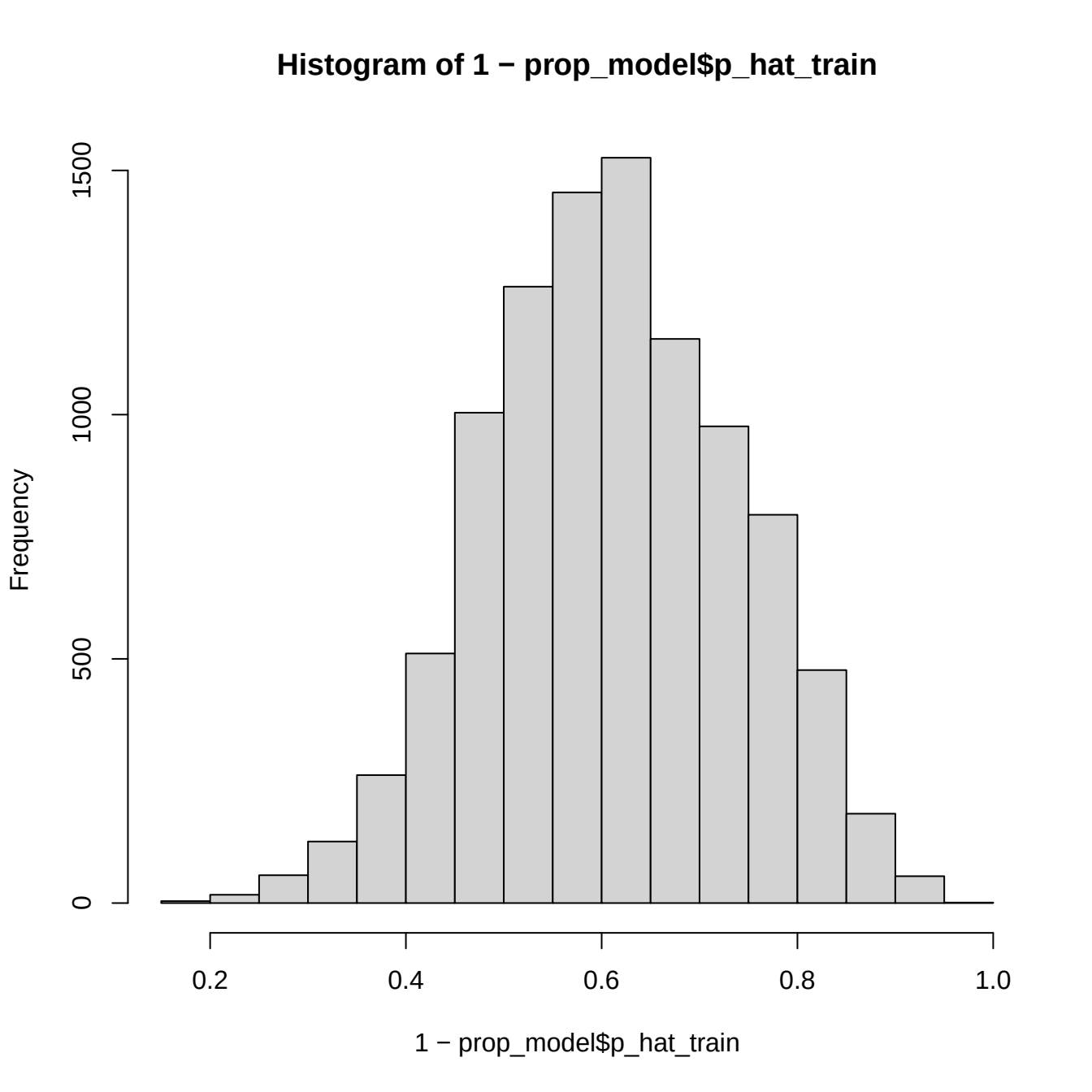
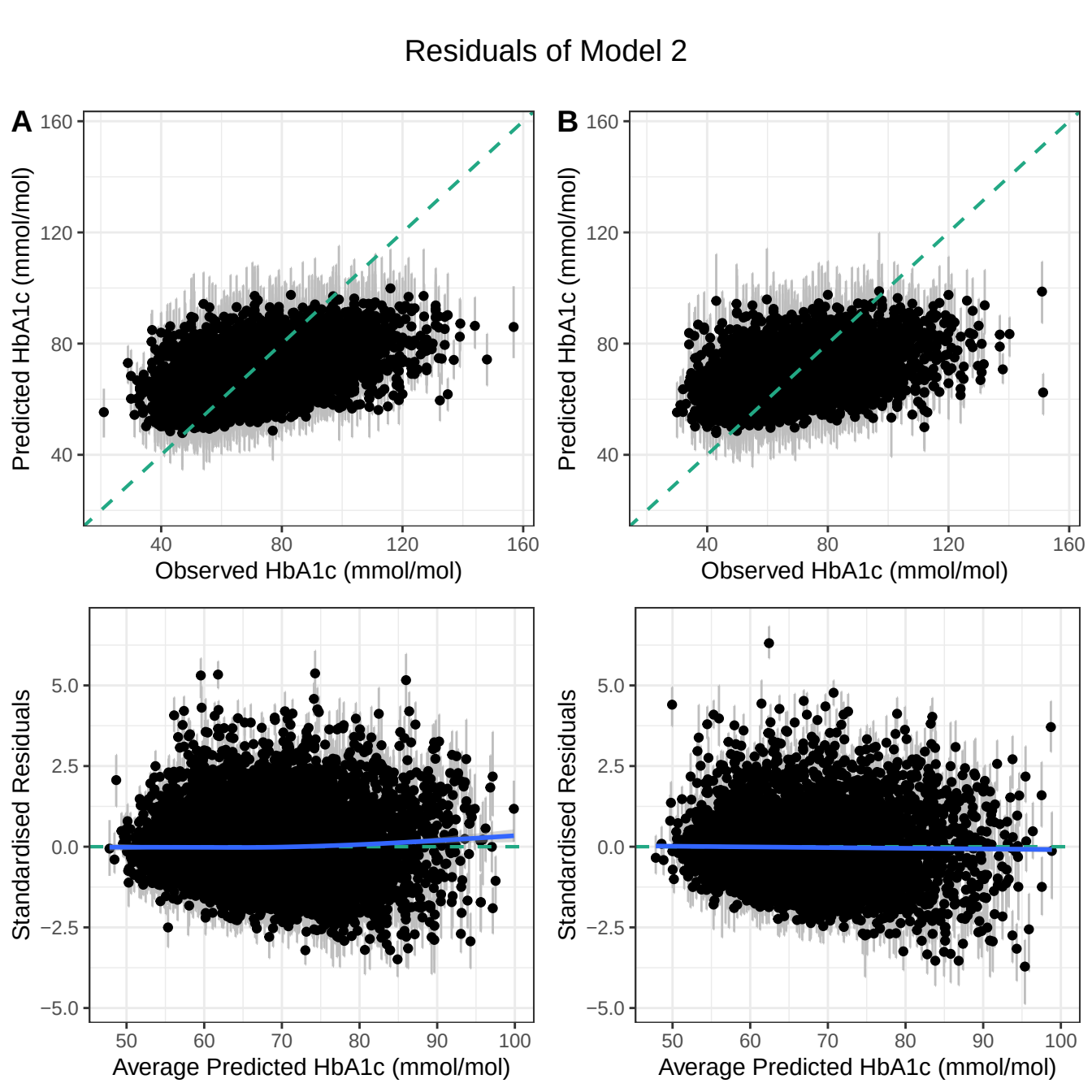
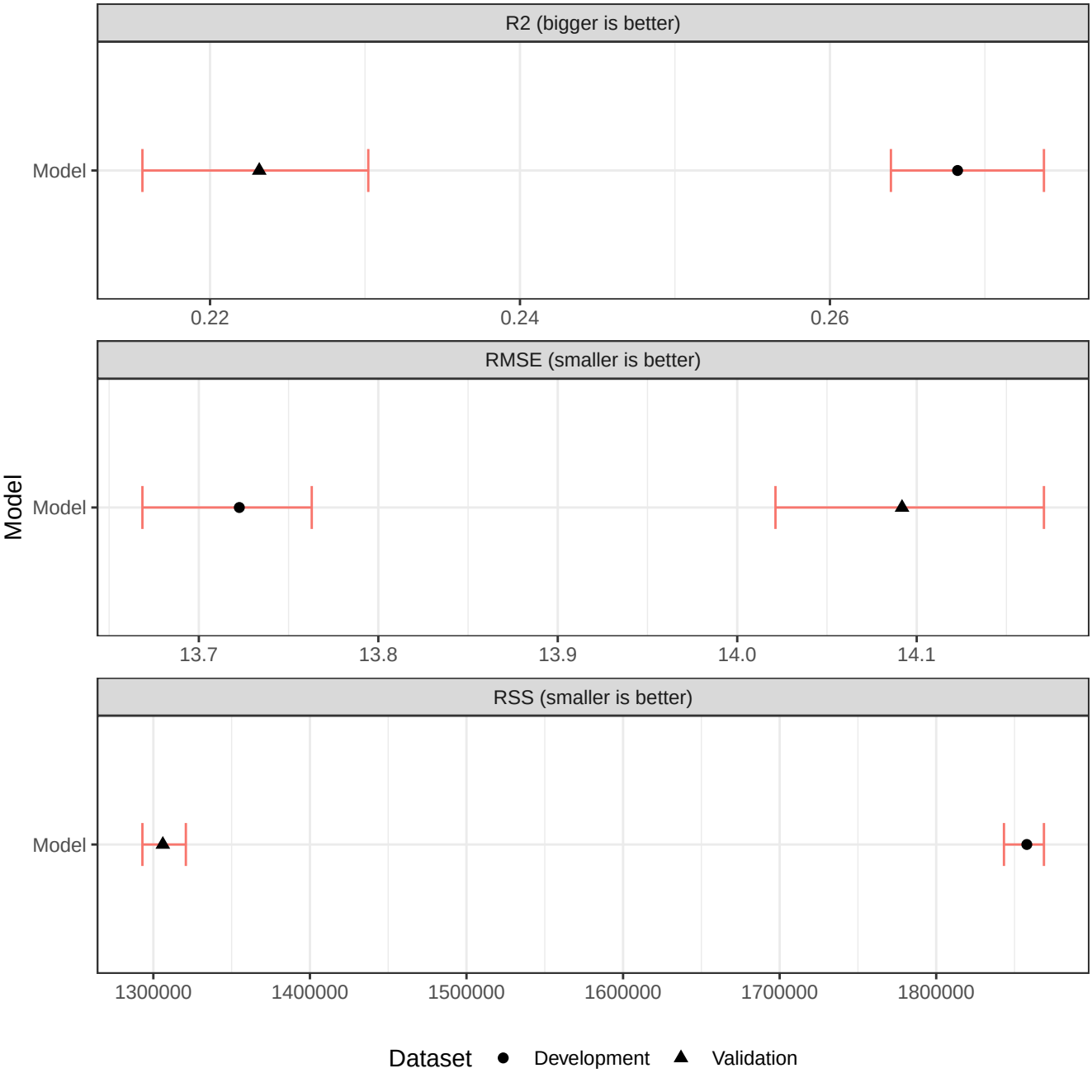
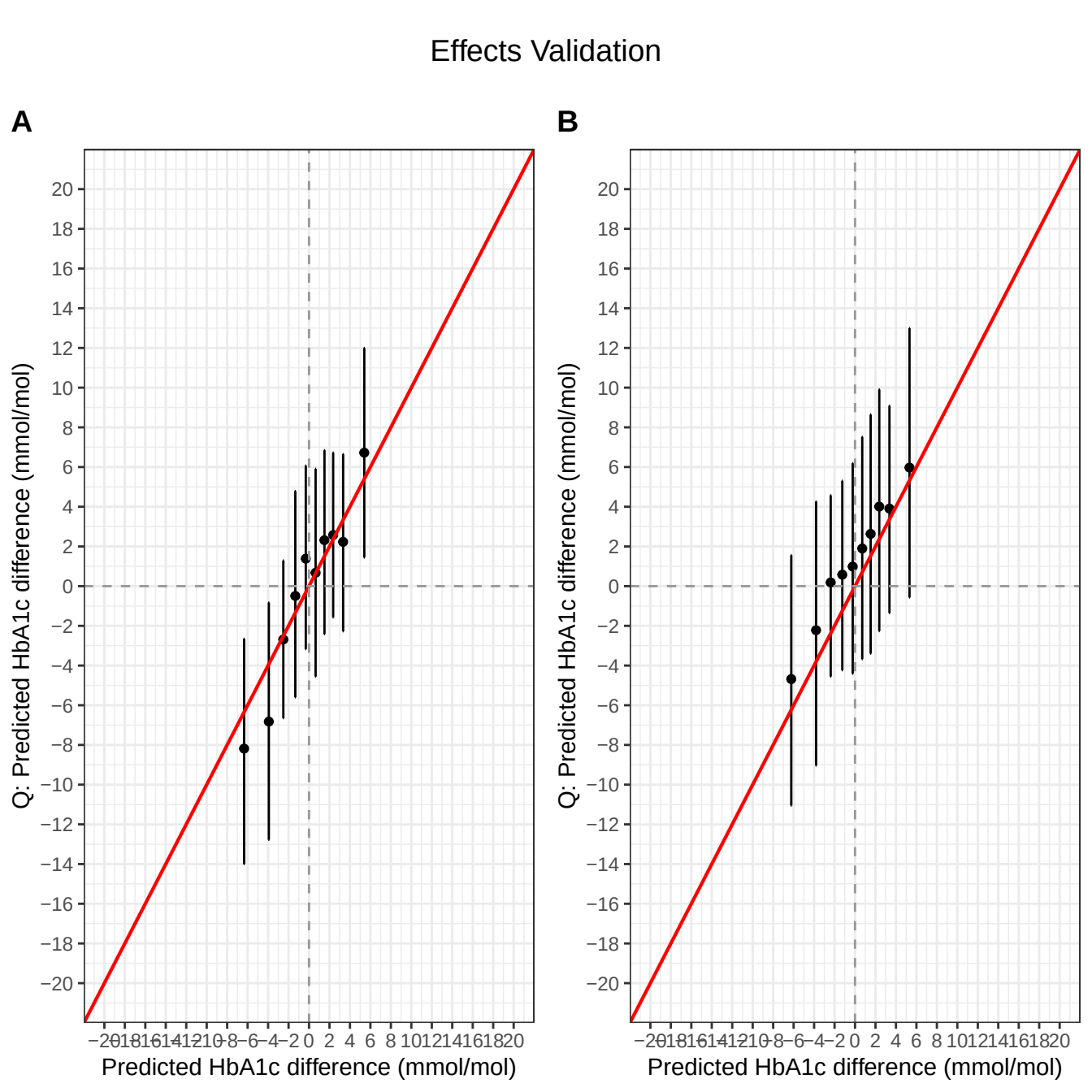
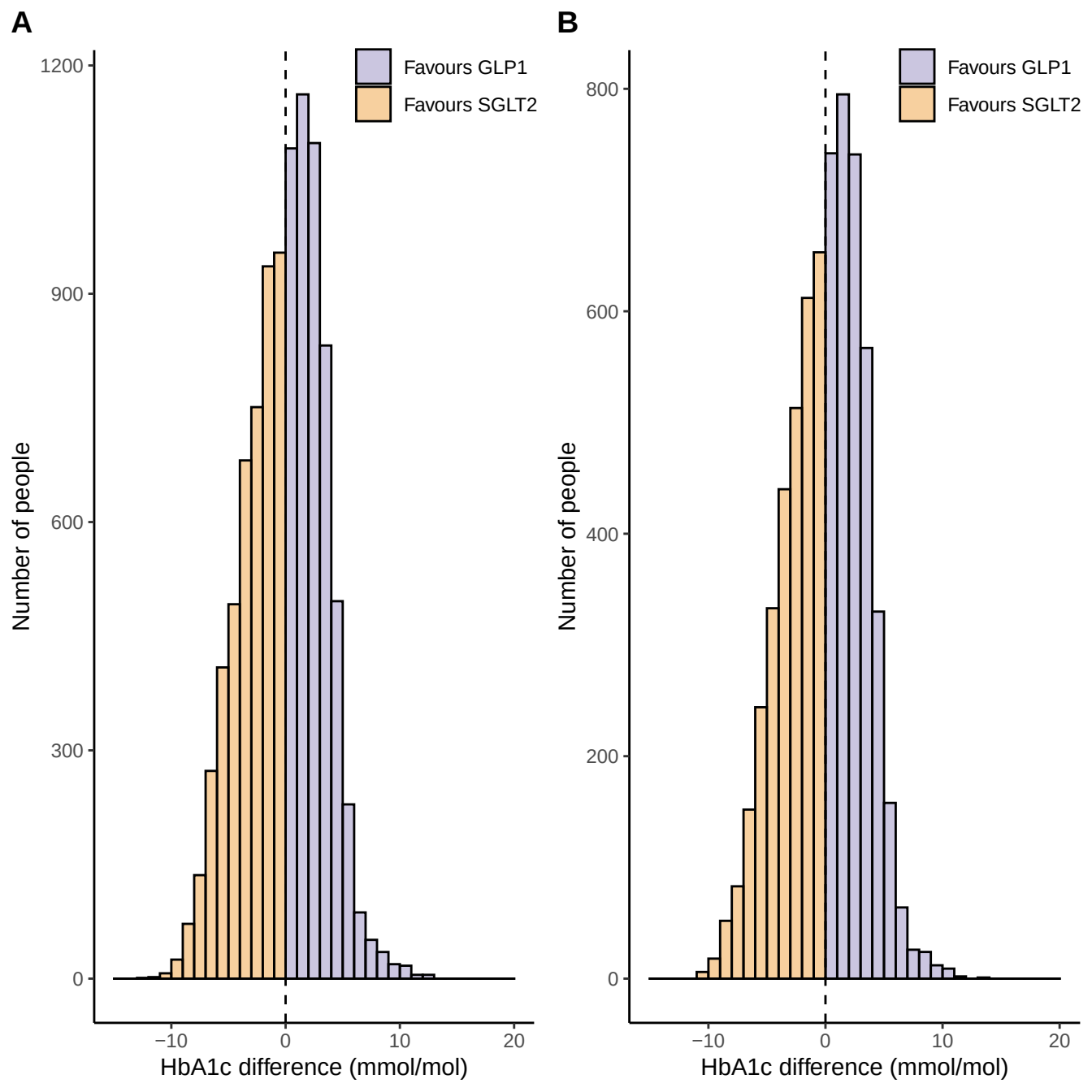

Histogram of 1 − prop_model$p_hat_train
Frequency
0
500
1000
1500
0.2
0.4
0.6
0.8
1.0
1 − prop_model$p_hat_train


Residuals of Model 2
A
B
Predicted HbA1c (mmol/mol)
Observed HbA1c (mmol/mol)
Standardised Residuals
Average Predicted HbA1c (mmol/mol)
40
80
120
160
−5.0
0.0
2.5
5.0
50
60
70
90
100


R2 (bigger is better)
Model
0.22
0.24
0.26
RMSE (smaller is better)
Model
13.7
13.8
13.9
14.0
14.1
RSS (smaller is better)
Model
1300000
1400000
1500000
1600000
1700000
1800000
Model
Dataset
Development
Validation


Effects Validation
A
B
Q: Predicted HbA1c difference (mmol/mol)
Predicted HbA1c difference (mmol/mol)
20
18
16
14
12
10
8
6
4
2
0
−2
−4
−6
−8
−10
−12
−14
−16
−18
−20


A
B
Favours GLP1
Favours SGLT2
Number of people
HbA1c difference (mmol/mol)
1200
900
600
300
0
800
400
200
−10
0
10
20