

TEWA 1: Advanced Data Analysis

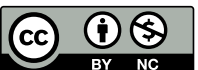
Lecture 12

Lei Zhang

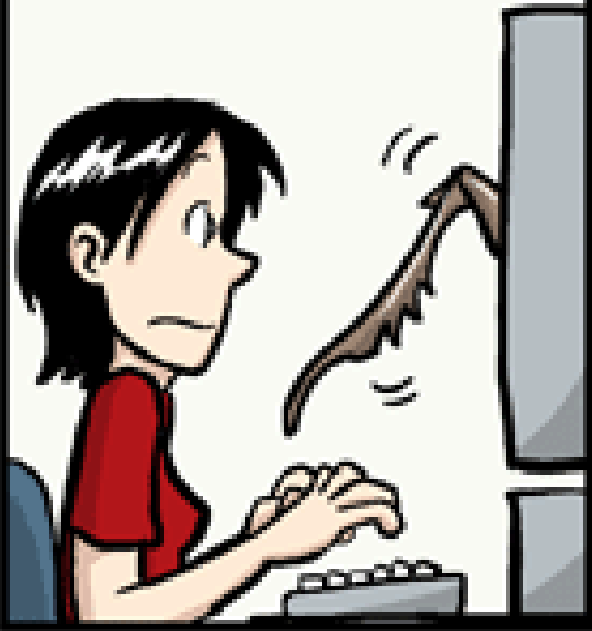
Social, Cognitive and Affective Neuroscience Unit (SCAN-Unit)
Department of Cognition, Emotion, and Methods in Psychology

https://github.com/lei-zhang/tewa1_univie

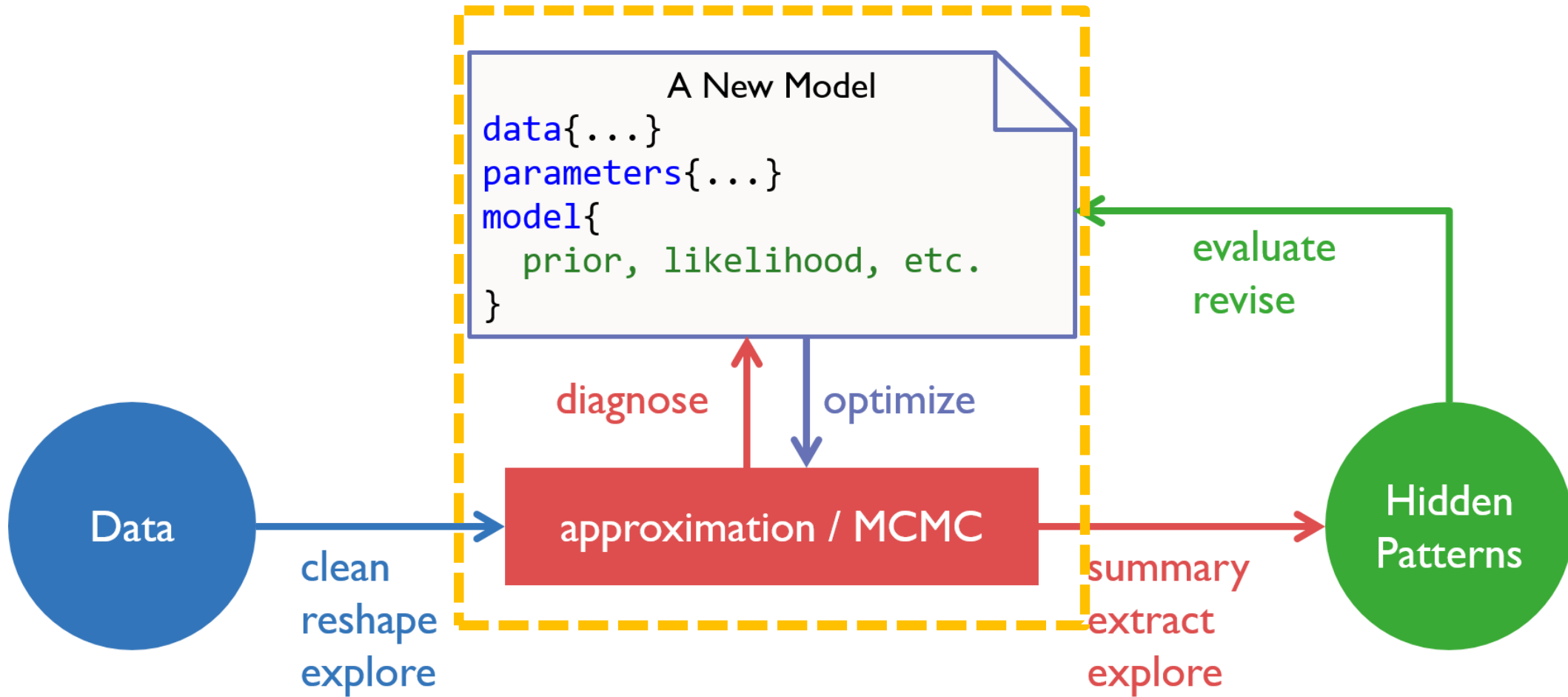
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STAN DEBUGGING



JORGE CHAM (C) 2005



Stan Style Tips

cognitive model

statistics

computing

Make it Reproducible

- Scripts are good documentations!
- Save your seed (not cross platform*)

Make it Readable

- Choose a consistent style
- Give meaningful variable names

Start with Simulated Data

Design Top-Down, Code Bottom-Up

Write Comments

- Code never lies!



* [Stan seed depends on hardware etc.](#)

The Editor of your Choice

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```
data {  
  int<lower=0> w;  
  int<lower=0> N;  
}
```

```
parameters {  
  real<lower=0,upper=1> p;  
}
```

```
model {  
  p ~ uniform(0,1);  
  w ~ binomial(N, p);  
}
```

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* Click on each logo to visit their homepage.

** [Comparison](#)

Common Error / Warning Types

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computing

ERRORS

- forget “ ; ”
- mis-indexing: mismatch or constant support mismatch
- improper constrain
- improper dimension declaration
- vectorizing when not supported
- wrong data type
- wrong distribution names
- forget priors
- miss spelling

WARNINGS


- forget last blank line
- use earlier version of Stan
- numerical problems
- divergent transitions
- hit max_treedepth
- BFMI too low
- improper prior

Debugging in Stan

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- always use a *.stan file
- press  in RStudio
- use `lookup()`
- start with simulated data
- be careful with copy/paste
- run 1 chain, 1 sample
- debugging by printing

```
for (s in 1:1) {  
  vector[2] v;  
  real pe;  
  v <- initV;  
  
  for (t in 1:nTrials) {  
    choice[s,t] ~ categorical_logit( tau[s] * v );  
  
    print("s = ", s, ", t = ", t, ", v = ", v);  
  
    pe <- reward[s,t] - v[choice[s,t]];  
    v[choice[s,t]] <- v[choice[s,t]] + lr[s] * pe;  
  }  
}
```

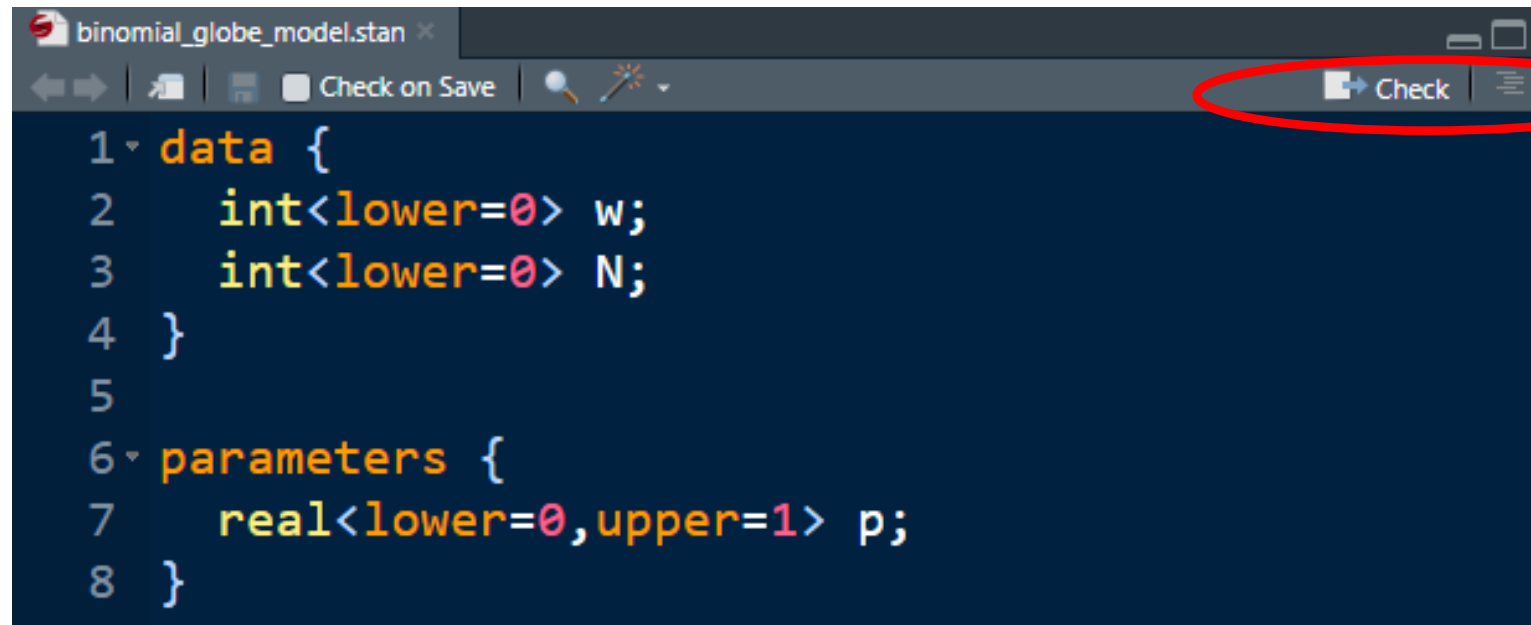
```
> lookup(dnorm)  
StanFunction Arguments ReturnType Page SamplingStatement  
344      normal      (reals mu, reals sigma)      real   369          TRUE  
348  normal_log (reals y, reals mu, reals sigma)      real   369          FALSE
```

Debugging Stan in RStudio

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```
1 data {  
2   int<lower=0> w;  
3   int<lower=0> N;  
4 }  
5  
6 parameters {  
7   real<lower=0,upper=1> p;  
8 }
```

```
rstan::rstudio_stanc("_scripts/binomial_globe_model.stan")
```

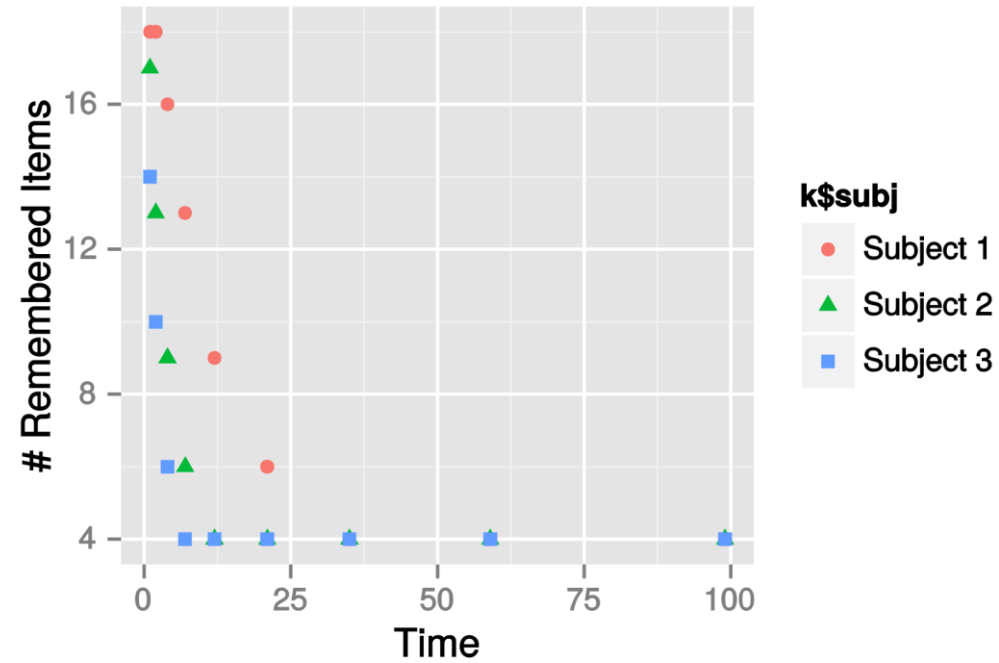



Example: Memory Retention

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Subject	Time Interval								
	1	2	4	7	12	21	35	59	99
1	18	18	16	13	9	6	4	4	4
2	17	13	9	6	4	4	4	4	4
3	14	10	6	4	4	4	4	4	4

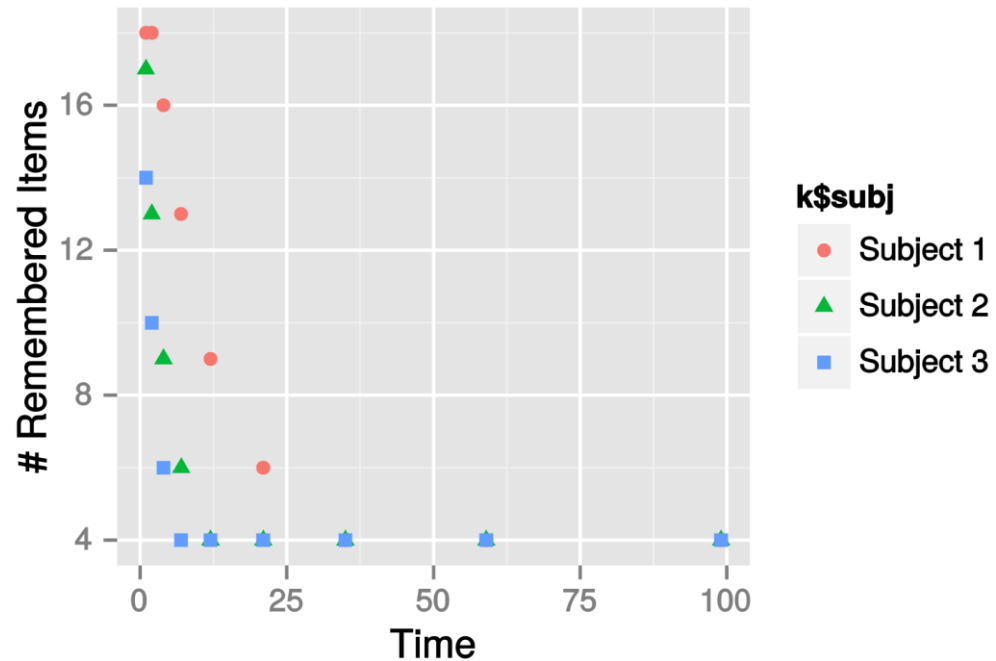


Simple Exponential Decay Model

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$$\theta_t = \min(1.0, \exp(-\alpha t) + \beta)$$

$p(\text{remember})$

decay rate

baseline

Exercise XIV

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.../09.debugging/_scripts/exp_decay_main.R

TASK: Debugging the Memory retention model

≥ 9 errors!

Viel Spaß!

```
> dataList
$`k`
      [,1] [,2] [,3] [,4] [,5] [,6] [,7] [,8] [,9]
[1,]   18   18   16   13    9    6    4    4    4
[2,]   17   13    9    6    4    4    4    4    4
[3,]   14   10    6    4    4    4    4    4    4

$nItem
[1] 18

$intervals
[1] 1 2 4 7 12 21 35 59 99

$ns
[1] 3

$nt
[1] 9
```

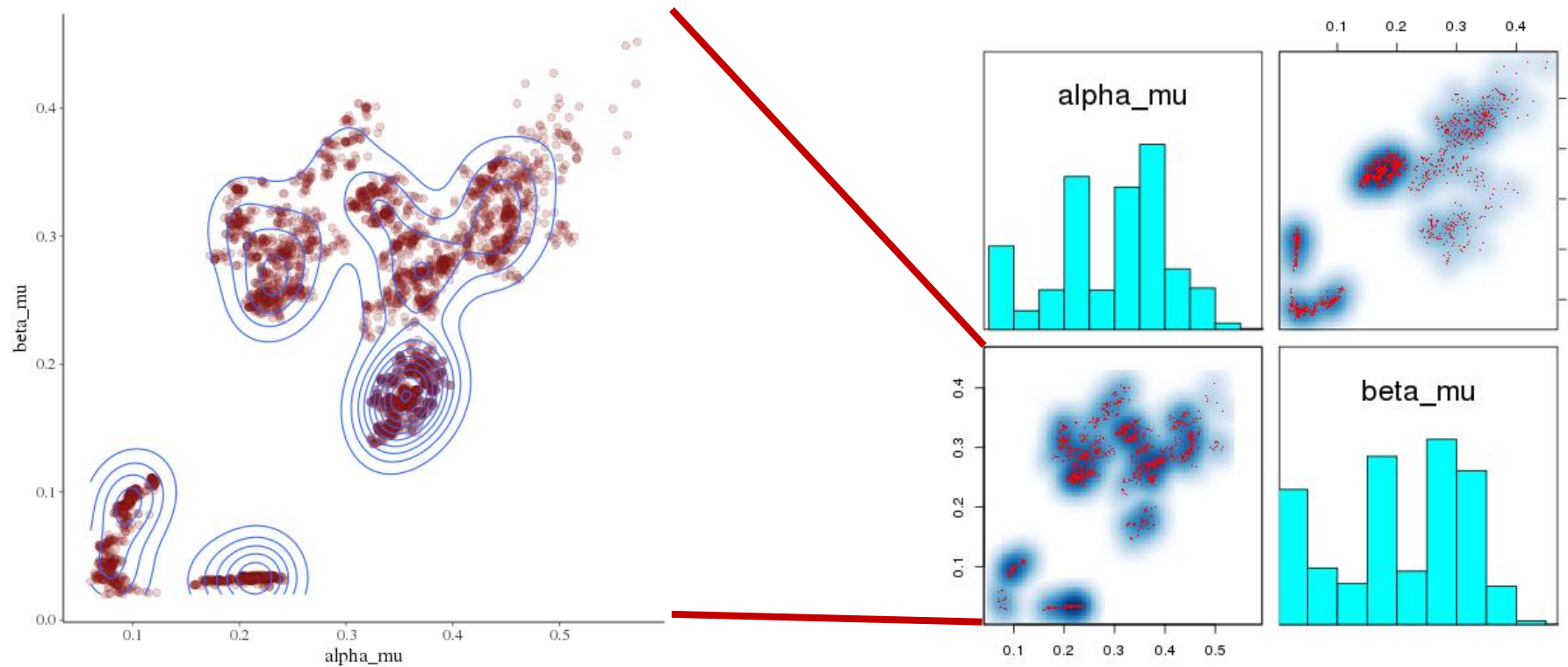
Satisfied with the results?

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```
Warning messages:  
1: There were 3998 divergent transitions after warmup. Increasing adapt_delta above 0.8 may help. See  
http://mc-stan.org/misc/warnings.html#divergent-transitions-after-warmup  
2: Examine the pairs() plot to diagnose sampling problems
```



Why Stan Fails?

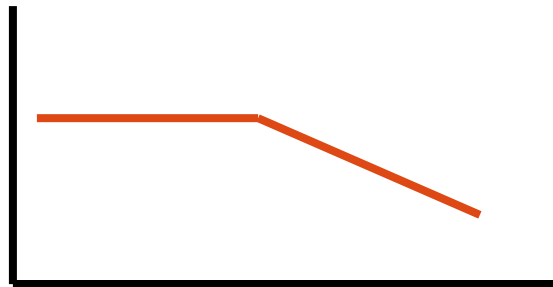
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```
for (s in 1:ns) {  
  for (t in 1:nt) {  
    theta[s,t] = fmin(1.0, exp(-alpha[s] * intervals[t]) + beta[s]);  
    k[s,t] ~ binomial(nItem, theta[s,t]);  
  }  
}
```

Non-differentiable link (likelihood) functions are bad news, particularly in Stan, which relies on derivatives.



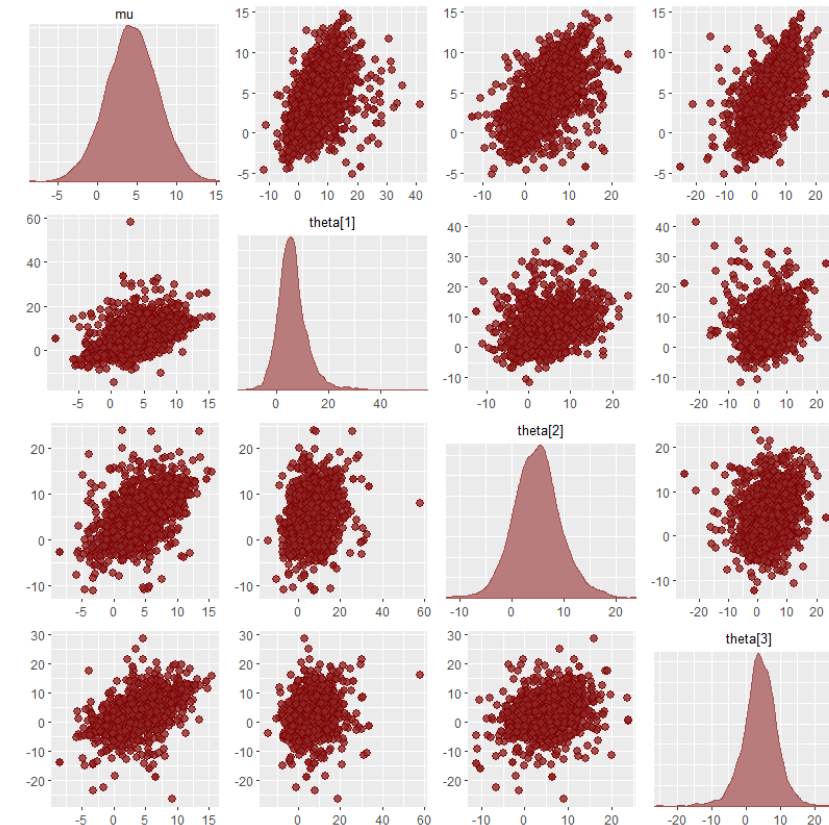
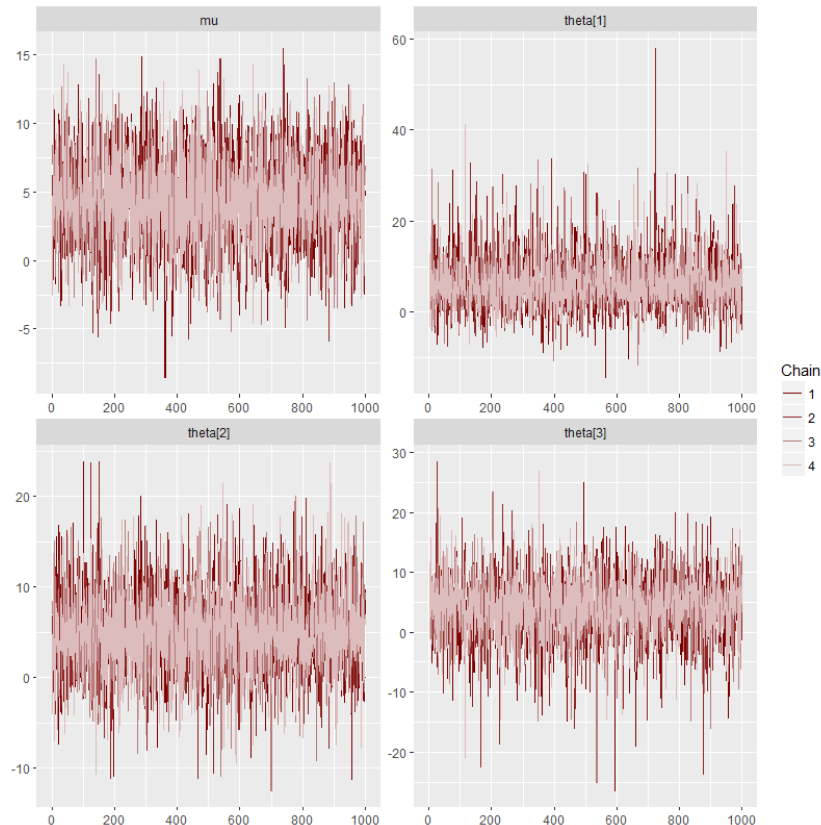
What to look for?

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```
> source('stan_utility.R')
> check_all_diagnostics(fit)
[1] "n_eff / iter looks reasonable for all parameters"
[1] "Rhat looks reasonable for all parameters"
[1] "0 of 4000 iterations ended with a divergence (0%)"
[1] "0 of 4000 iterations saturated the maximum tree depth of 10 (0%)"
[1] "E-BFMI indicated no pathological behavior"
```



ANY
QUESTIONS
?

Happy Computing!