

# Appendix for “Comparing methods for estimating patient-specific treatment effects in individual patient data meta-analysis”

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# 1 Descriptive statistics of the real datasets used in the paper

Table 1: Stents datasets. Separate summary is reported below for complete cases patients and patients with at least one missing covariate. *stable\_cad*: stable coronary artery disease (clinical presentation at the time of percutaneous coronary intervention); *ladtreated*: target vessel left anterior descending artery; *m\_dia\_above\_3*: mean diameter  $\geq 3$  mm; *num\_stents*: number of implanted stents.

Variable	Complete cases (n = 11106)	Patients with at least one missing covariate (n =327)
	Mean(SD) for continuous or % for binary	Mean(SD) for continuous or % for binary
binary outcome: cardiac death or myocardial infarction at 1-year after randomization	5%	7%
treatment (drug eluding stents)	56%	54%
age	68.60 (12.22)	70.47 (12.09)
gender	73%	78%
diabetes	23%	28%
stable_cad	34%	35%
multivessel	46%	52%
ladtreated	47%	48%
overlap	15%	12%
m_dia_above_3	97%	96%
num_stents	1.66 (1.05)	1.26 (0.77)

Table 2: Antidepressant dataset. Separate summary is reported below for complete cases patients and patients with at least one missing covariate. *HRSD*: 17-item Hamilton Rating Scale for Depression

Variable	Complete cases (n =1261)	Patients with at least one missing covariate (n =232)
	Mean(SD) for continuous or % for binary	Mean(SD) for continuous or % for binary
continuous outcome: depression severity at week 6 or 8	9.66 (6.19)	6.50 (4.41)
treatment (antidepressants)	71%	69%
baseline severity	20.92 (4.38)	21.70 (4.29)
age	36.72 (9.93)	39.48 (12.05)
female	50%	50%
age at onset	33.58 (10.40)	35.50 (12.37)
episode frequency over 3	16%	22%
episode duration week	42.04 (55.73)	53.53 (102.56)
guilty agitation HRSD	5.48 (2.06)	5.43 (2.12)
bodily symptoms HRSD	3.68 (1.50)	3.98 (1.59)
sleep problems HRSD	3.10 (1.68)	3.26 (1.65)
anhedonia retardation HRSD	7.48 (1.71)	7.65 (1.81)

## 2 Presentation of scenarios explored in simulations

In the following table, we present an overview of the scenarios we explored in our simulation study.

Table 3: Overview of the scenarios we explored in our simulations. In the column with the true values for effect modifier, we first present the main effect on the outcome and in the parenthesis the interaction with the treatment (i.e. effect modification). Scenario C1 models continuous outcome and scenario B1 models binary outcome.

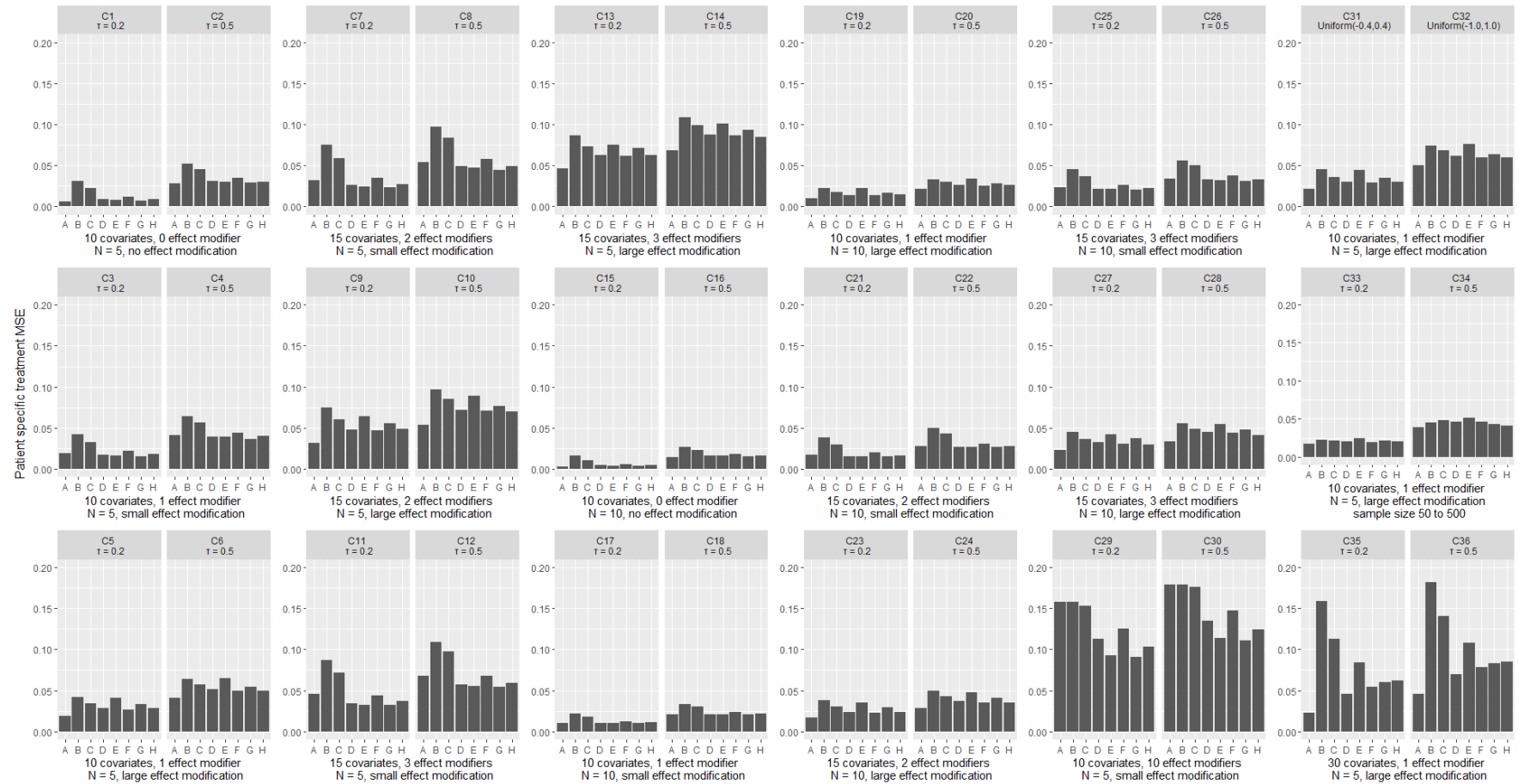
Scenarios	# of studies	# of covariates	# of nuisance covariates	# of prognostic factors	# of effect modifiers	True values for effect modifiers: main effect (effect modification)	Heterogeneity of treatment effect
C1	5	10	3 continuous 2 binary	3 continuous 2 binary	None	NA	$\tau=0.2$
C2							$\tau=0.5$
C3	5	10	3 continuous 2 binary	2 continuous 2 binary	1 continuous	continuous: 0.2 (0.1)	$\tau=0.2$
C4							$\tau=0.5$
C5	5	10	3 continuous 2 binary	2 continuous 2 binary	1 continuous	continuous: 0.2 (0.5)	$\tau=0.2$
C6							$\tau=0.5$
C7	5	15	5 continuous 3 binary	3 continuous 2 binary	1 continuous 1 binary	continuous: 0.2 (0.1) binary: 0.2 (0.1)	$\tau=0.2$
C8							$\tau=0.5$
C9	5	15	5 continuous 3 binary	3 continuous 2 binary	1 continuous 1 binary	continuous: 0.2 (0.5) binary: 0.2 (0.4)	$\tau=0.2$
C10							$\tau=0.5$
C11	5	15	5 continuous 3 binary	2 continuous 2 binary	2 continuous 1 binary	continuous: 0.2 (0.1) binary: 0.2 (0.1)	$\tau=0.2$
C12							$\tau=0.5$
C13	5	15	5 continuous 3 binary	2 continuous 2 binary	2 continuous 1 binary	continuous: 0.2 (0.5) binary: 0.2 (0.4)	$\tau=0.2$
C14							$\tau=0.5$
C15	10	10	3 continuous 2 binary	3 continuous 2 binary	None	NA	$\tau=0.2$
C16							$\tau=0.5$
C17	10	10	3 continuous 2 binary	2 continuous 2 binary	1 continuous	continuous: 0.2 (0.1)	$\tau=0.2$
C18							$\tau=0.5$
C19	10	10	3 continuous 2 binary	2 continuous 2 binary	1 continuous	continuous: 0.2 (0.5)	$\tau=0.2$
C20							$\tau=0.5$
C21	10	15	5 continuous 3 binary	3 continuous 2 binary	1 continuous 1 binary	continuous: 0.2 (0.1) binary: 0.2 (0.1)	$\tau=0.2$
C22							$\tau=0.5$
C23	10	15	5 continuous 3 binary	3 continuous 2 binary	1 continuous 1 binary	continuous: 0.2 (0.5) binary: 0.2 (0.4)	$\tau=0.2$
C24							$\tau=0.5$
C25	10	15	5 continuous 3 binary	2 continuous 2 binary	2 continuous 1 binary	continuous: 0.2 (0.1) binary: 0.2 (0.1)	$\tau=0.2$
C26							$\tau=0.5$
C27	10	15	5 continuous 3 binary	2 continuous 2 binary	2 continuous 1 binary	continuous: 0.2 (0.5) binary: 0.2 (0.4)	$\tau=0.2$
C28							$\tau=0.5$

Scenarios	# of studies	# of covariates	# of nuisance covariates	# of prognostic factors	# of effect modifiers	True values for effect modifiers: main effect (effect modification)	Heterogeneity of treatment effect
C29	5	10	None	None	5 continuous 5 binary	continuous: 0.2 (0.1) binary: 0.2 (0.1)	$\tau=0.2$
C30							$\tau=0.5$
C31	5	10	3 continuous 2 binary	2 continuous 2 binary	1 continuous	continuous: 0.2 (0.5)	Uniform(-0.4,0.4)
C32							Uniform(-1.0,1.0)
C33	5 (sample size from 50 to 500)	10	3 continuous 2 binary	2 continuous 2 binary	1 continuous	continuous: 0.2 (0.5)	$\tau=0.2$
C34							$\tau=0.5$
C35	5	30	8 continuous 7 binary	7 continuous 7 binary	1 continuous	continuous: 0.2 (0.5)	$\tau=0.2$
C36							$\tau=0.5$
B1	5	10	3 continuous 2 binary	3 continuous 2 binary	None	NA	$\tau=0.2$
B2							$\tau=0.5$
B3	5	10	3 continuous 2 binary	2 continuous 2 binary	1 continuous	continuous: 0.2 (0.1)	$\tau=0.2$
B4							$\tau=0.5$
B5	5	10	3 continuous 2 binary	2 continuous 2 binary	1 continuous	continuous: 0.2 (0.5)	$\tau=0.2$
B6							$\tau=0.5$
B7	5	15	5 continuous 3 binary	3 continuous 2 binary	1 continuous 1 binary	continuous: 0.2 (0.1) binary: 0.2 (0.1)	$\tau=0.2$
B8							$\tau=0.5$
B9	5	15	5 continuous 3 binary	3 continuous 2 binary	1 continuous 1 binary	continuous: 0.2 (0.5) binary: 0.2 (0.4)	$\tau=0.2$
B10							$\tau=0.5$
B11	5	15	5 continuous 3 binary	2 continuous 2 binary	2 continuous 1 binary	continuous: 0.2 (0.1) binary: 0.2 (0.1)	$\tau=0.2$
B12							$\tau=0.5$
B13	5	15	5 continuous 3 binary	2 continuous 2 binary	2 continuous 1 binary	continuous: 0.2 (0.5) binary: 0.2 (0.4)	$\tau=0.2$
B14							$\tau=0.5$
B15	10	10	3 continuous 2 binary	3 continuous 2 binary	None	NA	$\tau=0.2$
B16							$\tau=0.5$
B17	10	10	3 continuous 2 binary	2 continuous 2 binary	1 continuous	continuous: 0.2 (0.1)	$\tau=0.2$
B18							$\tau=0.5$
B19	10	10	3 continuous 2 binary	2 continuous 2 binary	1 continuous	continuous: 0.2 (0.5)	$\tau=0.2$
B20							$\tau=0.5$
B21	10	15	5 continuous 3 binary	3 continuous 2 binary	1 continuous 1 binary	continuous: 0.2 (0.1) binary: 0.2 (0.1)	$\tau=0.2$
B22							$\tau=0.5$
B23	10	15	5 continuous 3 binary	3 continuous 2 binary	1 continuous 1 binary	continuous: 0.2 (0.5) binary: 0.2 (0.4)	$\tau=0.2$
B24							$\tau=0.5$
B25	10	15	5 continuous 3 binary	2 continuous 2 binary	2 continuous 1 binary	continuous: 0.2 (0.1) binary: 0.2 (0.1)	$\tau=0.2$
B26							$\tau=0.5$
B27	10	15	5 continuous 3 binary	2 continuous 2 binary	2 continuous 1 binary	continuous: 0.2 (0.5) binary: 0.2 (0.4)	$\tau=0.2$
B28							$\tau=0.5$

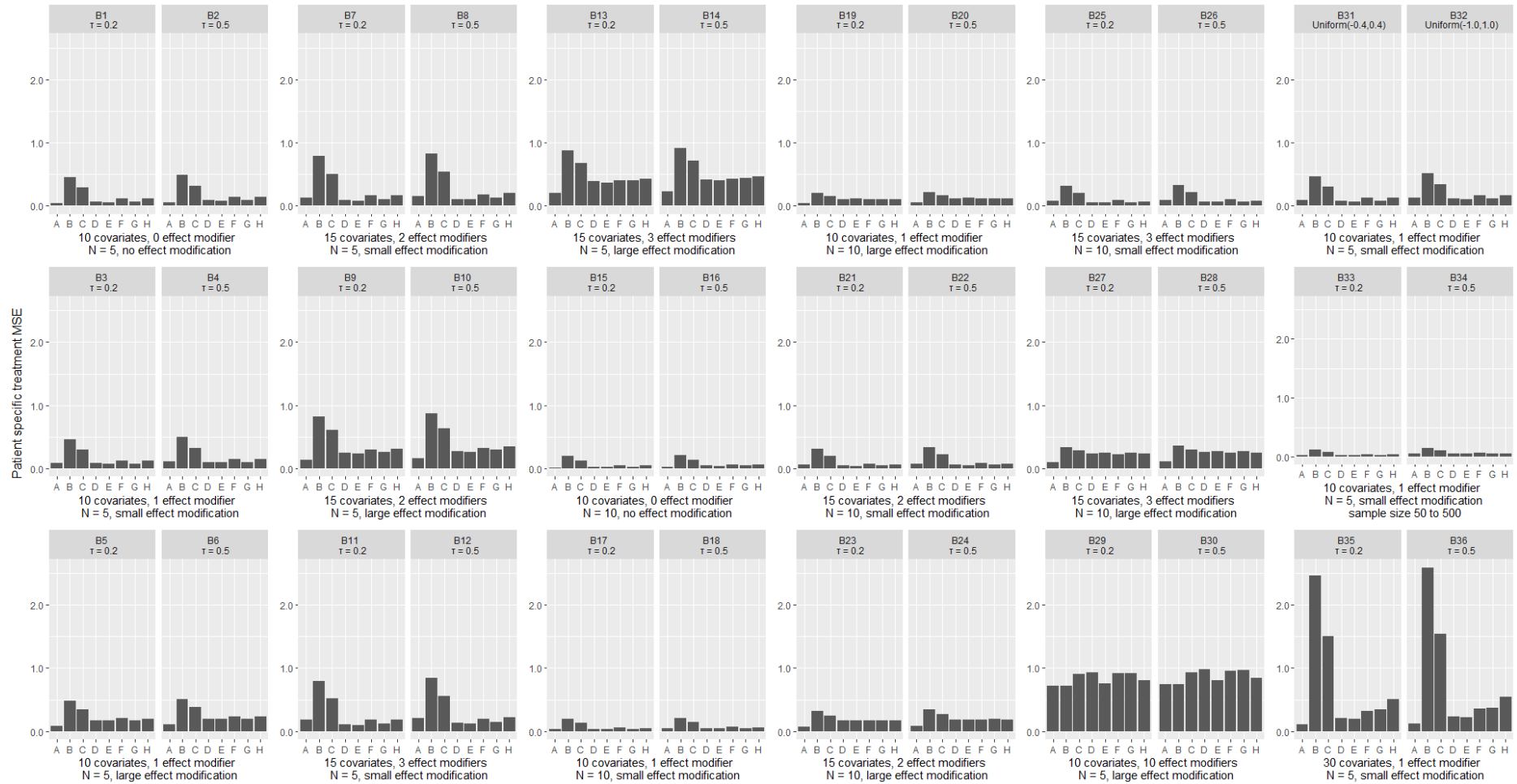
Scenarios	# of studies	# of covariates	# of nuisance covariates	# of prognostic factors	# of effect modifiers	True values for effect modifiers: main effect (effect modification)	Heterogeneity of treatment effect
B29	5	10	None	None	5 continuous 5 binary	continuous: 0.2 (0.5) binary: 0.2 (0.4)	$\tau=0.2$
B30							$\tau=0.5$
B31	5	10	3 continuous 2 binary	2 continuous 2 binary	1 continuous	continuous: 0.2 (0.1)	Uniform(-0.4,0.4)
B32							Uniform(-1.0,1.0)
B33	5 (sample size from 50 to 500)	10	3 continuous 2 binary	2 continuous 2 binary	1 continuous	continuous: 0.2 (0.1)	$\tau=0.2$
B34							$\tau=0.5$
B35	5	30	8 continuous 7 binary	7 continuous 7 binary	1 continuous	continuous: 0.2 (0.1)	$\tau=0.2$
B36							$\tau=0.5$

### 3 Results from simulations

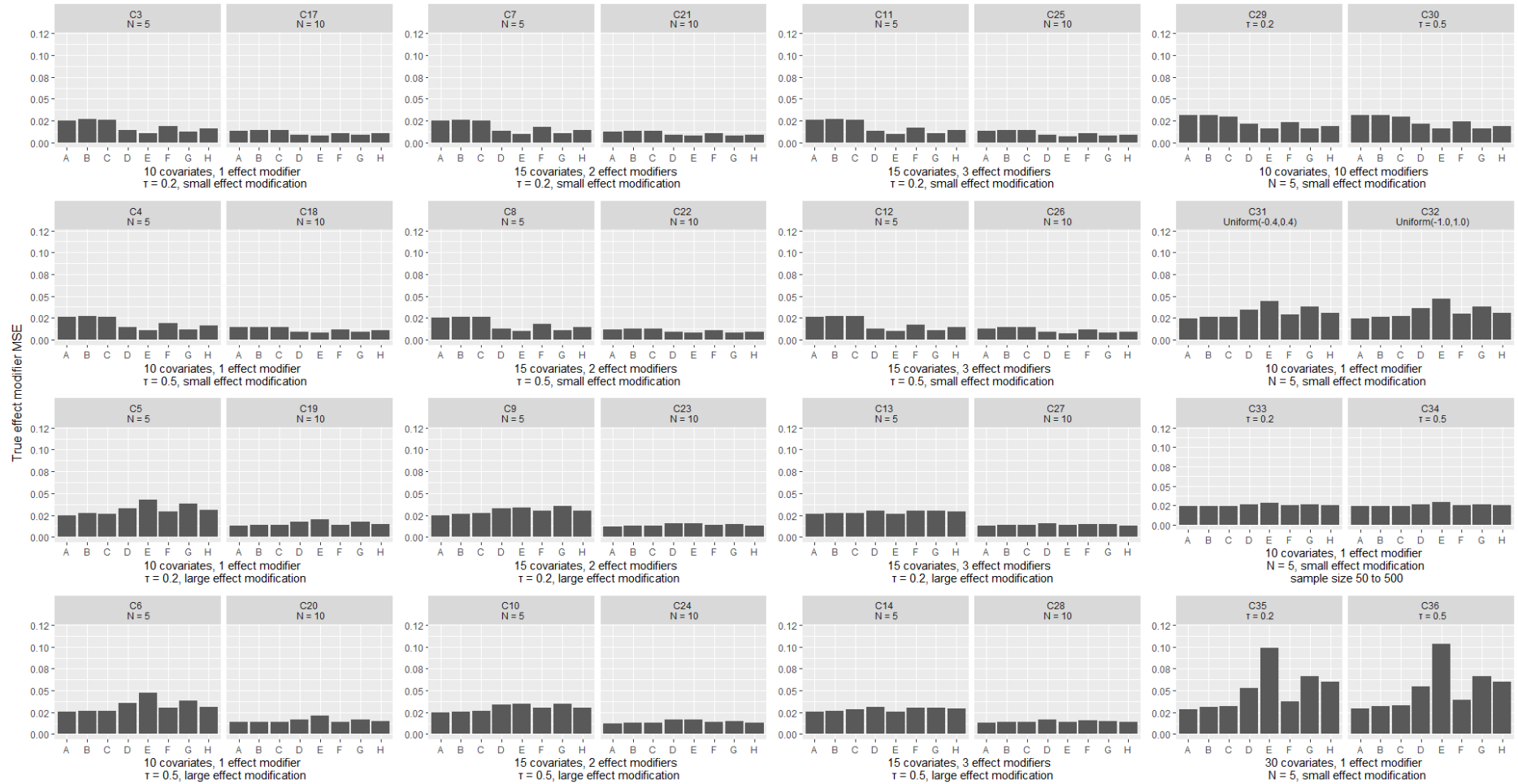
**Figure 1:** Results from simulations of a continuous outcome, comparing different methods in terms of the mean squared error (MSE) for the patient-specific treatment effect grouped by heterogeneity of the treatment effect. Scenarios C1-C36 are shown in pairs, differing only in the assumed heterogeneity ( $\tau$ ). Within each pair, the two scenarios explore the same type of outcome, have the same number of covariates and effect modifiers, equal heterogeneity and the number of studies. Scenarios are described in detail in Section 2 of the Appendix. A: GLMM-oracle; B: GLMM-full; C: STEP; D: LASSO; E: ridge; F: adaptive LASSO; G: Bayesian LASSO; H: SSVS.



**Figure 2:** Results from simulations of a binary outcome, comparing different methods in terms of the mean squared error (MSE) for the patient-specific treatment effect grouped by heterogeneity of the treatment effect. Scenarios B1-B36 are shown in pairs, differing only in the assumed heterogeneity ( $\tau$ ). Within each pair, the two scenarios explore the same type of outcome, have the same number of covariates and effect modifiers, equal heterogeneity and the number of studies. Scenarios are described in detail in Section 2 of the Appendix. A: GLMM-oracle; B: GLMM-full; C: STEP; D: LASSO; E: ridge; F: adaptive LASSO; G: Bayesian LASSO; H: SSVS.

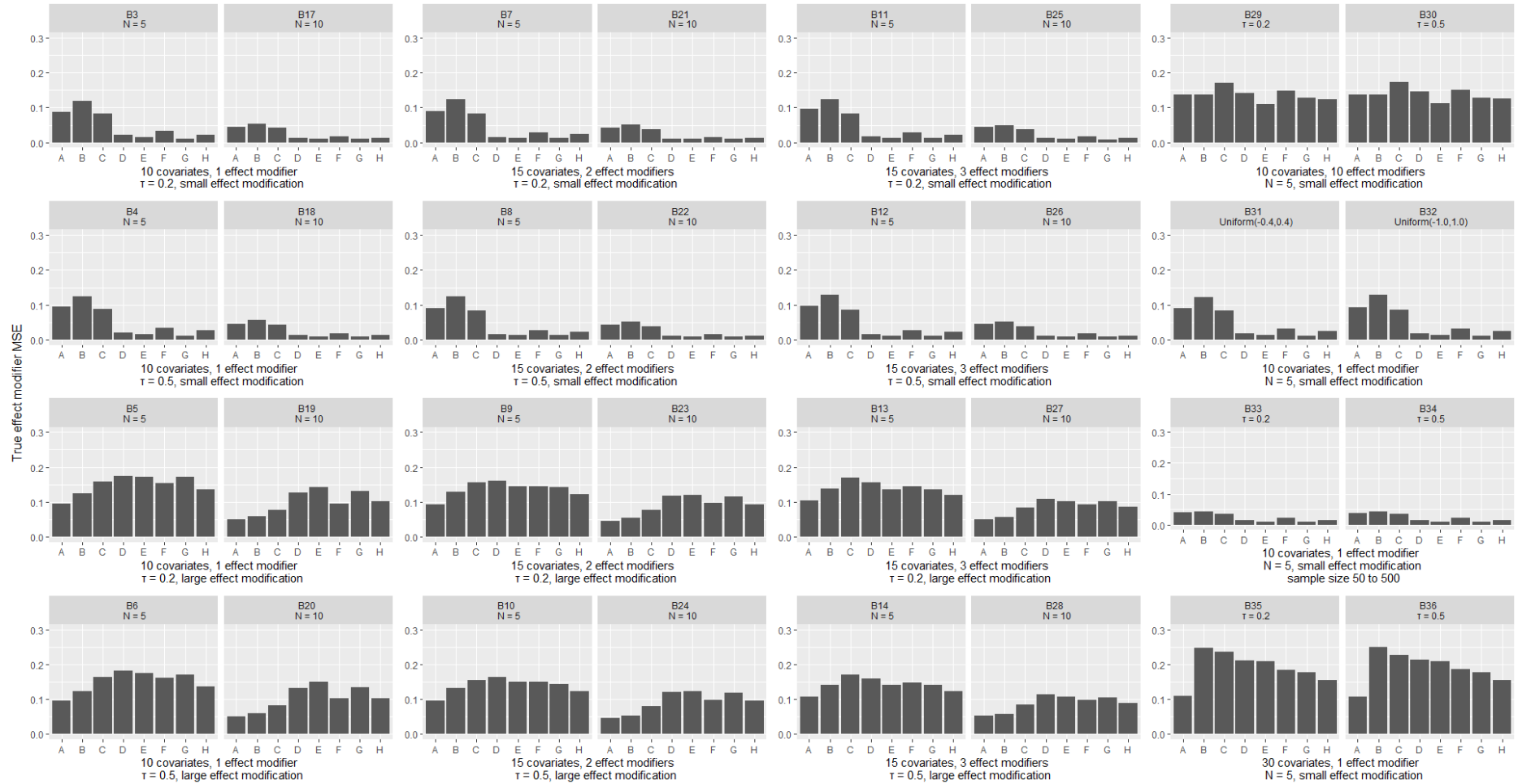


**Figure 3:** Simulation results of a continuous outcome comparing different methods in terms of true effect modifiers MSE. A: GLMM-oracle; B: GLMM-full; C: STEP; D: LASSO; E: ridge; F: adaptive LASSO; G: Bayesian LASSO; H: SSVS.

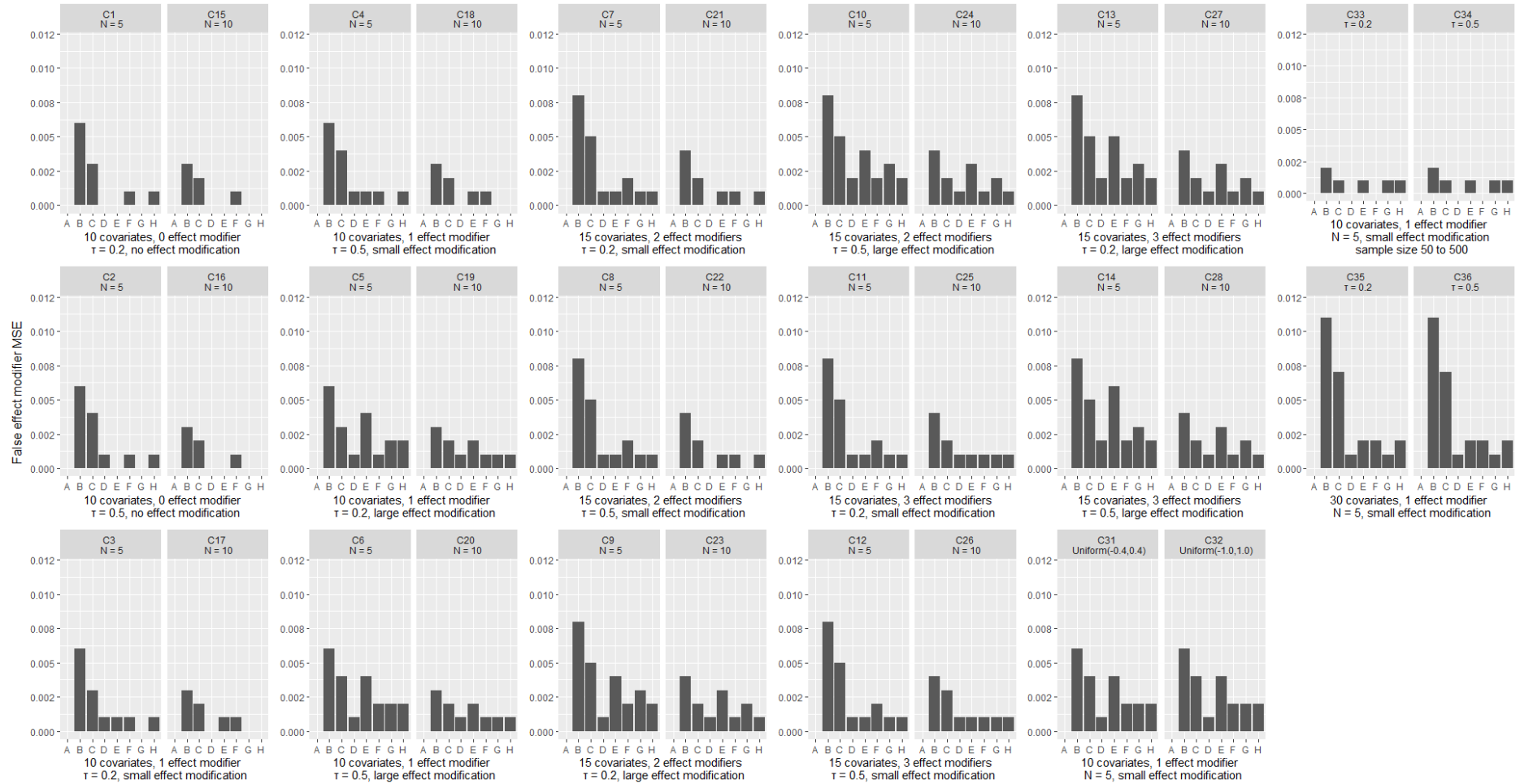




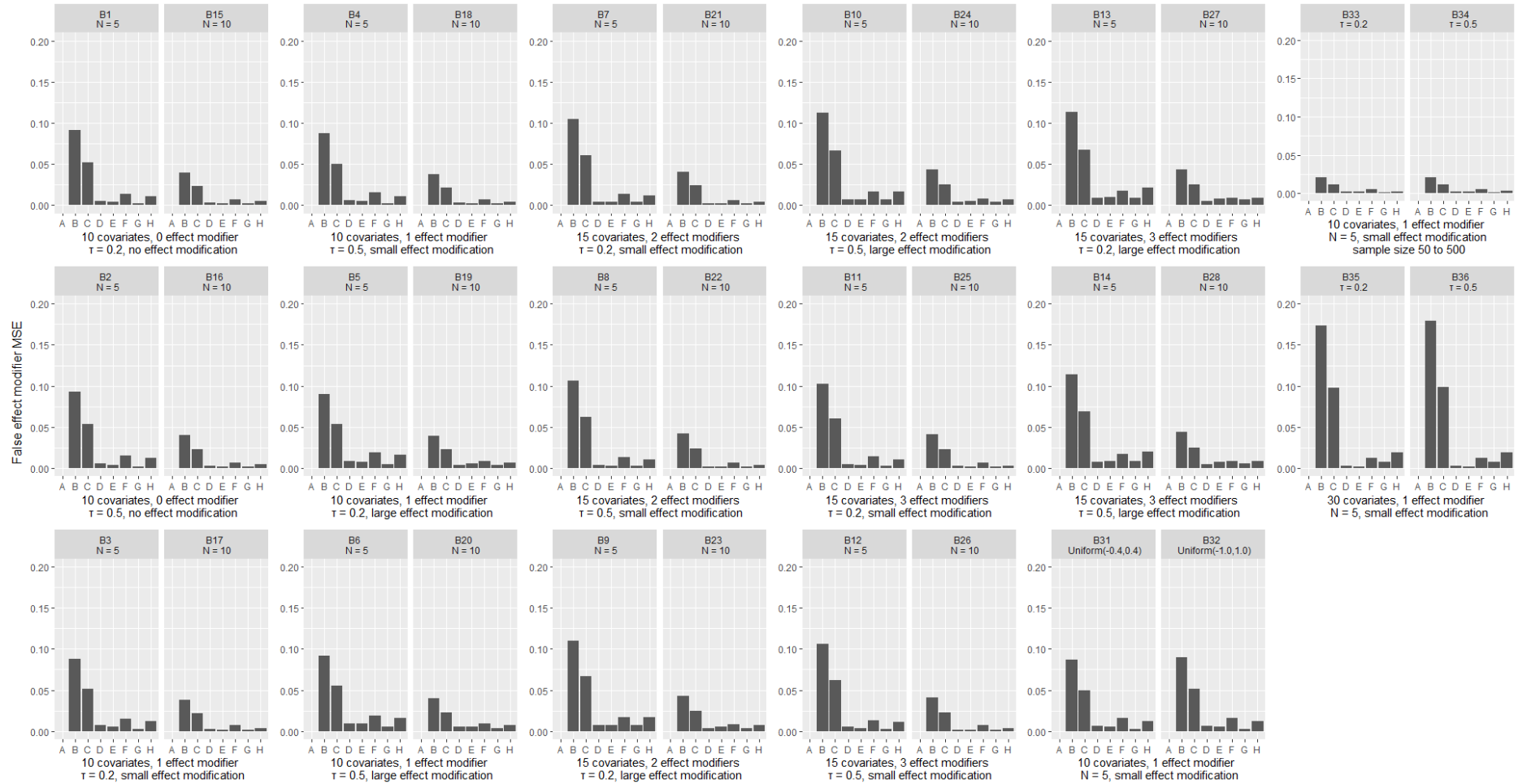
**Figure 4:** Simulation results of a binary outcome comparing different methods in terms of true effect modifiers MSE. A: GLMM-oracle; B: GLMM-full; C: STEP; D: LASSO; E: ridge; F: adaptive LASSO; G: Bayesian LASSO; H: SSVS.



**Figure 5:** Simulation results of a continuous outcome comparing different methods in terms of false effect modifier MSE. A: GLMM-oracle; B: GLMM-full; C: STEP; D: LASSO; E: ridge; F: adaptive LASSO; G: Bayesian LASSO; H: SSVS.



**Figure 6:** Simulation results of a binary outcome comparing different methods in terms of false effect modifier MSE. A: GLMM-oracle; B: GLMM-full; C: STEP; D: LASSO; E: ridge; F: adaptive LASSO; G: Bayesian LASSO; H: SSVS.



**Table 4:** Full results from the simulations. Model abbreviations as per Section 2 in the main paper. MSE: mean squared error. SE: standard error. To facilitate readers, the lowest patient-specific treatment-effect MSE is shown in bold (excluding MSE of GLMM-oracle)

Scenario	Model	False effect modifier MSE	True effect modifier MSE	Treatment MSE	Patient specific treatment effect MSE	Continuous effect modifier SE	Binary effect modifier SE	Treatment effect SE
C1	GLMM-oracle	0		0.012	0.006			0.109
	GLMM-full	0.006		0.013	0.031			0.109
	STEP	0.003		0.013	0.022			0.068
	LASSO	0		0.013	0.009			
	ridge	0		0.013	0.008			
	adaptive LASSO	0.001		0.013	0.012			
	Bayesian LASSO	0		0.012	<b>0.007</b>			0.167
	SSVS	0.001		0.013	0.009			0.167
C2	GLMM-oracle	0		0.055	0.028			0.223
	GLMM-full	0.006		0.055	0.052			0.223
	STEP	0.004		0.057	0.045			0.071
	LASSO	0.001		0.057	0.031			
	ridge	0		0.057	0.030			
	adaptive LASSO	0.001		0.057	0.035			
	Bayesian LASSO	0		0.055	<b>0.029</b>			0.305
	SSVS	0.001		0.055	0.030			0.305
C3	GLMM-oracle	0	0.025	0.013	0.019	0.070		0.111
	GLMM-full	0.006	0.027	0.013	0.042	0.078		0.111
	STEP	0.003	0.026	0.013	0.033	0.072		0.070
	LASSO	0.001	0.014	0.013	0.017			
	ridge	0.001	0.011	0.013	0.016			
	adaptive LASSO	0.001	0.019	0.013	0.022			
	Bayesian LASSO	0	0.012	0.013	<b>0.015</b>	0.050		0.169
	SSVS	0.001	0.016	0.013	0.018	0.059		0.169
C4	GLMM-oracle	0	0.026	0.055	0.041	0.070		0.224
	GLMM-full	0.006	0.027	0.056	0.064	0.078		0.225
	STEP	0.004	0.026	0.058	0.057	0.075		0.074
	LASSO	0.001	0.014	0.057	0.039			
	ridge	0.001	0.011	0.057	0.039			
	adaptive LASSO	0.001	0.019	0.057	0.044			

	Bayesian LASSO	0	0.012	0.055	<b>0.036</b>	0.051		0.307
	SSVS	0.001	0.016	0.055	0.040	0.059		0.306
C5	GLMM-oracle	0	0.025	0.013	0.019	0.070		0.111
	GLMM-full	0.006	0.027	0.013	0.042	0.078		0.111
	STEP	0.003	0.026	0.013	0.034	0.072		0.070
	LASSO	0.001	0.033	0.013	0.028			
	ridge	0.004	0.043	0.013	0.041			
	adaptive LASSO	0.001	0.029	0.013	<b>0.027</b>			
	Bayesian LASSO	0.002	0.038	0.013	0.033	0.079		0.169
	SSVS	0.002	0.031	0.013	0.028	0.077		0.170
C6	GLMM-oracle	0	0.026	0.055	0.041	0.070		0.224
	GLMM-full	0.006	0.027	0.056	0.064	0.078		0.225
	STEP	0.004	0.027	0.058	0.057	0.075		0.074
	LASSO	0.001	0.036	0.057	0.051			
	ridge	0.004	0.047	0.057	0.065			
	adaptive LASSO	0.002	0.03	0.057	<b>0.050</b>			
	Bayesian LASSO	0.002	0.038	0.055	0.054	0.079		0.306
	SSVS	0.002	0.031	0.055	<b>0.050</b>	0.077		0.306
C7	GLMM-oracle	0	0.025	0.015	0.032	0.078	0.078	0.116
	GLMM-full	0.008	0.026	0.015	0.075	0.088	0.080	0.118
	STEP	0.005	0.025	0.015	0.059	0.080	0.078	0.078
	LASSO	0.001	0.013	0.015	0.026			
	ridge	0.001	0.010	0.015	0.024			
	adaptive LASSO	0.002	0.018	0.015	0.035			
	Bayesian LASSO	0.001	0.011	0.015	<b>0.023</b>	0.054	0.053	0.176
	SSVS	0.001	0.014	0.015	0.027	0.061	0.060	0.176
C8	GLMM-oracle	0	0.025	0.057	0.054	0.078	0.078	0.227
	GLMM-full	0.008	0.026	0.057	0.097	0.088	0.080	0.228
	STEP	0.005	0.026	0.059	0.084	0.083	0.082	0.081
	LASSO	0.001	0.013	0.059	0.049			
	ridge	0.001	0.010	0.059	0.047			
	adaptive LASSO	0.002	0.018	0.059	0.058			
	Bayesian LASSO	0.001	0.011	0.057	<b>0.044</b>	0.054	0.053	0.307
	SSVS	0.001	0.014	0.057	0.049	0.061	0.060	0.308
C9	GLMM-oracle	0	0.025	0.015	0.032	0.078	0.078	0.116
	GLMM-full	0.008	0.026	0.015	0.075	0.088	0.080	0.118

	STEP	0.005	0.027	0.015	0.060	0.081	0.079	0.078
	LASSO	0.001	0.033	0.015	0.048			
	ridge	0.004	0.034	0.015	0.064			
	adaptive LASSO	0.002	0.03	0.01	<b>0.047</b>			
	Bayesian LASSO	0.003	0.035	0.015	0.056	0.087	0.080	0.177
	SSVS	0.002	0.030	0.015	0.049	0.086	0.080	0.177
C10	GLMM-oracle	0	0.025	0.057	0.054	0.078	0.078	0.227
	GLMM-full	0.008	0.026	0.057	0.097	0.088	0.080	0.228
	STEP	0.005	0.027	0.059	0.085	0.084	0.082	0.081
	LASSO	0.002	0.034	0.059	0.072			
	ridge	0.004	0.035	0.059	0.089			
	adaptive LASSO	0.002	0.03	0.059	0.071			
	Bayesian LASSO	0.003	0.035	0.057	0.077	0.087	0.081	0.308
	SSVS	0.002	0.030	0.057	<b>0.070</b>	0.086	0.081	0.308
C11	GLMM-oracle	0	0.026	0.016	0.046	0.085	0.080	0.118
	GLMM-full	0.008	0.027	0.016	0.087	0.091	0.083	0.118
	STEP	0.005	0.026	0.016	0.072	0.084	0.081	0.081
	LASSO	0.001	0.013	0.016	0.034			
	ridge	0.001	0.010	0.016	<b>0.032</b>			
	adaptive LASSO	0.002	0.017	0.016	0.044			
	Bayesian LASSO	0.001	0.011	0.016	<b>0.032</b>	0.059	0.057	0.178
	SSVS	0.001	0.014	0.016	0.037	0.066	0.063	0.177
C12	GLMM-oracle	0	0.026	0.059	0.068	0.085	0.081	0.227
	GLMM-full	0.008	0.027	0.060	0.109	0.091	0.083	0.227
	STEP	0.005	0.027	0.062	0.097	0.087	0.083	0.084
	LASSO	0.001	0.013	0.061	0.057			
	ridge	0.001	0.010	0.061	0.055			
	adaptive LASSO	0.002	0.017	0.061	0.068			
	Bayesian LASSO	0.001	0.011	0.060	<b>0.054</b>	0.059	0.057	0.307
	SSVS	0.001	0.014	0.060	0.059	0.066	0.063	0.307
C13	GLMM-oracle	0	0.026	0.016	0.046	0.085	0.080	0.118
	GLMM-full	0.008	0.027	0.016	0.087	0.091	0.083	0.118
	STEP	0.005	0.027	0.016	0.073	0.086	0.081	0.081
	LASSO	0.002	0.030	0.016	0.063			
	ridge	0.005	0.026	0.016	0.075			
	adaptive LASSO	0.002	0.03	0.016	<b>0.062</b>			

	Bayesian LASSO	0.003	0.030	0.016	0.071	0.090	0.083	0.178
	SSVS	0.002	0.029	0.016	0.063	0.089	0.083	0.178
C14	GLMM-oracle	0	0.026	0.059	0.068	0.085	0.081	0.227
	GLMM-full	0.008	0.027	0.060	0.109	0.091	0.083	0.227
	STEP	0.005	0.028	0.062	0.099	0.089	0.084	0.084
	LASSO	0.002	0.031	0.061	0.088			
	ridge	0.006	0.026	0.061	0.101			
	adaptive LASSO	0.002	0.03	0.061	0.087			
	Bayesian LASSO	0.003	0.030	0.060	0.093	0.090	0.083	0.307
	SSVS	0.002	0.029	0.060	<b>0.085</b>	0.089	0.084	0.307
C15	GLMM-oracle	0		0.006	0.003			0.078
	GLMM-full	0.003		0.006	0.016			0.078
	STEP	0.002		0.006	0.011			0.049
	LASSO	0		0.007	0.005			
	ridge	0		0.007	<b>0.004</b>			
	adaptive LASSO	0.001		0.007	0.006			
	Bayesian LASSO	0		0.006	<b>0.004</b>			0.092
SSVS	0		0.006	0.005			0.092	
C16	GLMM-oracle	0		0.028	0.014			0.161
	GLMM-full	0.003		0.028	0.027			0.161
	STEP	0.002		0.029	0.023			0.051
	LASSO	0		0.030	0.016			
	ridge	0		0.030	0.016			
	adaptive LASSO	0.001		0.03	0.018			
	Bayesian LASSO	0		0.028	<b>0.015</b>			0.187
SSVS	0		0.028	0.016			0.187	
C17	GLMM-oracle	0	0.013	0.007	0.010	0.051		0.079
	GLMM-full	0.003	0.014	0.007	0.022	0.056		0.080
	STEP	0.002	0.014	0.007	0.018	0.052		0.051
	LASSO	0	0.009	0.007	<b>0.010</b>			
	ridge	0.001	0.008	0.007	<b>0.010</b>			
	adaptive LASSO	0.001	0.011	0.007	0.012			
	Bayesian LASSO	0	0.009	0.007	<b>0.010</b>	0.041		0.093
SSVS	0	0.011	0.007	0.011	0.045		0.093	
C18	GLMM-oracle	0	0.014	0.028	0.021	0.051		0.161
	GLMM-full	0.003	0.014	0.028	0.033	0.056		0.161

	STEP	0.002	0.014	0.029	0.030	0.054		0.053
	LASSO	0	0.009	0.029	<b>0.021</b>			
	ridge	0.001	0.008	0.029	<b>0.021</b>			
	adaptive LASSO	0.001	0.012	0.029	0.024			
	Bayesian LASSO	0	0.009	0.028	<b>0.021</b>	0.041		0.188
	SSVS	0	0.011	0.028	0.022	0.045		0.188
C19	GLMM-oracle	0	0.013	0.007	0.010	0.051		0.079
	GLMM-full	0.003	0.014	0.007	0.022	0.056		0.080
	STEP	0.002	0.014	0.007	0.018	0.052		0.051
	LASSO	0.001	0.017	0.007	<b>0.014</b>			
	ridge	0.002	0.020	0.007	0.022			
	adaptive LASSO	0.001	0.014	0.007	<b>0.014</b>			
	Bayesian LASSO	0.001	0.017	0.007	0.017	0.056		0.093
SSVS	0.001	0.015	0.007	0.015	0.055		0.093	
C20	GLMM-oracle	0	0.014	0.028	0.021	0.051		0.161
	GLMM-full	0.003	0.014	0.028	0.033	0.056		0.161
	STEP	0.002	0.014	0.029	0.030	0.055		0.053
	LASSO	0.001	0.017	0.029	0.026			
	ridge	0.002	0.021	0.029	0.034			
	adaptive LASSO	0.001	0.014	0.029	<b>0.025</b>			
	Bayesian LASSO	0.001	0.017	0.028	0.028	0.056		0.188
SSVS	0.001	0.015	0.028	0.026	0.055		0.188	
C21	GLMM-oracle	0	0.012	0.008	0.017	0.057	0.056	0.083
	GLMM-full	0.004	0.013	0.008	0.038	0.063	0.057	0.083
	STEP	0.002	0.013	0.008	0.030	0.058	0.057	0.057
	LASSO	0	0.009	0.007	<b>0.015</b>			
	ridge	0.001	0.008	0.007	<b>0.015</b>			
	adaptive LASSO	0.001	0.011	0.007	0.020			
	Bayesian LASSO	0	0.008	0.008	<b>0.015</b>	0.044	0.043	0.097
SSVS	0.001	0.009	0.008	0.016	0.047	0.046	0.097	
C22	GLMM-oracle	0	0.012	0.031	0.028	0.057	0.057	0.163
	GLMM-full	0.004	0.013	0.031	0.050	0.063	0.057	0.163
	STEP	0.002	0.013	0.032	0.043	0.060	0.059	0.059
	LASSO	0	0.009	0.030	<b>0.027</b>			
	ridge	0.001	0.008	0.030	<b>0.027</b>			
	adaptive LASSO	0.001	0.011	0.03	0.031			



	Bayesian LASSO	0	0.008	0.031	<b>0.027</b>	0.044	0.043	0.189
	SSVS	0.001	0.009	0.031	0.028	0.047	0.046	0.189
C23	GLMM-oracle	0	0.012	0.008	0.017	0.057	0.056	0.083
	GLMM-full	0.004	0.013	0.008	0.038	0.063	0.057	0.083
	STEP	0.002	0.013	0.008	0.030	0.058	0.057	0.057
	LASSO	0.001	0.016	0.007	0.024			
	ridge	0.003	0.016	0.007	0.035			
	adaptive LASSO	0.001	0.014	0.007	<b>0.023</b>			
	Bayesian LASSO	0.002	0.015	0.008	0.029	0.062	0.058	0.097
	SSVS	0.001	0.013	0.008	0.024	0.061	0.057	0.097
C24	GLMM-oracle	0	0.012	0.031	0.028	0.057	0.057	0.163
	GLMM-full	0.004	0.013	0.031	0.050	0.063	0.057	0.163
	STEP	0.002	0.013	0.032	0.043	0.060	0.059	0.059
	LASSO	0.001	0.017	0.030	0.037			
	ridge	0.003	0.017	0.030	0.048			
	adaptive LASSO	0.001	0.014	0.03	<b>0.035</b>			
	Bayesian LASSO	0.002	0.015	0.031	0.041	0.062	0.058	0.189
	SSVS	0.001	0.013	0.031	<b>0.035</b>	0.061	0.057	0.189
C25	GLMM-oracle	0	0.013	0.008	0.023	0.062	0.059	0.084
	GLMM-full	0.004	0.014	0.008	0.045	0.065	0.059	0.085
	STEP	0.002	0.014	0.008	0.037	0.061	0.059	0.059
	LASSO	0.001	0.009	0.008	0.021			
	ridge	0.001	0.007	0.008	0.021			
	adaptive LASSO	0.001	0.011	0.008	0.026			
	Bayesian LASSO	0.001	0.008	0.008	<b>0.020</b>	0.048	0.047	0.099
	SSVS	0.001	0.009	0.008	0.022	0.051	0.049	0.099
C26	GLMM-oracle	0	0.013	0.030	0.034	0.062	0.059	0.164
	GLMM-full	0.004	0.014	0.030	0.056	0.065	0.060	0.164
	STEP	0.003	0.014	0.031	0.050	0.063	0.061	0.061
	LASSO	0.001	0.009	0.030	0.033			
	ridge	0.001	0.007	0.030	0.032			
	adaptive LASSO	0.001	0.012	0.030	0.038			
	Bayesian LASSO	0.001	0.008	0.030	<b>0.031</b>	0.048	0.047	0.190
	SSVS	0.001	0.009	0.030	0.033	0.051	0.049	0.190
C27	GLMM-oracle	0	0.013	0.008	0.023	0.062	0.059	0.084
	GLMM-full	0.004	0.014	0.008	0.045	0.065	0.059	0.085

	STEP	0.002	0.014	0.008	0.036	0.063	0.059	0.059
	LASSO	0.001	0.016	0.008	0.033			
	ridge	0.003	0.014	0.008	0.042			
	adaptive LASSO	0.001	0.015	0.008	0.031			
	Bayesian LASSO	0.002	0.015	0.008	0.037	0.065	0.060	0.099
	SSVS	0.001	0.013	0.008	<b>0.030</b>	0.063	0.060	0.099
C28	GLMM-oracle	0	0.013	0.030	0.034	0.062	0.059	0.164
	GLMM-full	0.004	0.014	0.030	0.056	0.065	0.060	0.164
	STEP	0.002	0.014	0.031	0.049	0.065	0.061	0.061
	LASSO	0.001	0.017	0.030	0.045			
	ridge	0.003	0.014	0.030	0.055			
	adaptive LASSO	0.001	0.016	0.03	0.044			
	Bayesian LASSO	0.002	0.015	0.030	0.048	0.065	0.060	0.190
	SSVS	0.001	0.014	0.030	<b>0.041</b>	0.063	0.060	0.190
C29	GLMM-oracle		0.031	0.019	0.158	0.111	0.103	0.135
	GLMM-full		0.031	0.019	0.158	0.111	0.103	0.135
	STEP		0.030	0.019	0.153	0.105	0.102	0.101
	LASSO		0.021	0.018	0.113			
	ridge		0.016	0.018	0.093			
	adaptive LASSO		0.023	0.018	0.125			
	Bayesian LASSO		0.016	0.019	<b>0.091</b>	0.087	0.083	0.199
	SSVS		0.019	0.019	0.103	0.090	0.085	0.199
C30	GLMM-oracle		0.031	0.061	0.179	0.111	0.103	0.235
	GLMM-full		0.031	0.061	0.179	0.111	0.103	0.235
	STEP		0.030	0.062	0.176	0.107	0.104	0.104
	LASSO		0.021	0.062	0.135			
	ridge		0.016	0.062	0.114			
	adaptive LASSO		0.024	0.062	0.147			
	Bayesian LASSO		0.016	0.061	<b>0.111</b>	0.087	0.083	0.314
	SSVS		0.019	0.061	0.124	0.090	0.085	0.314
C31	GLMM-oracle	0.000	0.024	0.017	0.021	0.070		0.121
	GLMM-full	0.006	0.026	0.017	0.045	0.079		0.122
	STEP	0.004	0.026	0.017	0.036	0.073		0.071
	LASSO	0.001	0.034	0.017	0.030			
	ridge	0.004	0.044	0.017	0.044			
	adaptive LASSO	0.002	0.029	0.017	<b>0.029</b>			

	Bayesian LASSO	0.002	0.038	0.017	0.035	0.080		0.184
	SSVS	0.002	0.031	0.017	0.030	0.078		0.184
C32	GLMM-oracle	0.000	0.024	0.076	0.050	0.071		0.258
	GLMM-full	0.006	0.026	0.076	0.074	0.079		0.258
	STEP	0.004	0.027	0.078	0.068	0.077		0.075
	LASSO	0.001	0.036	0.078	0.062			
	ridge	0.004	0.047	0.078	0.076			
	adaptive LASSO	0.002	0.030	0.078	<b>0.060</b>			
	Bayesian LASSO	0.002	0.038	0.076	0.064	0.080		0.343
	SSVS	0.002	0.031	0.076	<b>0.060</b>	0.078		0.343
C33	GLMM-oracle	0.000	0.024	0.010	0.017	0.036		0.095
	GLMM-full	0.002	0.024	0.010	0.023	0.040		0.095
	STEP	0.001	0.024	0.011	0.021	0.037		0.037
	LASSO	0.000	0.026	0.011	0.020			
	ridge	0.001	0.028	0.011	0.025			
	adaptive LASSO	0.000	0.025	0.011	<b>0.019</b>			
	Bayesian LASSO	0.001	0.026	0.010	0.021	0.040		0.153
	SSVS	0.001	0.025	0.010	0.020	0.039		0.153
C34	GLMM-oracle	0.000	0.024	0.053	0.039	0.036		0.219
	GLMM-full	0.002	0.024	0.053	0.045	0.040		0.219
	STEP	0.001	0.024	0.063	0.048	0.039		0.038
	LASSO	0.000	0.026	0.063	0.046			
	ridge	0.001	0.029	0.063	0.051			
	adaptive LASSO	0.000	0.025	0.063	0.046			
	Bayesian LASSO	0.001	0.026	0.053	0.043	0.040		0.303
	SSVS	0.001	0.025	0.053	<b>0.041</b>	0.039		0.303
C35	GLMM-oracle	0.000	0.028	0.018	0.023	0.093		0.128
	GLMM-full	0.011	0.031	0.019	0.159	0.109		0.131
	STEP	0.007	0.032	0.019	0.113	0.096		0.093
	LASSO	0.001	0.053	0.019	<b>0.046</b>			
	ridge	0.002	0.099	0.019	0.084			
	adaptive LASSO	0.002	0.037	0.019	0.054			
	Bayesian LASSO	0.001	0.066	0.019	0.060	0.103		0.192
	SSVS	0.002	0.060	0.019	0.062	0.104		0.192
C36	GLMM-oracle	0.000	0.029	0.062	0.046	0.093		0.231

	GLMM-full	0.011	0.032	0.063	0.182	0.109		0.231
	STEP	0.007	0.033	0.064	0.141	0.109		0.231
	LASSO	0.001	0.055	0.064	<b>0.070</b>			
	ridge	0.002	0.103	0.064	0.108			
	adaptive LASSO	0.002	0.039	0.064	0.078			
	Bayesian LASSO	0.001	0.066	0.063	0.083	0.103		0.310
	SSVS	0.002	0.060	0.063	0.085	0.104		0.309
B1	GLMM-oracle	0		0.076	0.038			0.259
	GLMM-full	0.091		0.110	0.459			0.286
	STEP	0.052		0.096	0.289			0.268
	LASSO	0.005		0.080	0.065			
	ridge	0.004		0.080	<b>0.059</b>			
	adaptive LASSO	0.014		0.082	0.111			
	Bayesian LASSO	0.002		0.106	0.065			0.395
SSVS	0.011		0.116	0.114			0.401	
B2	GLMM-oracle	0		0.116	0.058			0.281
	GLMM-full	0.093		0.159	0.491			0.308
	STEP	0.054		0.140	0.317			0.270
	LASSO	0.006		0.128	0.093			
	ridge	0.004		0.127	<b>0.084</b>			
	adaptive LASSO	0.015		0.131	0.139			
	Bayesian LASSO	0.002		0.155	0.090			0.439
SSVS	0.012		0.165	0.141			0.444	
B3	GLMM-oracle	0	0.087	0.077	0.082	0.260		0.261
	GLMM-full	0.088	0.118	0.106	0.461	0.297		0.285
	STEP	0.051	0.082	0.094	0.297	0.277		0.267
	LASSO	0.007	0.021	0.087	0.084			
	ridge	0.005	0.016	0.086	0.075			
	adaptive LASSO	0.015	0.033	0.091	0.128			
	Bayesian LASSO	0.003	0.011	0.107	<b>0.072</b>	0.109		0.394
SSVS	0.012	0.021	0.115	0.122	0.165		0.398	
B4	GLMM-oracle	0	0.094	0.125	0.109	0.262		0.282
	GLMM-full	0.088	0.125	0.162	0.493	0.300		0.306
	STEP	0.050	0.088	0.147	0.323	0.279		0.269
	LASSO	0.006	0.020	0.131	0.102			

	ridge	0.005	0.015	0.130	<b>0.095</b>			
	adaptive LASSO	0.016	0.033	0.136	0.155			
	Bayesian LASSO	0.002	0.011	0.165	0.100	0.109		0.440
	SSVS	0.011	0.026	0.173	0.150	0.165		0.446
B5	GLMM-oracle	0	0.096	0.079	0.087	0.273		0.267
	GLMM-full	0.090	0.124	0.107	0.476	0.310		0.291
	STEP	0.054	0.159	0.098	0.345	0.287		0.272
	LASSO	0.009	0.175	0.096	0.171			
	ridge	0.008	0.171	0.096	0.167			
	adaptive LASSO	0.019	0.154	0.099	0.203			
	Bayesian LASSO	0.005	0.171	0.123	<b>0.166</b>	0.164		0.401
	SSVS	0.016	0.135	0.130	0.200	0.241		0.407
B6	GLMM-oracle	0	0.095	0.130	0.112	0.274		0.288
	GLMM-full	0.092	0.123	0.167	0.512	0.312		0.312
	STEP	0.055	0.163	0.154	0.379	0.287		0.274
	LASSO	0.009	0.181	0.144	0.198			
	ridge	0.009	0.174	0.143	<b>0.194</b>			
	adaptive LASSO	0.019	0.162	0.149	0.234			
	Bayesian LASSO	0.005	0.171	0.182	0.195	0.163		0.441
	SSVS	0.016	0.137	0.189	0.230	0.240		0.448
B7	GLMM-oracle	0	0.089	0.080	0.129	0.265	0.260	0.268
	GLMM-full	0.105	0.123	0.143	0.786	0.312	0.282	0.306
	STEP	0.061	0.083	0.119	0.507	0.286	0.277	0.280
	LASSO	0.004	0.016	0.087	0.088			
	ridge	0.004	0.013	0.086	<b>0.082</b>			
	adaptive LASSO	0.014	0.029	0.093	0.164			
	Bayesian LASSO	0.004	0.013	0.128	0.101	0.111	0.109	0.411
	SSVS	0.012	0.024	0.140	0.170	0.158	0.152	0.417
B8	GLMM-oracle	0	0.09	0.128	0.154	0.267	0.262	0.291
	GLMM-full	0.106	0.125	0.208	0.828	0.314	0.284	0.331
	STEP	0.062	0.083	0.176	0.538	0.289	0.277	0.281
	LASSO	0.004	0.016	0.136	0.109			
	ridge	0.003	0.013	0.134	<b>0.104</b>			
	adaptive LASSO	0.013	0.028	0.142	0.184			
	Bayesian LASSO	0.003	0.013	0.187	0.129	0.111	0.109	0.455
	SSVS	0.011	0.023	0.204	0.199	0.158	0.153	0.46

B9	GLMM-oracle	0	0.094	0.086	0.137	0.279	0.267	0.275
	GLMM-full	0.110	0.130	0.149	0.825	0.328	0.291	0.314
	STEP	0.067	0.157	0.134	0.611	0.298	0.281	0.288
	LASSO	0.007	0.160	0.103	0.255			
	ridge	0.007	0.146	0.102	<b>0.239</b>			
	adaptive LASSO	0.017	0.146	0.105	0.302			
	Bayesian LASSO	0.007	0.143	0.165	0.266	0.169	0.154	0.419
	SSVS	0.017	0.122	0.175	0.316	0.238	0.212	0.427
B10	GLMM-oracle	0	0.096	0.135	0.164	0.281	0.269	0.298
	GLMM-full	0.112	0.132	0.215	0.867	0.330	0.293	0.339
	STEP	0.066	0.154	0.192	0.634	0.299	0.284	0.289
	LASSO	0.007	0.163	0.145	0.275			
	ridge	0.007	0.150	0.143	<b>0.262</b>			
	adaptive LASSO	0.017	0.15	0.148	0.329			
	Bayesian LASSO	0.007	0.144	0.226	0.298	0.168	0.153	0.460
	SSVS	0.017	0.122	0.239	0.349	0.237	0.212	0.469
B11	GLMM-oracle	0	0.097	0.093	0.185	0.277	0.261	0.270
	GLMM-full	0.103	0.124	0.173	0.798	0.311	0.282	0.305
	STEP	0.060	0.084	0.140	0.514	0.285	0.272	0.280
	LASSO	0.005	0.017	0.100	0.110			
	ridge	0.004	0.013	0.100	<b>0.096</b>			
	adaptive LASSO	0.014	0.029	0.104	0.183			
	Bayesian LASSO	0.003	0.012	0.153	0.115	0.112	0.109	0.408
	SSVS	0.011	0.022	0.168	0.184	0.159	0.151	0.415
B12	GLMM-oracle	0	0.098	0.135	0.207	0.279	0.263	0.287
	GLMM-full	0.106	0.128	0.233	0.846	0.314	0.284	0.324
	STEP	0.062	0.085	0.195	0.553	0.288	0.274	0.282
	LASSO	0.005	0.016	0.152	0.132			
	ridge	0.004	0.012	0.151	<b>0.121</b>			
	adaptive LASSO	0.013	0.027	0.157	0.199			
	Bayesian LASSO	0.003	0.012	0.206	0.143	0.112	0.110	0.447
	SSVS	0.011	0.022	0.227	0.217	0.160	0.154	0.452
B13	GLMM-oracle	0	0.105	0.094	0.198	0.295	0.274	0.282
	GLMM-full	0.113	0.139	0.173	0.871	0.333	0.298	0.317
	STEP	0.067	0.169	0.155	0.673	0.308	0.287	0.293
	LASSO	0.009	0.156	0.133	0.392			

	ridge	0.010	0.137	0.132	<b>0.370</b>			
	adaptive LASSO	0.018	0.146	0.132	0.406			
	Bayesian LASSO	0.009	0.137	0.231	0.408	0.194	0.168	0.424
	SSVS	0.021	0.121	0.234	0.434	0.257	0.222	0.431
B14	GLMM-oracle	0	0.107	0.137	0.224	0.297	0.275	0.301
	GLMM-full	0.114	0.142	0.228	0.912	0.335	0.299	0.338
	STEP	0.069	0.171	0.204	0.711	0.309	0.288	0.294
	LASSO	0.008	0.159	0.186	0.419			
	ridge	0.009	0.14	0.184	<b>0.398</b>			
	adaptive LASSO	0.017	0.147	0.187	0.431			
	Bayesian LASSO	0.009	0.141	0.278	0.438	0.190	0.167	0.457
	SSVS	0.020	0.124	0.286	0.464	0.255	0.221	0.466
B15	GLMM-oracle	0		0.037	0.018			0.178
	GLMM-full	0.040		0.045	0.202			0.188
	STEP	0.023		0.042	0.129			0.181
	LASSO	0.003		0.037	0.031			
	ridge	0.002		0.037	<b>0.028</b>			
	adaptive LASSO	0.007		0.038	0.053			
	Bayesian LASSO	0.002		0.045	0.031			0.236
	SSVS	0.005		0.046	0.047			0.237
B16	GLMM-oracle	0		0.061	0.031			0.190
	GLMM-full	0.040		0.072	0.216			0.200
	STEP	0.023		0.069	0.142			0.183
	LASSO	0.003		0.059	0.044			
	ridge	0.002		0.059	<b>0.040</b>			
	adaptive LASSO	0.007		0.061	0.066			
	Bayesian LASSO	0.002		0.074	0.046			0.276
	SSVS	0.005		0.076	0.062			0.277
B17	GLMM-oracle	0	0.044	0.034	0.039	0.180		0.180
	GLMM-full	0.038	0.054	0.041	0.197	0.200		0.188
	STEP	0.022	0.042	0.038	0.130	0.187		0.182
	LASSO	0.003	0.012	0.037	0.037			
	ridge	0.002	0.010	0.037	<b>0.034</b>			
	adaptive LASSO	0.007	0.018	0.038	0.059			
	Bayesian LASSO	0.002	0.010	0.045	<b>0.034</b>	0.086		0.239
	SSVS	0.004	0.014	0.046	0.049	0.111		0.241

B18	GLMM-oracle	0	0.044	0.057	0.051	0.181		0.193
	GLMM-full	0.038	0.056	0.067	0.213	0.202		0.202
	STEP	0.021	0.042	0.063	0.140	0.187		0.183
	LASSO	0.003	0.013	0.060	0.049			
	ridge	0.002	0.010	0.060	<b>0.045</b>			
	adaptive LASSO	0.007	0.018	0.061	0.072			
	Bayesian LASSO	0.002	0.010	0.072	0.048	0.086		0.277
	SSVS	0.004	0.014	0.074	0.063	0.112		0.278
B19	GLMM-oracle	0	0.050	0.037	0.044	0.189		0.183
	GLMM-full	0.039	0.060	0.044	0.207	0.209		0.191
	STEP	0.023	0.078	0.042	0.151	0.195		0.185
	LASSO	0.004	0.127	0.042	<b>0.102</b>			
	ridge	0.006	0.143	0.043	0.114			
	adaptive LASSO	0.009	0.095	0.042	0.104			
	Bayesian LASSO	0.004	0.132	0.057	0.109	0.156		0.242
	SSVS	0.007	0.101	0.055	0.108	0.192		0.244
B20	GLMM-oracle	0	0.051	0.057	0.054	0.190		0.195
	GLMM-full	0.040	0.060	0.067	0.222	0.210		0.203
	STEP	0.023	0.082	0.064	0.165	0.195		0.186
	LASSO	0.005	0.133	0.063	<b>0.117</b>			
	ridge	0.005	0.150	0.063	0.127			
	adaptive LASSO	0.009	0.102	0.063	0.119			
	Bayesian LASSO	0.004	0.134	0.081	0.122	0.155		0.278
	SSVS	0.007	0.102	0.080	0.121	0.191		0.281
B21	GLMM-oracle	0	0.042	0.039	0.062	0.181	0.179	0.182
	GLMM-full	0.041	0.051	0.052	0.311	0.204	0.185	0.194
	STEP	0.024	0.038	0.048	0.203	0.191	0.183	0.185
	LASSO	0.002	0.011	0.038	0.045			
	ridge	0.002	0.010	0.038	<b>0.040</b>			
	adaptive LASSO	0.006	0.016	0.040	0.077			
	Bayesian LASSO	0.002	0.010	0.055	0.049	0.083	0.083	0.242
	SSVS	0.004	0.012	0.055	0.065	0.099	0.099	0.244
B22	GLMM-oracle	0	0.043	0.062	0.074	0.183	0.180	0.193
	GLMM-full	0.042	0.052	0.079	0.333	0.206	0.186	0.205
	STEP	0.024	0.038	0.074	0.220	0.191	0.183	0.186
	LASSO	0.002	0.011	0.060	0.057			



	ridge	0.002	0.010	0.060	<b>0.052</b>			
	adaptive LASSO	0.007	0.016	0.062	0.090			
	Bayesian LASSO	0.002	0.010	0.083	0.063	0.083	0.084	0.278
	SSVS	0.004	0.012	0.085	0.080	0.100	0.100	0.281
B23	GLMM-oracle	0	0.046	0.041	0.067	0.191	0.183	0.186
	GLMM-full	0.043	0.054	0.053	0.325	0.214	0.191	0.198
	STEP	0.025	0.077	0.050	0.246	0.199	0.187	0.190
	LASSO	0.004	0.118	0.049	<b>0.168</b>			
	ridge	0.005	0.120	0.049	0.174			
	adaptive LASSO	0.008	0.098	0.048	0.169			
	Bayesian LASSO	0.004	0.115	0.073	0.175	0.148	0.136	0.249
	SSVS	0.007	0.094	0.071	0.171	0.181	0.165	0.251
B24	GLMM-oracle	0	0.047	0.065	0.079	0.192	0.184	0.197
	GLMM-full	0.043	0.054	0.082	0.343	0.215	0.192	0.209
	STEP	0.025	0.080	0.078	0.267	0.199	0.188	0.191
	LASSO	0.004	0.121	0.069	<b>0.180</b>			
	ridge	0.005	0.124	0.069	0.187			
	adaptive LASSO	0.008	0.098	0.068	0.183			
	Bayesian LASSO	0.004	0.118	0.104	0.193	0.146	0.135	0.282
	SSVS	0.007	0.096	0.103	0.189	0.180	0.164	0.284
B25	GLMM-oracle	0	0.045	0.041	0.084	0.190	0.179	0.183
	GLMM-full	0.041	0.050	0.052	0.314	0.204	0.185	0.194
	STEP	0.023	0.038	0.048	0.207	0.190	0.182	0.185
	LASSO	0.003	0.012	0.044	0.057			
	ridge	0.002	0.010	0.043	<b>0.052</b>			
	adaptive LASSO	0.007	0.018	0.044	0.089			
	Bayesian LASSO	0.002	0.009	0.056	0.054	0.084	0.084	0.241
	SSVS	0.003	0.012	0.058	0.070	0.102	0.101	0.244
B26	GLMM-oracle	0	0.046	0.060	0.095	0.191	0.180	0.195
	GLMM-full	0.041	0.051	0.075	0.330	0.205	0.186	0.205
	STEP	0.023	0.038	0.070	0.219	0.190	0.183	0.186
	LASSO	0.002	0.012	0.065	0.067			
	ridge	0.002	0.010	0.065	<b>0.062</b>			
	adaptive LASSO	0.007	0.017	0.066	0.100			
	Bayesian LASSO	0.002	0.010	0.080	0.066	0.084	0.084	0.277
	SSVS	0.004	0.012	0.082	0.082	0.103	0.101	0.278

B27	GLMM-oracle	0	0.051	0.043	0.097	0.202	0.187	0.191
	GLMM-full	0.043	0.056	0.053	0.340	0.217	0.194	0.202
	STEP	0.025	0.084	0.053	0.282	0.205	0.191	0.194
	LASSO	0.005	0.109	0.058	0.242			
	ridge	0.008	0.102	0.06	0.253			
	adaptive LASSO	0.009	0.093	0.054	<b>0.223</b>			
	Bayesian LASSO	0.007	0.101	0.085	0.255	0.168	0.148	0.254
	SSVS	0.009	0.086	0.081	0.232	0.196	0.173	0.255
B28	GLMM-oracle	0	0.052	0.064	0.109	0.202	0.188	0.203
	GLMM-full	0.044	0.057	0.078	0.357	0.218	0.195	0.214
	STEP	0.025	0.084	0.076	0.296	0.205	0.191	0.195
	LASSO	0.005	0.113	0.081	0.264			
	ridge	0.008	0.107	0.082	0.274			
	adaptive LASSO	0.009	0.099	0.078	<b>0.246</b>			
	Bayesian LASSO	0.006	0.105	0.112	0.273	0.166	0.147	0.284
	SSVS	0.009	0.088	0.107	0.248	0.196	0.172	0.287
B29	GLMM-oracle		0.137	0.130	0.715	0.335	0.305	0.325
	GLMM-full		0.137	0.130	<b>0.715</b>	0.335	0.305	0.325
	STEP		0.172	0.152	0.900	0.319	0.298	0.301
	LASSO		0.141	0.178	0.930			
	ridge		0.110	0.157	0.753			
	adaptive LASSO		0.148	0.170	0.914			
	Bayesian LASSO		0.127	0.281	0.923	0.227	0.202	0.440
	SSVS		0.124	0.257	0.811	0.277	0.240	0.443
B30	GLMM-oracle		0.138	0.179	0.746	0.336	0.306	0.342
	GLMM-full		0.138	0.179	<b>0.746</b>	0.336	0.306	0.342
	STEP		0.173	0.198	0.936	0.319	0.298	0.302
	LASSO		0.145	0.219	0.984			
	ridge		0.113	0.198	0.802			
	adaptive LASSO		0.150	0.214	0.960			
	Bayesian LASSO		0.129	0.324	0.964	0.223	0.199	0.462
	SSVS		0.125	0.302	0.846	0.275	0.238	0.470
B31	GLMM-oracle	0.000	0.091	0.083	0.086	0.261		0.263
	GLMM-full	0.087	0.122	0.116	0.464	0.299		0.288
	STEP	0.050	0.083	0.104	0.300	0.276		0.268
	LASSO	0.006	0.018	0.088	0.079			

	ridge	0.005	0.014	0.088	<b>0.072</b>			
	adaptive LASSO	0.016	0.031	0.092	0.130			
	Bayesian LASSO	0.003	0.011	0.115	0.076	0.109		0.402
	SSVS	0.012	0.024	0.125	0.130	0.166		0.407
B32	GLMM-oracle	0.000	0.093	0.156	0.124	0.264		0.294
	GLMM-full	0.090	0.128	0.203	0.521	0.302		0.320
	STEP	0.051	0.086	0.182	0.344	0.277		0.271
	LASSO	0.006	0.019	0.162	0.117			
	Ridge	0.005	0.014	0.160	<b>0.109</b>			
	adaptive LASSO	0.016	0.031	0.166	0.167			
	Bayesian LASSO	0.003	0.011	0.199	0.118	0.110		0.460
	SSVS	0.012	0.025	0.210	0.172	0.167		0.464
B33	GLMM-oracle	0.000	0.040	0.024	0.032	0.134		0.140
	GLMM-full	0.021	0.043	0.027	0.119	0.147		0.145
	STEP	0.012	0.036	0.026	0.082	0.136		0.134
	LASSO	0.002	0.016	0.025	0.029			
	ridge	0.002	0.011	0.025	<b>0.026</b>			
	adaptive LASSO	0.005	0.022	0.025	0.044			
	Bayesian LASSO	0.001	0.011	0.028	<b>0.026</b>	0.076		0.250
	SSVS	0.002	0.015	0.029	0.033	0.090		0.250
B34	GLMM-oracle	0.000	0.039	0.067	0.053	0.135		0.200
	GLMM-full	0.021	0.043	0.072	0.143	0.149		0.204
	STEP	0.012	0.036	0.073	0.106	0.136		0.135
	LASSO	0.002	0.015	0.071	0.053			
	ridge	0.002	0.011	0.071	0.049			
	adaptive LASSO	0.005	0.022	0.072	0.069			
	Bayesian LASSO	0.001	0.011	0.072	<b>0.048</b>	0.077		0.338
	SSVS	0.003	0.015	0.072	0.055	0.091		0.339
B35	GLMM-oracle	0.000	0.110	0.094	0.102	0.283		0.280
	GLMM-full	0.173	0.248	0.401	2.465	0.381		0.394
	STEP	0.098	0.235	0.258	1.501	0.330		0.323
	LASSO	0.003	0.212	0.124	0.208			
	ridge	0.002	0.208	0.123	<b>0.199</b>			
	adaptive LASSO	0.012	0.185	0.133	0.324			
	Bayesian LASSO	0.008	0.178	0.270	0.341	0.176		0.465

	SSVS	0.019	0.155	0.307	0.508	0.235		0.477
B36	GLMM-oracle	0.000	0.107	0.136	0.121	0.284		0.303
	GLMM-full	0.179	0.250	0.491	2.584	0.394		0.424
	STEP	0.099	0.227	0.317	1.543	0.384		0.325
	LASSO	0.003	0.213	0.168	0.233			
	ridge	0.002	0.209	0.168	<b>0.222</b>			
	adaptive LASSO	0.012	0.186	0.179	0.352			
	Bayesian LASSO	0.008	0.177	0.334	0.373	0.176		0.506
	SSVS	0.019	0.154	0.379	0.547	0.235		0.519

## 4 Example of R code used for fitting the various meta-analysis models

### 4.1 Generate example datasets

Here we generate two mock datasets. The first has a continuous outcome ( $y$ ) and two continuous covariates ( $z_1, z_2$ ). The second dataset has a binary outcome ( $x$ ), one continuous covariate ( $w_1$ ) and one binary ( $w_2$ ).

```
# The github library ("bipd") contains functions for generating sample data and
running Bayesian IPD-MA methods.
library(devtools)
devtools::install_github("MikeJSeo/bipd") #parallel packages take a while to
install
library(bipd)

##load data
ds <- generate_ipdma_example(type = "continuous")
ds2 <- generate_ipdma_example(type = "binary")
head(ds)
head(ds2)
```

### 4.2 Code for fitting GLMM

```
# continuous outcome
library(lme4) #for fitting glmm
m1 <- lmer(y ~ studyid + (z1+z2)*treat + (-1 + treat|studyid), data = ds)
summary(m1)

# estimating treatment effect for specific values of the covariates
contr <- c(rep(0, 8), 1, 1, 0.5) #if covariates are standardized this needs to be
modified
v1 <- vcov(m1)
sel <- c(sqrt(contr %*% v1 %*% contr))
mean1 <- c(contr %*% summary(m1)$coefficients[, "Estimate"])
mean1 + qnorm(c(.025, 0.5, .975)) * as.vector(sel[[1]])

# binary outcome
m2 <- glmer(y ~ studyid + (w1+w2)*treat + (-1 + treat|studyid), data = ds2, family
= binomial)
summary(m2)

# estimating treatment effect for specific values of the covariates
contr <- c(rep(0, 8), 1, 1, 0.5) #if covariates are standardized this needs to be
modified
v1 <- vcov(m2)
sel <- c(sqrt(contr %*% v1 %*% contr))
mean1 <- c(contr %*% summary(m2)$coefficients[, "Estimate"])
exp(mean1 + qnorm(c(.025, 0.5, .975)) * as.vector(sel[[1]])) #calculate odds ratio
```

### 4.3 Code for fitting STEP

```
# continuous outcome
m3 <- glm(y ~ studyid + (z1+z2)*treat, data = ds) #glm model without mixed effects
s1 <- step(m3, scope=list(lower = ~ z1+z2+treat), direction = "both")
summary(s1)

# binary outcome
m4 <- glm(y ~ studyid + (w1+w2)*treat, family = binomial(link = "logit"), data =
ds2)
s2 <- step(m4, scope=list(lower = ~ w1+w2+treat), direction = "both")
```

```
summary(s2)
```

#### 4.4 Code for fitting LASSO

```
# continuous outcome
library(glmnet)
p.fac <- c(rep(0, 5), rep(0, 2), 0, rep(1,2)) # Shrinkage is only on effect
modifiers
lambdas <- 10^seq(3, -3, by = -.1) # manually specify lambda value to cross
validate

data_glmnet <- model.matrix(y~ studyid + (z1+z2)*treat, data = ds)
data_glmnet <- data_glmnet[,-1]
data_glmnet <- cbind(y = ds$y, data_glmnet = data_glmnet)
cvfit <- cv.glmnet(as.matrix(data_glmnet[,-1]), as.matrix(data_glmnet[,1]),
penalty.factor = p.fac, family = "gaussian", type.measure = "deviance", lambda =
lambdas)
coef(cvfit, s = "lambda.min")

# binary outcome
data_glmnet <- model.matrix(y~ studyid + (w1+w2)*treat, data = ds2)
data_glmnet <- data_glmnet[,-1]
data_glmnet <- cbind(y = ds2$y, data_glmnet = data_glmnet)
cvfit <- cv.glmnet(as.matrix(data_glmnet[,-1]), as.matrix(data_glmnet[,1]),
penalty.factor = p.fac, family = "binomial", type.measure = "deviance", lambda =
lambdas)
coef(cvfit, s = "lambda.min")
```

#### 4.5 Code for fitting ridge

```
# continuous outcome
data_glmnet <- model.matrix(y~ studyid + (z1+z2)*treat, data = ds)
data_glmnet <- data_glmnet[,-1]
data_glmnet <- cbind(y = ds$y, data_glmnet = data_glmnet)
cvfit.ridge <- cv.glmnet(as.matrix(data_glmnet[,-1]), as.matrix(data_glmnet[,1]),
penalty.factor = p.fac, family = "gaussian", alpha = 0, type.measure = "deviance",
lambda = lambdas)
coef(cvfit.ridge, s = "lambda.min")

# binary outcome
data_glmnet <- model.matrix(y~ studyid + (w1+w2)*treat, data = ds2)
data_glmnet <- data_glmnet[,-1]
data_glmnet <- cbind(y = ds2$y, data_glmnet = data_glmnet)
cvfit.ridge2 <- cv.glmnet(as.matrix(data_glmnet[,-1]), as.matrix(data_glmnet[,1]),
penalty.factor = p.fac, family = "binomial", alpha = 0, type.measure = "deviance",
lambda = lambdas)
coef(cvfit.ridge2, s = "lambda.min")
```

#### 4.6 Code for fitting adaptive LASSO

```
# continuous outcome
ridge_result <- coef(cvfit.ridge, s = "lambda.min")[-1]
p.fac2 <- p.fac/ abs(ridge_result)

data_glmnet <- model.matrix(y~ studyid + (z1+z2)*treat, data = ds)
data_glmnet <- data_glmnet[,-1]
data_glmnet <- cbind(y = ds$y, data_glmnet = data_glmnet)
cvfit <- cv.glmnet(as.matrix(data_glmnet[,-1]), as.matrix(data_glmnet[,1]),
penalty.factor = p.fac2, family = "gaussian", type.measure = "deviance", lambda =
lambdas)
coef(cvfit, s = "lambda.min")

# binary outcome
```

```

ridge_result <- coef(cvfit.ridge2, s = "lambda.min")[-1]
p.fac2 <- p.fac/ abs(ridge_result)

data_glmnet <- model.matrix(y~ studyid + (w1+w2)*treat, data = ds2)
data_glmnet <- data_glmnet[,-1]
data_glmnet <- cbind(y = ds2$y, data_glmnet = data_glmnet)
cvfit <- cv.glmnet(as.matrix(data_glmnet[,-1]), as.matrix(data_glmnet[,1]),
penalty.factor = p.fac2, family = "binomial", type.measure = "deviance", lambda =
lambdas)
coef(cvfit, s = "lambda.min")

```

#### 4.7 Code for fitting Bayesian LASSO

```

# Stored variable names are as follows:
#"beta" - coefficients for main effects of the covariates
#"gamma" - coefficients for effect modifiers
#"delta" - average treatment effect
#"lambda" - shrinkage parameter

# continuous outcome
ipd <- with(ds, ipdma.model.onestage(y = y, study = studyid, treat = treat, X =
cbind(z1, z2), response = "normal", shrinkage = "laplace", lambda.prior =
list("dgamma",2,0.1)))
##To see the JAGS code used to run the model use the command:
cat(ipd$code)
samples <- ipd.run(ipd, pars.save = c("lambda", "beta", "gamma", "delta"), n.chains
= 3, n.burnin = 500, n.iter = 5000)

samples <- samples[,-3] #remove delta[1] which is 0
summary(samples)
plot(samples) #traceplot and posterior of parameters
coda::gelman.plot(samples) #gelman diagnostic plot

# can also find treatment effect
treatment.effect(ipd, samples, newpatient = c(1,0.5))

# binary outcome
ipd <- with(ds2, ipdma.model.onestage(y = y, study = studyid, treat = treat, X =
cbind(w1, w2), response = "binomial", shrinkage = "laplace"))
samples <- ipd.run(ipd, pars.save = c("lambda", "beta", "gamma", "delta"))
summary(samples)

# can also run methods in parallel using dclone package
samples2 <- ipd.run.parallel(ipd, pars.save = c("lambda", "beta", "gamma",
"delta"))
summary(samples2)

```

#### 4.8 Code for fitting SSVS

```

# Stored variable names are as follows:
#"beta" - coefficients for main effects of the covariates
#"gamma" - coefficients for effect modifiers
#"delta" - coefficient of average treatment effect
#"Ind" - Indicator for assigning a slab prior (instead of a spike prior) i.e.
indicator for including a covariate
#"eta" - Standard deviation of the slab prior

# continuous outcome
ipd <- with(ds, ipdma.model.onestage(y = y, study = studyid, treat = treat, X =
cbind(z1, z2), response = "normal", shrinkage = "SSVS", hy.prior.eta =
list("dunif", 0, 5), g = 1000))
samples <- ipd.run(ipd, pars.save = c("beta", "gamma", "delta", "Ind", "eta"))

samples <- samples[,-5]
summary(samples)
plot(samples)

```

```
coda::gelman.plot(samples)

# binary outcome
ipd <- with(ds2, ipdma.model.onestage(y = y, study = studyid, treat = treat, X =
cbind(w1, w2), response = "binomial", shrinkage = "SSVS"))
samples <- ipd.run(ipd, pars.save = c("beta", "gamma", "delta", "Ind", "eta"))
summary(samples)
treatment.effect(ipd, samples, newpatient = c(1,0.5)) # binary outcome reports odds
ratio
```