Hierarchical Normal Model

Congulation time (seconds) for randomly drawn blood

Diet	Measurements
A	62, 60, 63, 59
В	63, 67, 71, 64, 65, 66
\mathbf{C}	68, 66, 71, 67, 68, 68
D	56, 62, 60, 61, 63, 64, 63, 59

Posterior from 10 Gibbs sequences of length 100

Estimand	Posterior quantiles					\widehat{R}
	2.5%	25%	median	75%	97.5%	
θ_1	58.9	60.6	61.3	62.1	63.5	1.01
$ heta_{2}$	63.9	65.3	65.9	66.6	67.7	1.01
$ heta_3$	66.0	67.1	67.8	68.5	69.5	1.01
$ heta_4$	59.5	60.6	61.1	61.7	62.8	1.01
$oldsymbol{\mu}$	56.9	62.2	63.9	65.5	73.4	1.04
σ	1.8	2.2	2.4	2.6	3.3	1.00
au	2.1	3.6	4.9	7.6	26.6	1.05
$\frac{1}{\log p(\mu, \log \sigma, \log \tau y)}$	-67.6	-64.3	-63.4	-62.6	-62.0	1.02
$\log p(heta, \mu, \log \sigma, \log au y)$	-70.6	-66.5	-65.1	-64.0	-62.4	1.01

Hierarchical Normal Model

		Stepwise ascent			
Parameter	Crude estimate	First iteration	Second iteration	Third iteration	
$\overline{\hspace{0.1cm}}^{\hspace{0.1cm}} \overline{\hspace{0.1cm}}_{\hspace{0.1cm} 1}$	61.00	61.28	61.29	61.29	
$ heta_2$	66.00	65.87	65.87	65.87	
$ heta_3$	68.00	67.74	67.73	67.73	
$ heta_4$	61.00	61.15	61.15	61.15	
μ	64.00	64.01	64.01	64.01	
σ	2.29	2.17	2.17	2.17	
au	3.56	3.32	3.31	3.31	
$\log p(heta, \mu, \log \sigma, \log au y)$	-63.70	-61.42	-61.42	-61.42	

		EM algorithm		
Parameter	Value at joint mode	First iteration	Second iteration	Third iteration
μ	64.01	64.01	64.01	64.01
σ	2.17	2.33	2.36	2.36
au	3.31	3.46	3.47	3.47
$\log p(\mu, \log \sigma, \log \tau y)$	-61.99	-61.835	-61.832	-61.832

Parameter	r Posterior quantiles				
	2.5%	25%	median	75%	97.5%
θ_1	59.15	60.63	61.38	62.18	63.87
$oldsymbol{ heta_2}$	63.83	65.20	65.78	66.42	67.79
$ heta_3$	65.46	66.95	67.65	68.32	69.64
$ heta_4$	59.51	60.68	61.21	61.77	62.99
${m \mu}$	60.43	62.73	64.05	65.29	67.69
σ	1.75	2.12	2.37	2.64	3.21
au	1.44	2.62	3.43	4.65	8.19