

# Signal detection theory

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## A psychological model: Signal detection theory

$$\mathcal{M}_{sdt} : \begin{cases} \delta \sim N(1, 1) & \beta \sim N(0, 1) \\ \phi_h = \Phi(\delta/2 - \beta) & \phi_f = \Phi(-\delta/2 - \beta) \\ h \sim B(\phi_h, n_s) & f \sim B(\phi_f, n_n) \end{cases}$$

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```
model {  
  d ~ dnorm(1, 1)  
  b ~ dnorm(0, 1)  
  
  phih <- phi( d / 2 - b)  
  phif <- phi(-d / 2 - b)  
  
  h ~ dbin(phih, sigtrials)  
  f ~ dbin(phif, noistrials)  
}
```

## Signal detection theory ~ implementation

```
library(rjags)
data <- list( h = 60 ,   sigtrials  = 100 ,
              f = 11 ,   noistrials = 100 )
modelString = "
  model {
    d ~ dnorm(1, 1)
    b ~ dnorm(0, 1)

    phih <- phi( d / 2 - b)
    phif <- phi(-d / 2 - b)

    h ~ dbin(phih, sigtrials)
    f ~ dbin(phif, noistrials)
  }
"
```

## Signal detection theory ~ implementation

```
writeLines( modelString , con = "sdt.txt" )

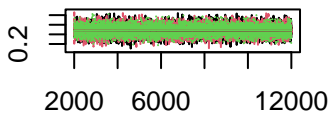
jagsModel = jags.model( file      = "sdt.txt" ,
                        data      =      data ,
                        n.chains =      3 ,
                        n.adapt  =     1000 )

set.seed(0)
update( jagsModel , n.iter = 1000 ) # burn-in

samples = coda.samples( jagsModel ,
                        variable.names = c("d", "b") ,
                        n.iter        =     10000 )
```

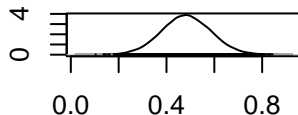
## Signal detection theory ~ results

**Trace of  $b$**



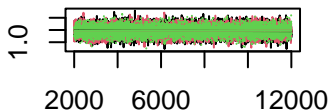
Iterations

**Density of  $b$**



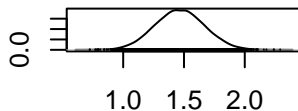
$N = 10000$  Bandwidth = 0.0139

**Trace of  $d$**



Iterations

**Density of  $d$**



$N = 10000$  Bandwidth = 0.0275

# Signal detection theory ~ summary statistics

```
summary(samples)$statistics
```

##	Mean	SD	Naive SE	Time-series SE
## b	0.4831641	0.1035147	0.0005976422	0.0007923464
## d	1.4660096	0.2038841	0.0011771255	0.0016408567

```
summary(samples)$quantiles
```

##	2.5%	25%	50%	75%	97.5%
## b	0.2799827	0.4131269	0.482196	0.5526179	0.6884915
## d	1.0710899	1.3263457	1.465063	1.6026530	1.8658863

# Signal detection theory ~ convergence

```
effectiveSize(samples)
```

```
##           b           d  
## 17112.68 15441.89
```

```
gelman.diag(samples)
```

```
## Potential scale reduction factors:
```

```
##
```

```
##   Point est. Upper C.I.
```

```
## b           1           1
```

```
## d           1           1
```

```
##
```

```
## Multivariate psrf
```

```
##
```

```
## 1
```