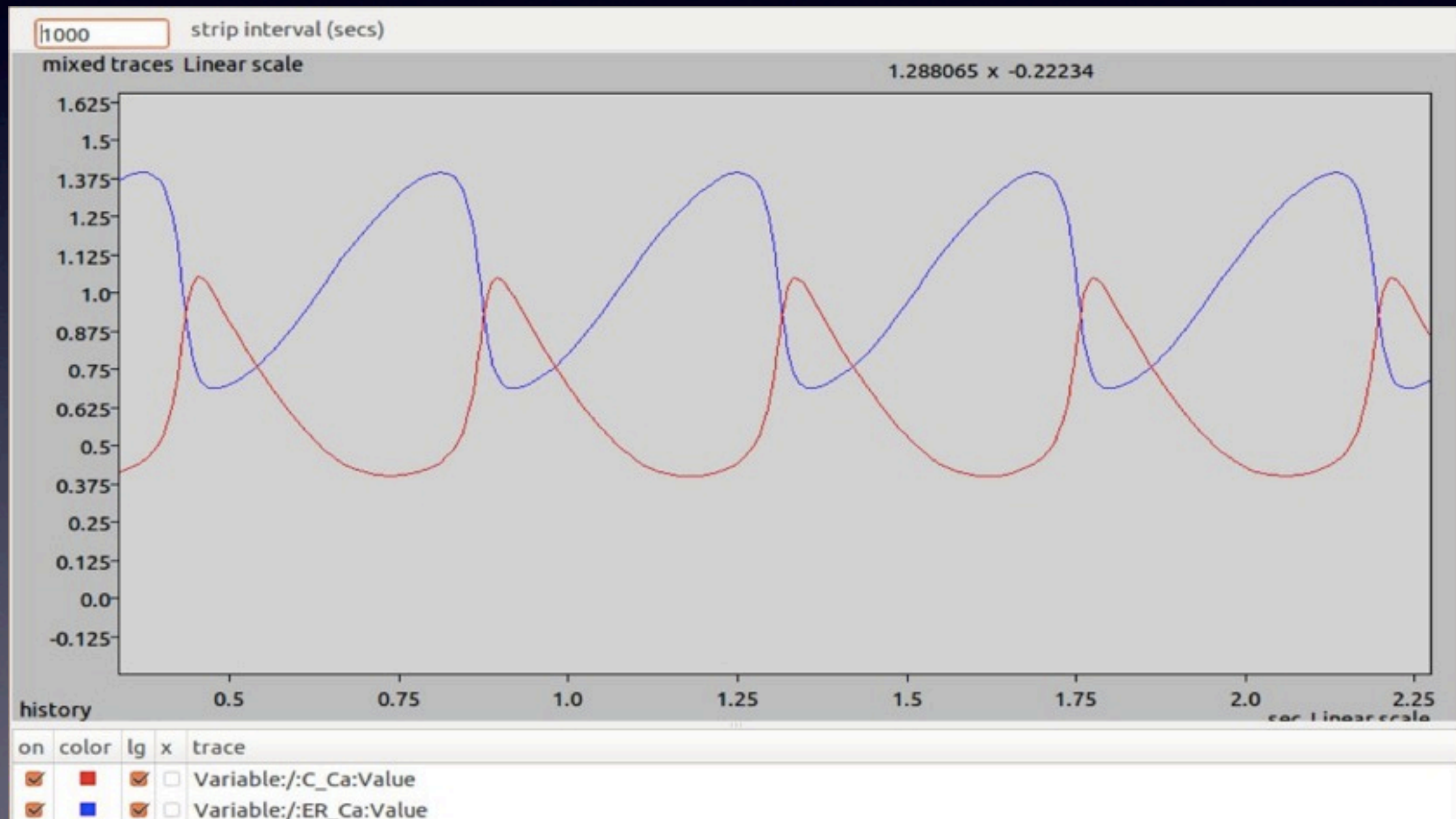


Goldbeter, A., Dupont, G., & Berridge, M. J. Minimal model for signal-induced  $\text{Ca}^{2+}$  oscillations and for their frequency encoding through protein phosphorylation. *Proc. Natl. Acad. Sci. USA* 87, 1461-1465(1990).



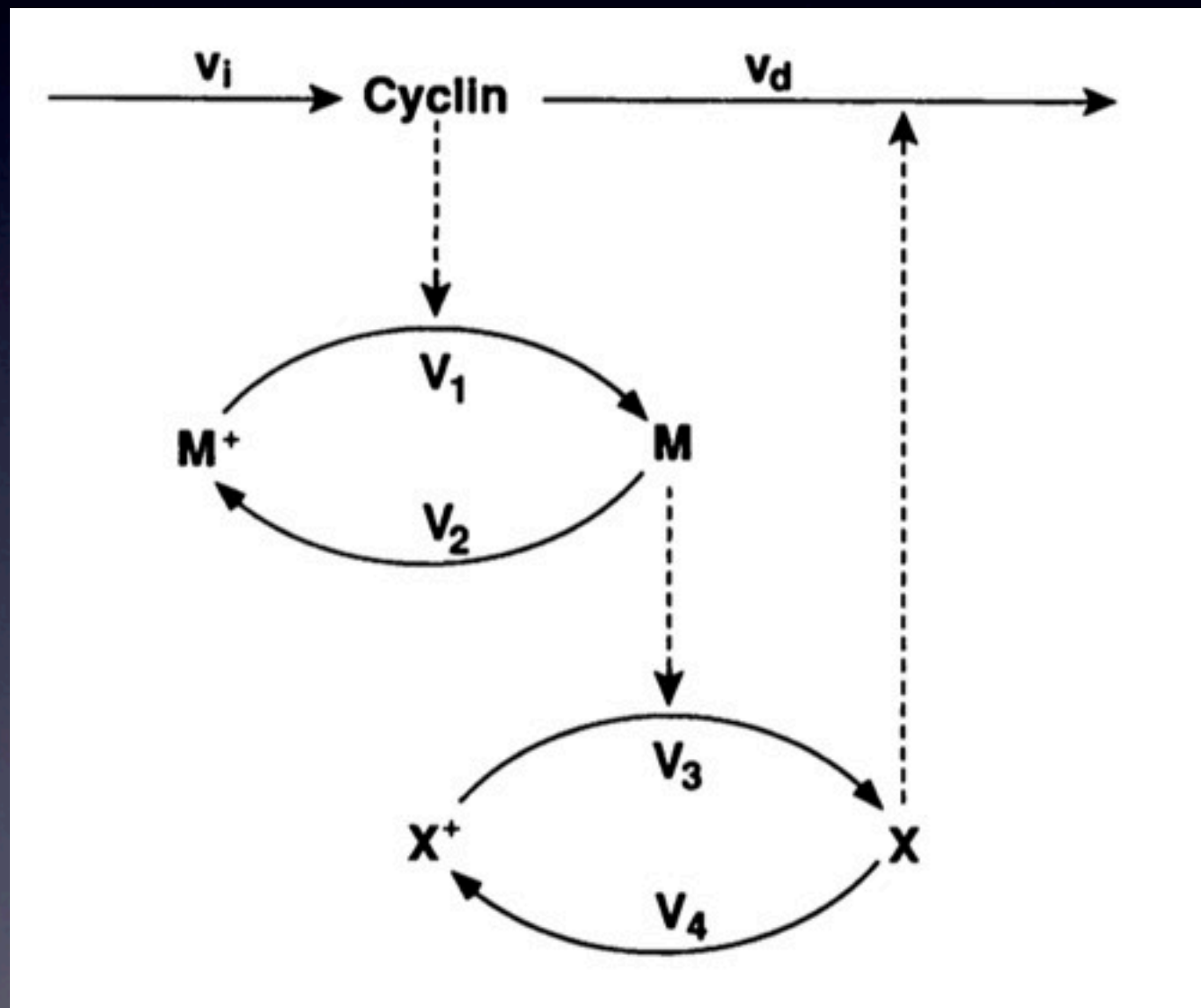
# Modelings

- Simple Cyclin Oscillation Model



# Modelings

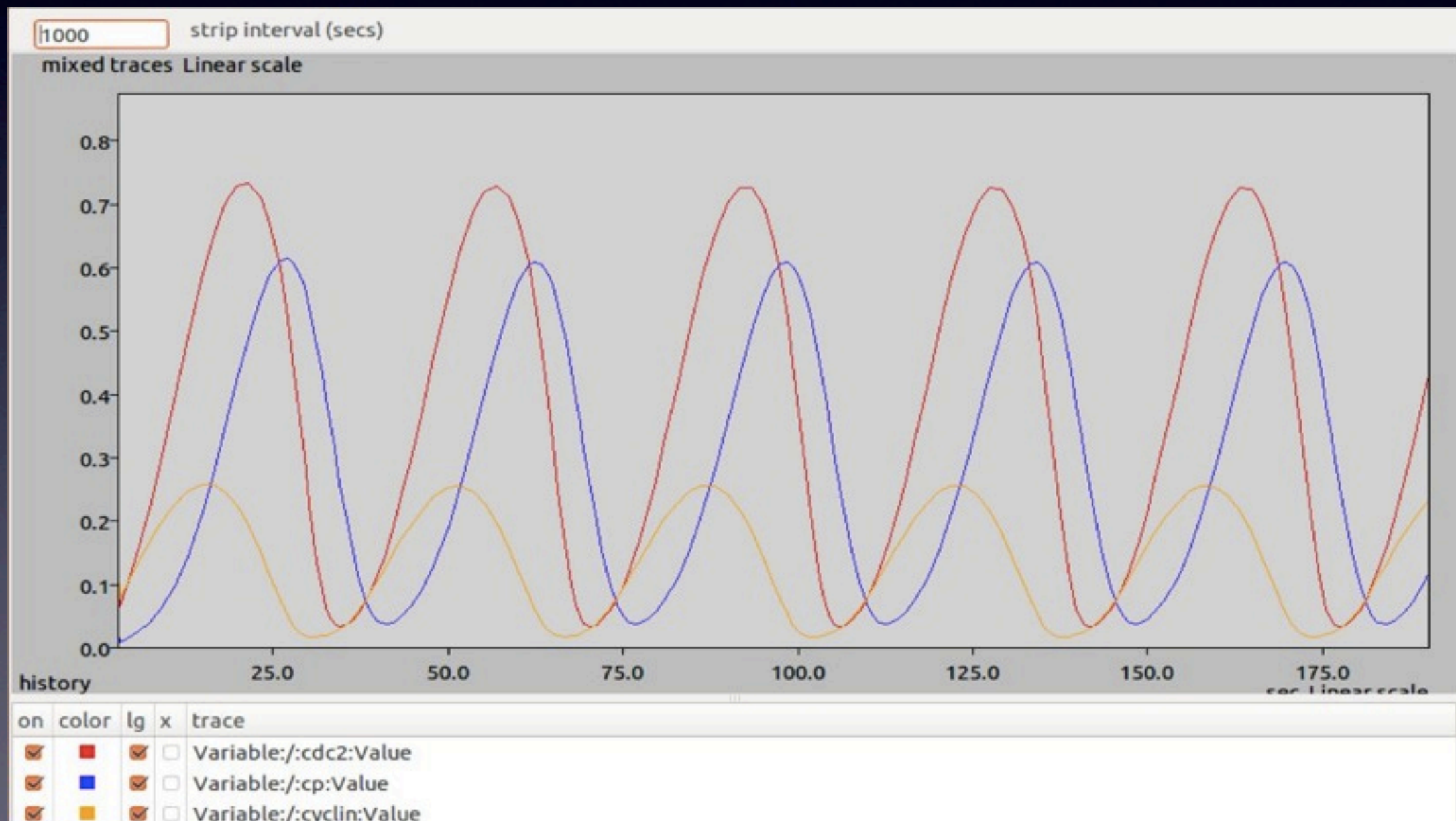
- Simple Cyclin Oscillation Model



Goldbeter, A. A minimal cascade model for the mitotic oscillator involving cyclin and cdc2 kinase. *Proc. Natl. Acad. Sci. USA* 88, 9107-9111(1991).

# Modelings

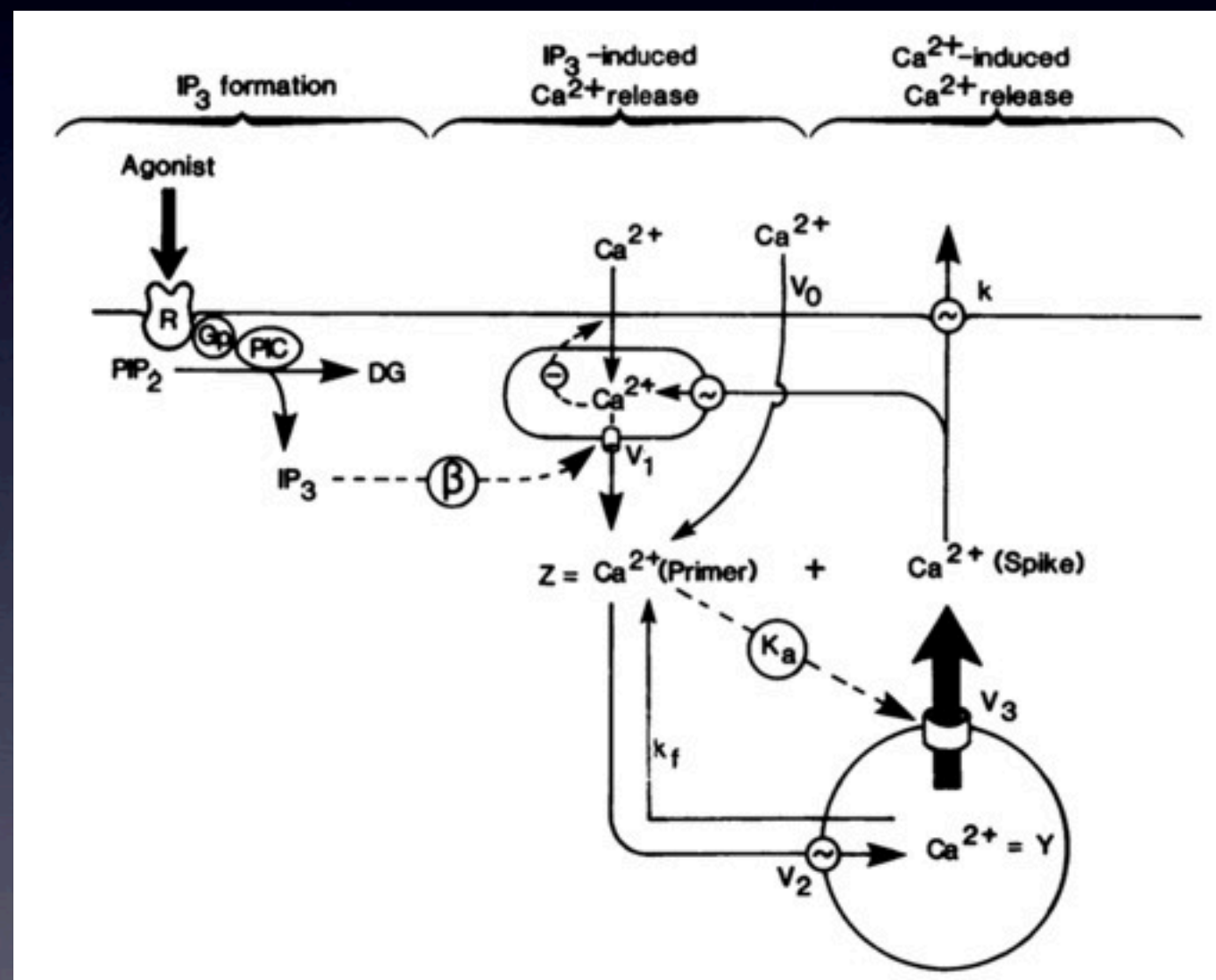
- Simple Cyclin Oscillation Model





# Modelings

## ● Simple $\text{Ca}^{2+}$ Oscillation Model



Goldbeter, A., Dupont, G., & Berridge, M. J. Minimal model for signal-induced  $\text{Ca}^{2+}$  oscillations and for their frequency encoding through protein phosphorylation. *Proc. Natl. Acad. Sci. USA* 87, 1461-1465(1990).

# Pairwise Alignment Program by Python

- Aligning Amino Acid Sequences
- Scoring Matrix: BLOSUM50
- Using Dynamic Programming

```
> demo1
K---SMRPDNRQTGGFSSIGPPPTLFNVYGR TSGIQNL-G---GGLNLMTHSMKSSSGIVWDDANLMRFDFKAI IKKTF RYMKNPM
> demo2
KQCHSMRPDNRQTGGFSSIG---TLFNVYGR TSGISGISGIQNLGGLNLMTHSMK-SSGIVWDDANLMRFDFKA-I-KTF RYMKN--
```



Thank You For Your Listening