# **Multinomial Processing Tree with JAGS**

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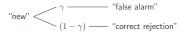
Probability of a correct rejection = correct rejection rate =  $1-\theta^{\rm f}$ 

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The parameters  $\rho$  and  $\gamma$  together determine the hit rate  $\theta^{\rm h}$  and false alarm rate  $\theta^{\rm f}$ 

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Those rates tell us how often old items lead to hits (the data are  $k^{\rm h}$  hits out of  $n_o$  old items) and how often new items lead to false alarms ( $k^{\rm f}$  false alarms out of  $n_n$  new items):

$$k^{\rm h} \sim {\rm binomial}\left(\theta^{\rm h}, n_o\right)$$
  
 $k^{\rm f} \sim {\rm binomial}\left(\theta^{\rm f}, n_n\right)$ 

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\begin{array}{lll} \rho & \sim & \mathrm{uniform}\left(0,1\right) \\ \gamma & \sim & \mathrm{uniform}\left(0,1\right) \\ \theta^{\mathrm{h}} & = & \rho + \left(1-\rho\right)\gamma \\ \theta^{\mathrm{f}} & = & \gamma \\ k^{\mathrm{h}} & \sim & \mathrm{binomial}\left(\theta^{\mathrm{h}}, n_{o}\right) \\ k^{\mathrm{f}} & \sim & \mathrm{binomial}\left(\theta^{\mathrm{f}}, n_{n}\right) \end{array}
```

$\rho$	$\sim$	$\mathrm{uniform}(0,1)$
$\gamma$	$\sim$	uniform $(0,1)$
$ heta^{ m h}$	=	$\rho + (1 - \rho) \gamma$
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$k^{\mathrm{h}}$	$\sim$	binomial $(\theta^{\rm h}, n_o)$
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Amyloid Status	Hits	False Alarms
negative	13	0
positive	8	4
negative	12	1
negative	14	0
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Amyloid Status	Hits	False Alarms
positive	8	4
positive	9	4
positive	14	0
positive	14	1
positive	13	2

```
\begin{array}{lll} & \operatorname{model}\{ \\ \rho & \sim & \operatorname{uniform}\left(0,1\right) \\ \gamma & \sim & \operatorname{uniform}\left(0,1\right) \\ \theta^{\mathrm{h}} & = & \rho + (1-\rho)\gamma \\ \theta^{\mathrm{f}} & = & \gamma \\ & & \forall p \in (1,\dots,P) \\ k_p^{\mathrm{h}} & \sim & \operatorname{binomial}\left(\theta^{\mathrm{h}}, n_o\right) \\ k_p^{\mathrm{f}} & \sim & \operatorname{binomial}\left(\theta^{\mathrm{f}}, n_n\right) \\ k_p^{\mathrm{f}} & \sim & \operatorname{binomial}\left(\theta^{\mathrm{f}}, n_n\right) \\ \end{array} \right. \left. \begin{array}{ll} \operatorname{model}\{ \\ \text{rho} & \tilde{} & \operatorname{dunif}(0, 1) \\ \text{gamma} & \tilde{} & \operatorname{dunif}(0, 1) \\ \text{thetaHit} & = & \operatorname{rho} + (1-\operatorname{rho}) * \operatorname{gamma} \\ \text{thetaFA} & = & \operatorname{gamma} \\ \text{for (p in 1:nPeople)}\{ \\ \text{hit[p]} & \tilde{} & \operatorname{dbin(thetaHit, nOld)} \\ \text{fa[p]} & \tilde{} & \operatorname{dbin(thetaFA, nNew)} \\ \} \\ \end{array} \right. \right\}
```

Patients remember around 60-70% of the items, and guess "old" 5-10% of the time when they do not remember

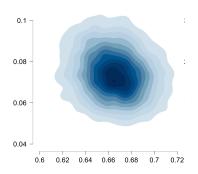
	Posterior			95% Cr	red. Int.
Parameter	Mean	Median	SD	Lower	Upper
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"Convergence of the MCMC procedure was good, with all  $\hat{R} < 1.01$ ."

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