TEST

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Contents

0.1 Functions

Table 1: CAPTION FOR TABLE

Parameter	Estimate	Est.Error	l-95% CI	u-95% CI	Rhat	Bulk_ESS	Tail_ESS
Group-Level Effects:							
fkClientID (Number of levels: 1034)							
sd(Intercept)	0.10	0.01	0.08	0.12	1.00	599	1 106
Population-Level Effects:							
Intercept	-0.03	0.04	-0.11	0.04	1.00	4 700	5 335
PHQ9_baseline	-3.86	0.08	-4.03	-3.70	1.00	3 647	4 904
PHQ9_change	-2.45	0.13	-2.70	-2.20	1.00	1 342	3 300
SOFAS_baseline	0.01	0.06	-0.10	0.12	1.00	4 767	5 393
SOFAS_change	0.41	0.07	0.27	0.55	1.00	6 621	6 049
Family Specific Parameters:							
sigma	0.10	0.00	0.09	0.10	1.01	661	1 473

Formula: aqol6d_total_w ~PHQ9_baseline + PHQ9_change + SOFAS_baseline + SOFAS_change + (1 | fkClientID)

Family: gaussian

Links: mu = log; sigma = identity

Data: data_tb (Number of observations: 1651)

Samples: 4 chains, each with iter = 4000; warmup = 2000; thin = 1;

total post-warmup samples = 8000

Samples were drawn using sampling(NUTS). For each parameter, Bulk_ESS

and Tail_ESS are effective sample size measures, and Rhat is the potential

scale reduction factor on split chains (at convergence, Rhat = 1).

```
#rlang::exec(print_table_xx,!!!brms_mdl_print_ls[[5]])
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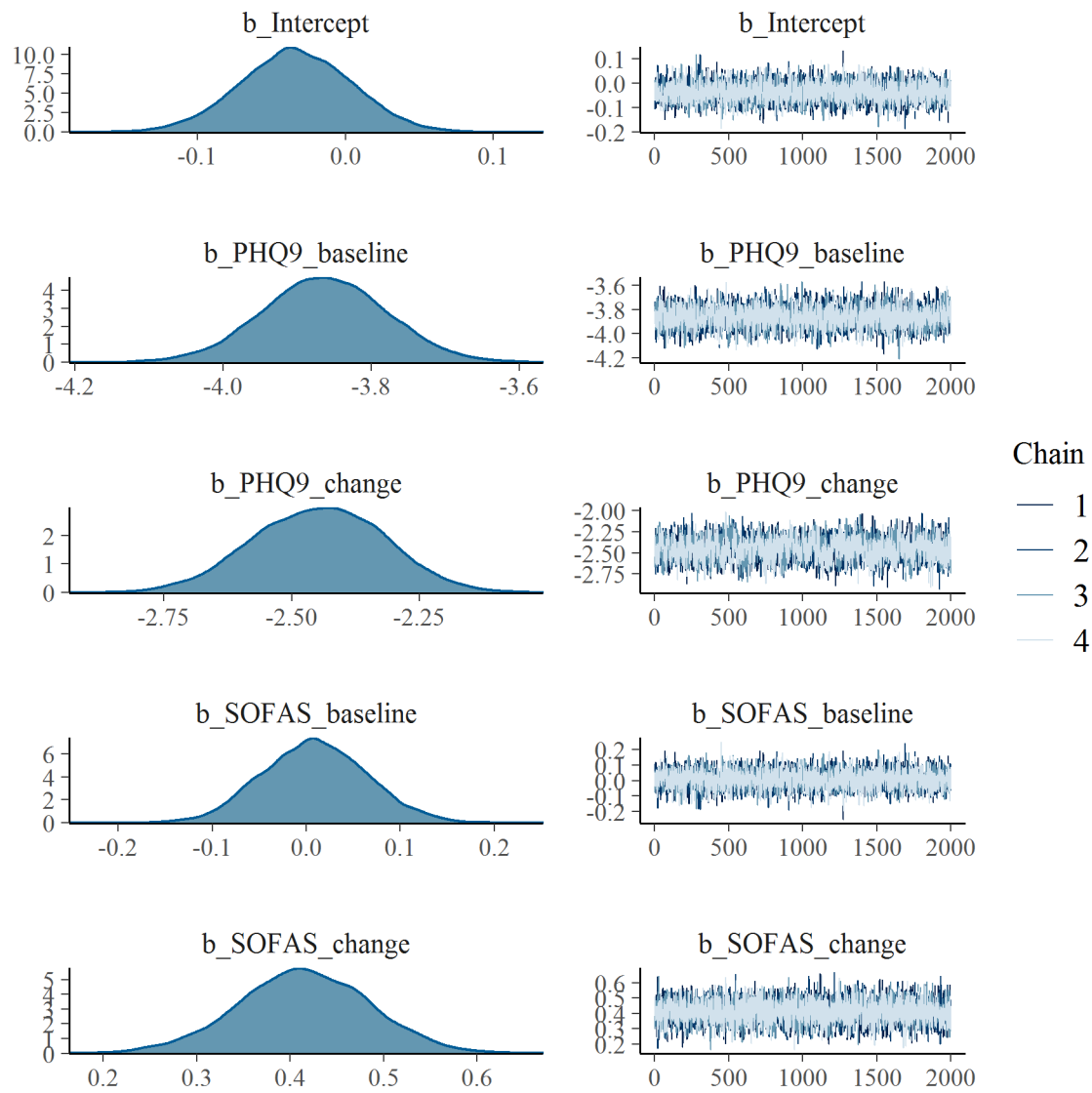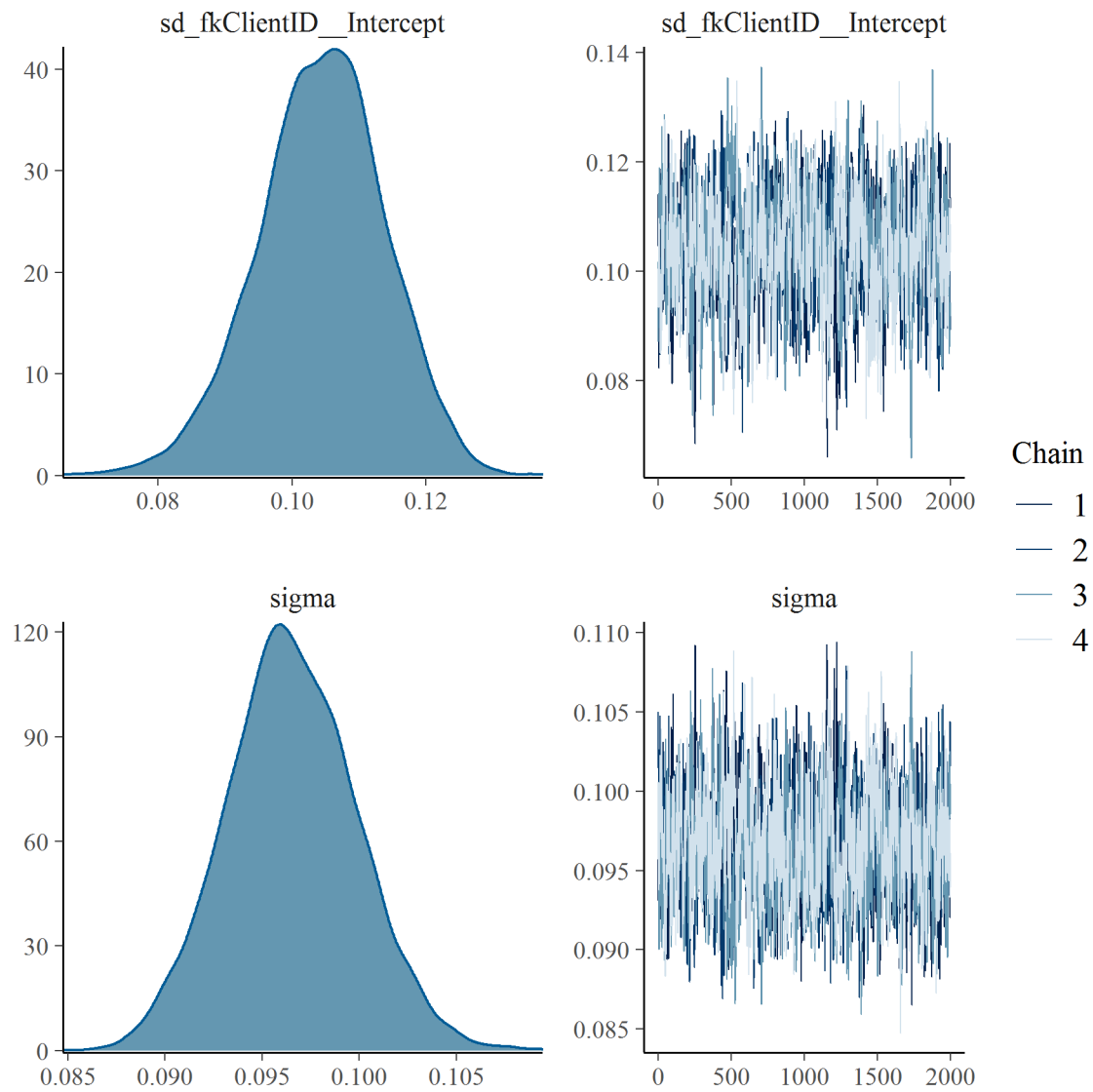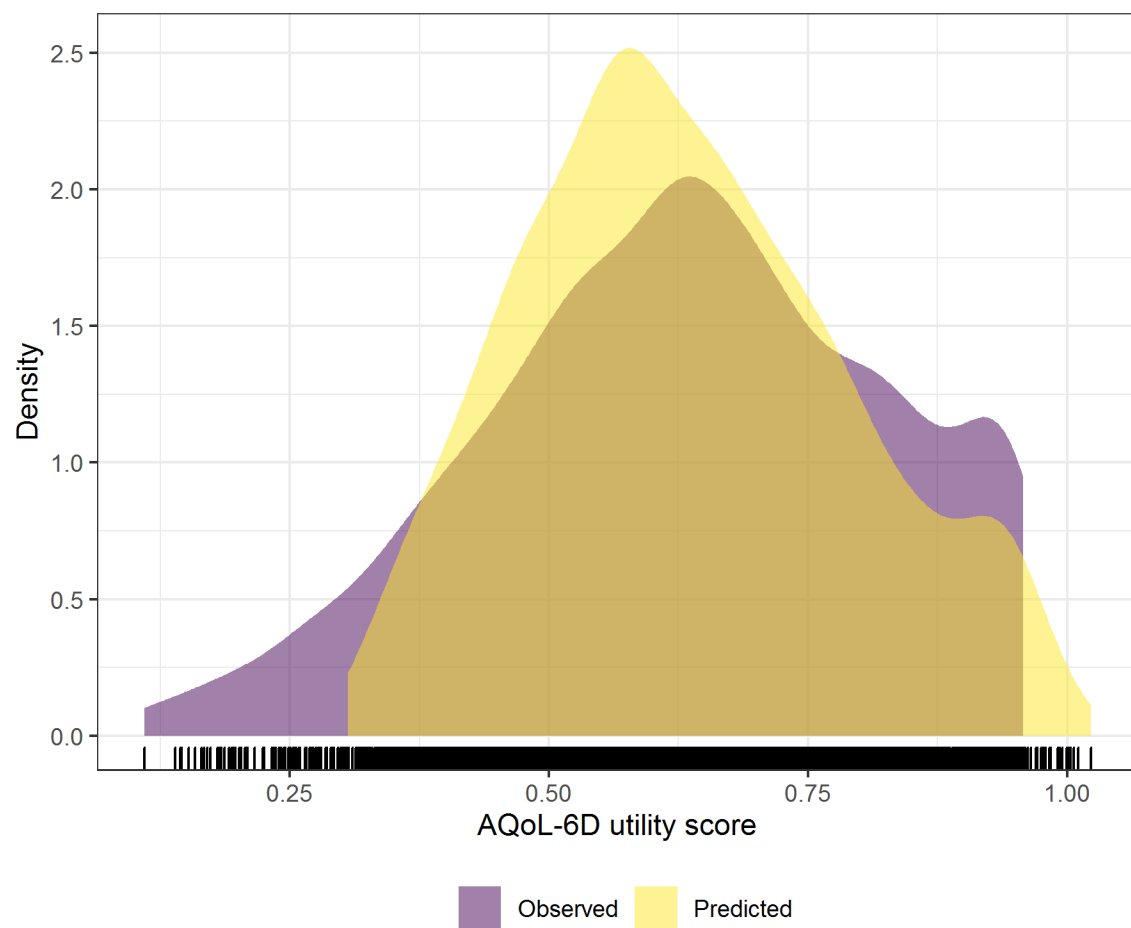
Figure 1

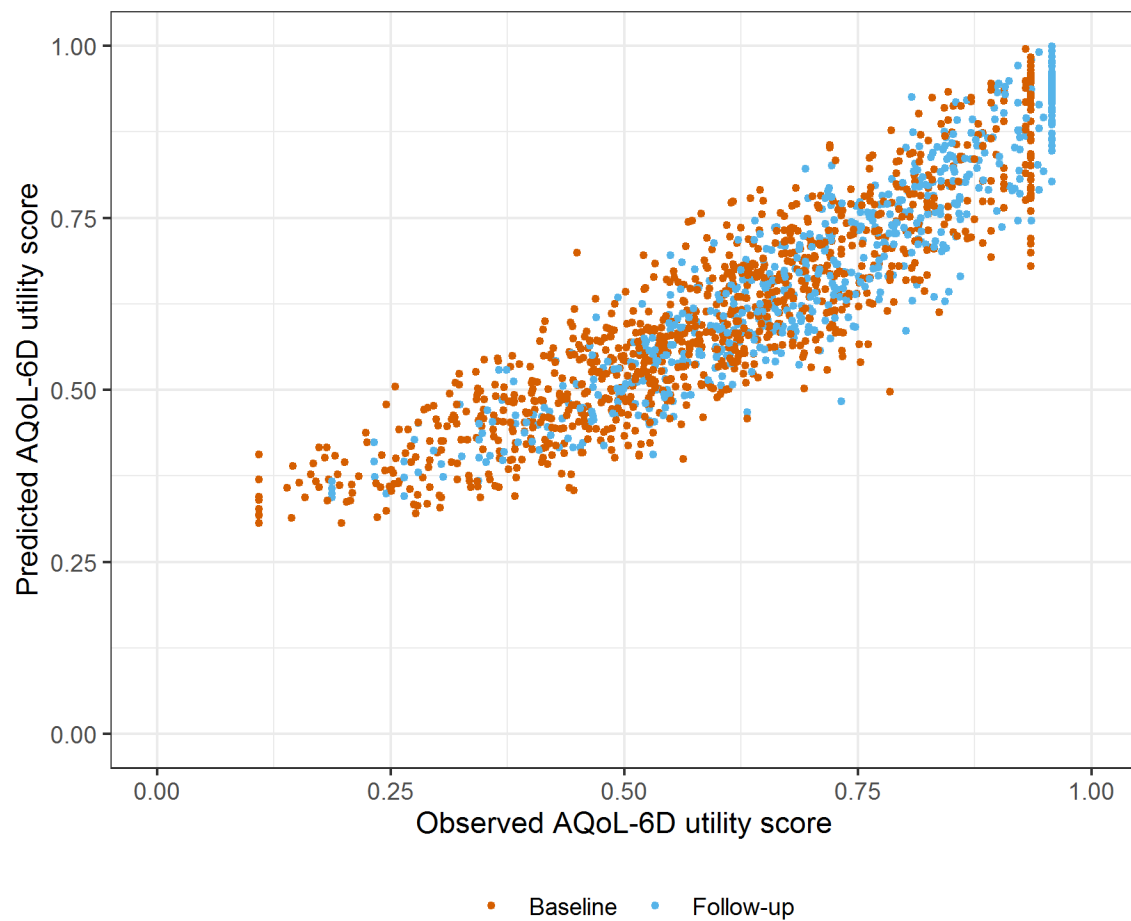Figure 1: Caption 1

Figure 2: Caption 2

Figure 3: Caption 3

Figure 4: Caption 4

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### TEST
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