* The description part.
  + Welcome:
    - It is not a “cox cluster model of pulse location pattern”
    - It Is not “smooth changing baseline” in the third paragraph.
  + General
    - The description of the default value of “Rate of console refresh” is 5000, and Actually it is 1000.
    - The description of “number of subject is “ is corresponding to “rate of writing to file rate”, and the default number is 11. **It is very important to correctly specify the number of subject in this version of the program, or the BAPHL will just keep on running with wrong number of subjects and wrong number of observations per subject without reporting an error.**
    - The description of the default value of “Constant birth rate” is 12, and Actually it is 1.0.
    - The description of the default value of “MCMC Iteration Count” is 500,000, and Actually it is 100,000
  + Baseline
    - The whole description is about half-life
    - It says the prior is an exponential distribution.
    - It says the default value for prior on population SD is 10, but actually it is 3.
    - It says the initial value for population SD is 3, but actually it is 0.5.
  + Half-life
    - It says the prior is an exponential distribution.
    - It says the default value for prior on population SD is 3, but actually it is 10.
    - It says the initial value for population mean is 45, but actually it is 2.5.
    - It says the proposal covariance for subject baseline and half-life is -0.85, but shows 0.9
  + Model error
    - It says the default value for shape of prior and rate of prior is 0.01 but actually it is 0.001.
  + Pulse Mass and Pulse width
    - The description says” the distribution of each mass and width is a bivariate t with 4 degree of freedom”.
    - The default values are not consistent with they are claimed to be.
  + Pulse Location
    - The description is about the variance for pulse width and mass.