Simulation3\_Calculation with Single Parameter.

No.5 is the transcription rate for Bmal1

No.1 is the transcription rate for Per1

Simulation2r\_Calculation with Multiple Parameter.

Three parameters: No.63\_No.30\_No.19

No.63: Degradation rate for Bmal1

No.30: Unbinding rate for Rev-erb to NPAS2

No.19: Unbinding rate for PER to CRY

Categorize\_bifurcation

By calculation with numbers of Hopf and SNIC

