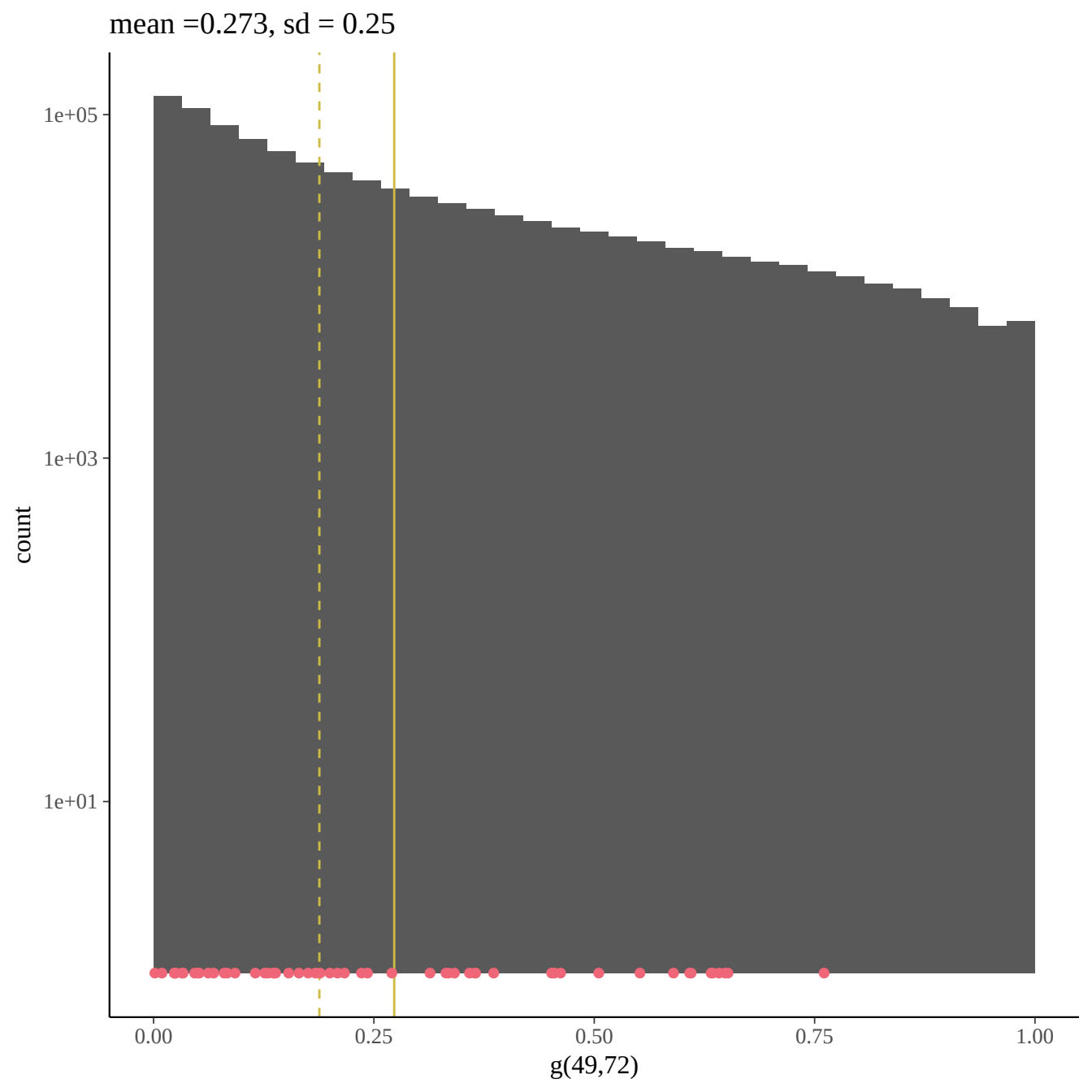
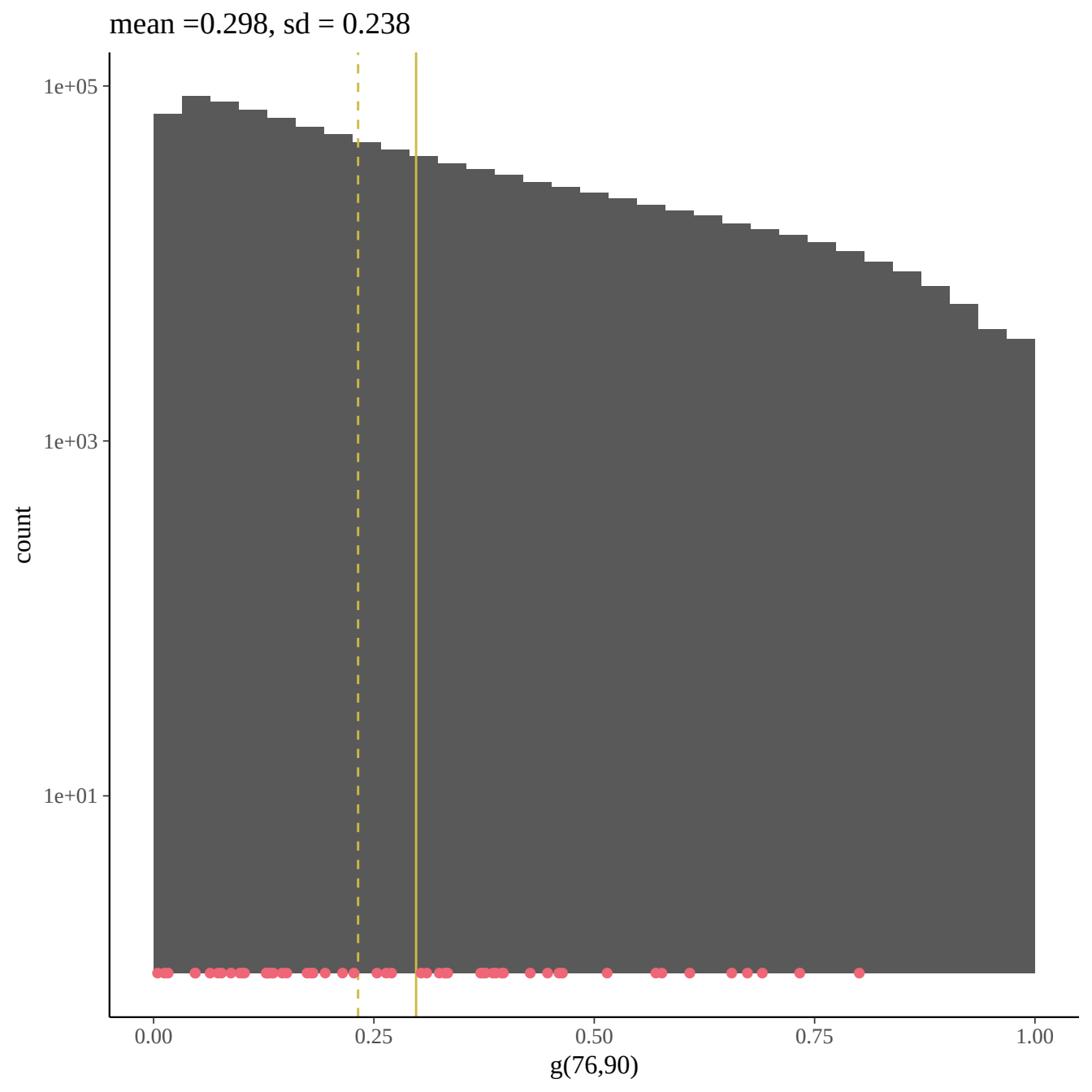
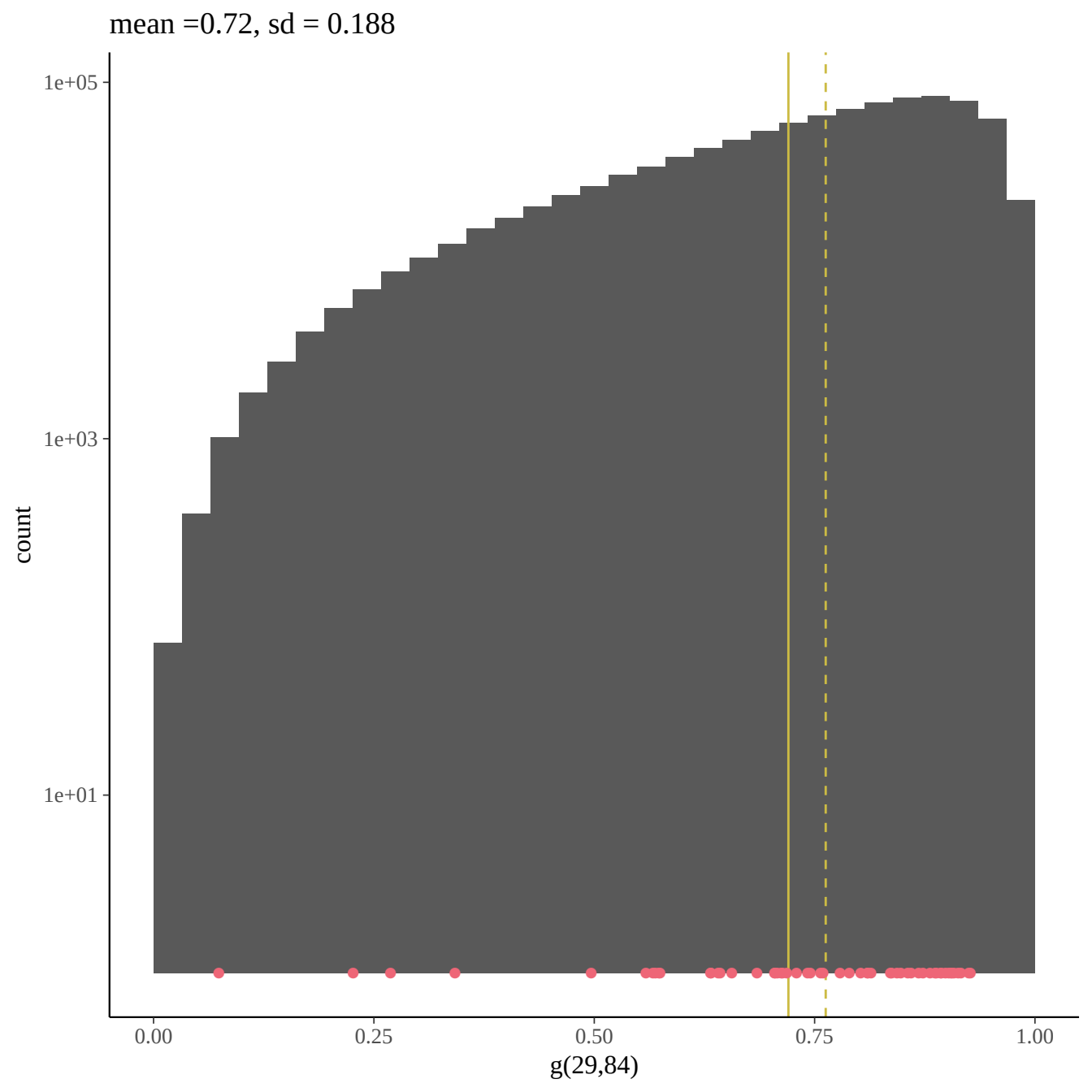
logit−uniform model, Healthy all (RDMH)

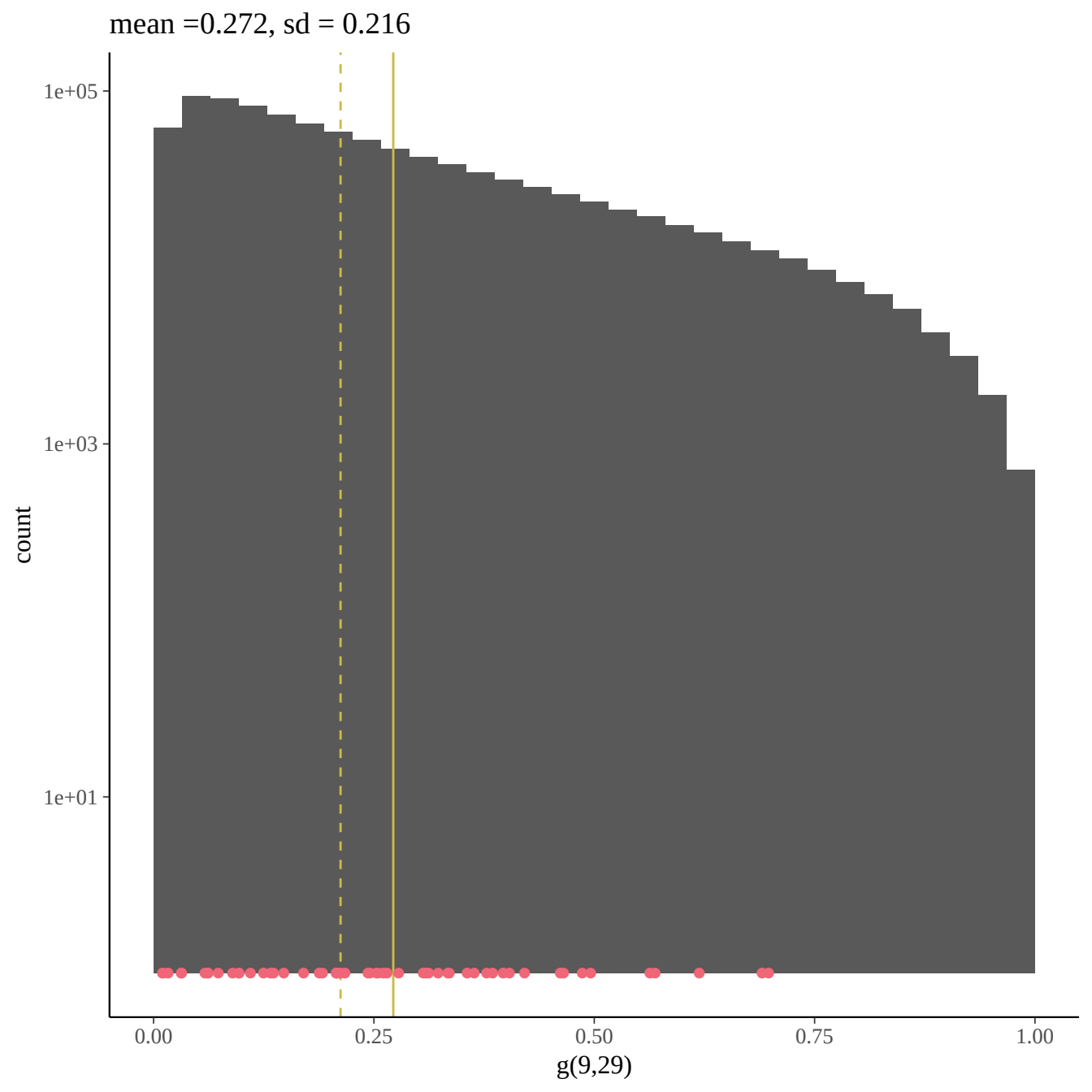
10 graph prop, 55 indiv

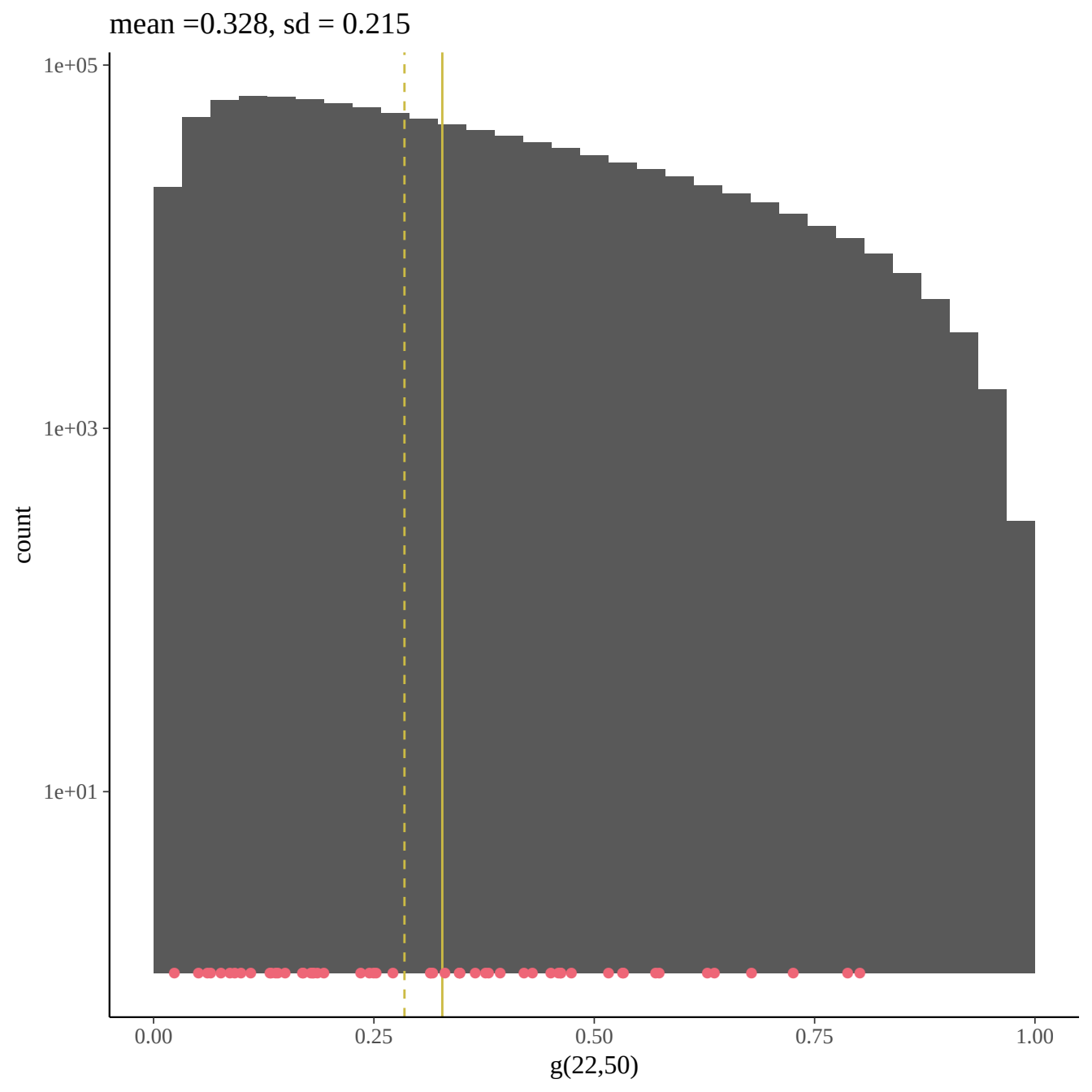
LML = −976.62353793386

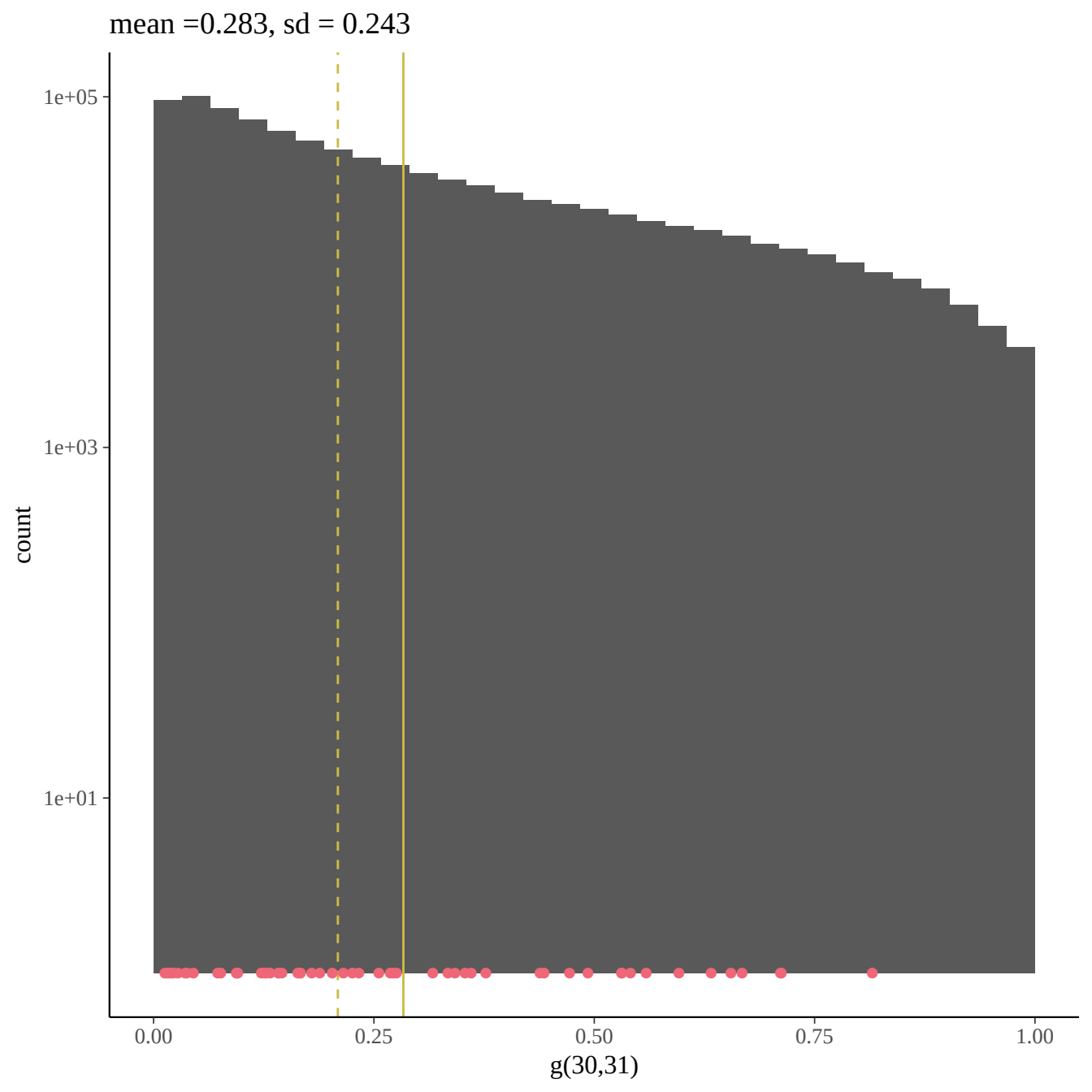

mean =0.273, sd = 0.25
1e+05
1e+03
1e+01
count
0.00
0.25
0.50
0.75
1.00
g(49,72)

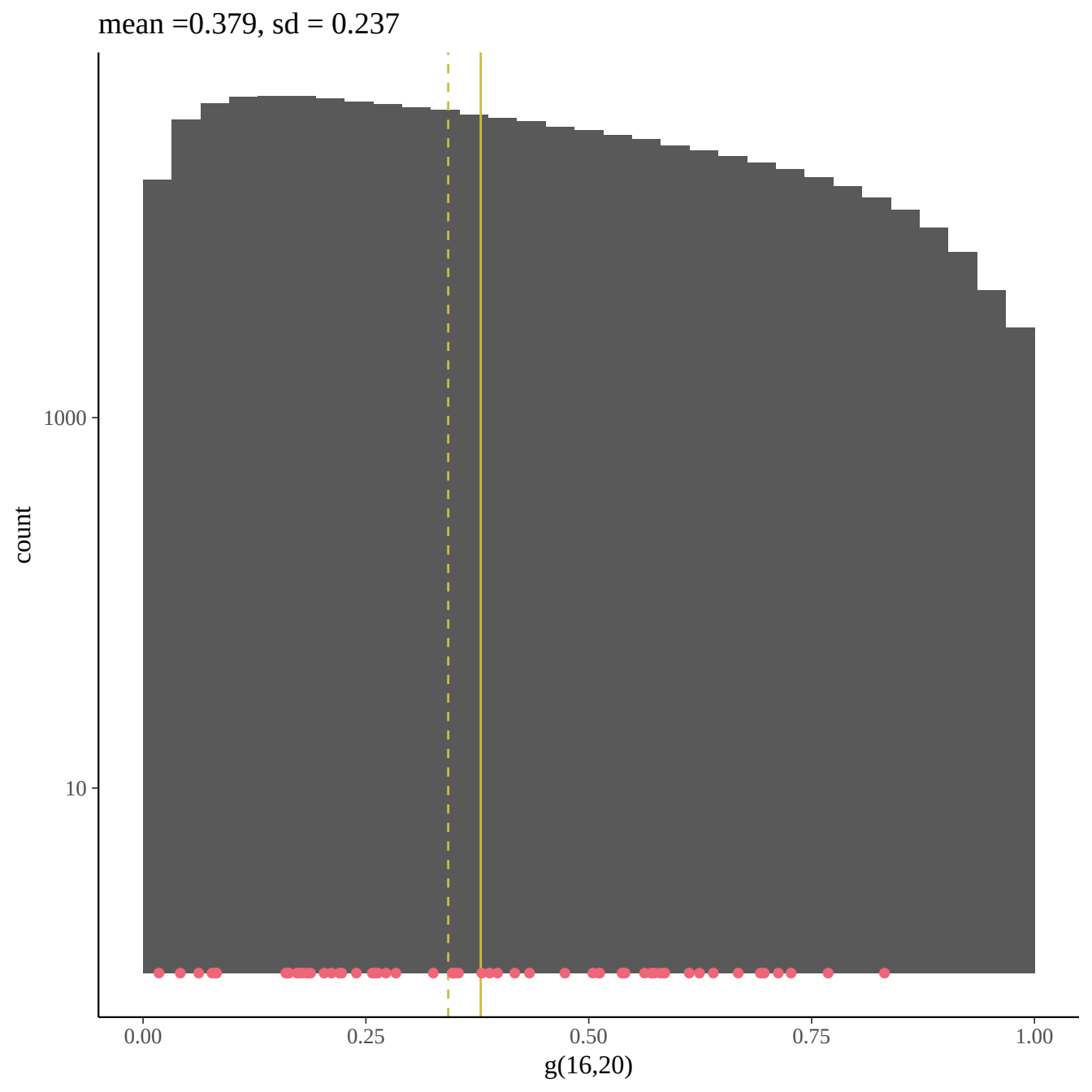

mean =0.298, sd = 0.238
1e+05
1e+03
1e+01
count
0.00
0.25
0.50
0.75
1.00
g(76,90)

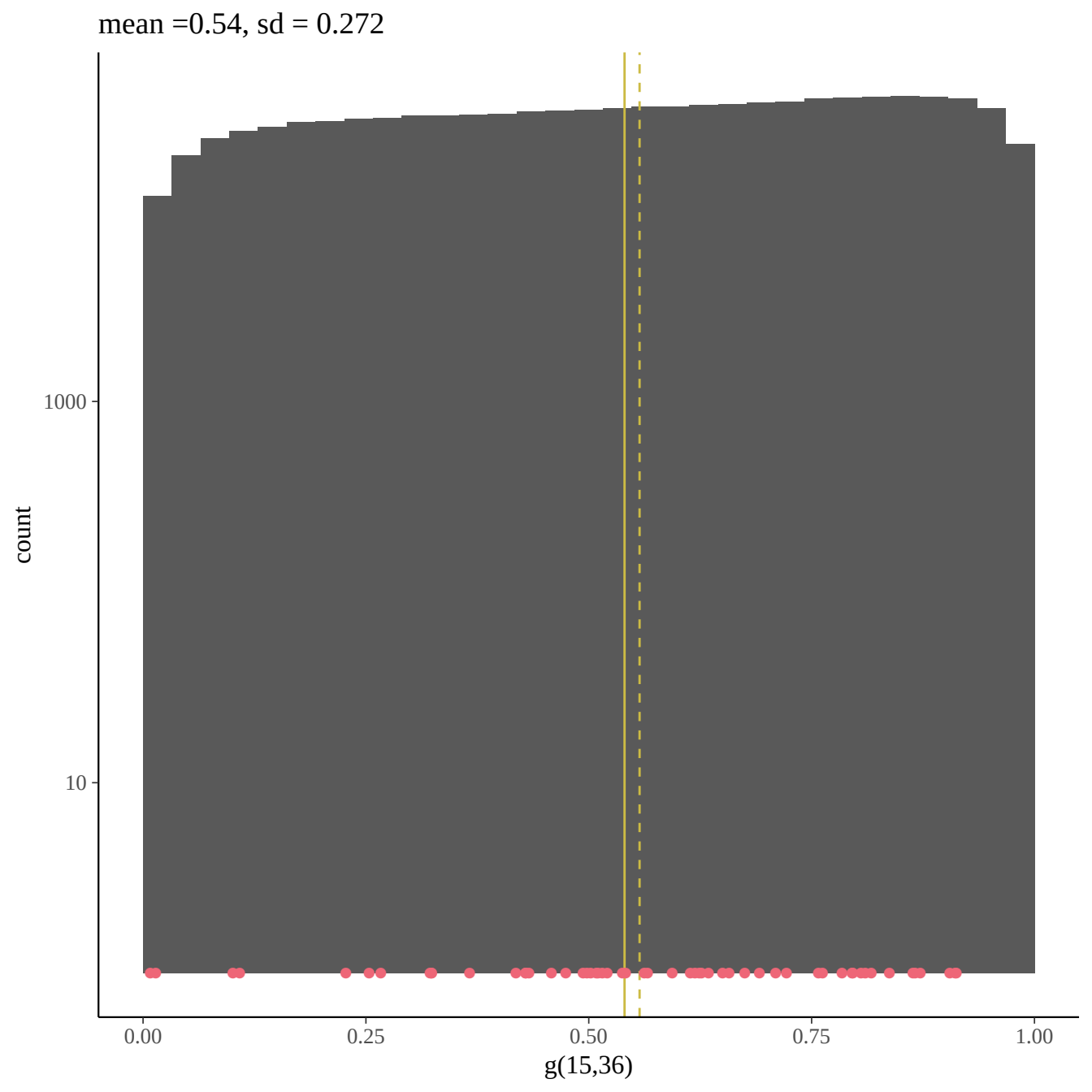

mean =0.72, sd = 0.188
1e+05
1e+03
1e+01
count
0.00
0.25
0.50
0.75
1.00
g(29,84)

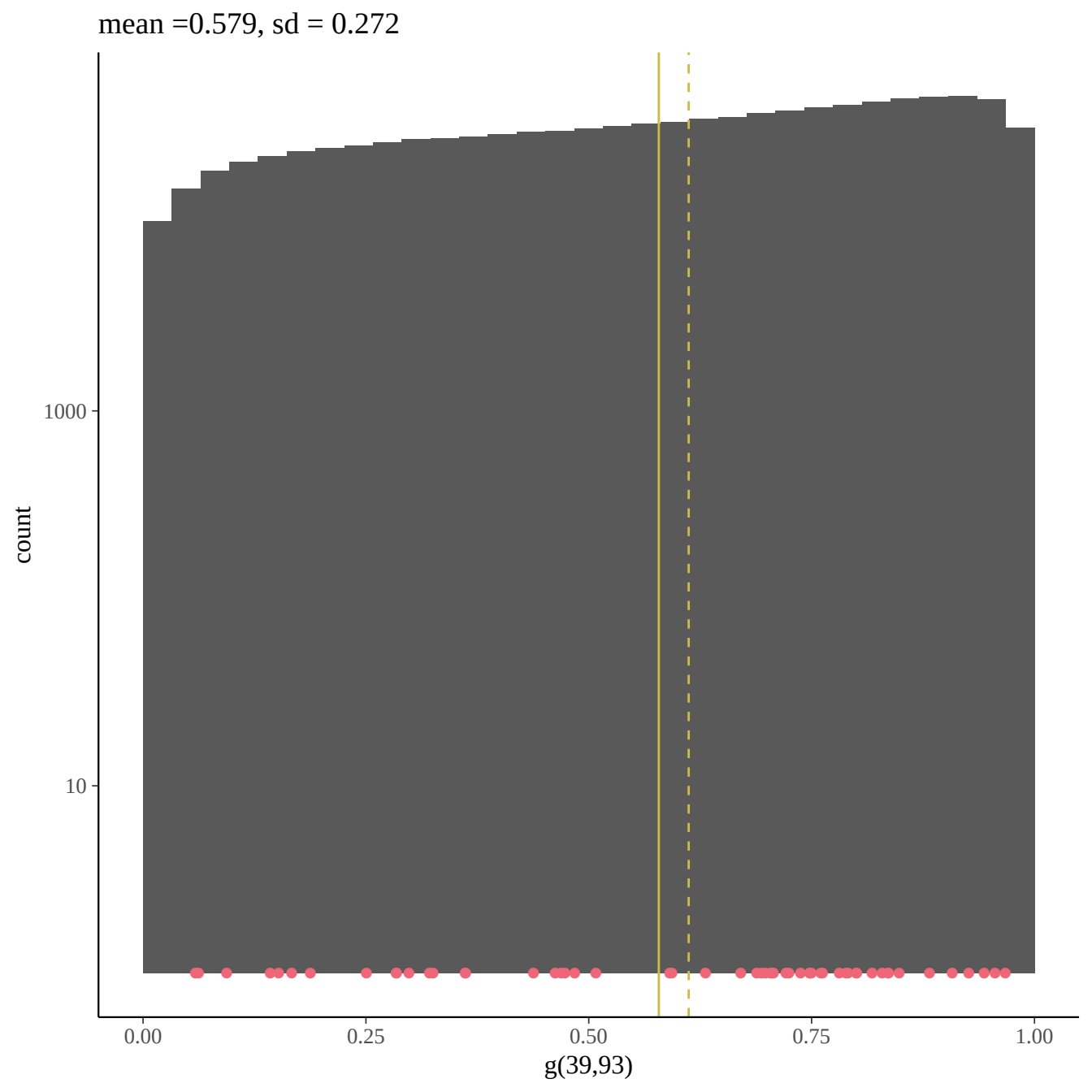

mean =0.272, sd = 0.216
1e+05
1e+03
1e+01
count
0.00
0.25
0.50
0.75
1.00
g(9,29)

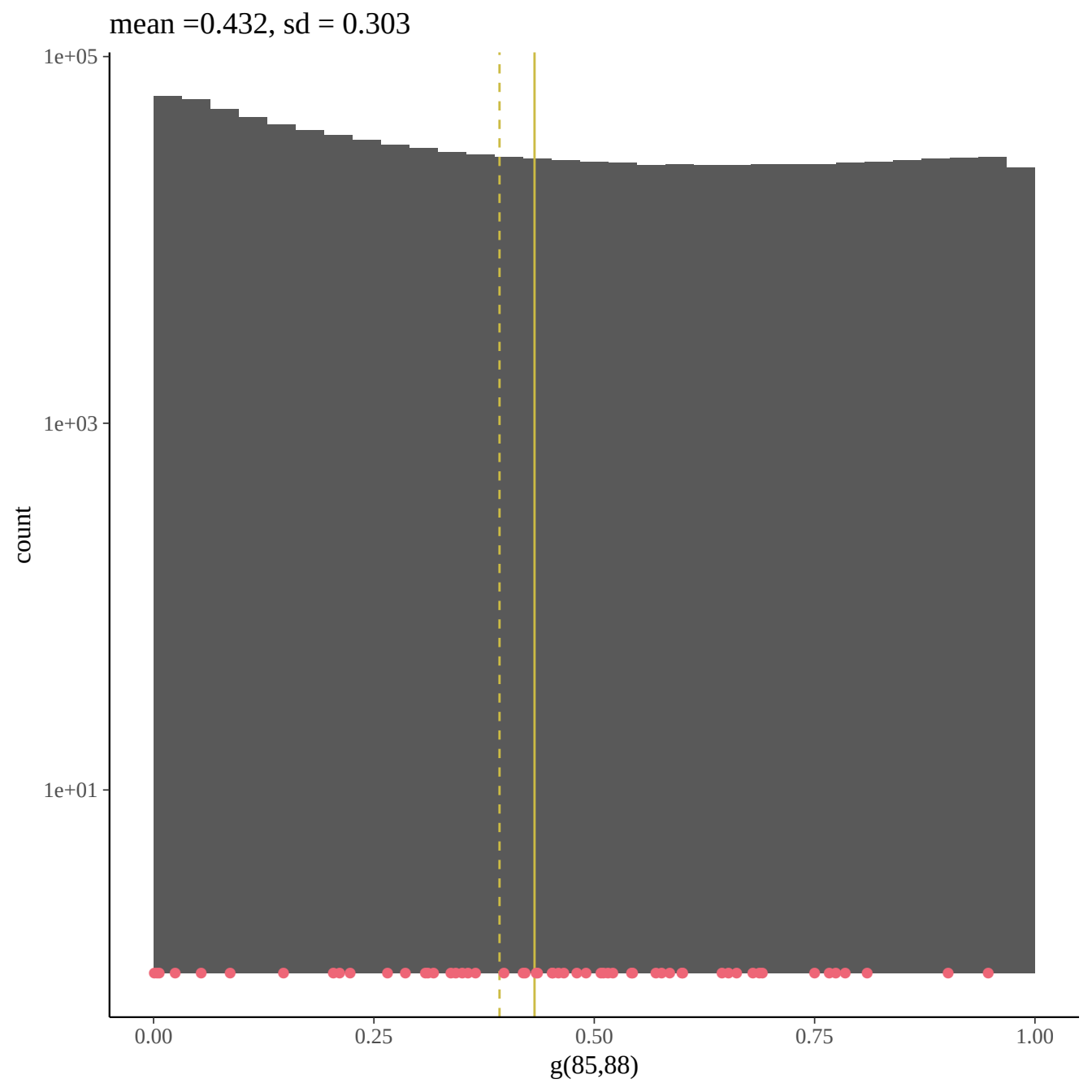

mean =0.328, sd = 0.215
1e+05
1e+03
1e+01
count
0.00
0.25
0.50
0.75
1.00
g(22,50)

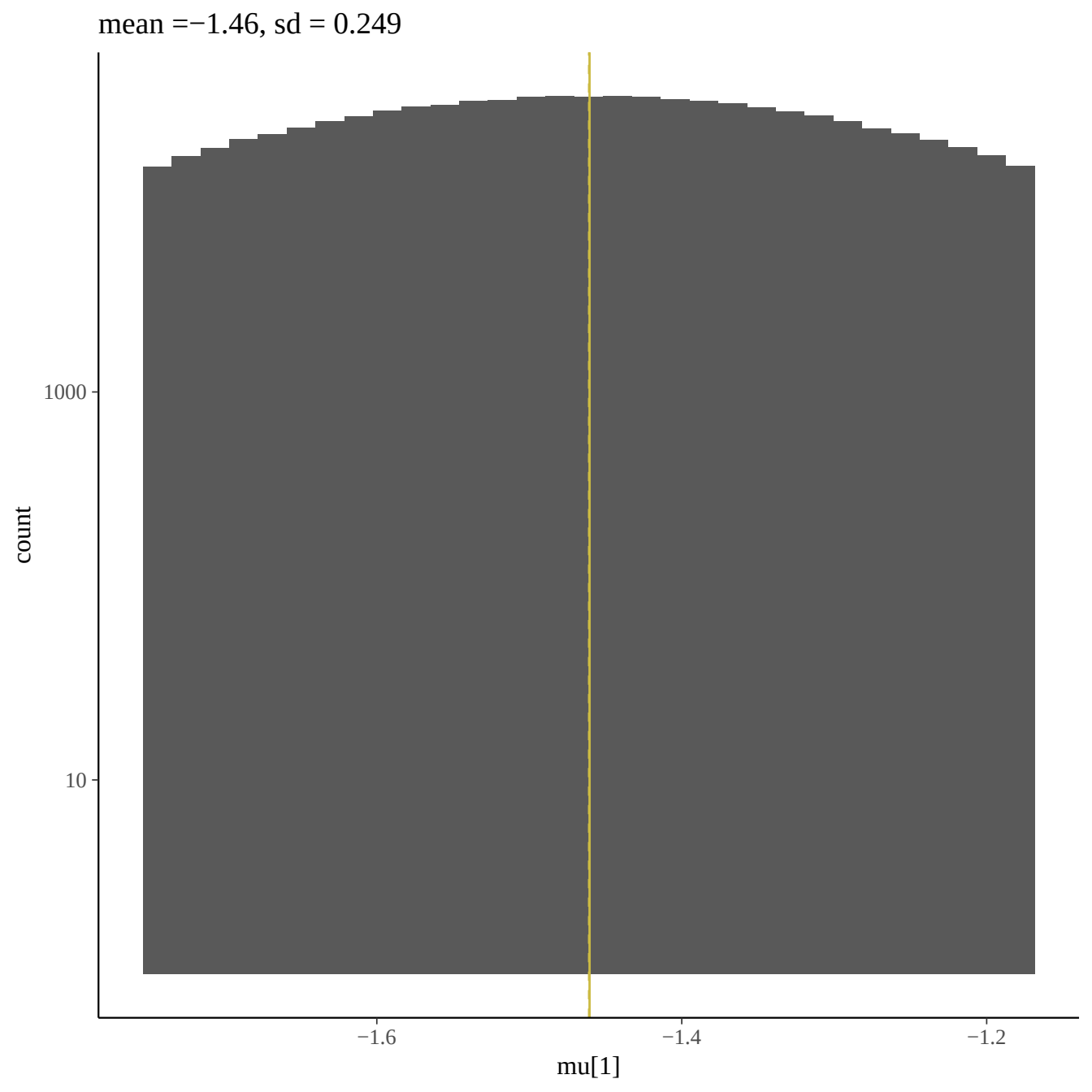

mean =0.283, sd = 0.243
1e+05
1e+03
1e+01
count
0.00
0.25
0.50
0.75
1.00
g(30,31)

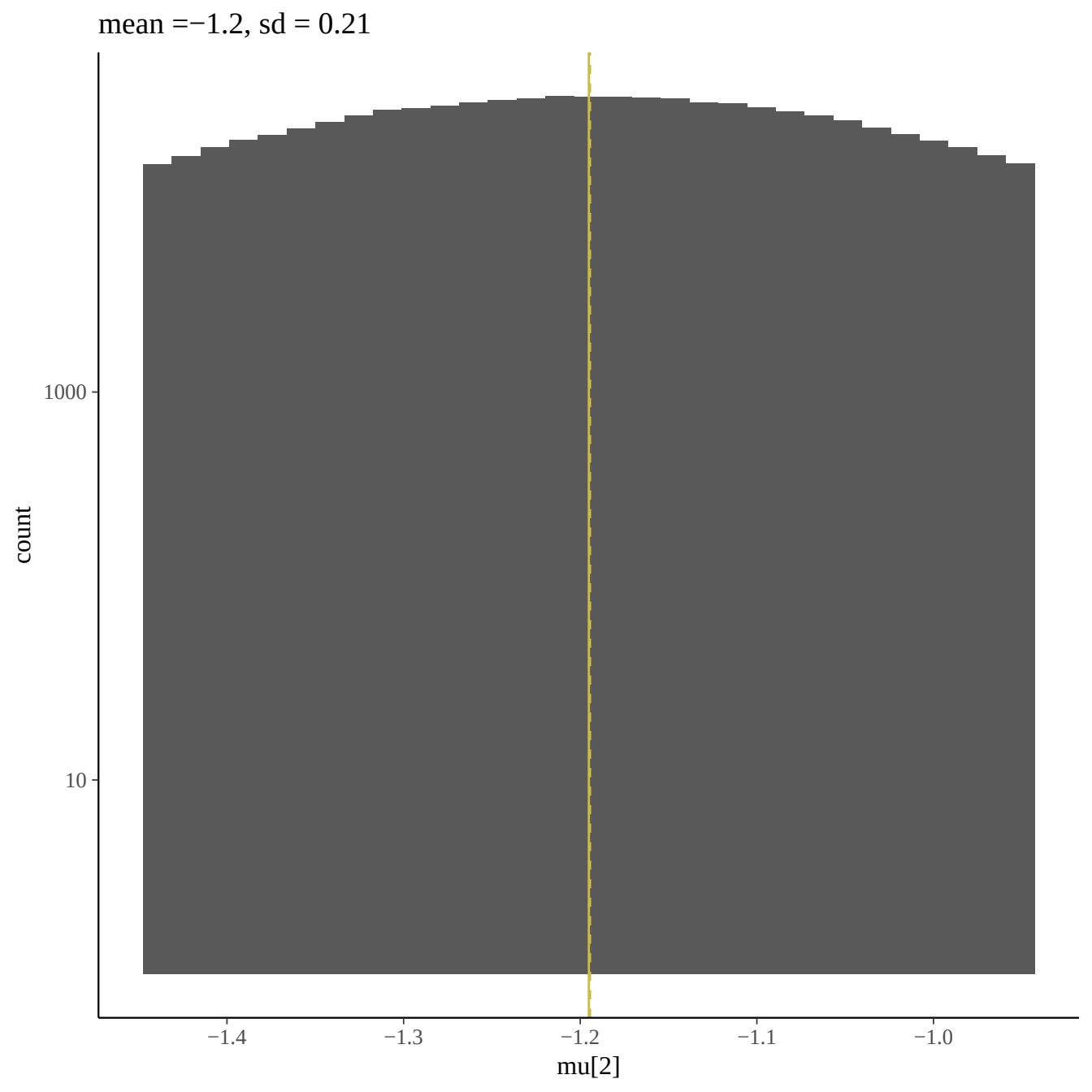

mean =0.379, sd = 0.237
1000
10
count
0.00
0.25
0.50
0.75
1.00
g(16,20)

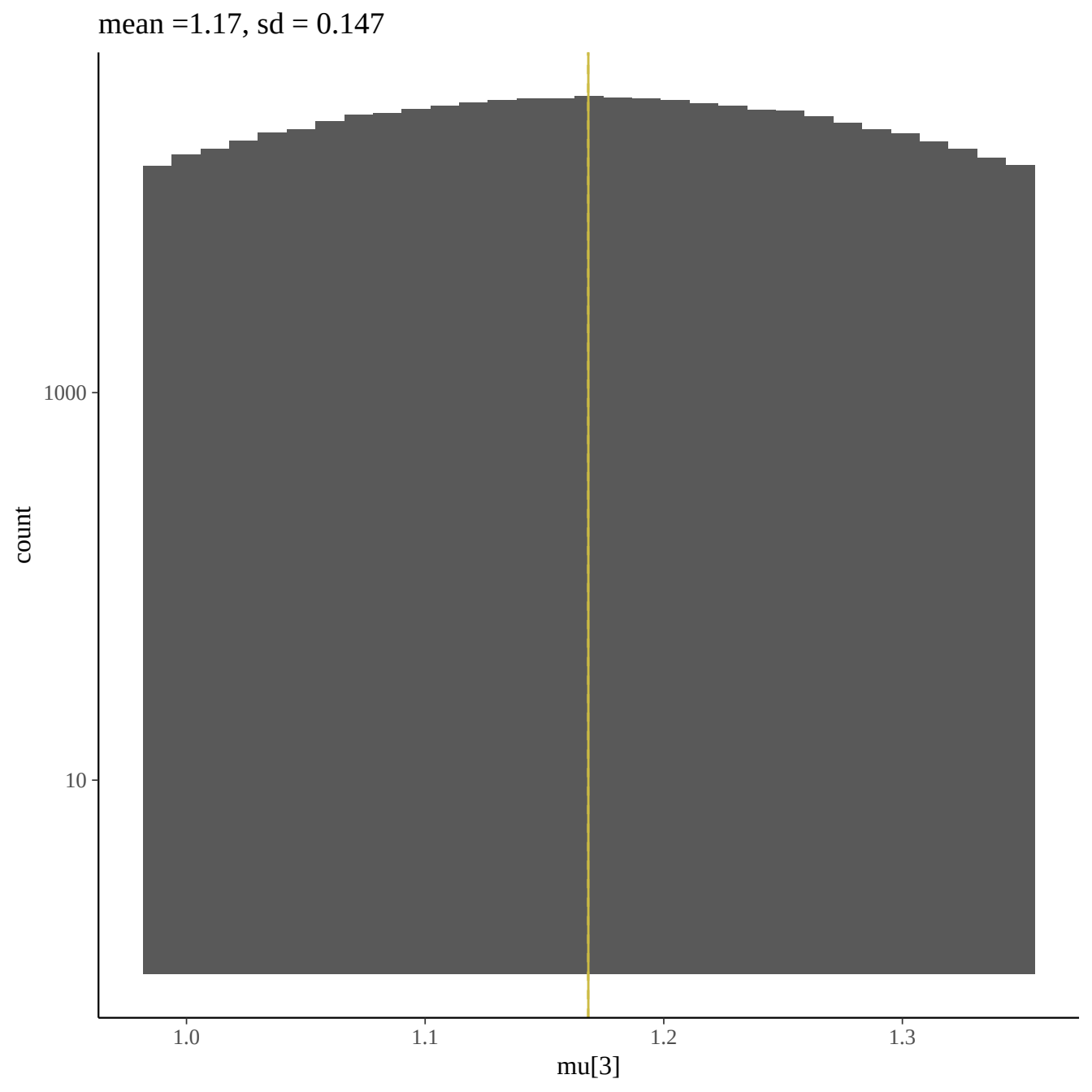

mean =0.54, sd = 0.272
1000
10
count
0.00
0.25
0.50
0.75
1.00
g(15,36)

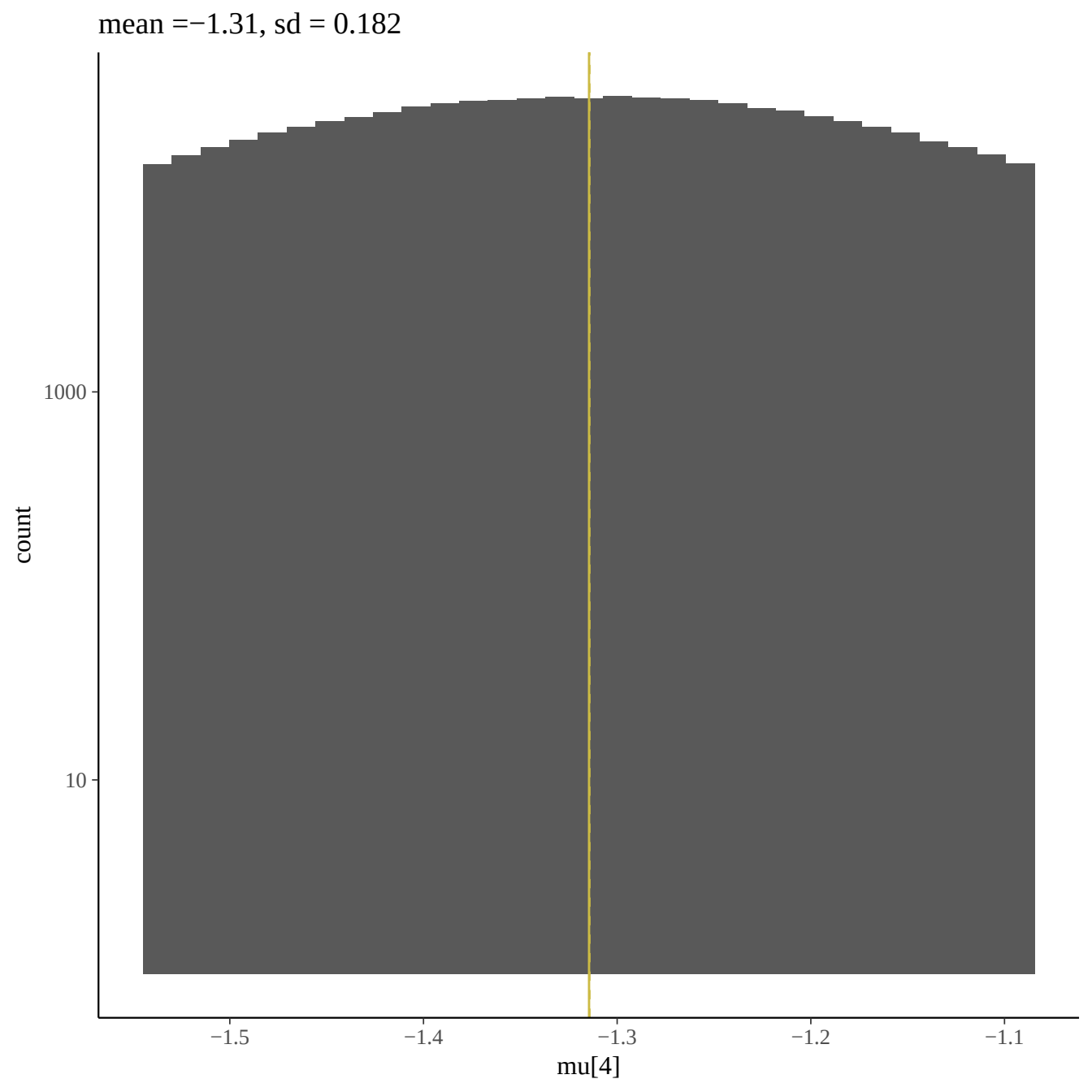

mean =0.579, sd = 0.272
1000
10
count
0.00
0.25
0.50
0.75
1.00
g(39,93)

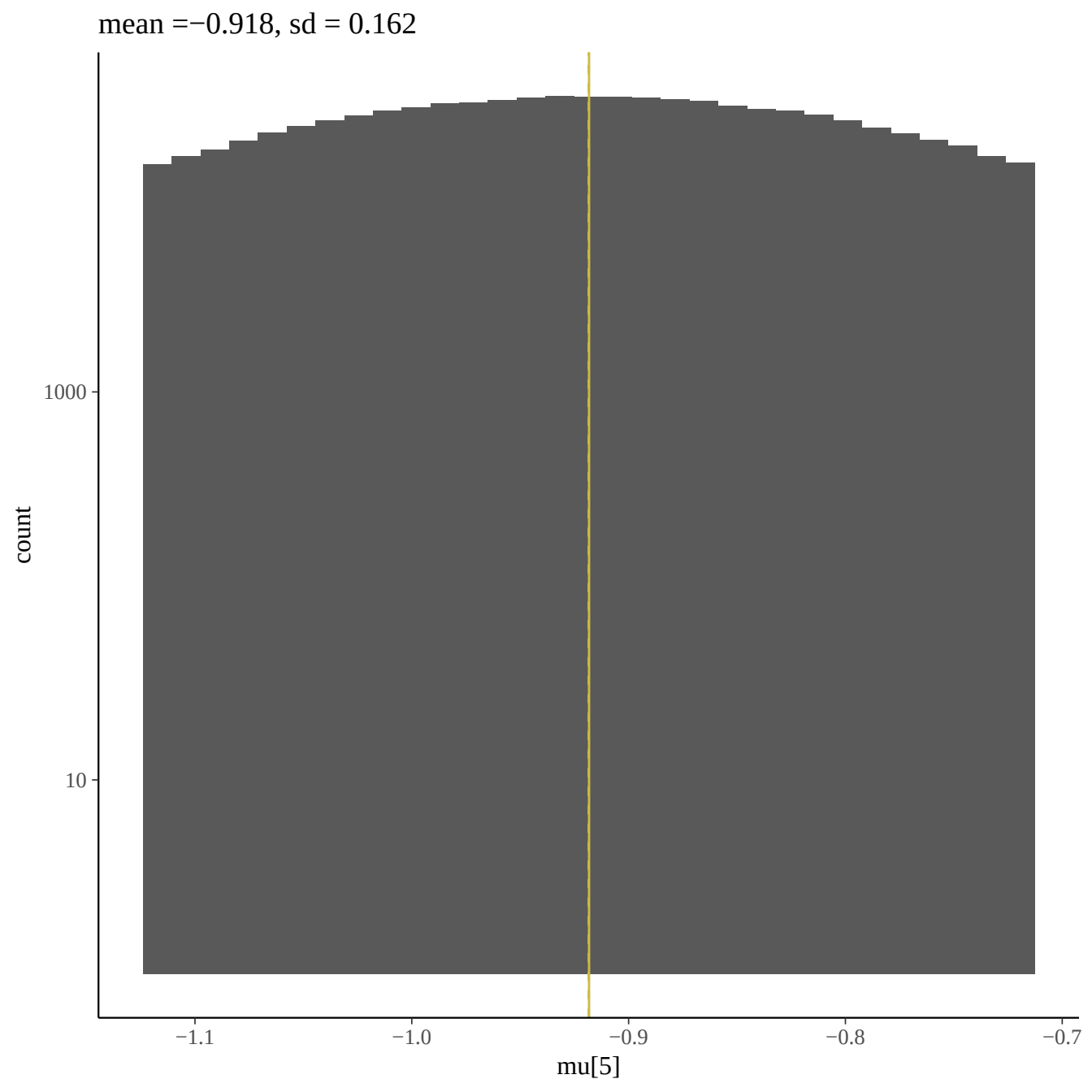

mean =0.432, sd = 0.303
1e+05
1e+03
1e+01
count
0.00
0.25
0.50
0.75
1.00
g(85,88)

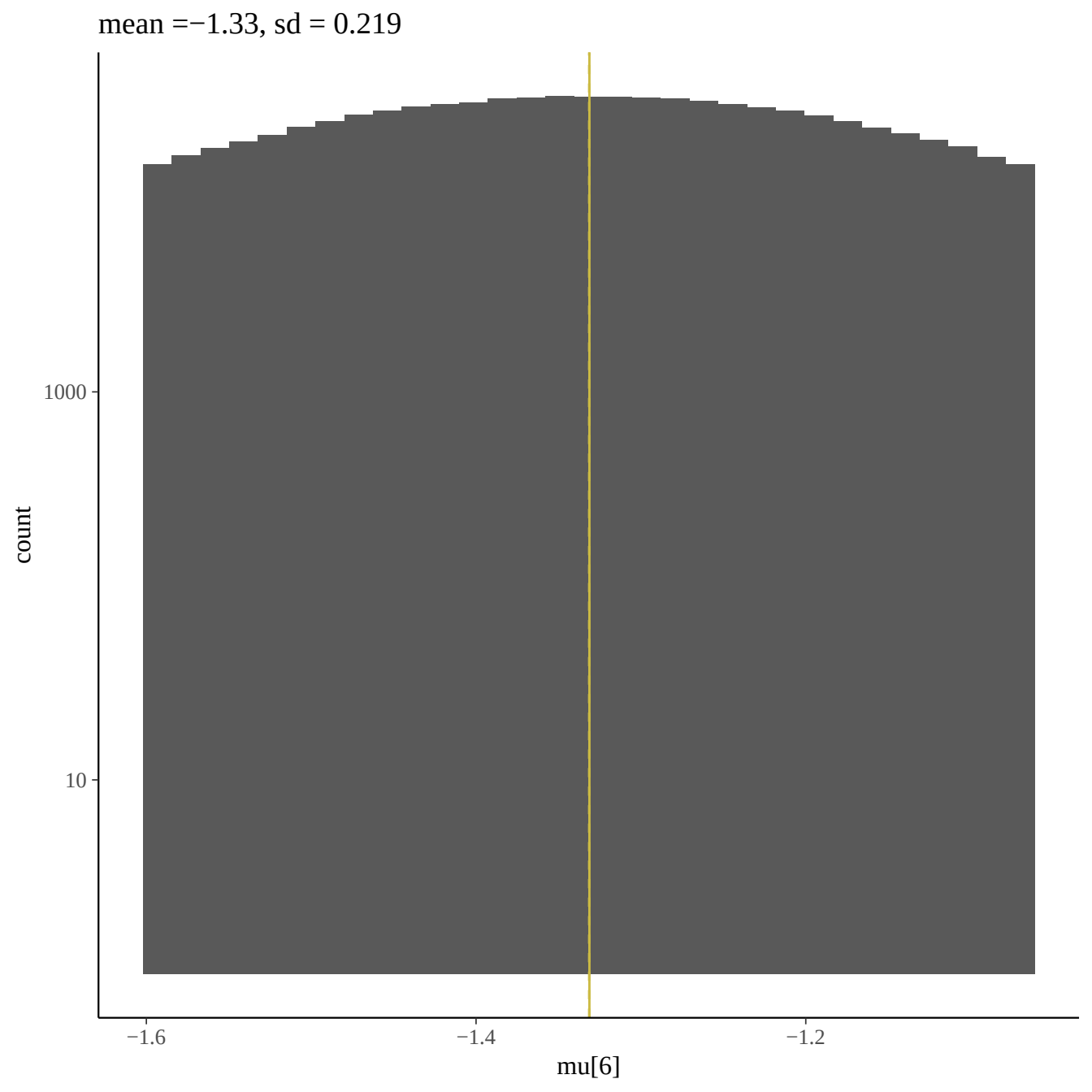

mean =−1.46, sd = 0.249
1000
10
count
−1.6
−1.4
−1.2
mu[1]

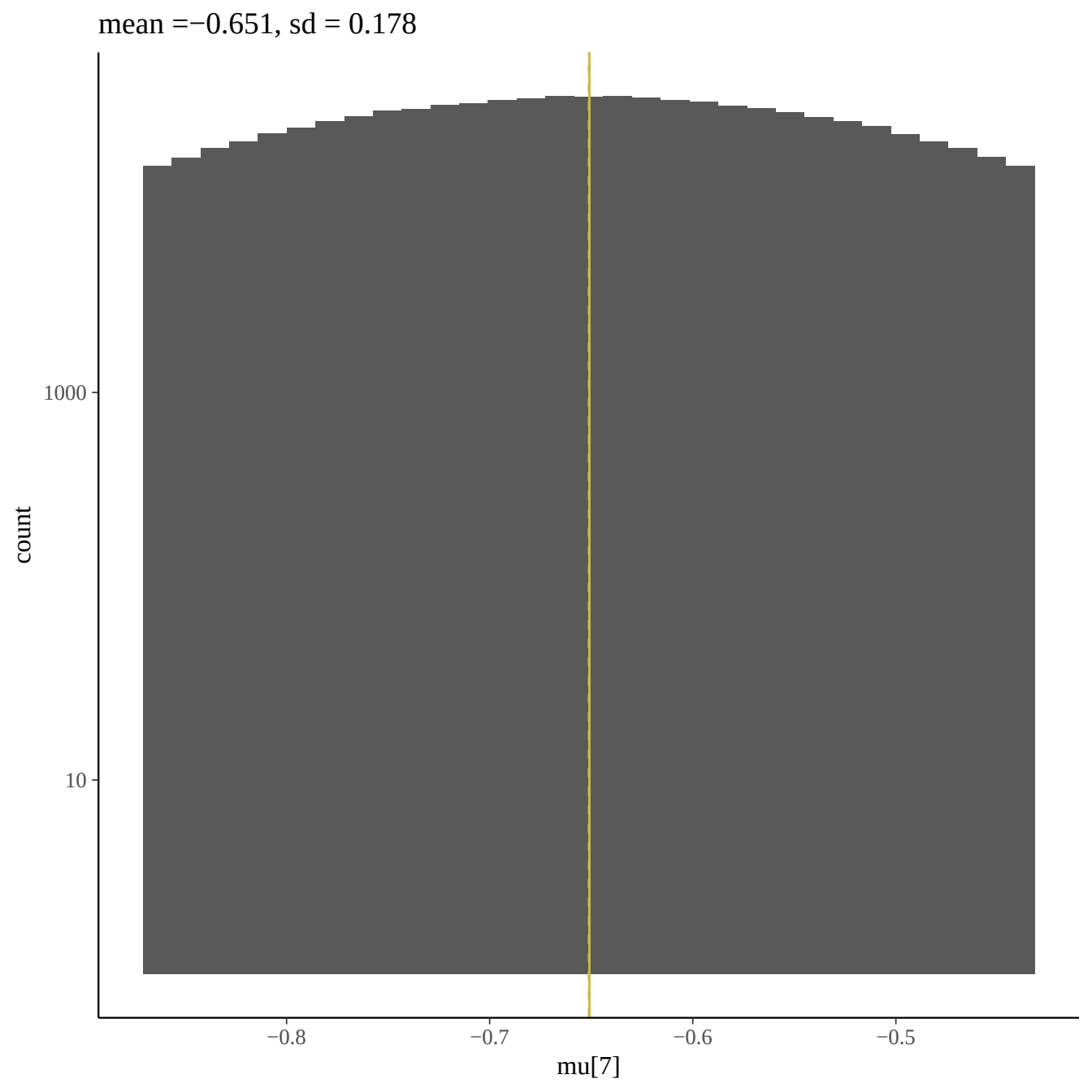

mean =−1.2, sd = 0.21
1000
10
count
−1.4
−1.3
−1.2
−1.1
−1.0
mu[2]

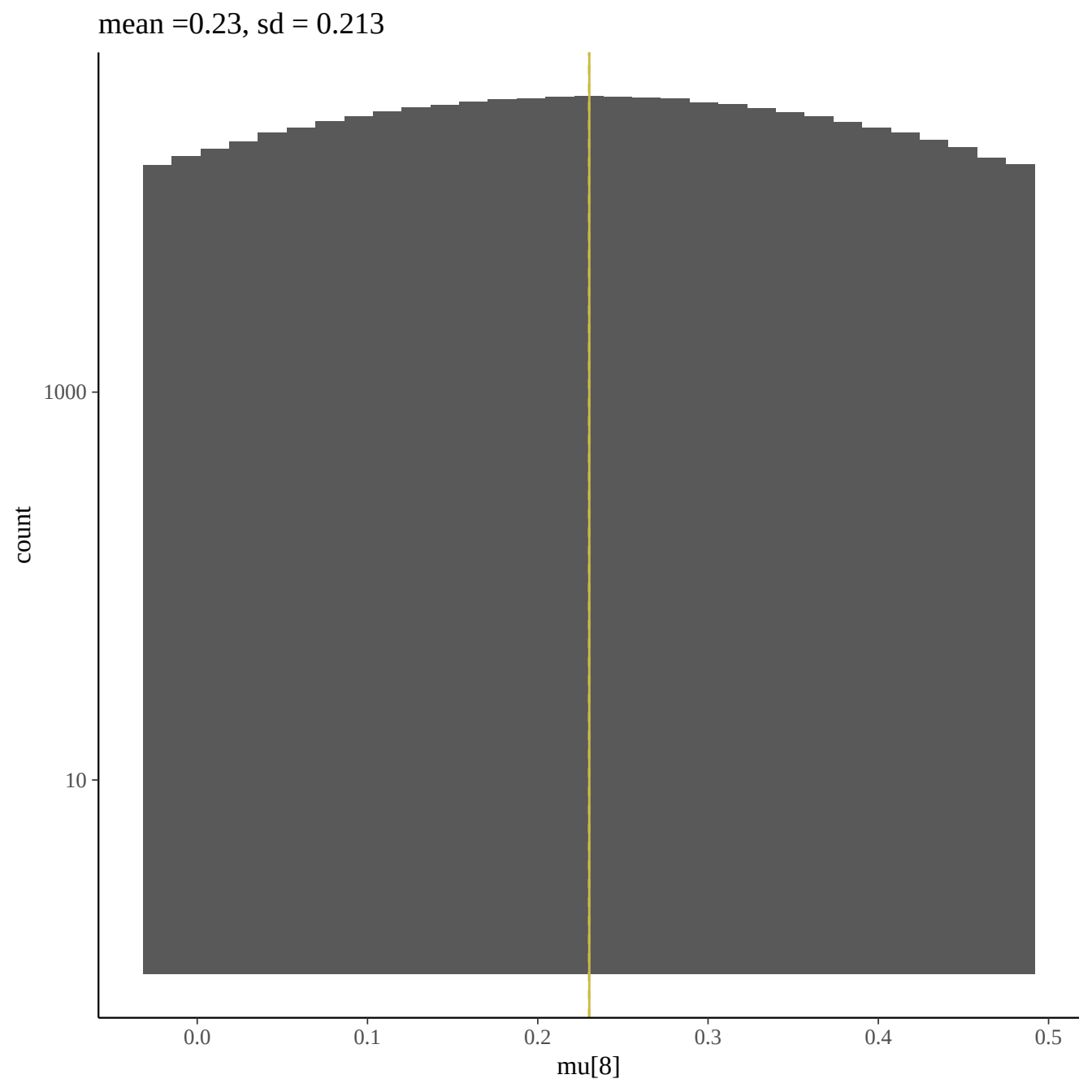

mean =1.17, sd = 0.147
1000
10
count
1.0
1.1
1.2
1.3
mu[3]

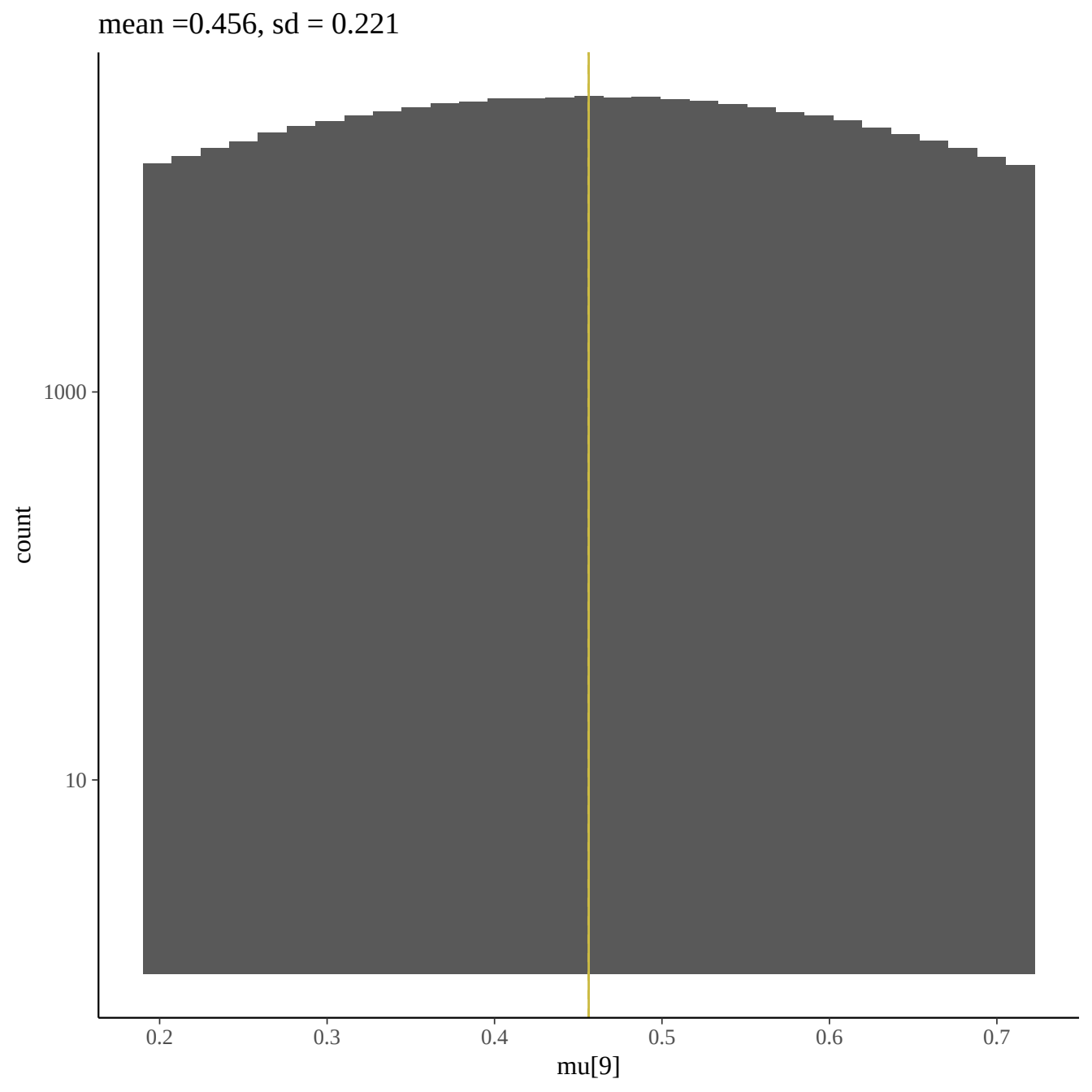

mean =−1.31, sd = 0.182
1000
10
count
−1.5
−1.4
−1.3
−1.2
−1.1
mu[4]

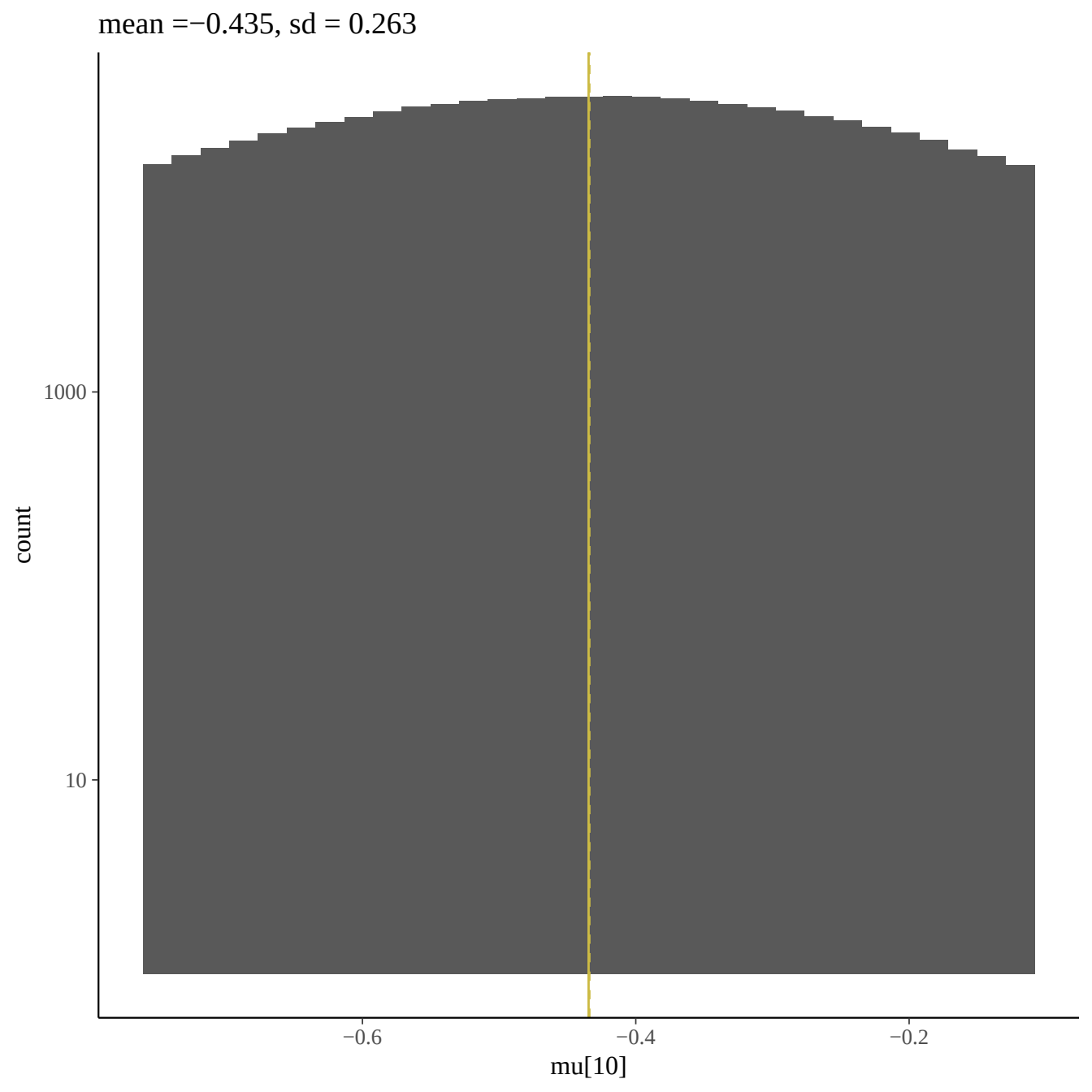

mean =−0.918, sd = 0.162
1000
10
count
−1.1
−1.0
−0.9
−0.8
−0.7
mu[5]

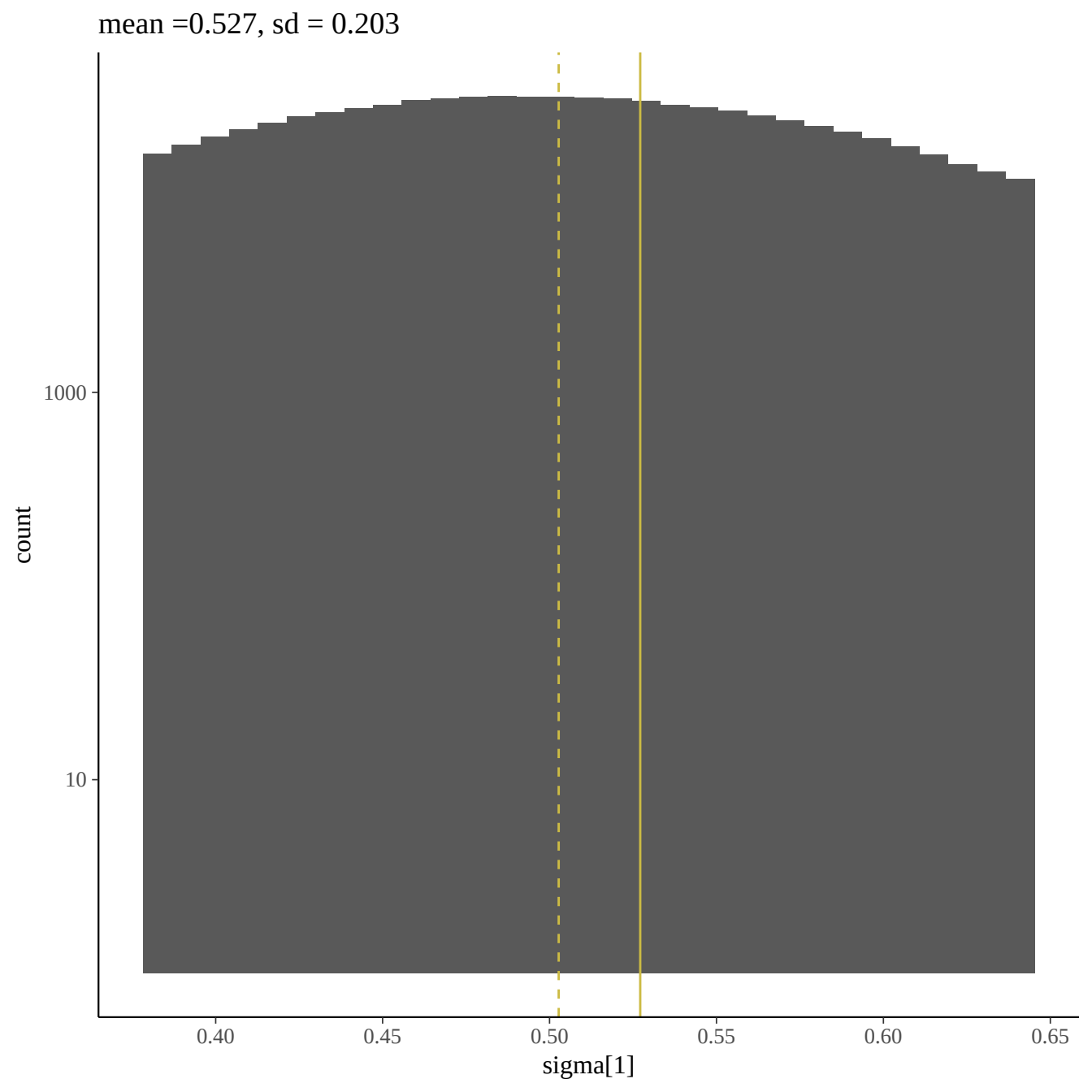

mean =−1.33, sd = 0.219
1000
10
count
−1.6
−1.4
−1.2
mu[6]

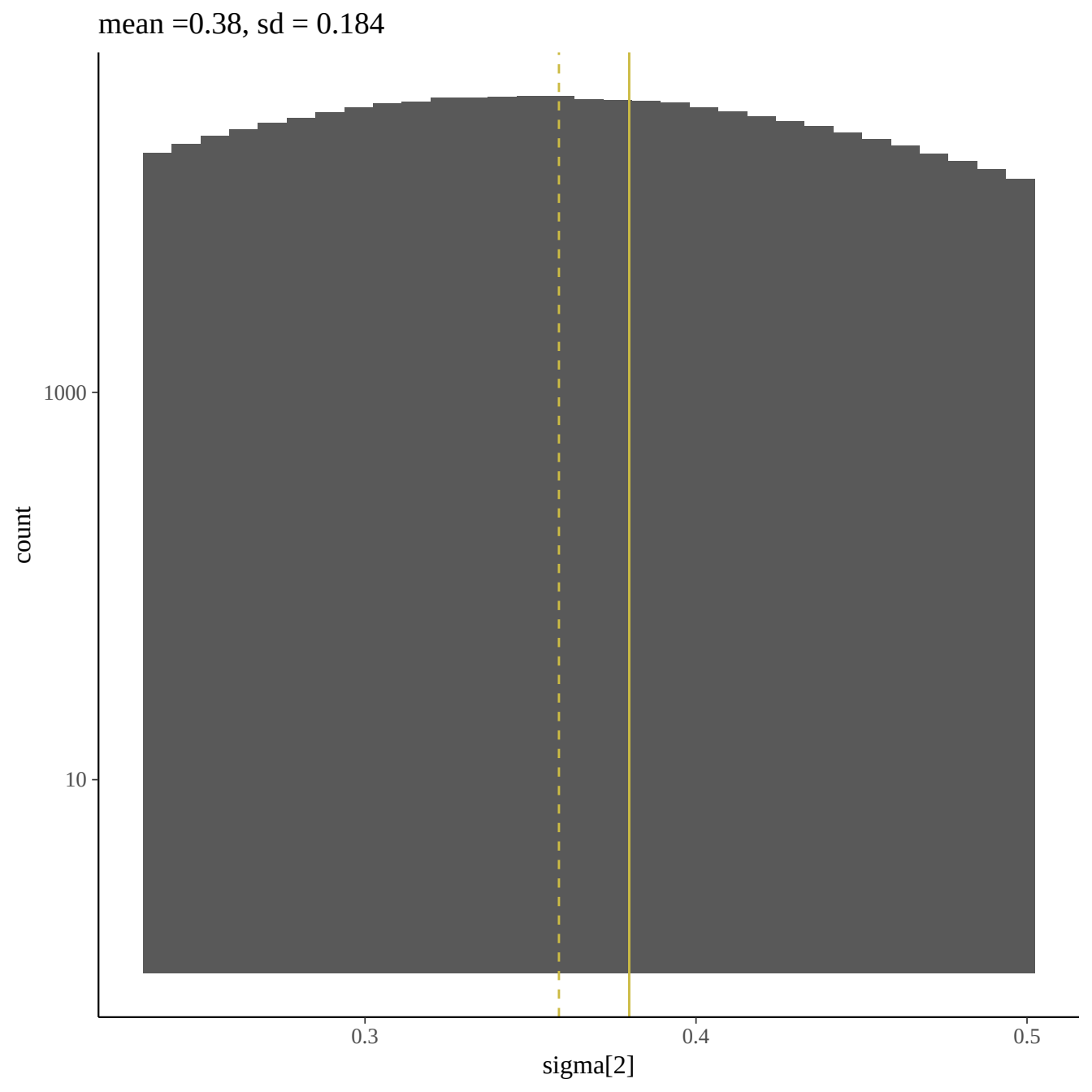

mean =−0.651, sd = 0.178
1000
10
count
−0.8
−0.7
−0.6
−0.5
mu[7]

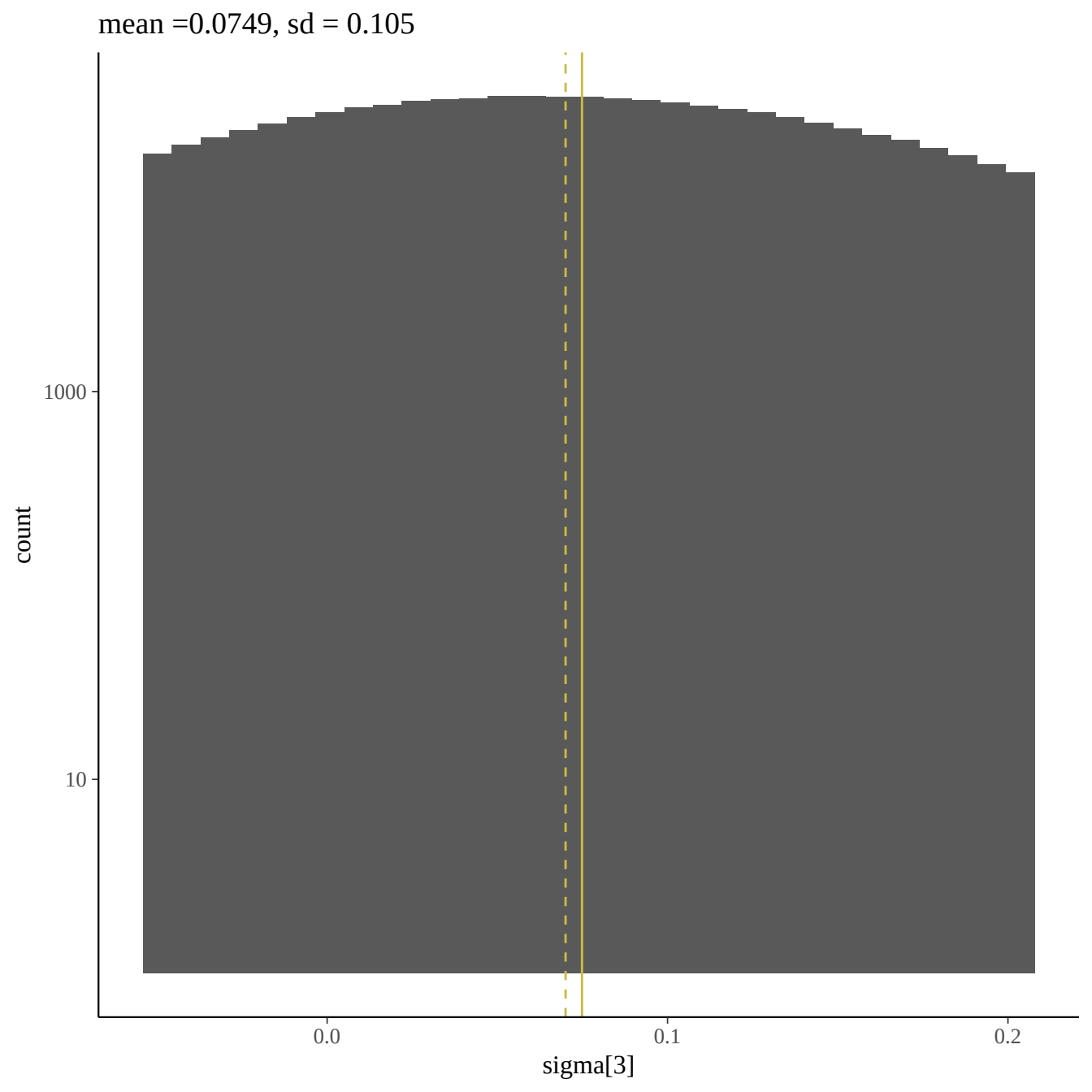

mean =0.23, sd = 0.213
1000
10
count
0.0
0.1
0.2
0.3
0.4
0.5
mu[8]

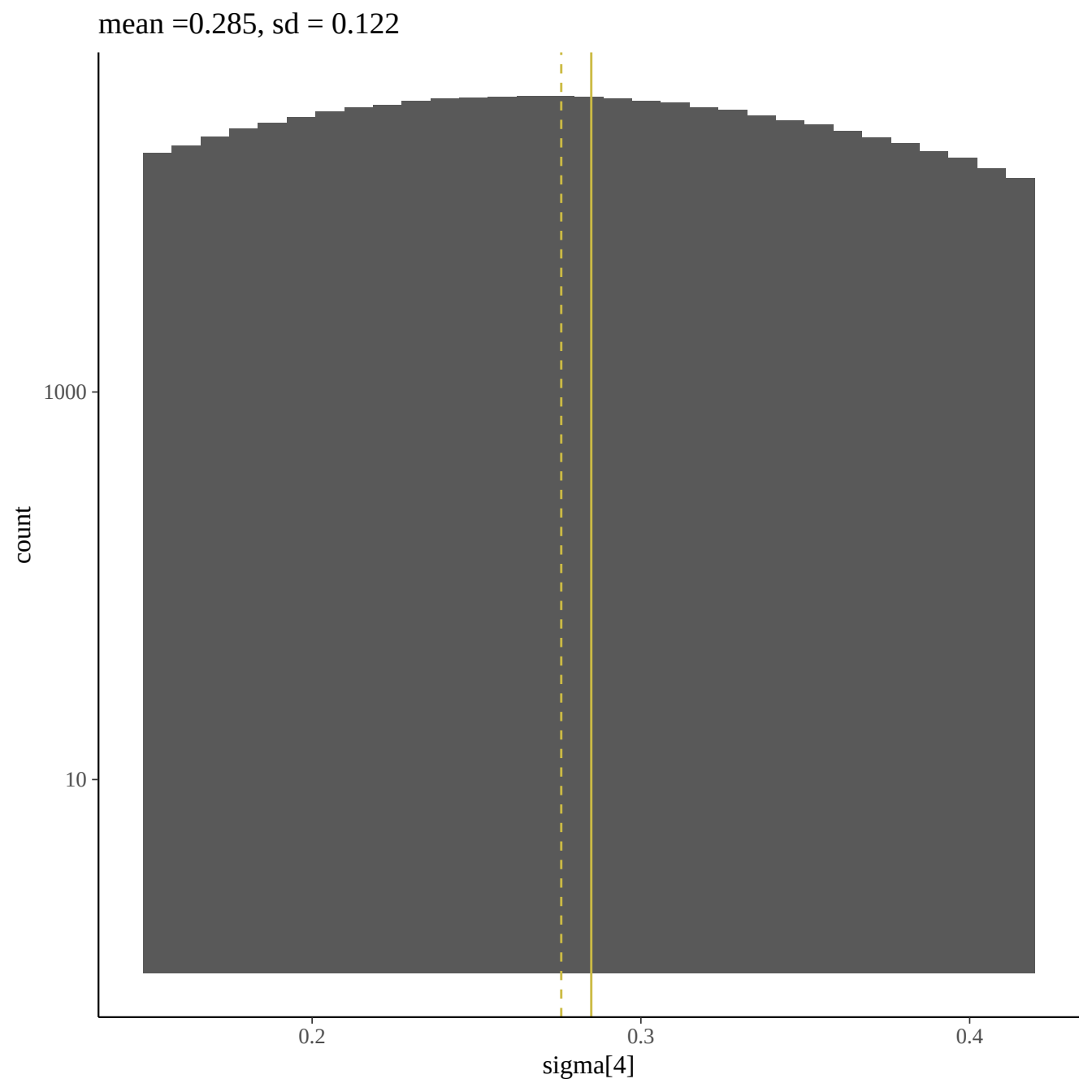

mean =0.456, sd = 0.221
count
1000
10
0.2
0.3
0.4
0.5
0.6
0.7
mu[9]

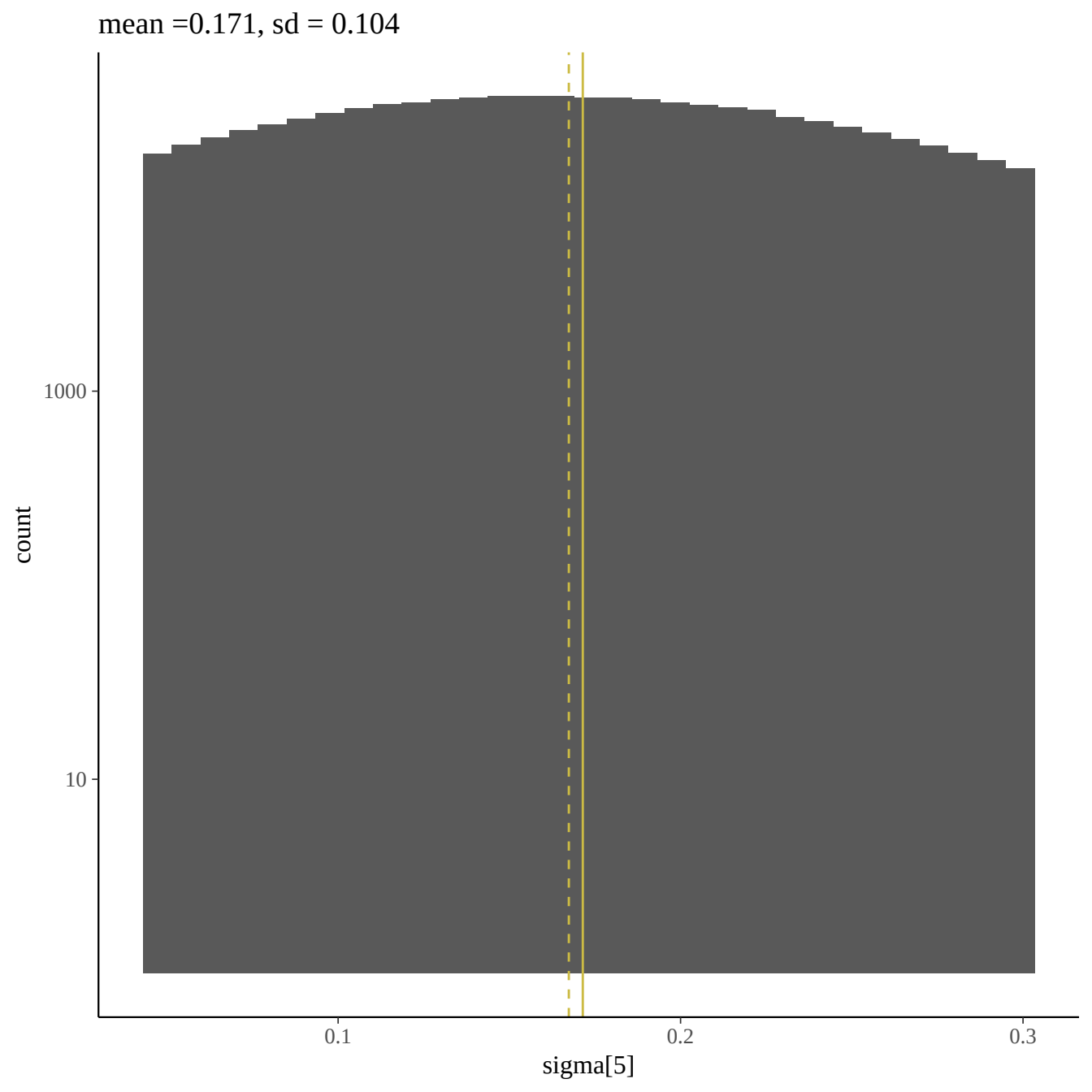

mean =−0.435, sd = 0.263
1000
10
count
−0.6
−0.4
−0.2
mu[10]

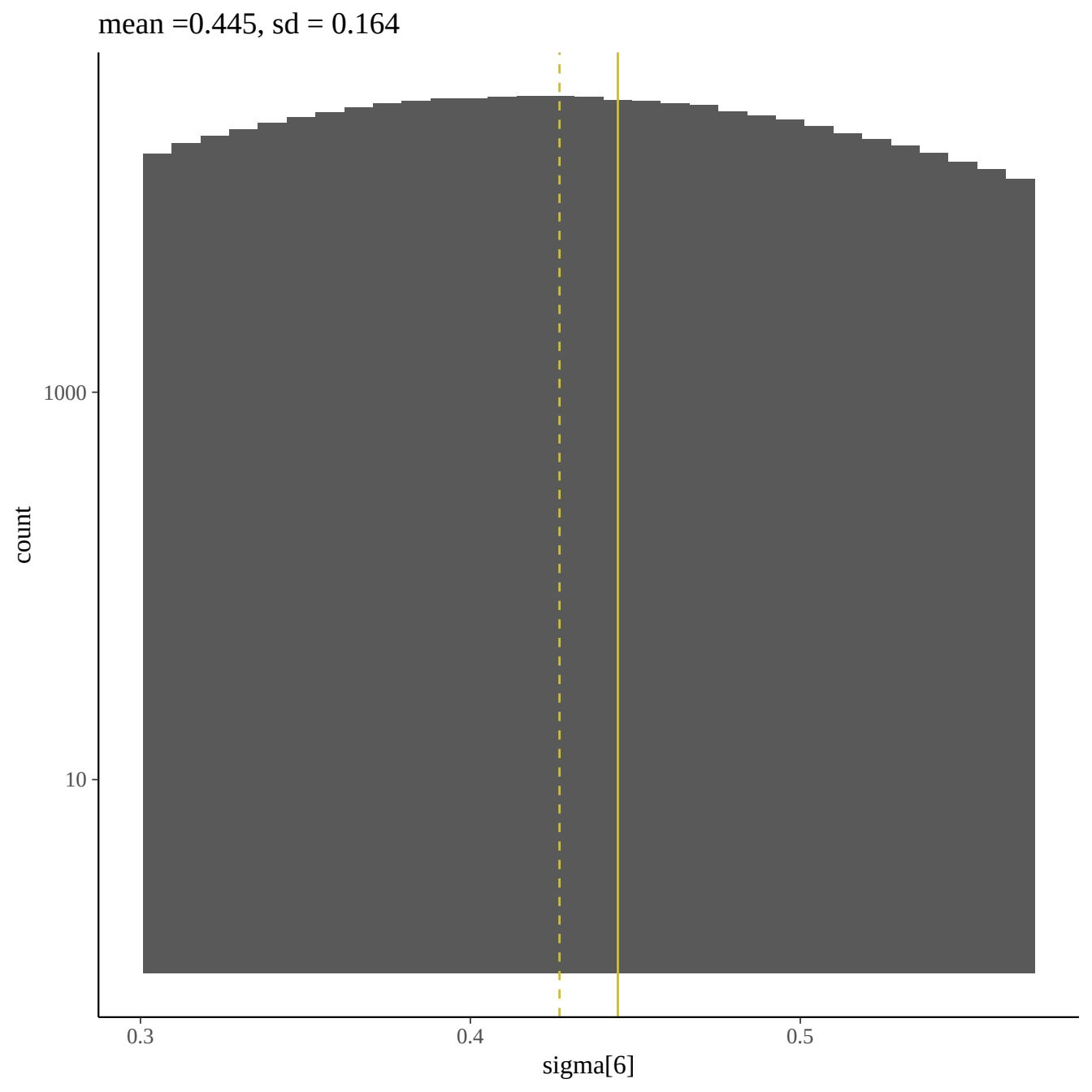

mean =0.527, sd = 0.203
1000
10
count
0.40
0.45
0.50
0.55
0.60
0.65
sigma[1]

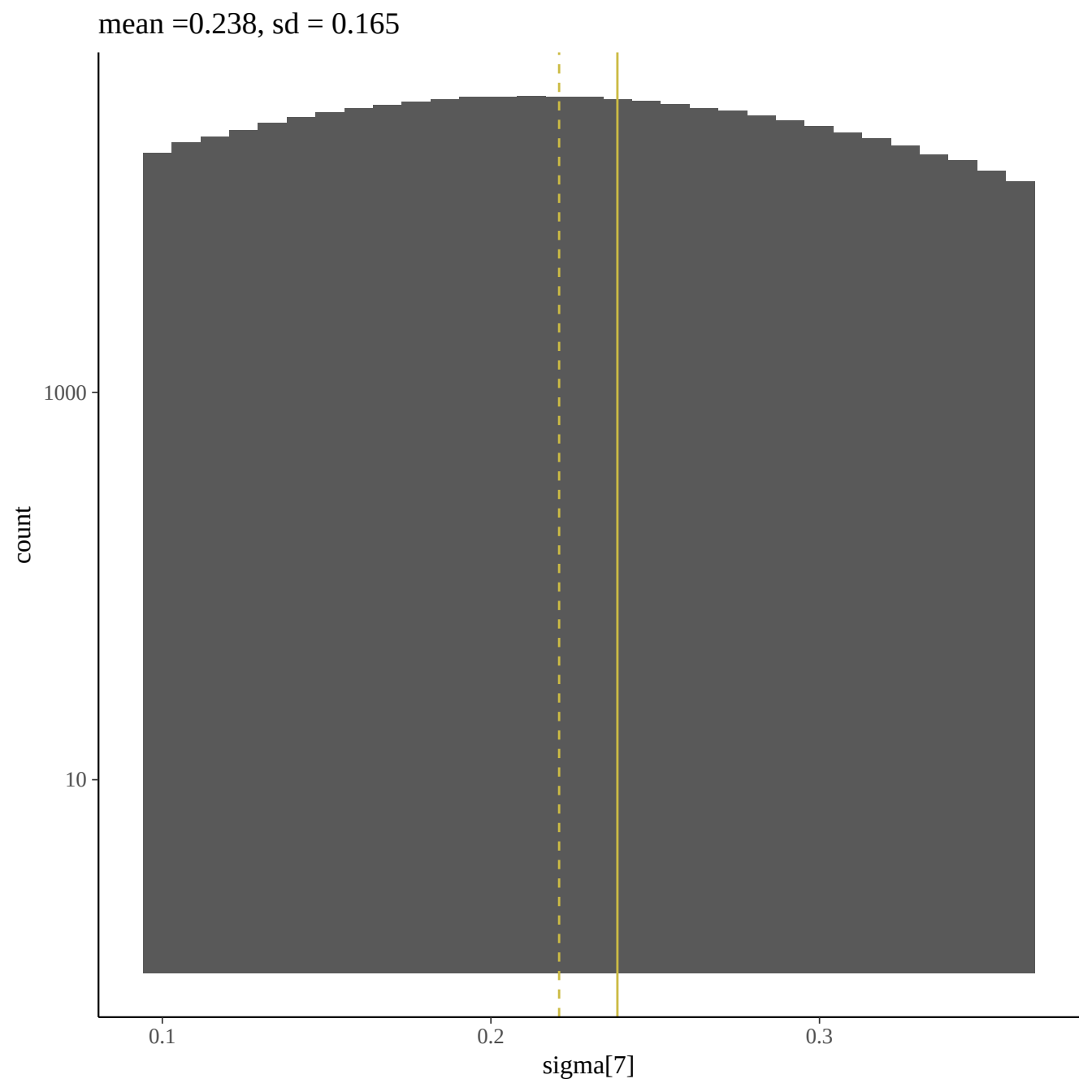

mean =0.38, sd = 0.184
1000
10
count
0.3
0.4
0.5
sigma[2]

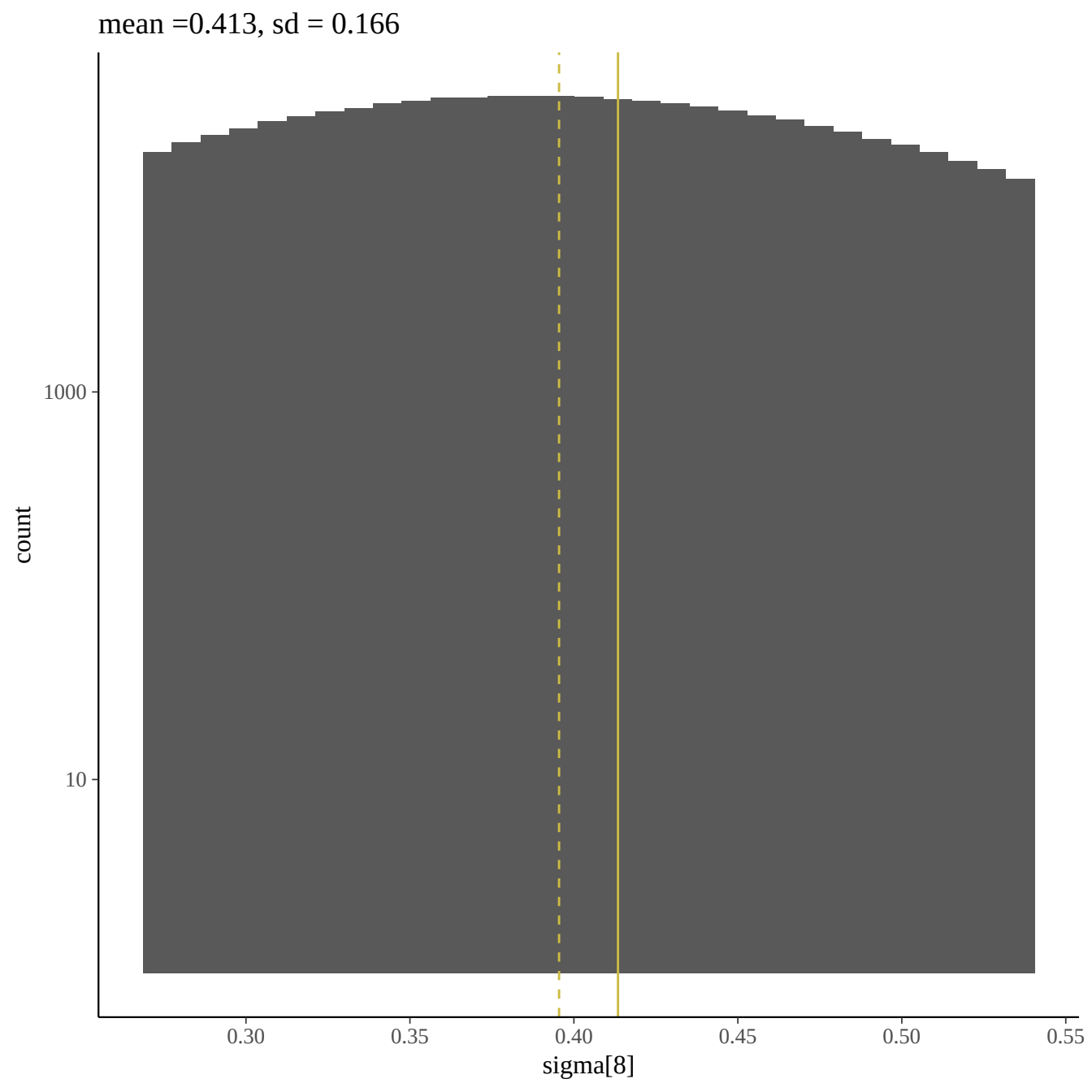

mean =0.0749, sd = 0.105
1000
10
count
0.0
0.1
0.2
sigma[3]

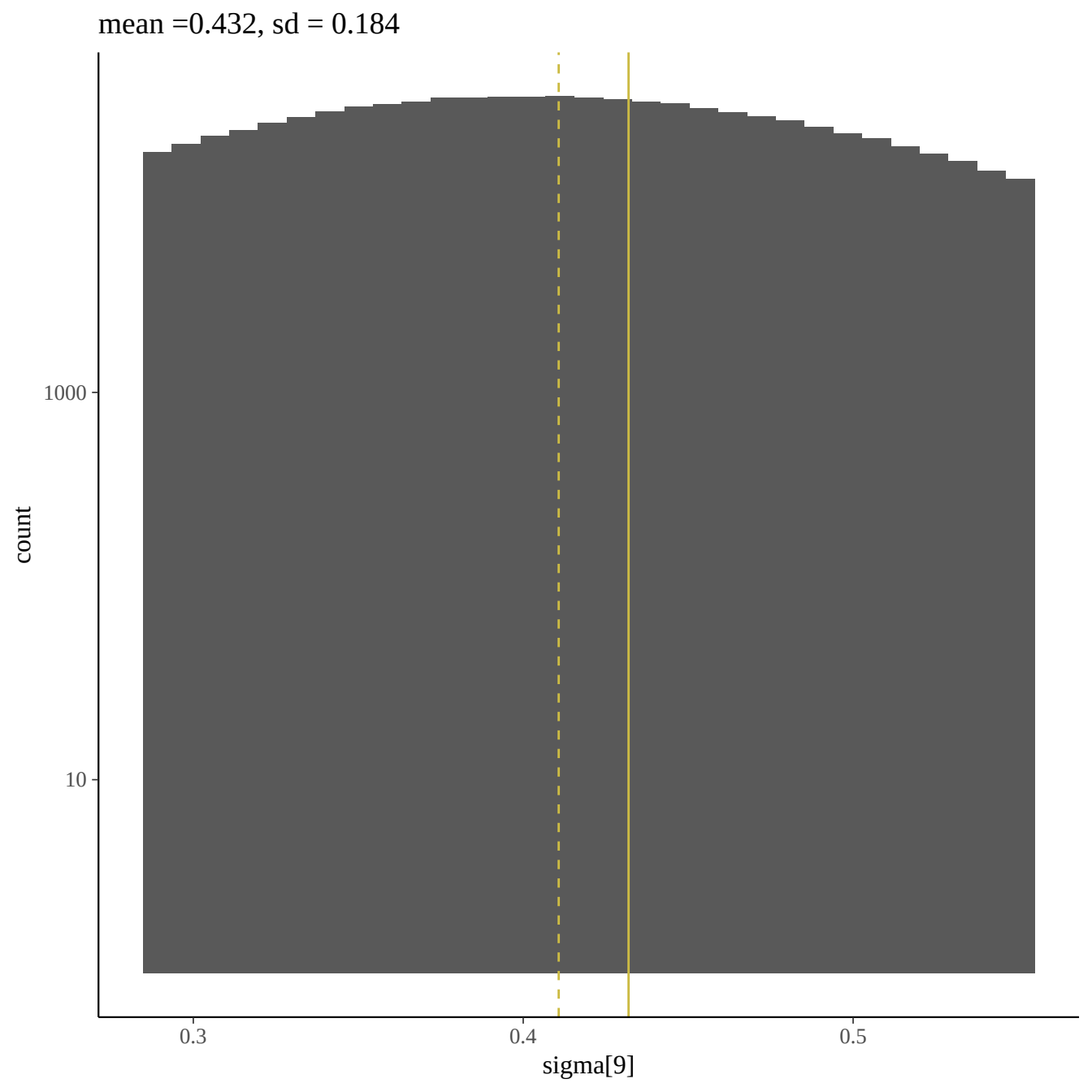

mean =0.285, sd = 0.122
1000
10
count
0.2
0.3
0.4
sigma[4]

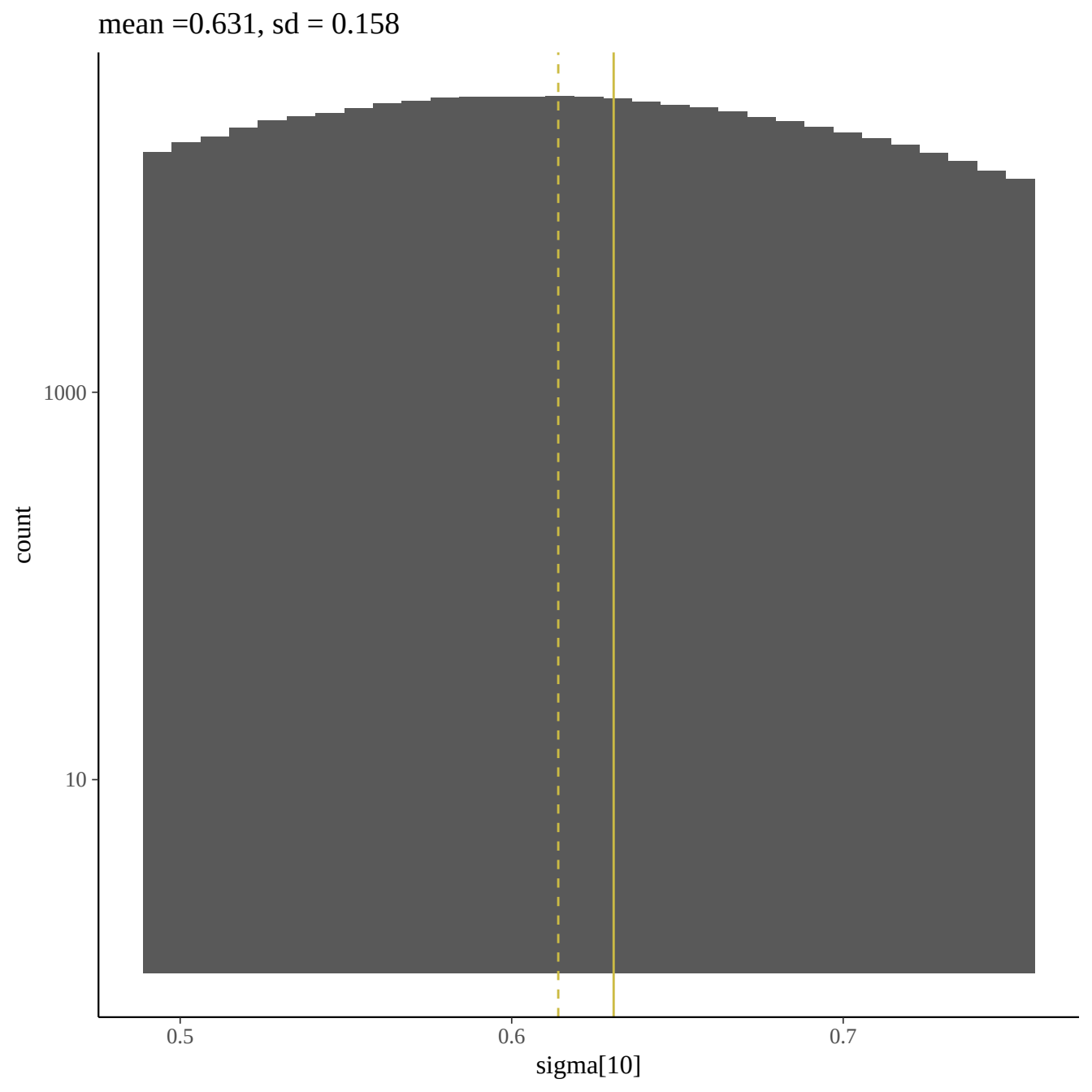

mean =0.171, sd = 0.104
1000
10
count
0.1
0.2
0.3
sigma[5]

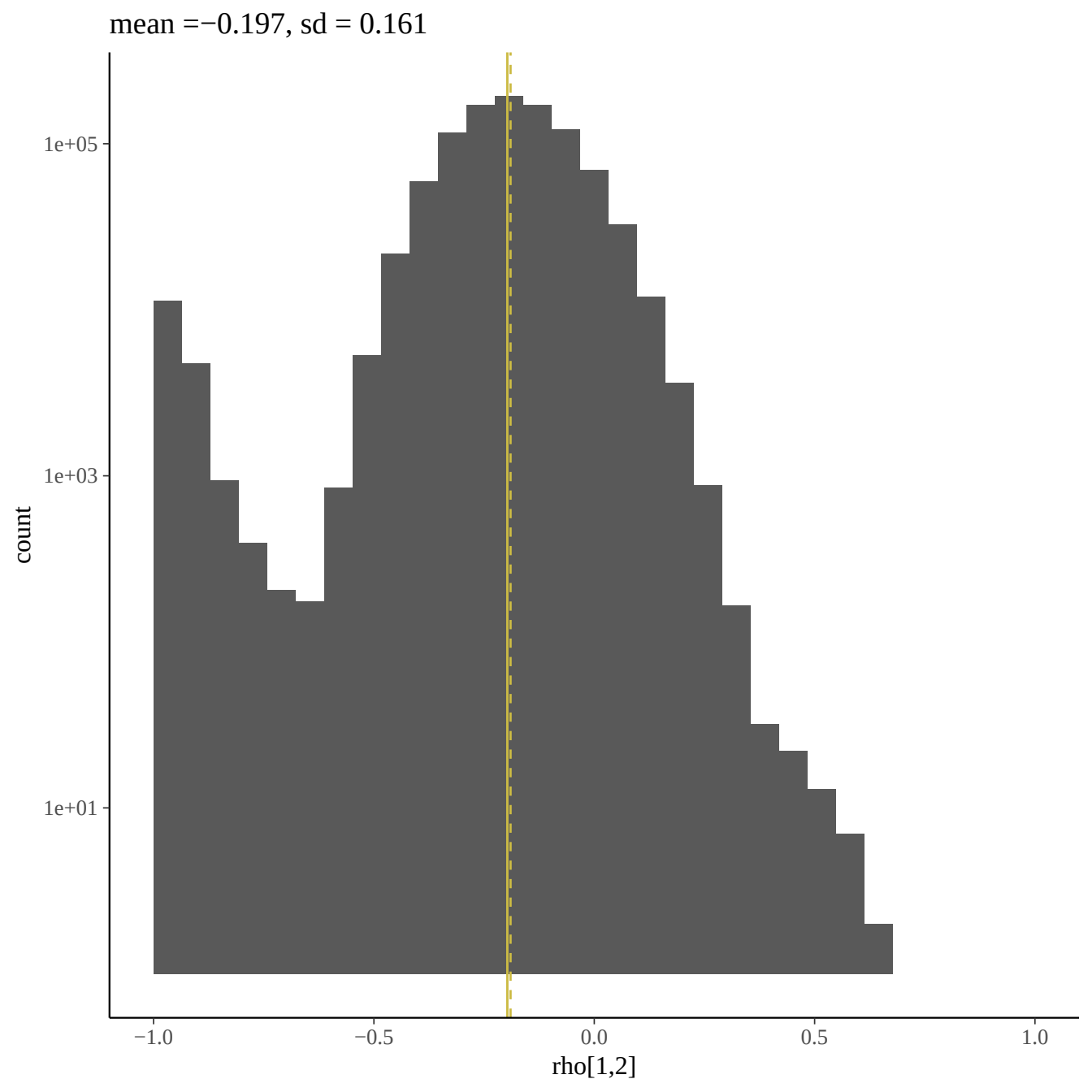

mean =0.445, sd = 0.164
1000
10
count
0.3
0.4
0.5
sigma[6]

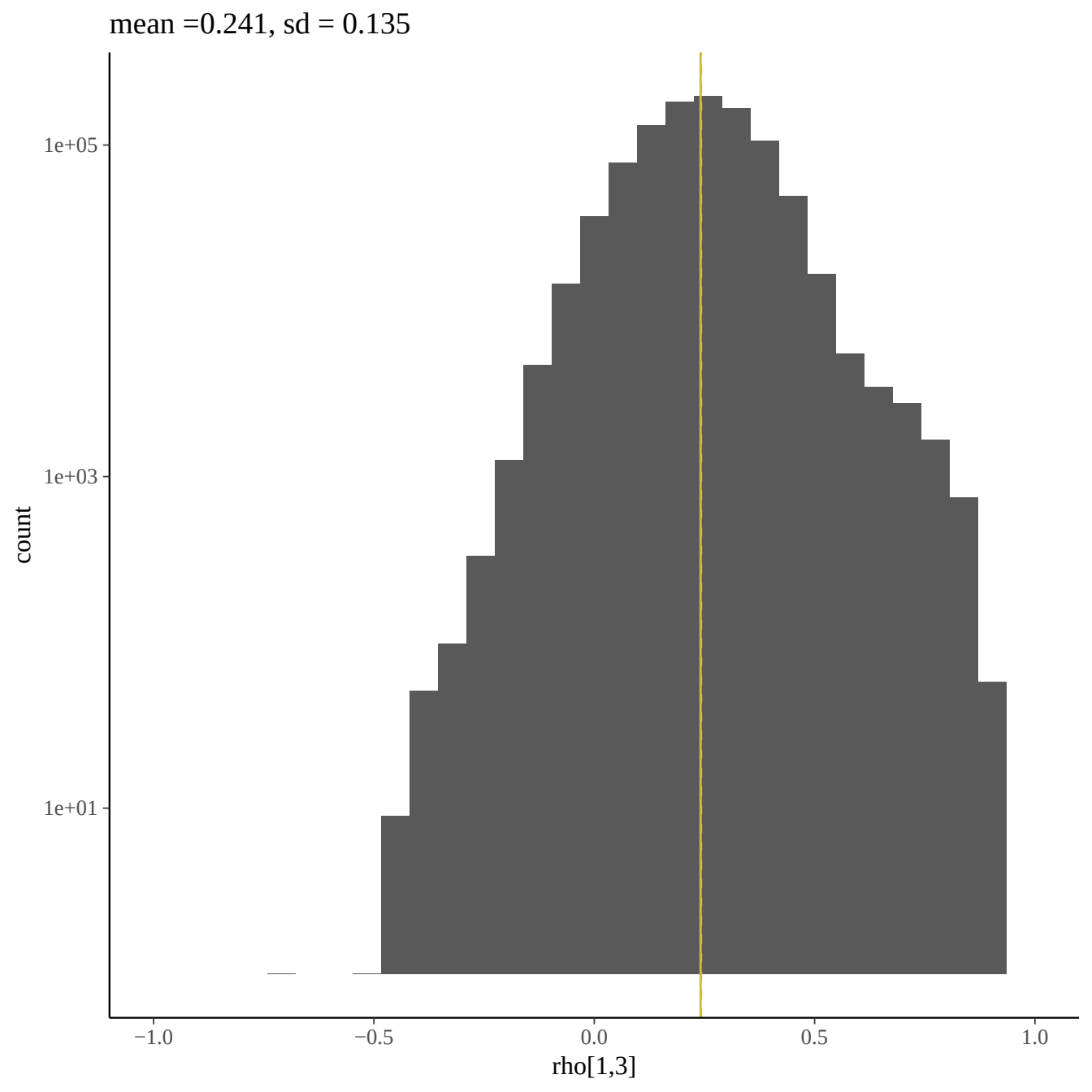

mean =0.238, sd = 0.165
1000
10
count
0.1
0.2
0.3
sigma[7]

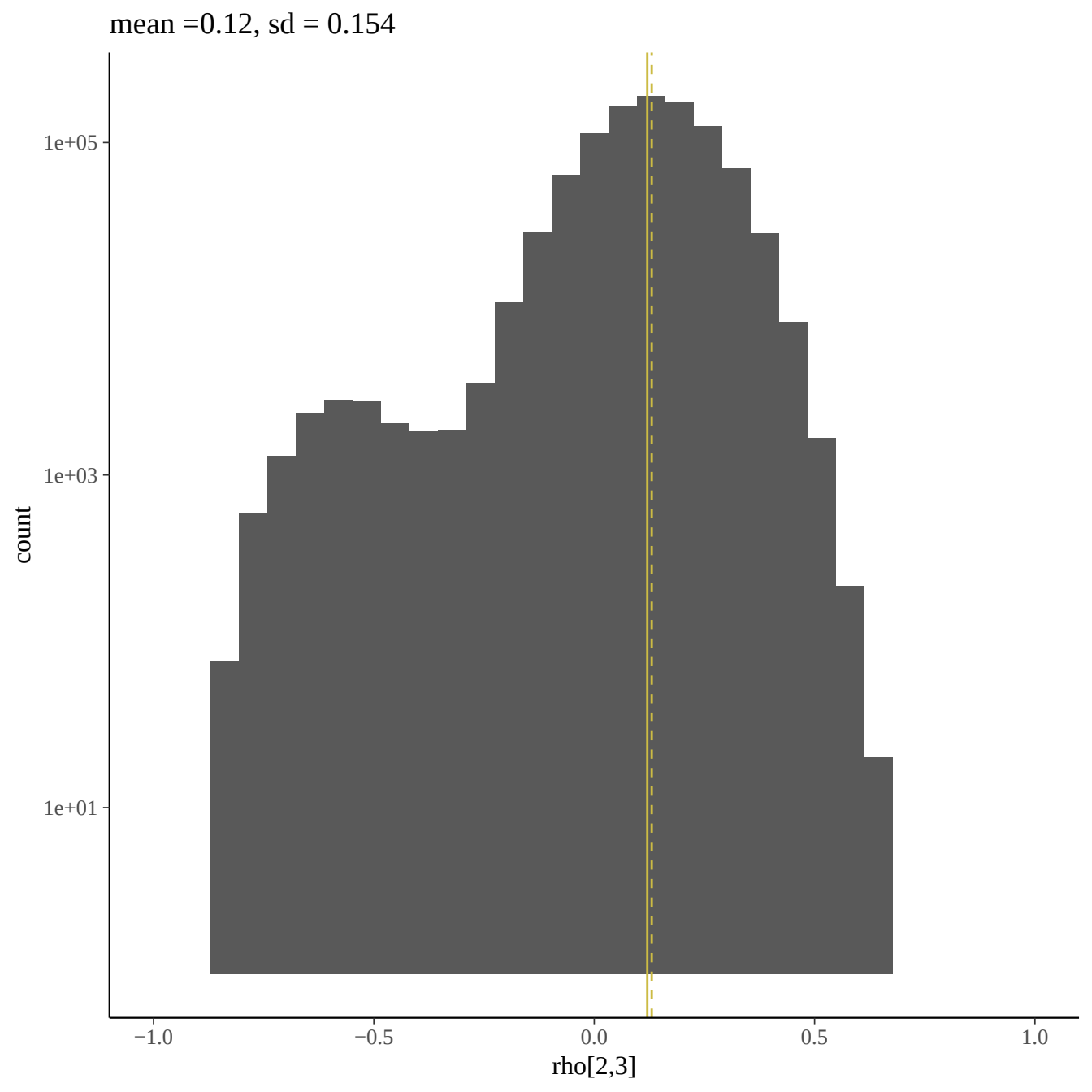

mean =0.413, sd = 0.166
count
1000
10
0.30
0.35
0.40
0.45
0.50
0.55
sigma[8]

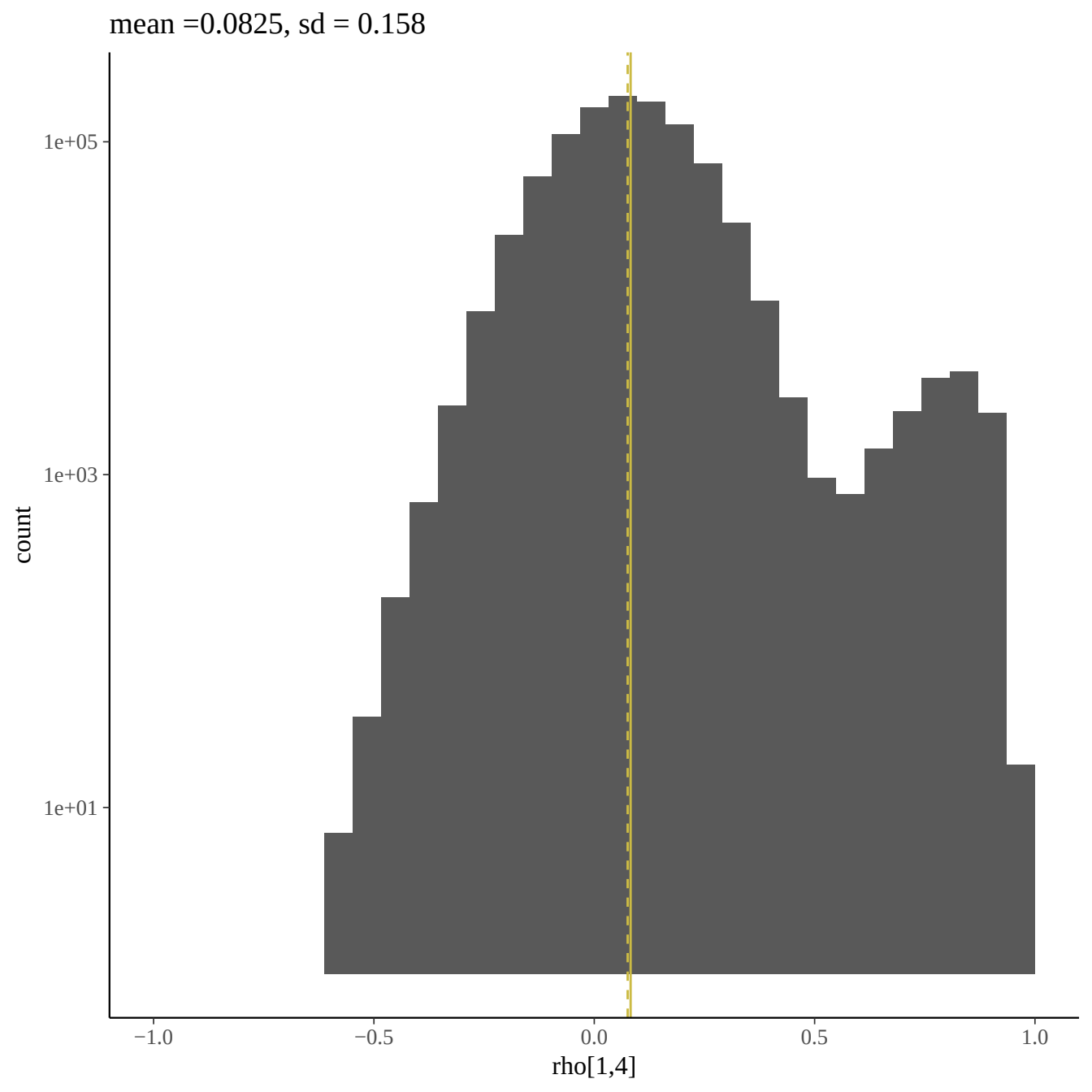

mean =0.432, sd = 0.184
1000
10
count
0.3
0.4
0.5
sigma[9]

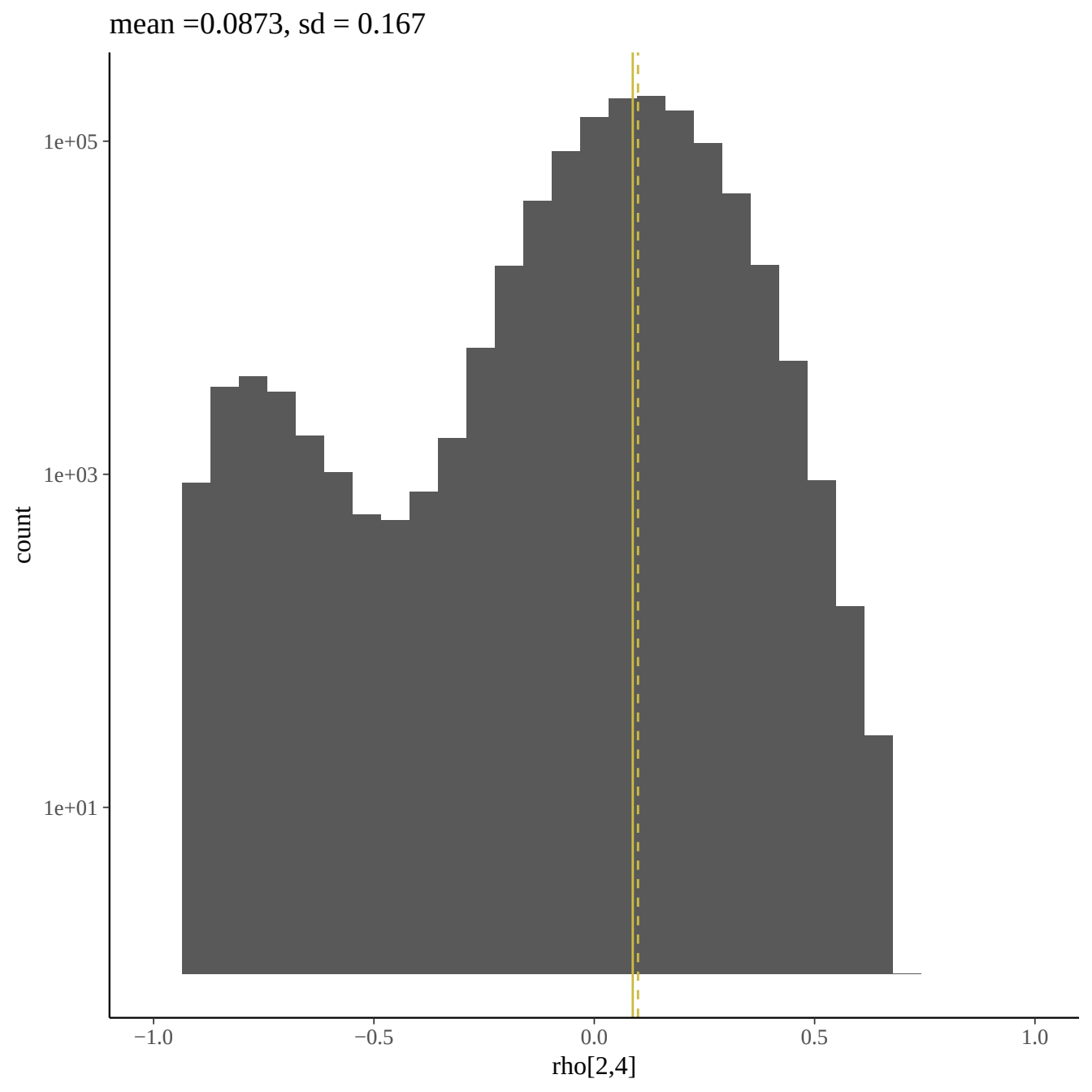

mean =0.631, sd = 0.158
1000
10
count
0.5
0.6
0.7
sigma[10]

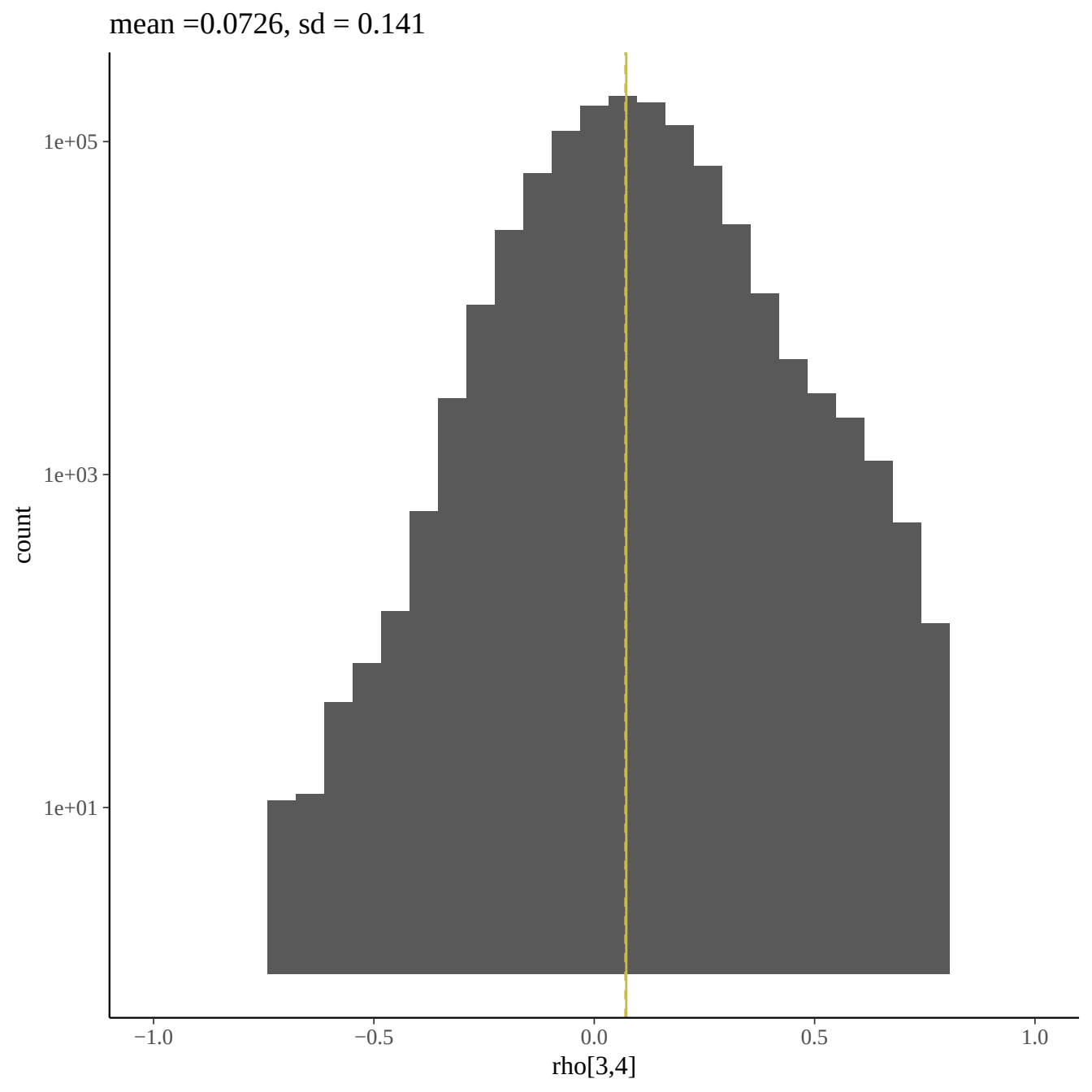

mean =−0.197, sd = 0.161
1e+05
1e+03
1e+01
count
−1.0
−0.5
0.0
0.5
1.0
rho[1,2]

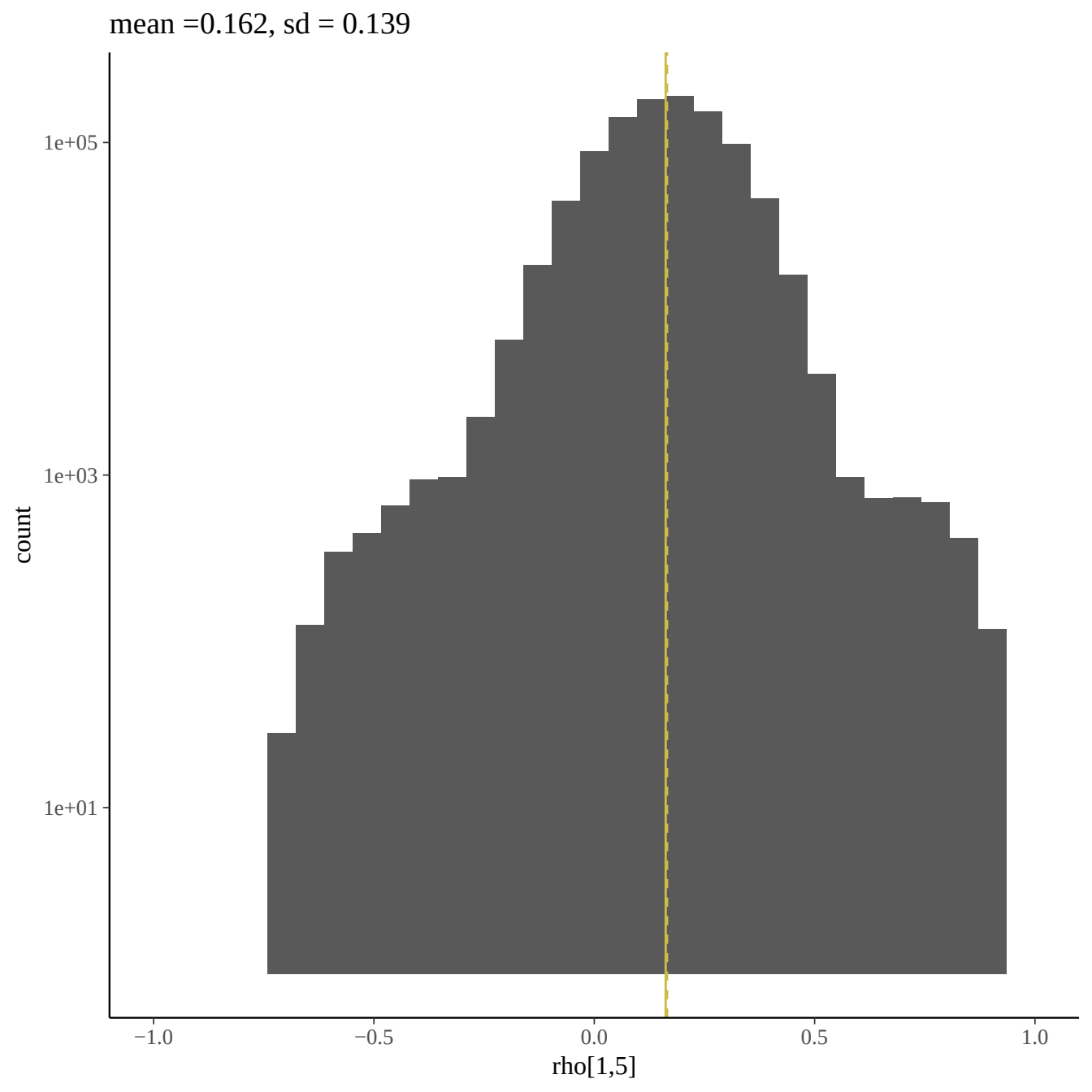

mean =0.241, sd = 0.135
1e+05
1e+03
1e+01
count
−1.0
−0.5
0.0
0.5
1.0
rho[1,3]

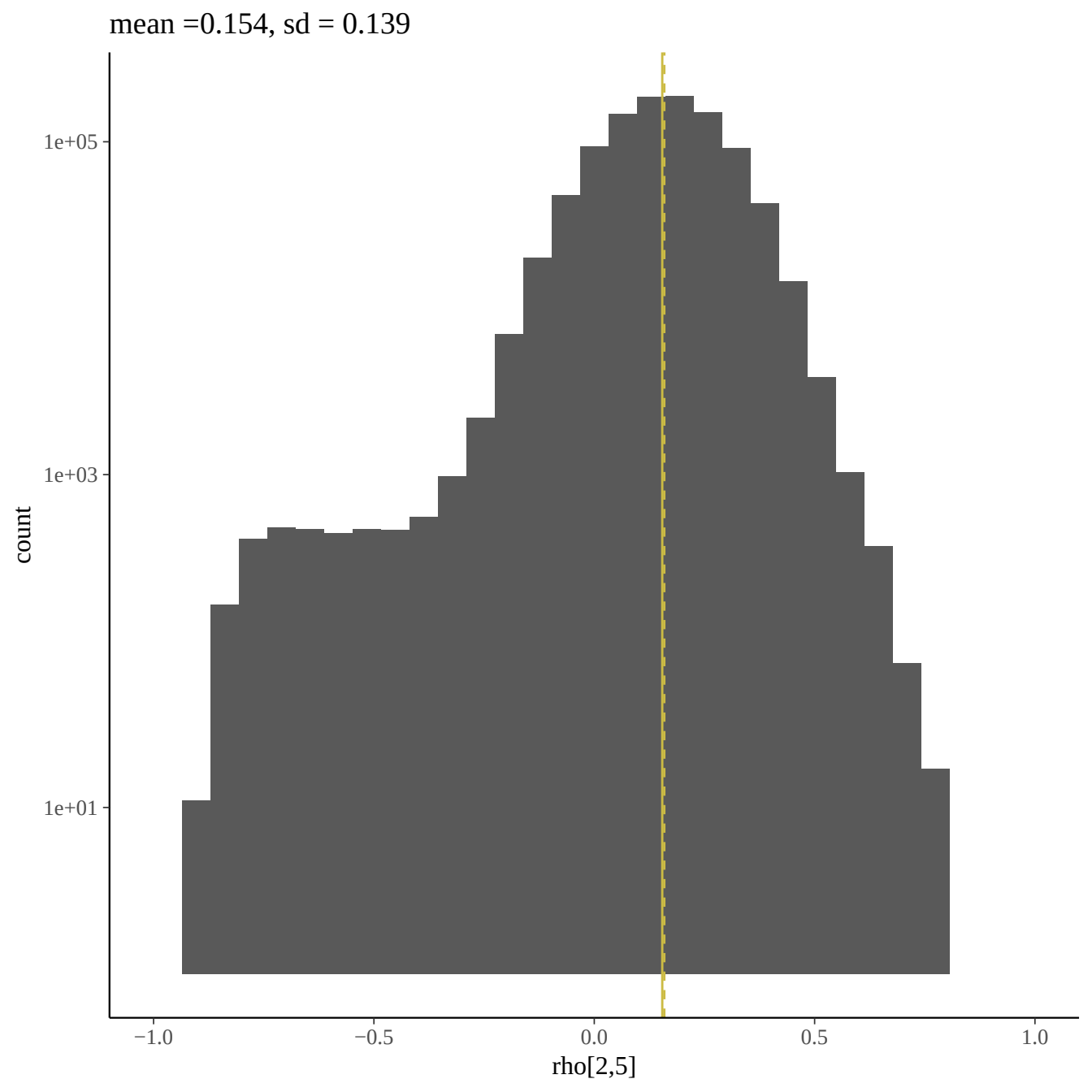

mean =0.12, sd = 0.154
1e+05
1e+03
1e+01
count
−1.0
−0.5
0.0
0.5
1.0
rho[2,3]

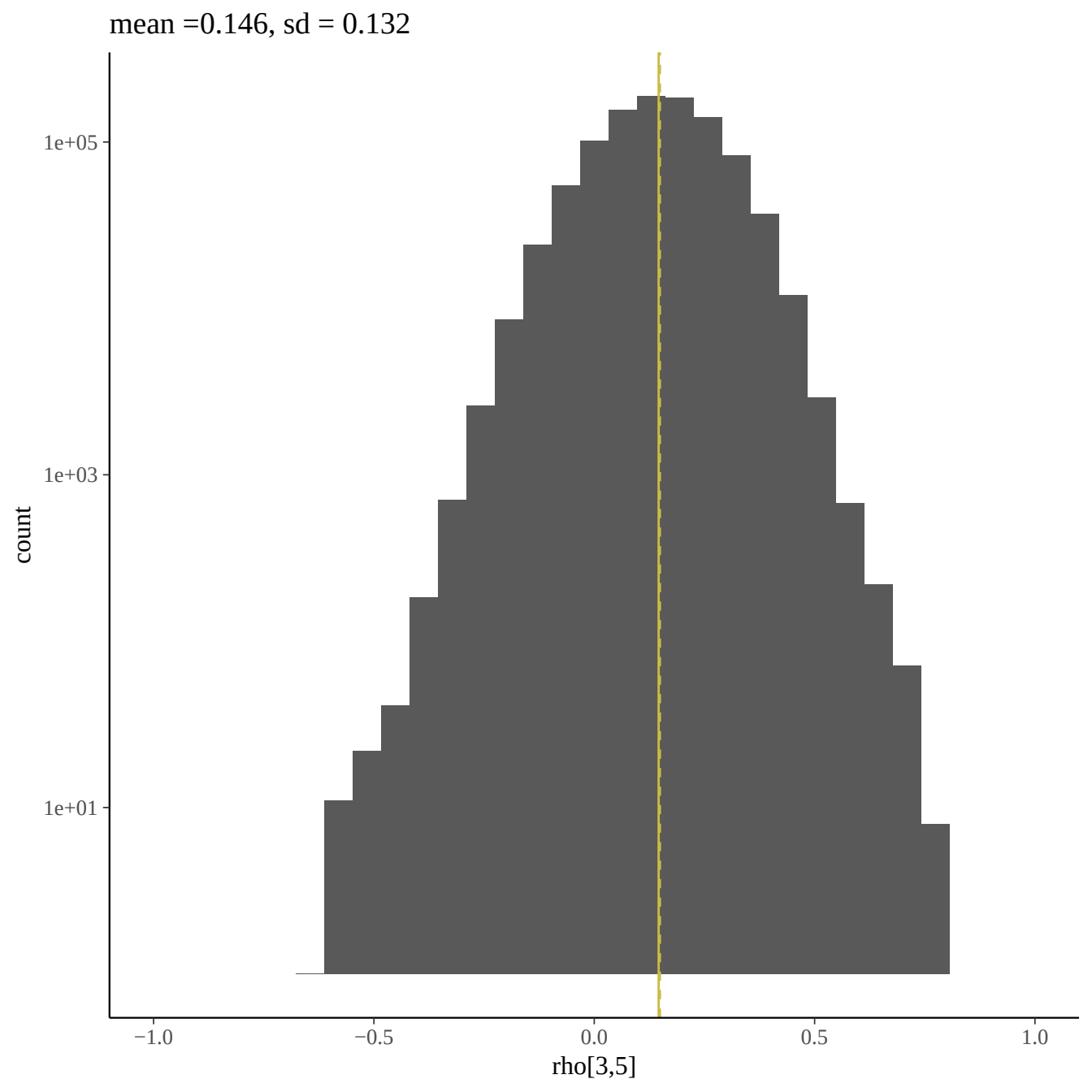

mean =0.0825, sd = 0.158
1e+05
1e+03
1e+01
count
−1.0
−0.5
0.0
0.5
1.0
rho[1,4]

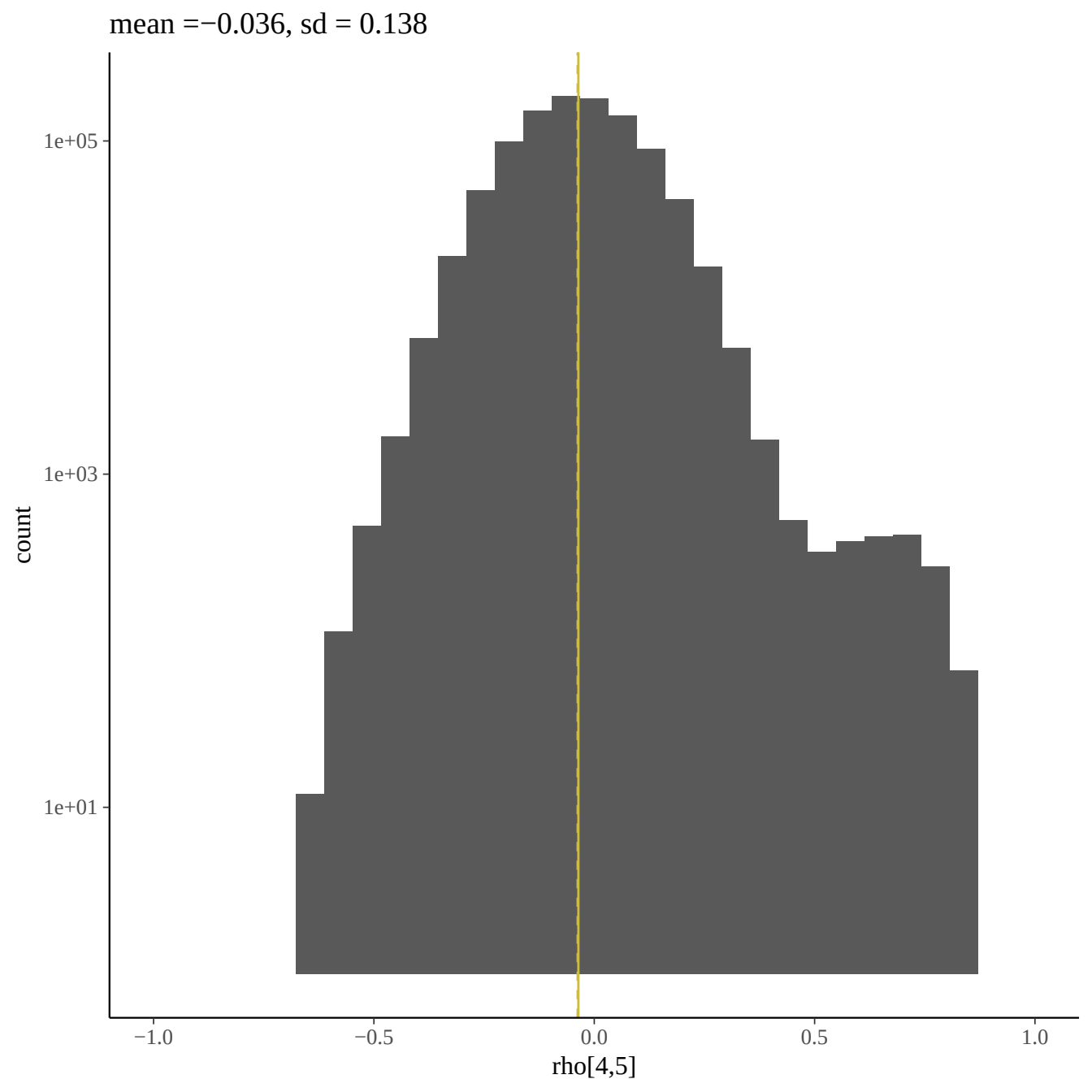

mean =0.0873, sd = 0.167
1e+05
1e+03
1e+01
count
−1.0
−0.5
0.0
0.5
1.0
rho[2,4]

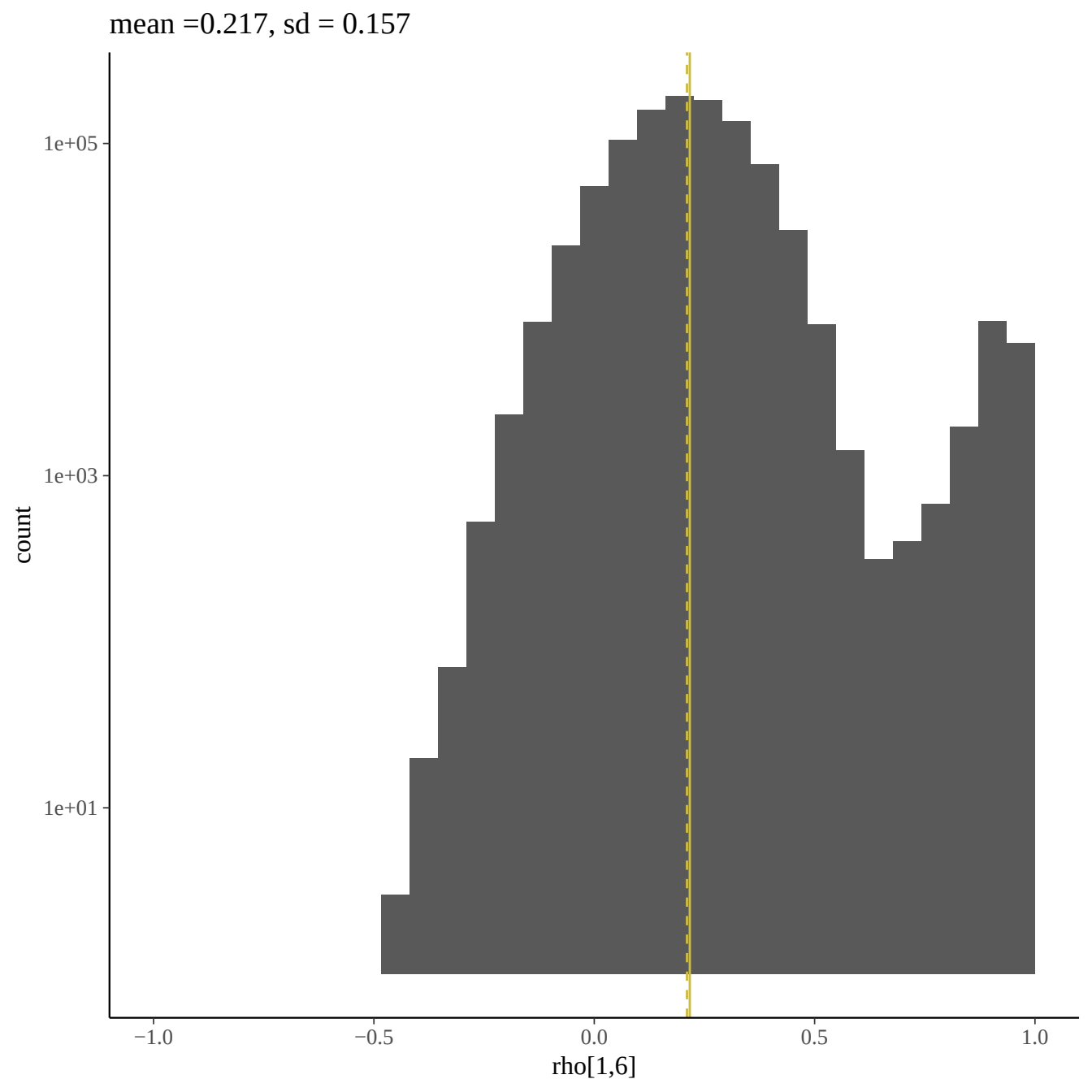

mean =0.0726, sd = 0.141
1e+05
1e+03
1e+01
count
−1.0
−0.5
0.0
0.5
1.0
rho[3,4]

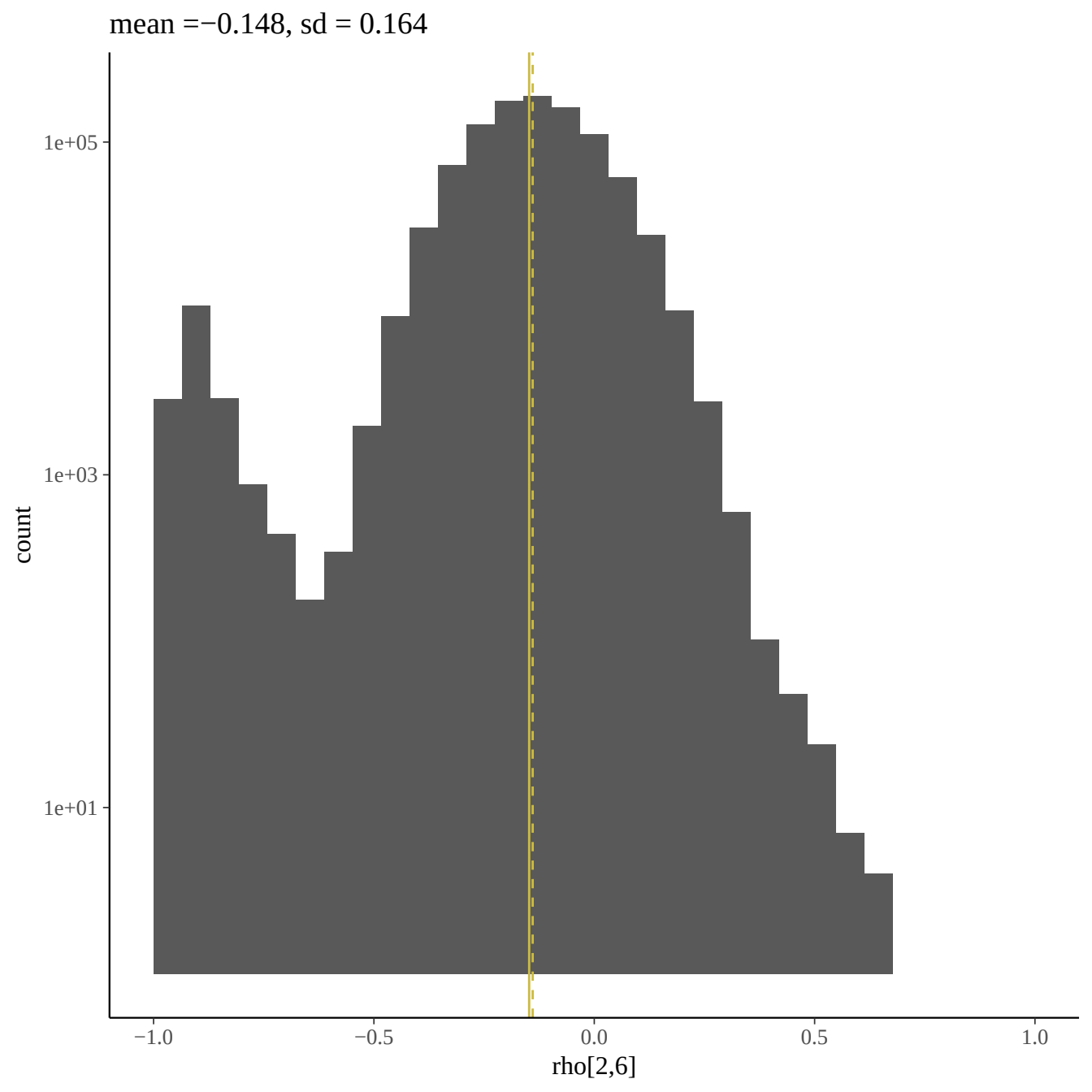

mean =0.162, sd = 0.139
1e+05
1e+03
1e+01
count
−1.0
−0.5
0.0
0.5
1.0
rho[1,5]

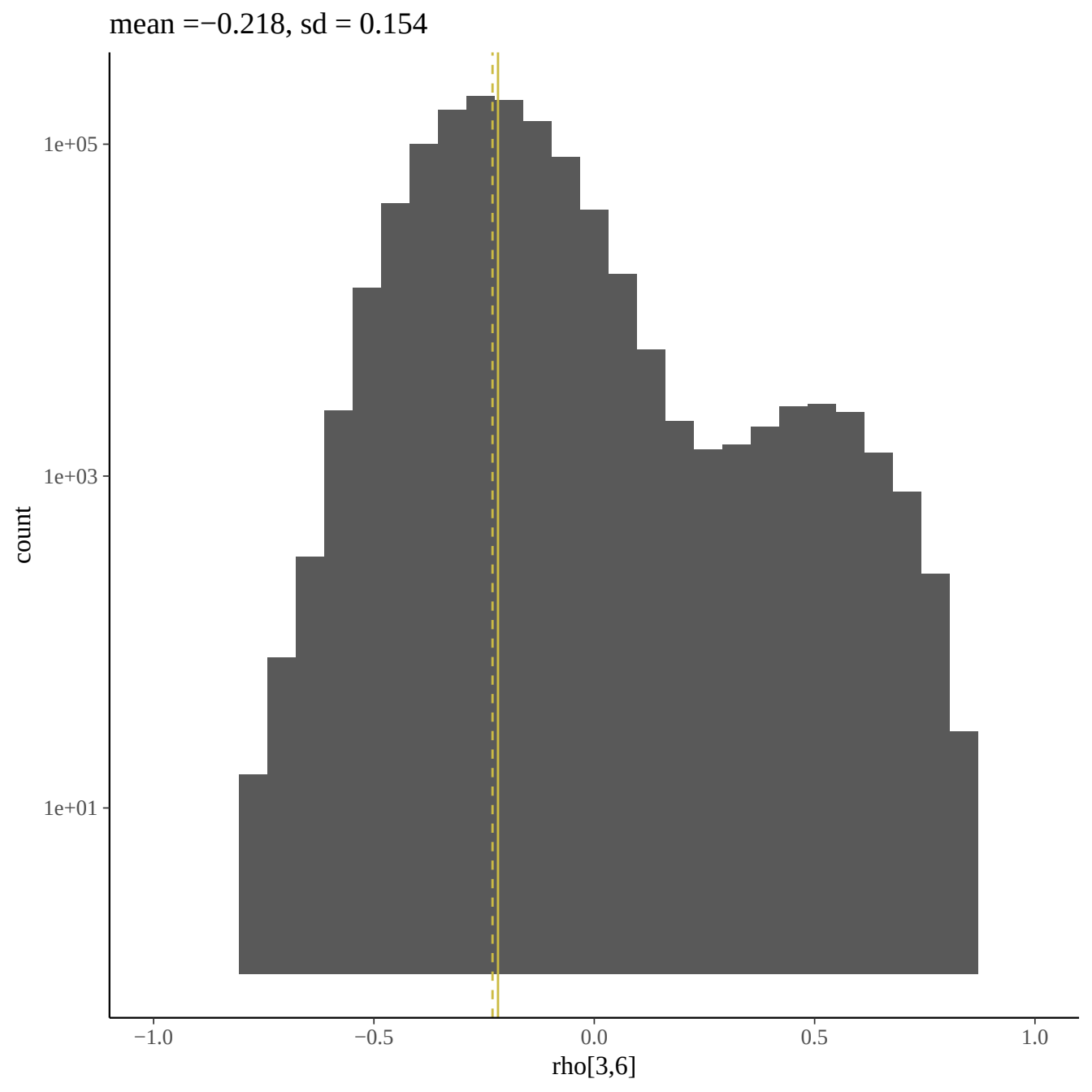

mean =0.154, sd = 0.139
1e+05
1e+03
1e+01
count
−1.0
−0.5
0.0
0.5
1.0
rho[2,5]

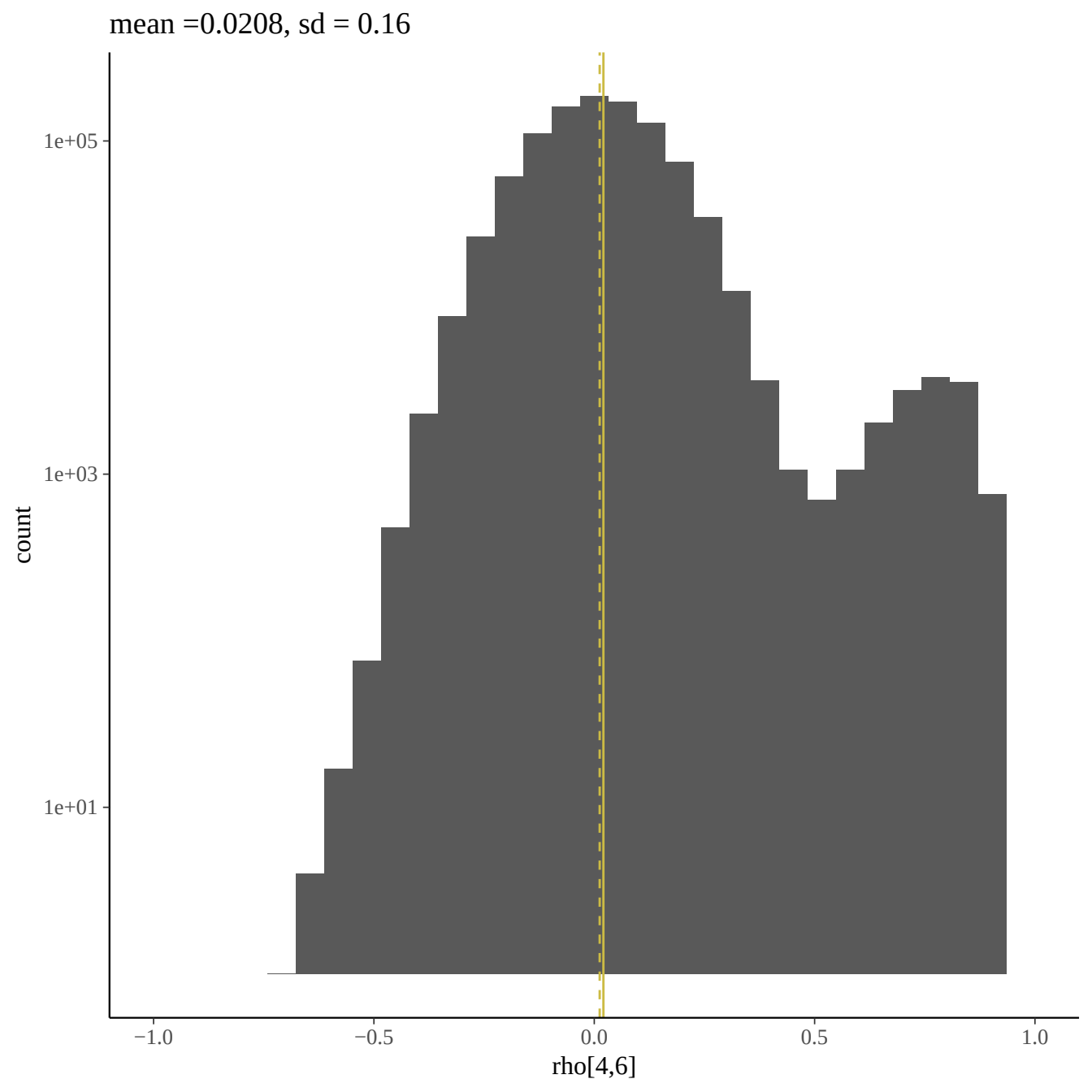

mean =0.146, sd = 0.132
1e+05
1e+03
1e+01
count
−1.0
−0.5
0.0
0.5
1.0
rho[3,5]

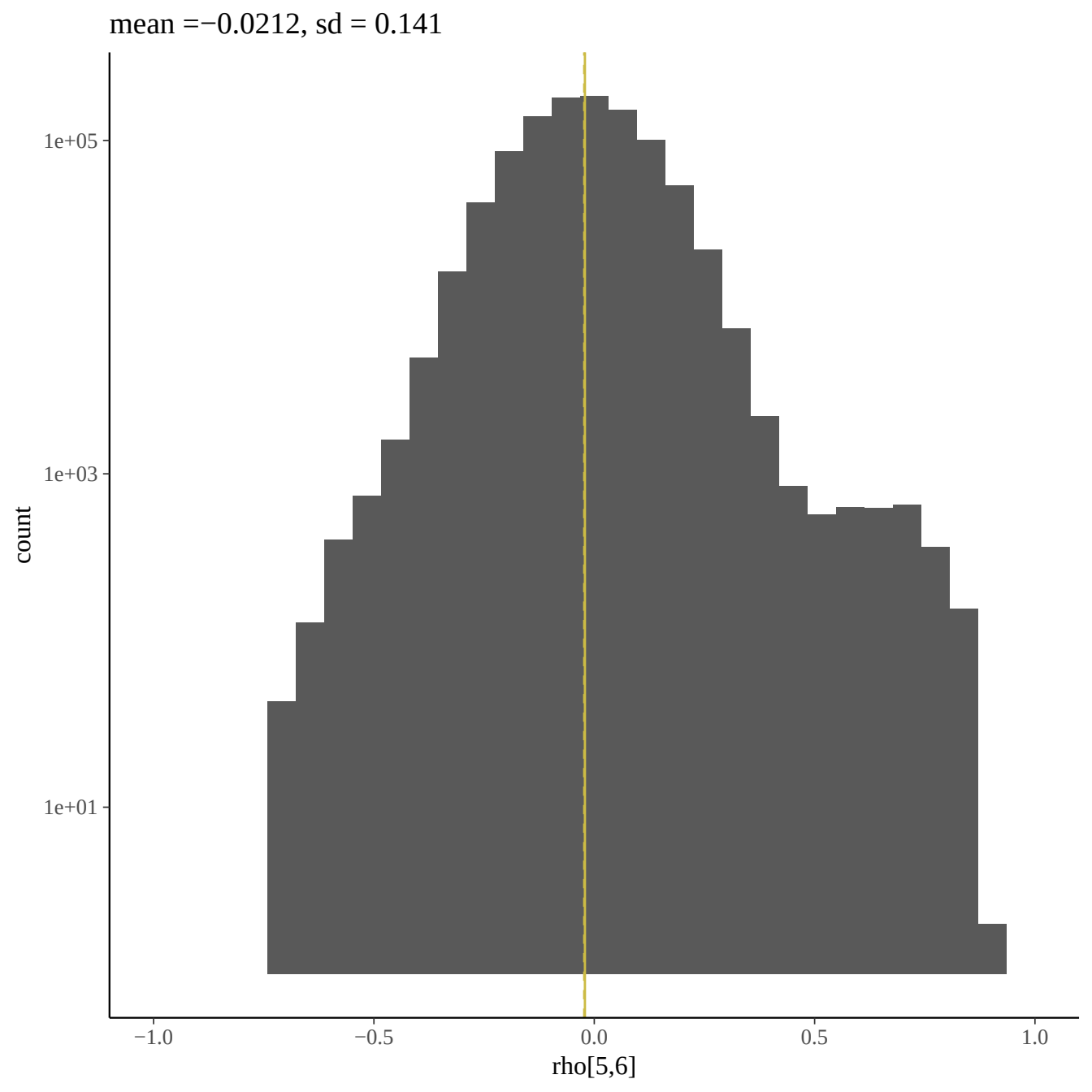

mean =−0.036, sd = 0.138
1e+05
1e+03
1e+01
count
−1.0
−0.5
0.0
0.5
1.0
rho[4,5]

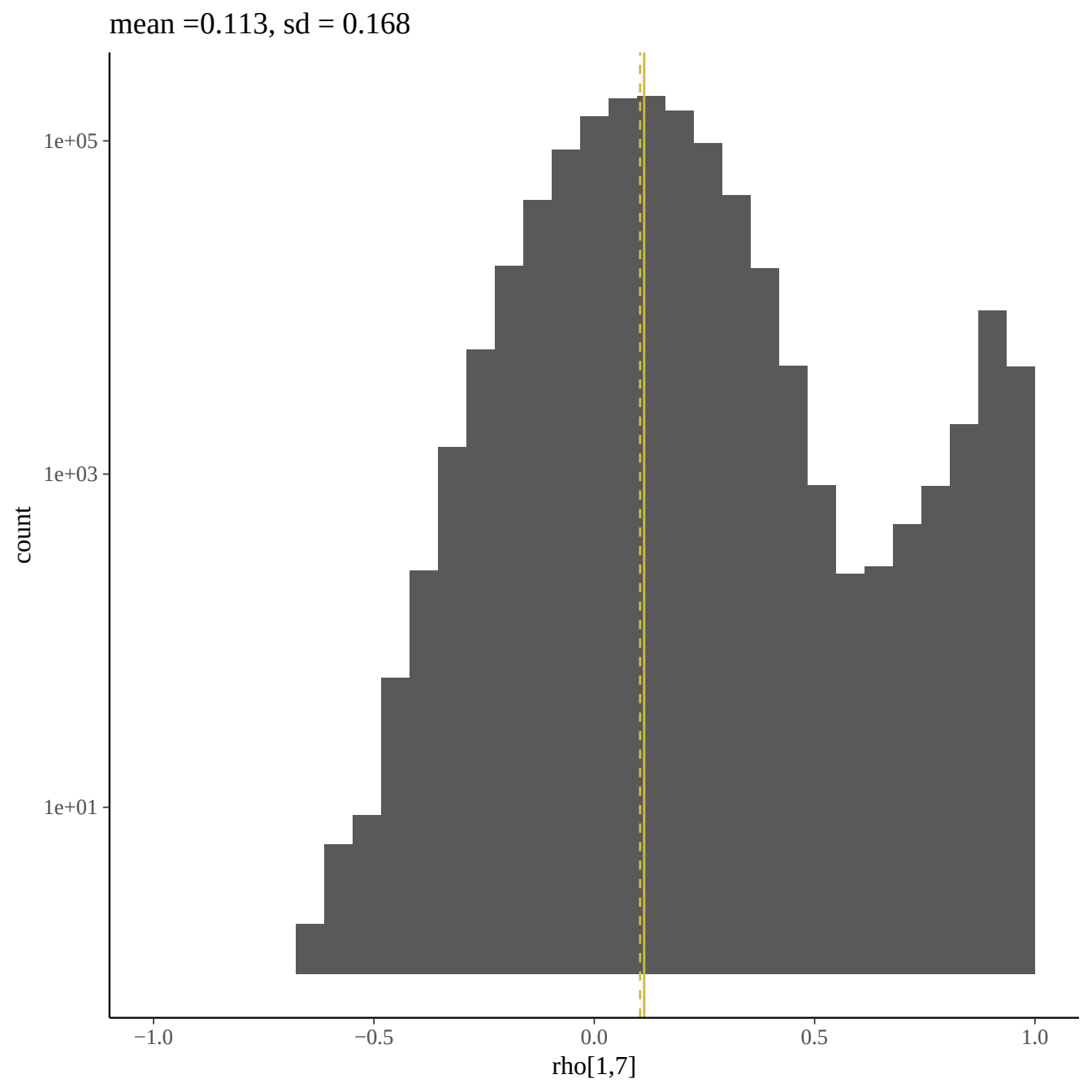

mean =0.217, sd = 0.157
1e+05
1e+03
1e+01
count
−1.0
−0.5
0.0
0.5
1.0
rho[1,6]

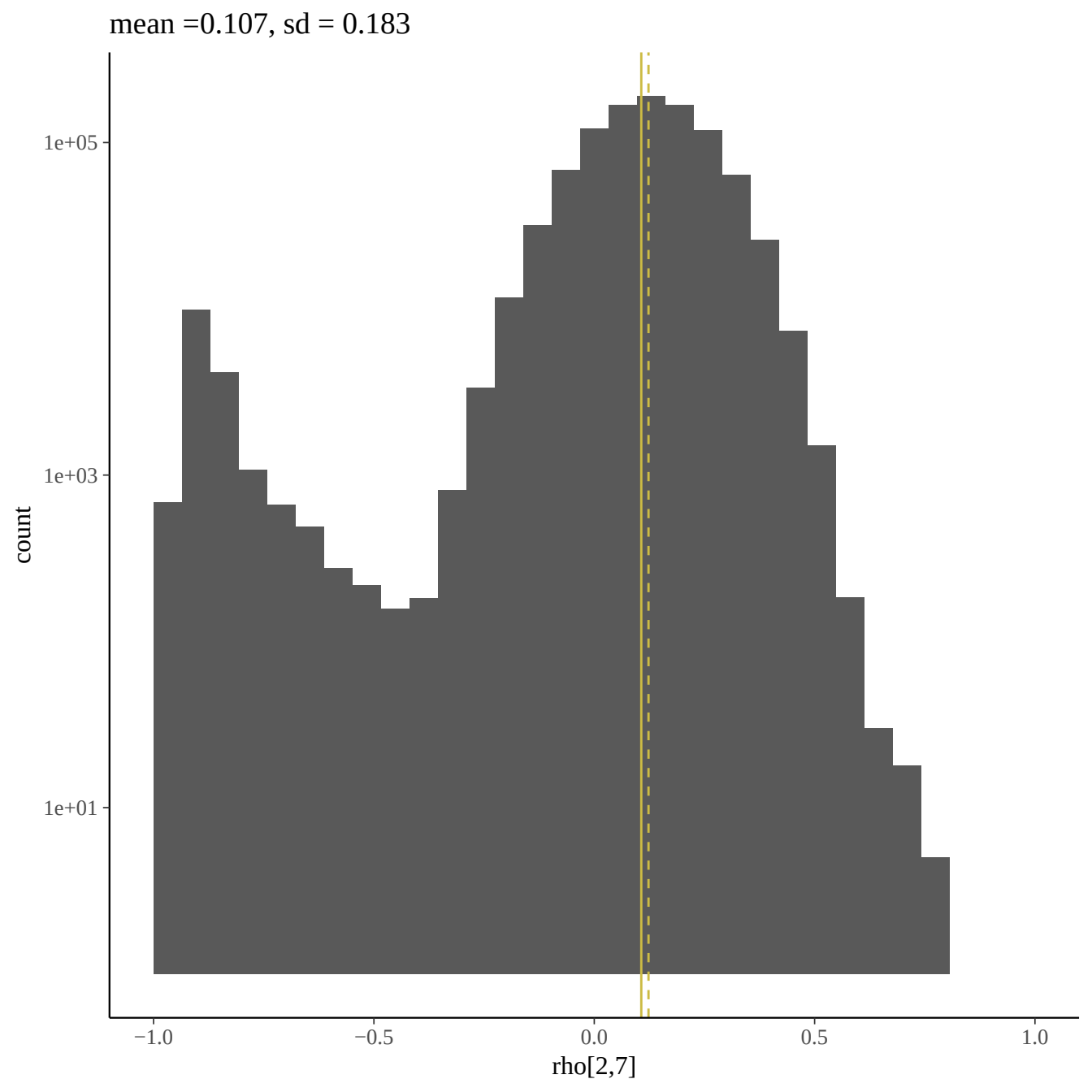

mean =−0.148, sd = 0.164
1e+05
1e+03
1e+01
count
−1.0
−0.5
0.0
0.5
1.0
rho[2,6]

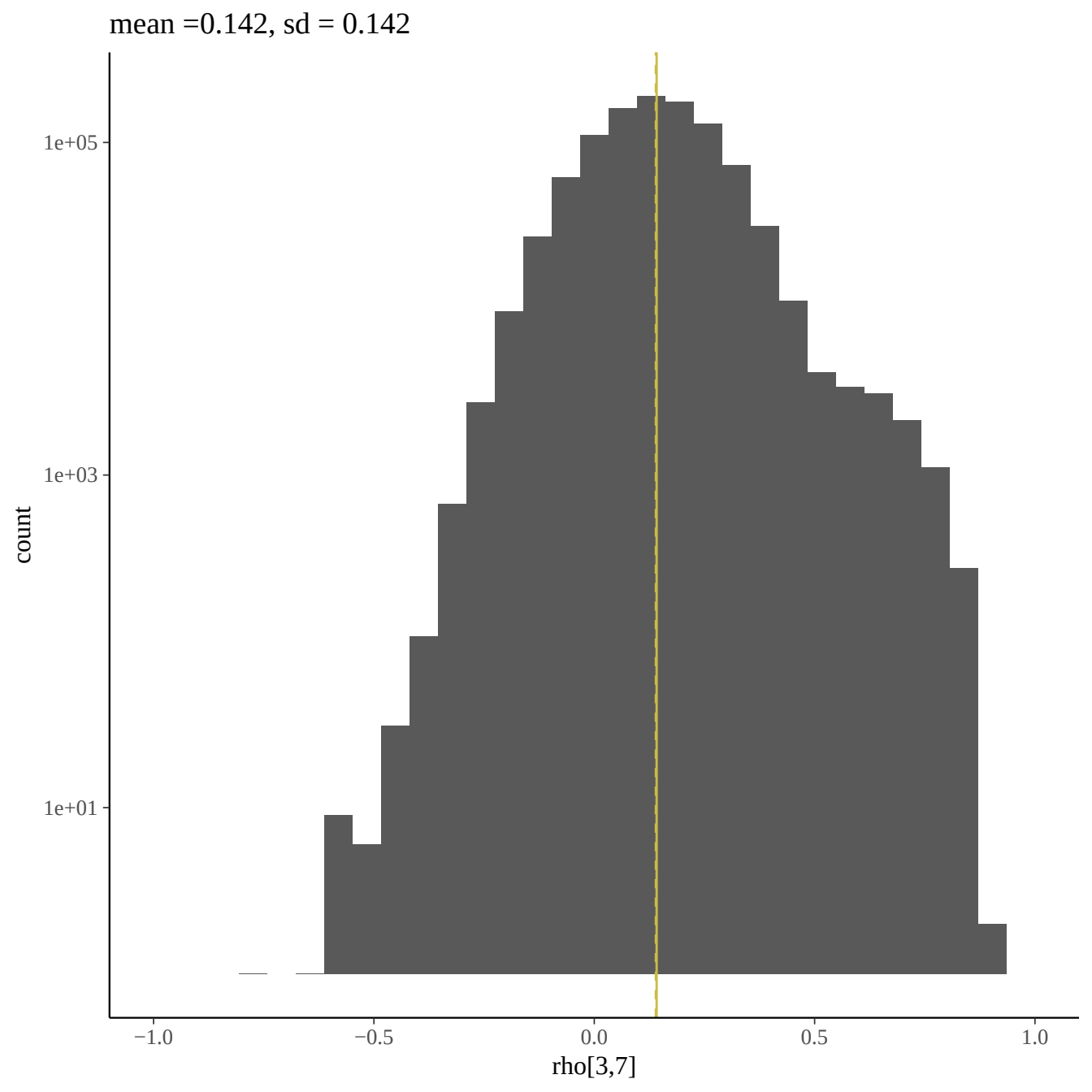

mean =−0.218, sd = 0.154
1e+05
1e+03
1e+01
count
−1.0
−0.5
0.0
0.5
1.0
rho[3,6]

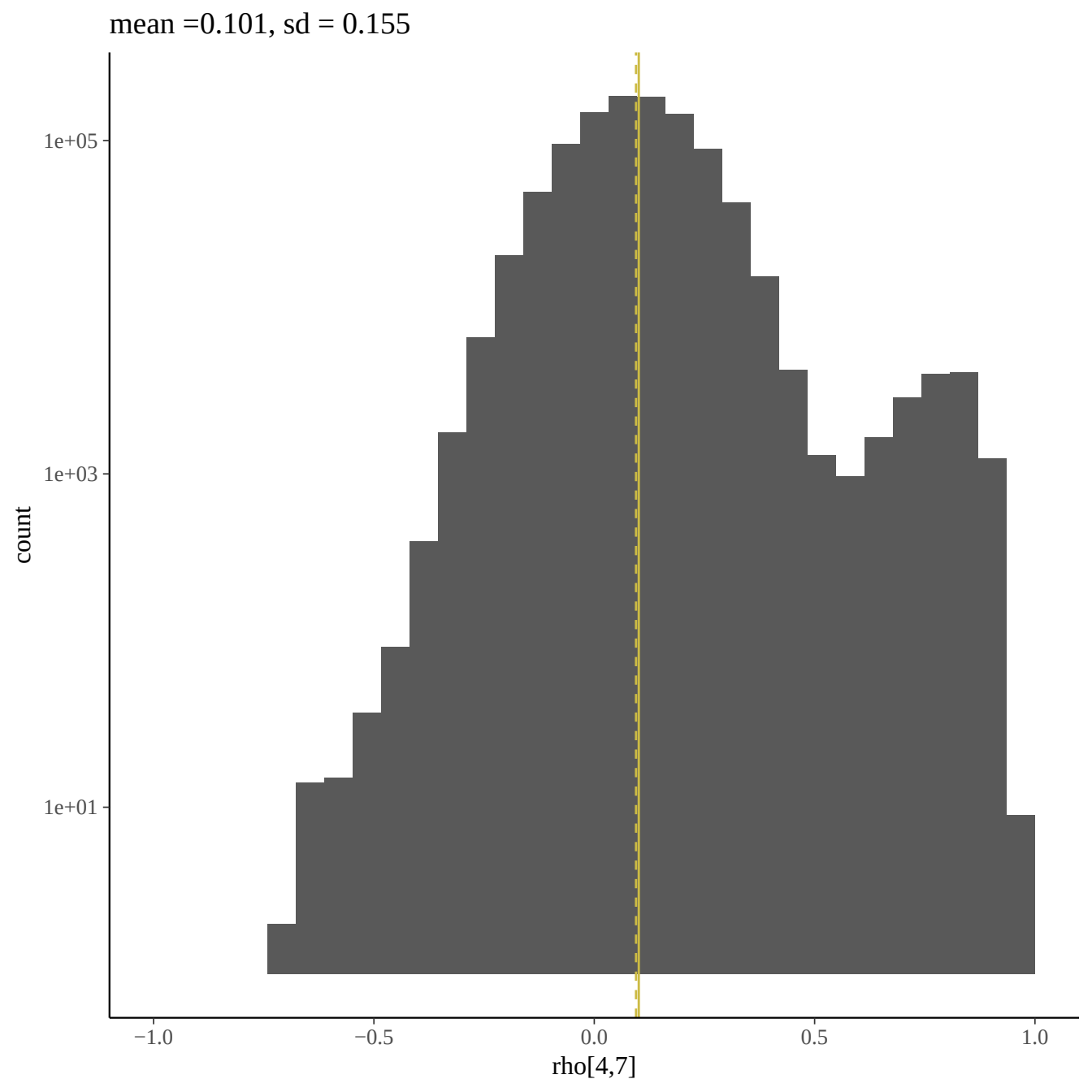

mean =0.0208, sd = 0.16
1e+05
1e+03
1e+01
count
−1.0
−0.5
0.0
0.5
1.0
rho[4,6]

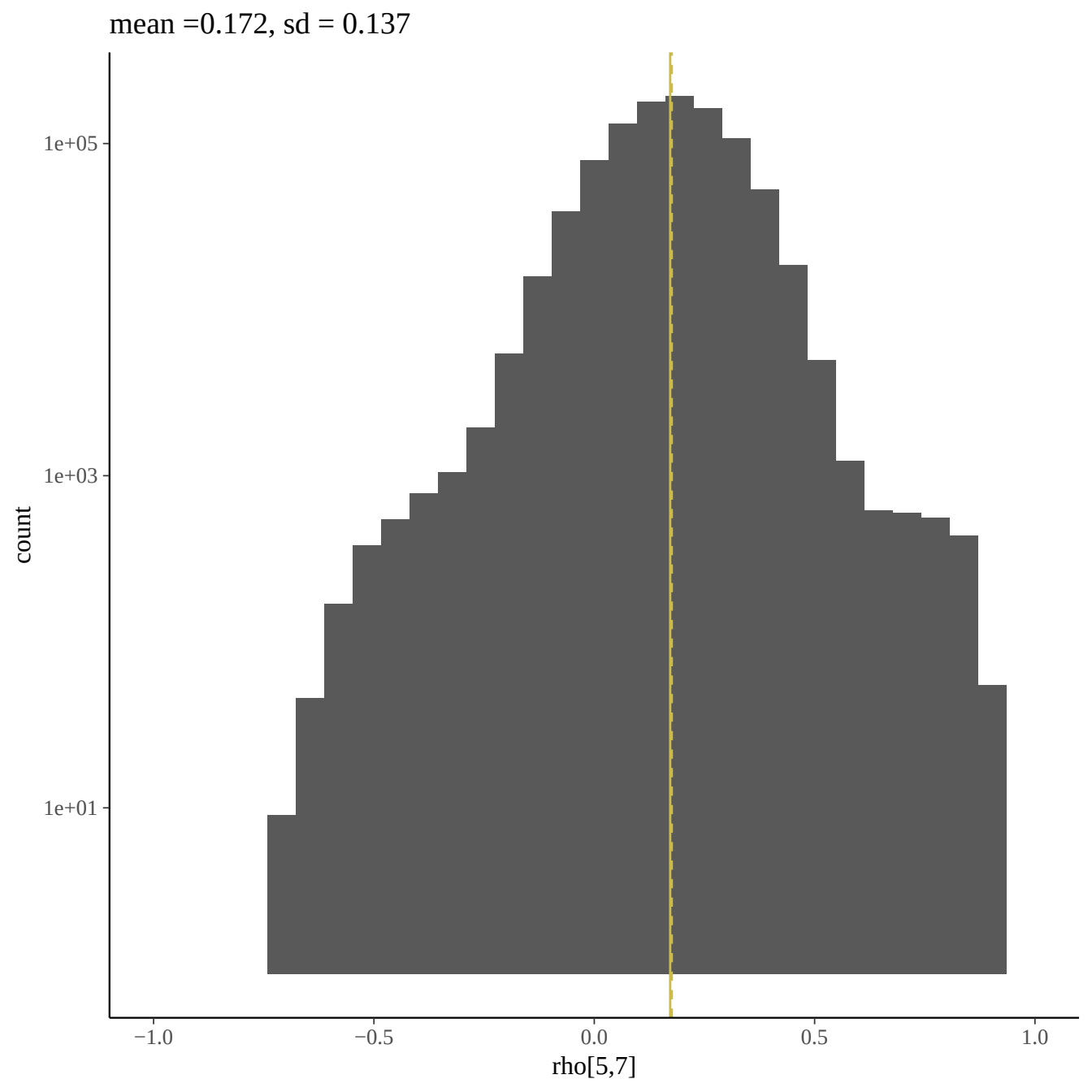

mean =−0.0212, sd = 0.141
1e+05
1e+03
1e+01
count
−1.0
−0.5
0.0
0.5
1.0
rho[5,6]

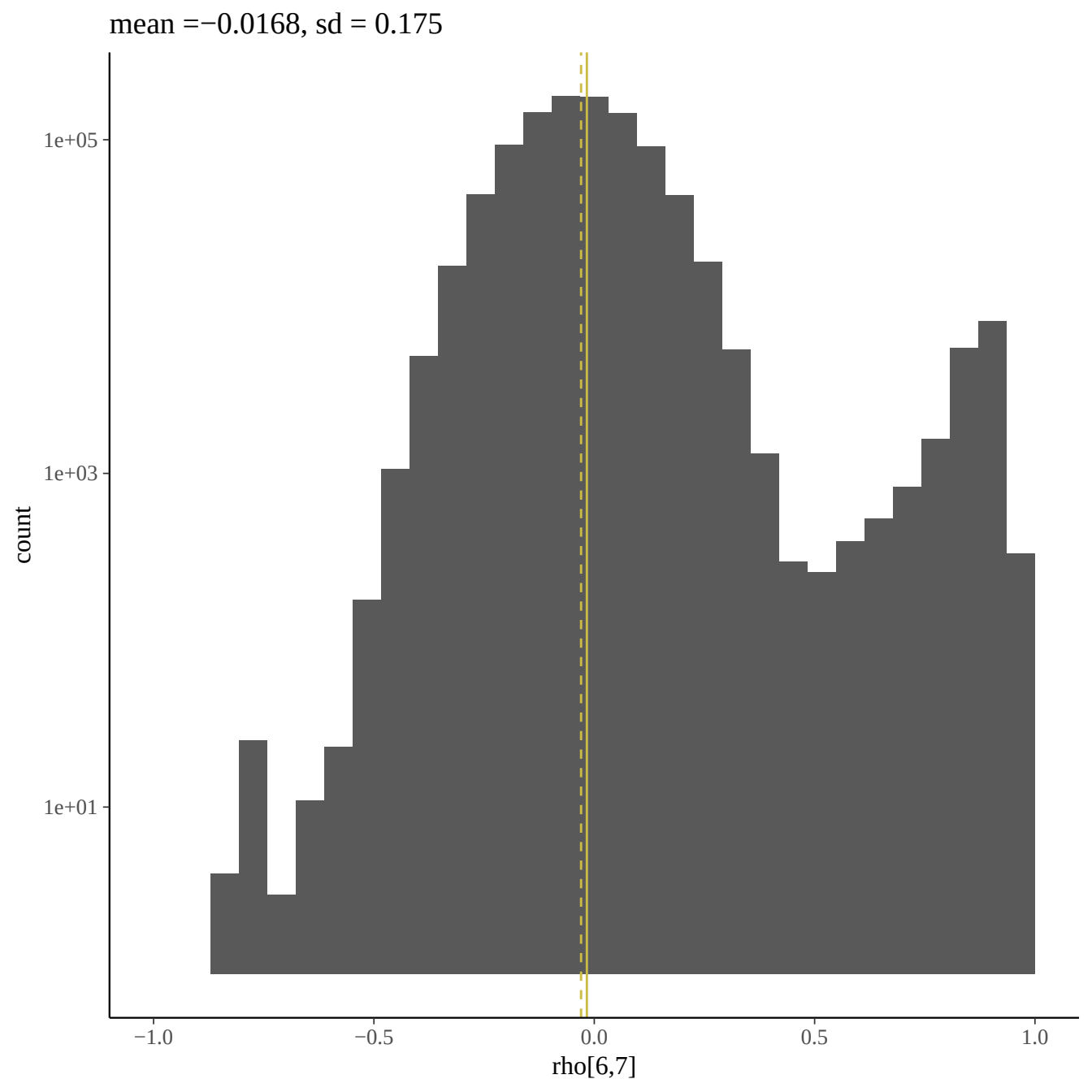

mean =0.113, sd = 0.168
1e+05
1e+03
1e+01
count
−1.0
−0.5
0.0
0.5
1.0
rho[1,7]

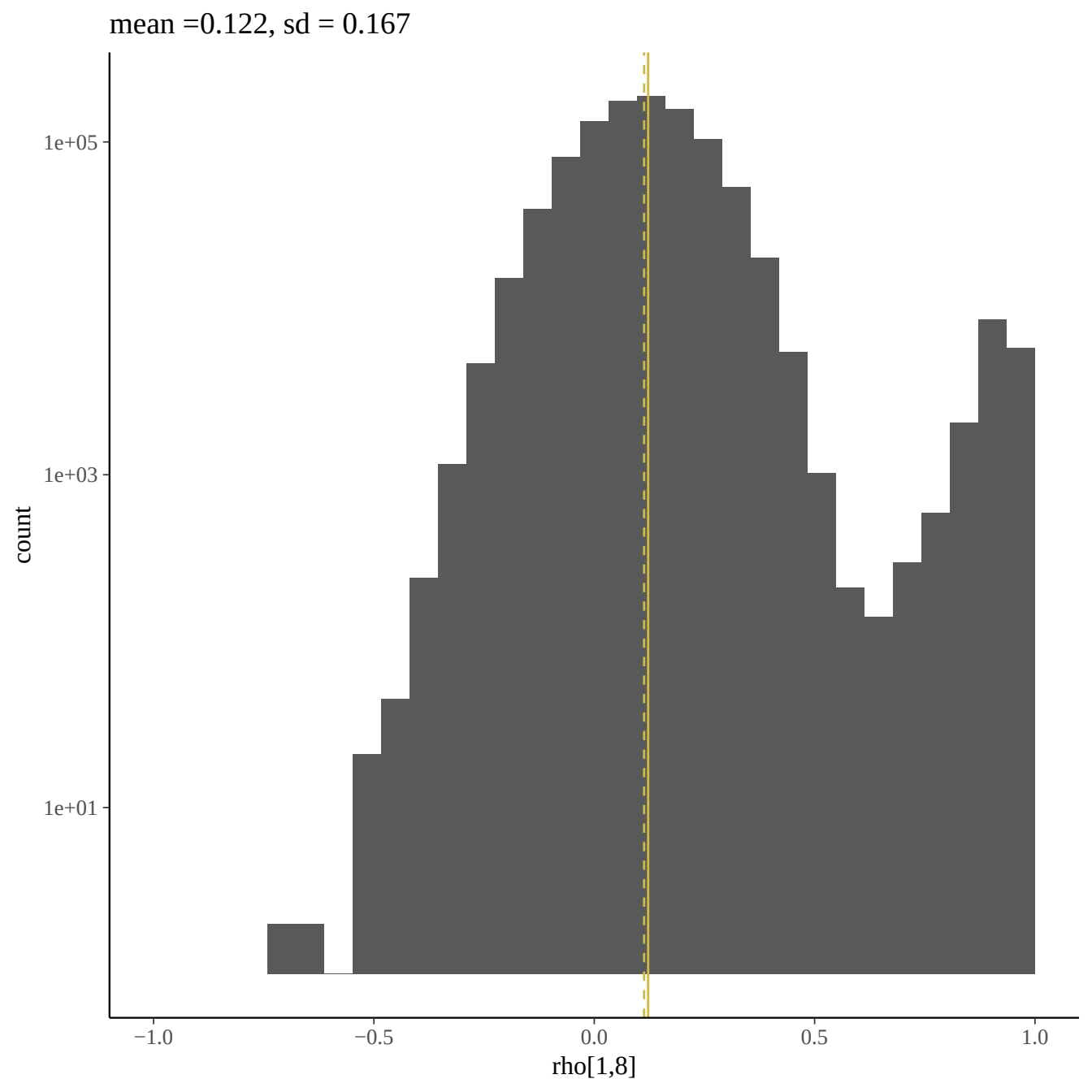

mean =0.107, sd = 0.183
1e+05
1e+03
1e+01
count
−1.0
−0.5
0.0
0.5
1.0
rho[2,7]

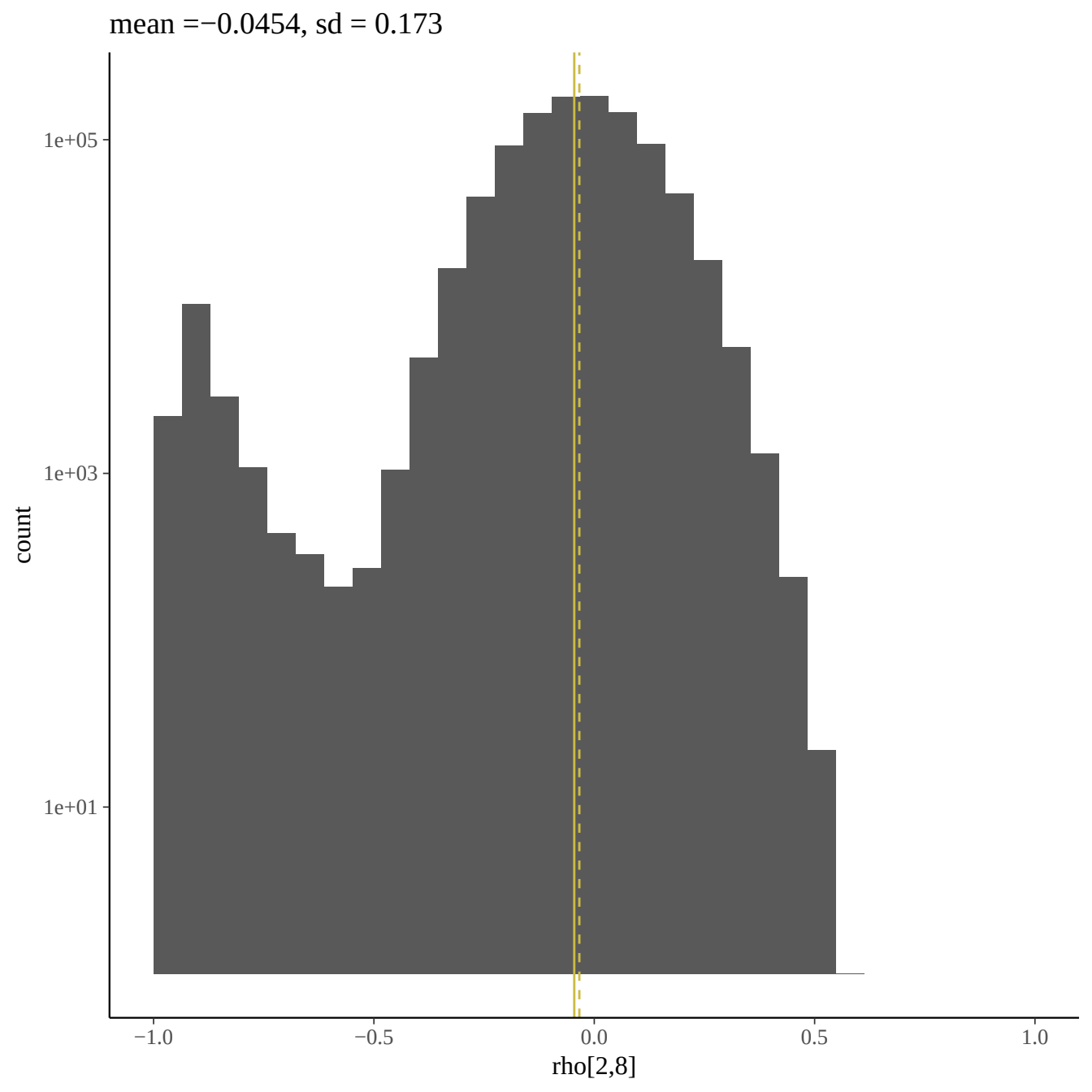

mean =0.142, sd = 0.142
1e+05
1e+03
1e+01
count
−1.0
−0.5
0.0
0.5
1.0
rho[3,7]

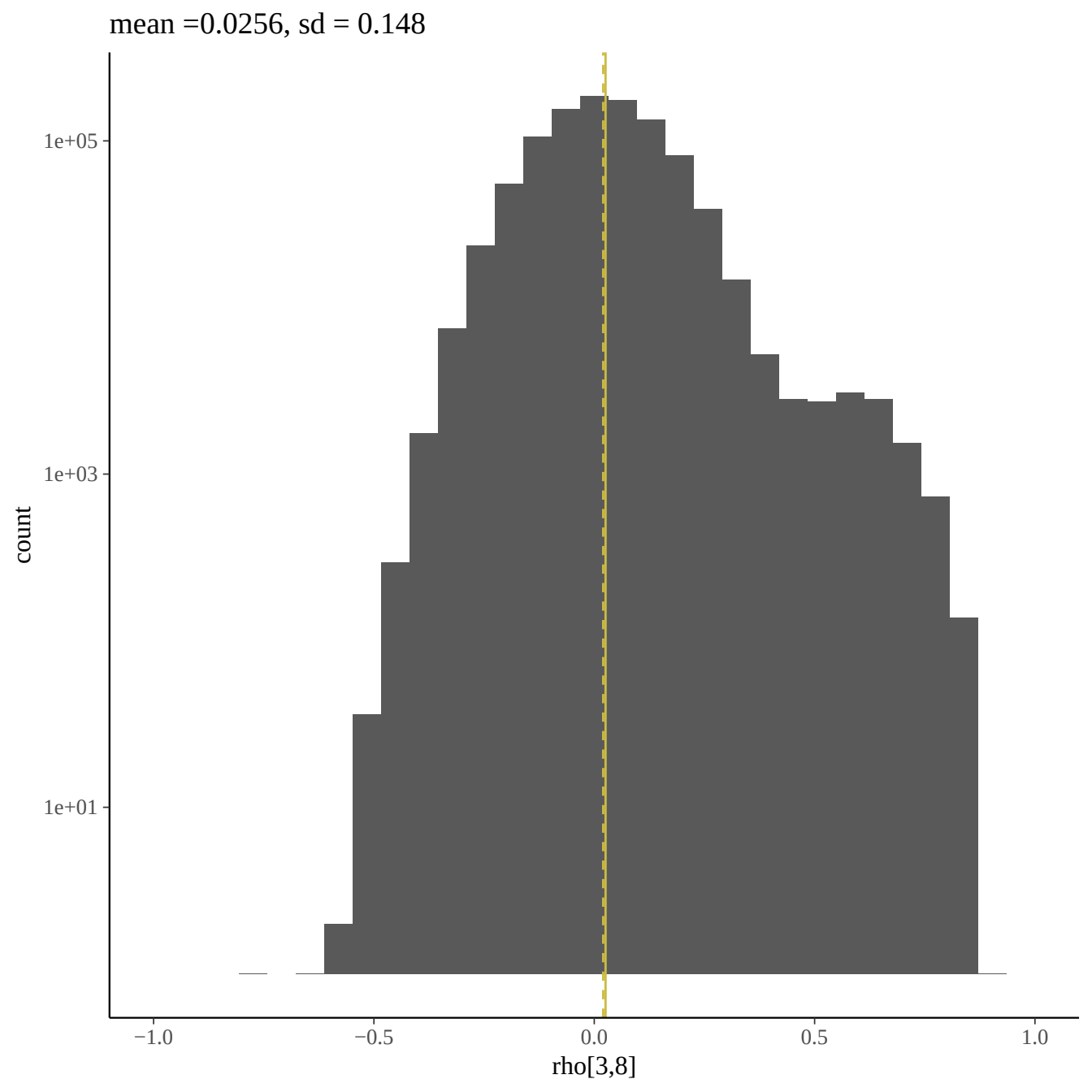

mean =0.101, sd = 0.155
1e+05
1e+03
1e+01
count
−1.0
−0.5
0.0
0.5
1.0
rho[4,7]

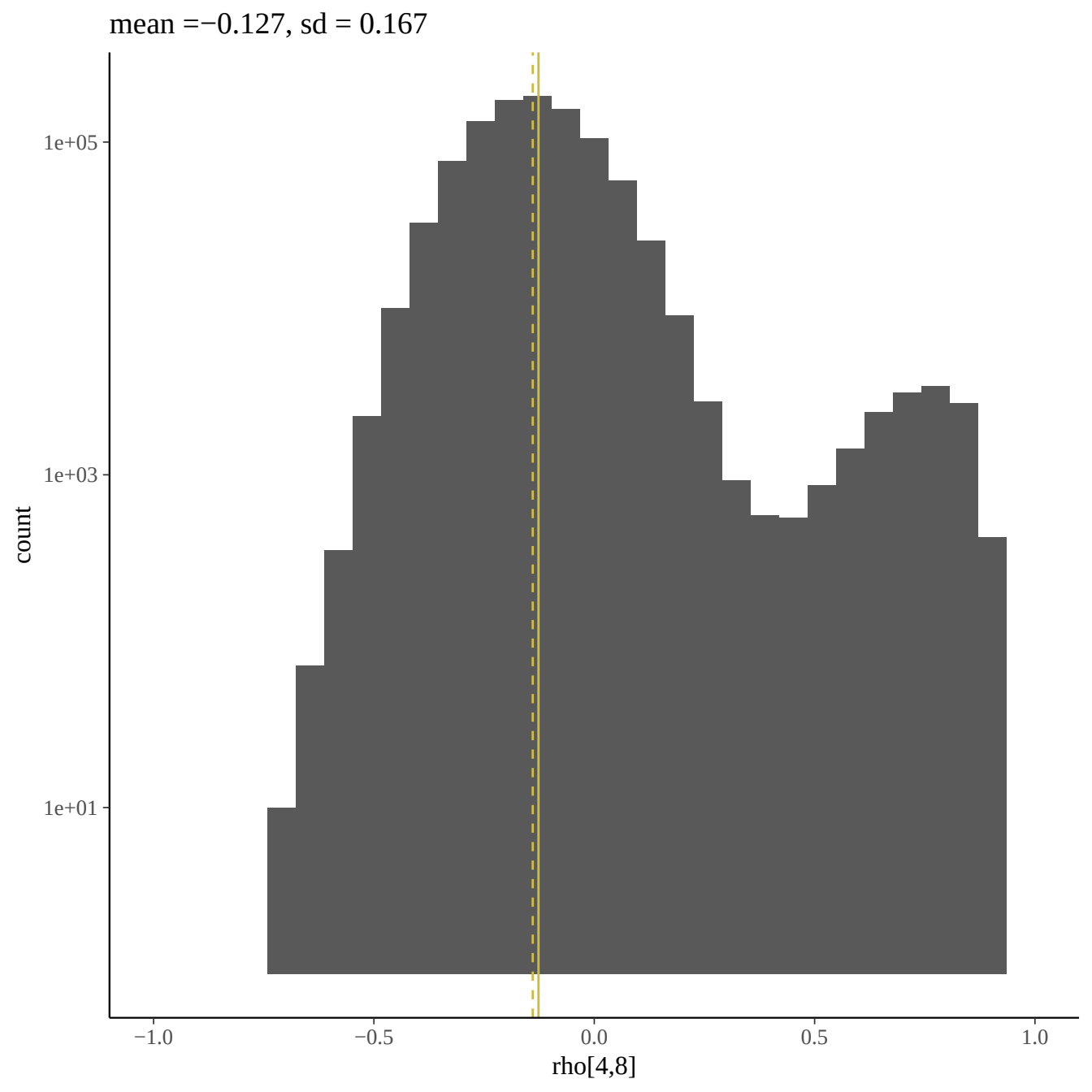

mean =0.172, sd = 0.137
1e+05
1e+03
1e+01
count
−1.0
−0.5
0.0
0.5
1.0
rho[5,7]

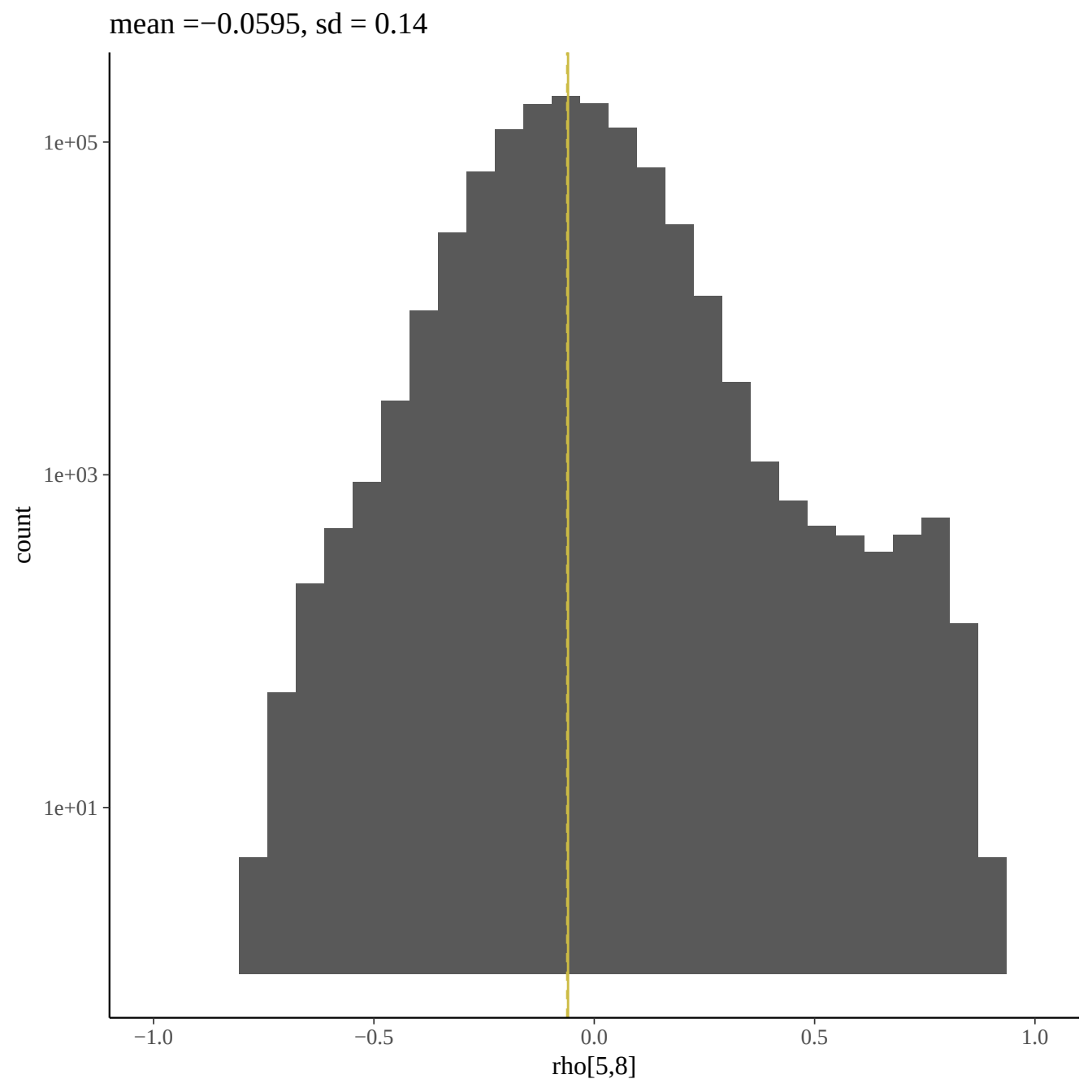

mean =−0.0168, sd = 0.175
1e+05
1e+03
1e+01
count
−1.0
−0.5
0.0
0.5
1.0
rho[6,7]

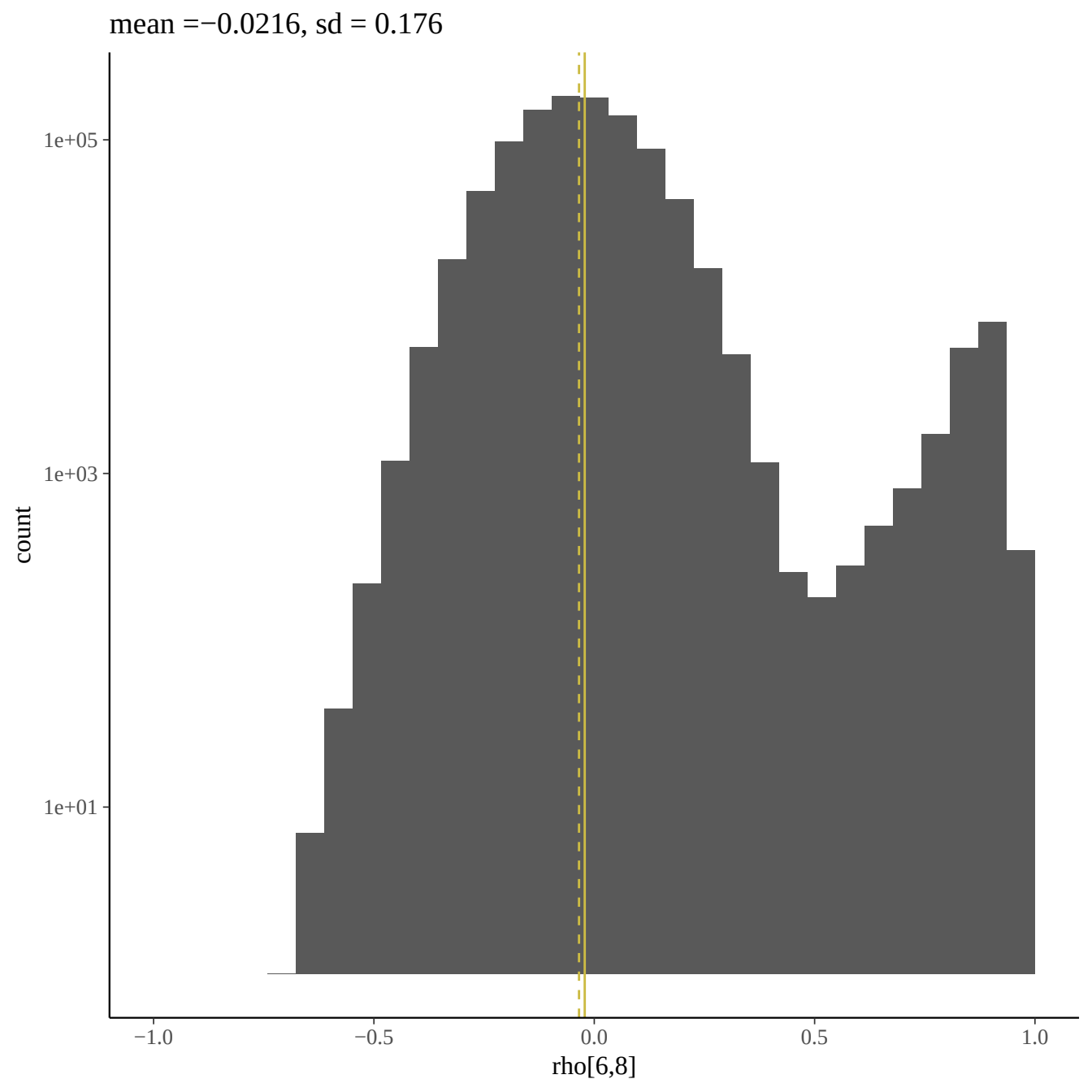

mean =0.122, sd = 0.167
1e+05
1e+03
1e+01
count
−1.0
−0.5
0.0
0.5
1.0
rho[1,8]

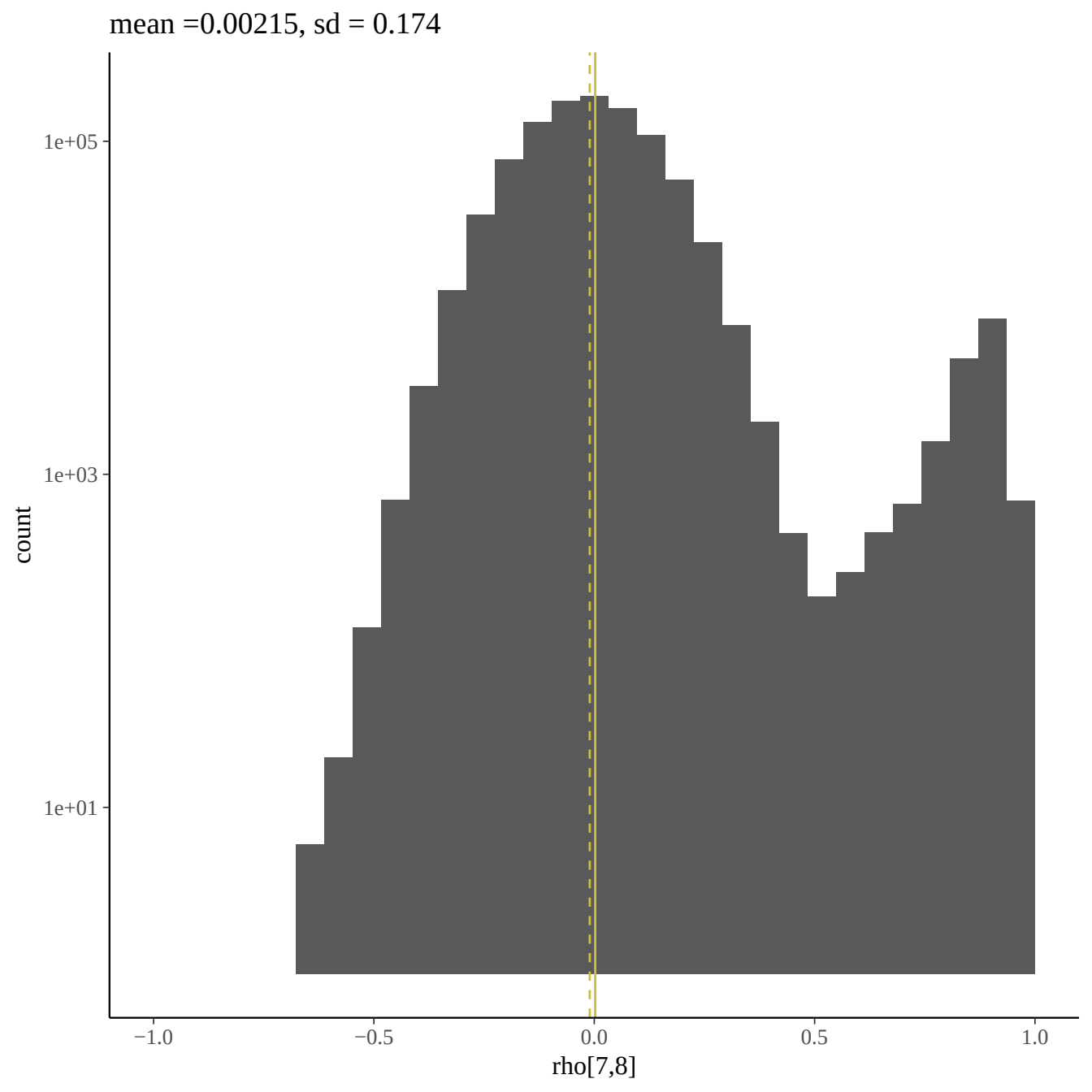

mean =−0.0454, sd = 0.173
1e+05
1e+03
1e+01
count
−1.0
−0.5
0.0
0.5
1.0
rho[2,8]

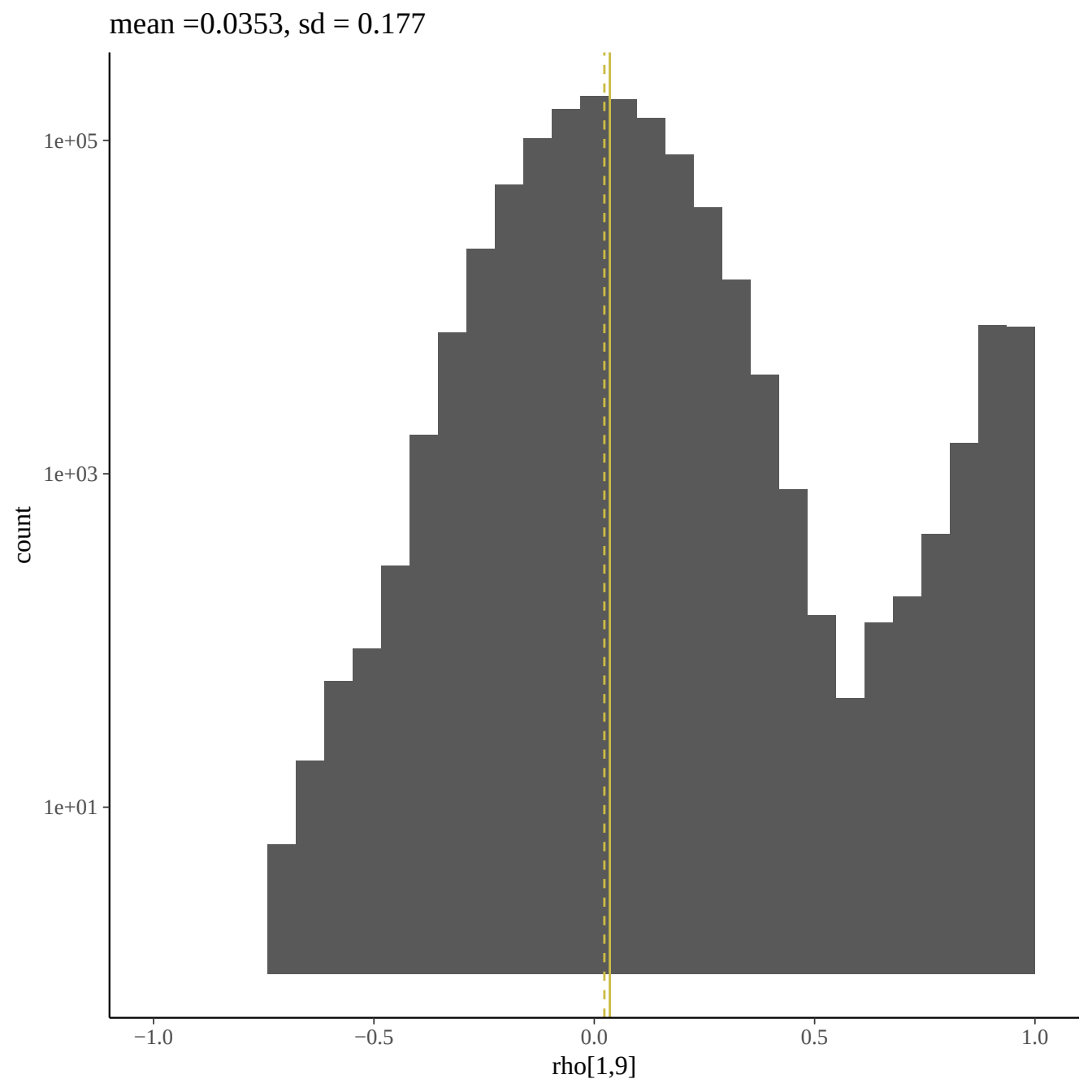

mean =0.0256, sd = 0.148
1e+05
1e+03
1e+01
count
−1.0
−0.5
0.0
0.5
1.0
rho[3,8]

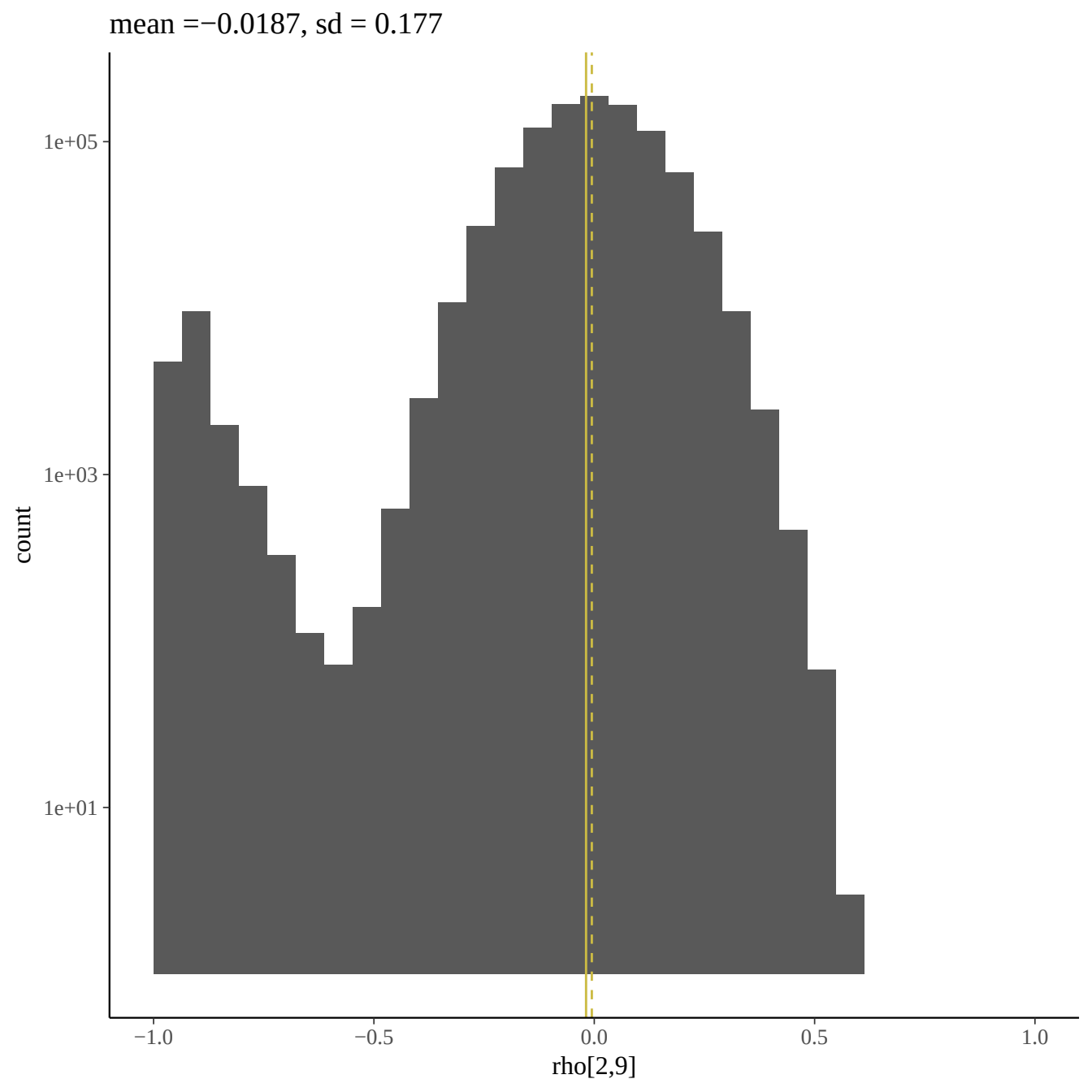

mean =−0.127, sd = 0.167
1e+05
1e+03
1e+01
count
−1.0
−0.5
0.0
0.5
1.0
rho[4,8]

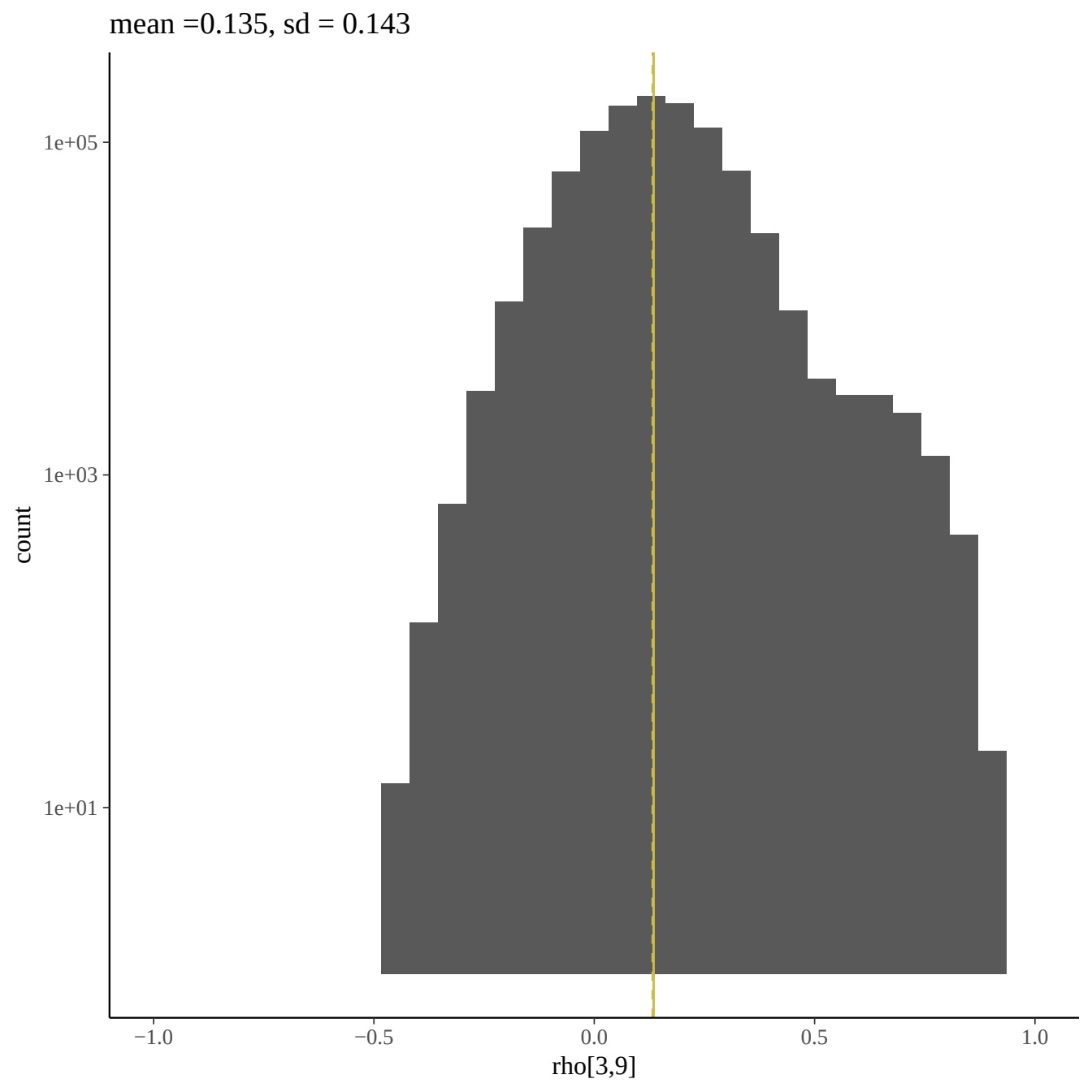

mean =−0.0595, sd = 0.14
1e+05
1e+03
1e+01
count
−1.0
−0.5
0.0
0.5
1.0
rho[5,8]

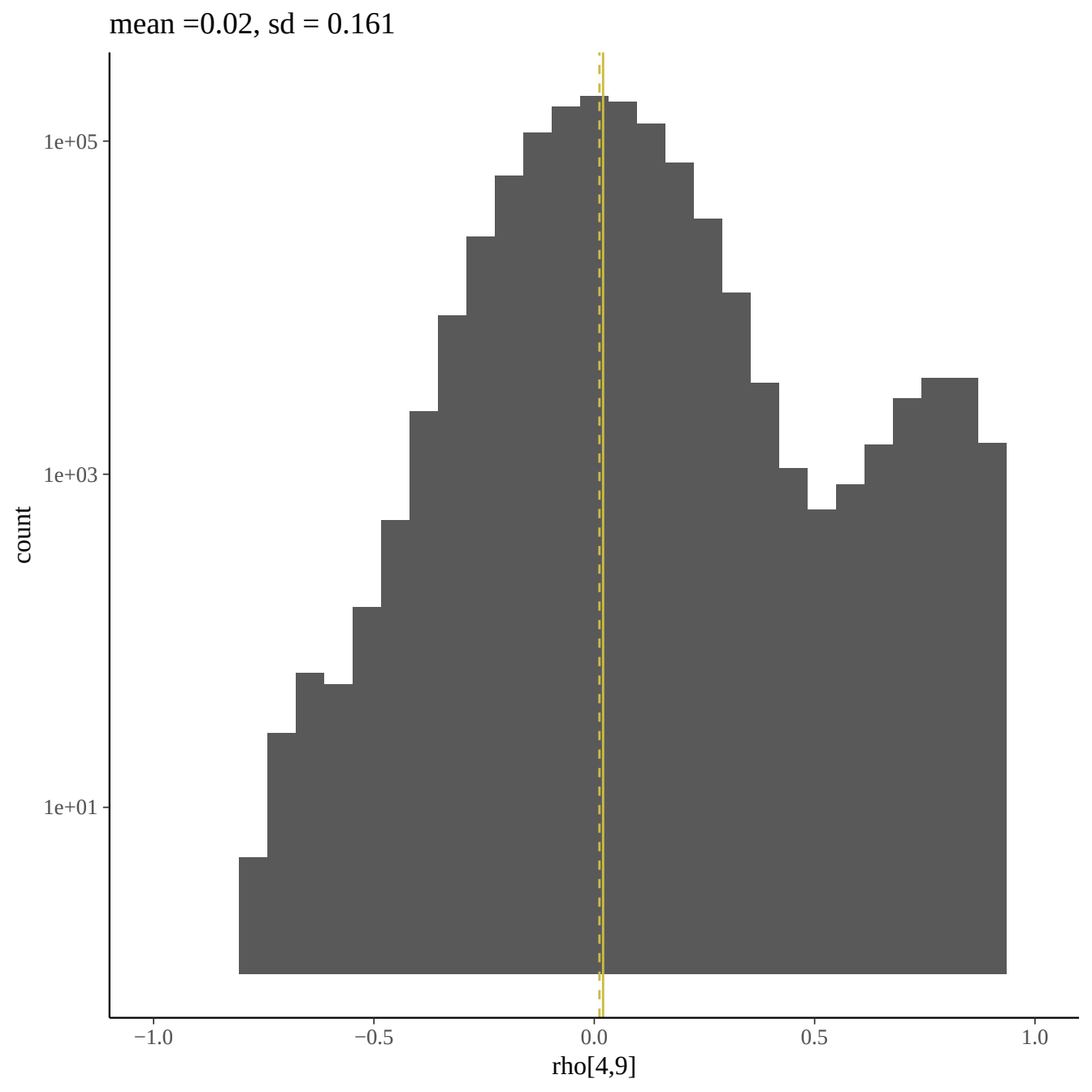

mean =−0.0216, sd = 0.176
1e+05
1e+03
1e+01
count
−1.0
−0.5
0.0
0.5
1.0
rho[6,8]

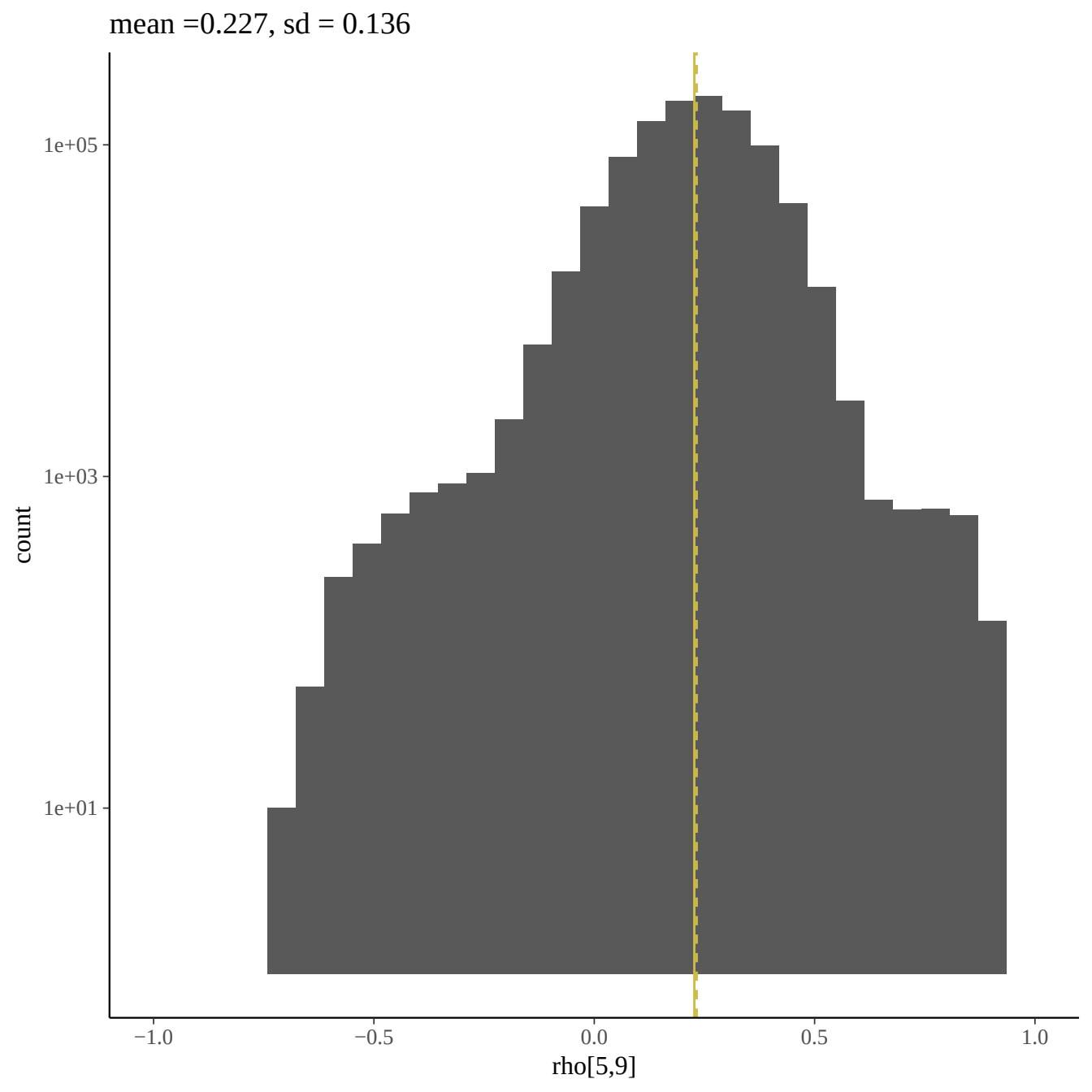

mean =0.00215, sd = 0.174
1e+05
1e+03
1e+01
count
−1.0
−0.5
0.0
0.5
1.0
rho[7,8]

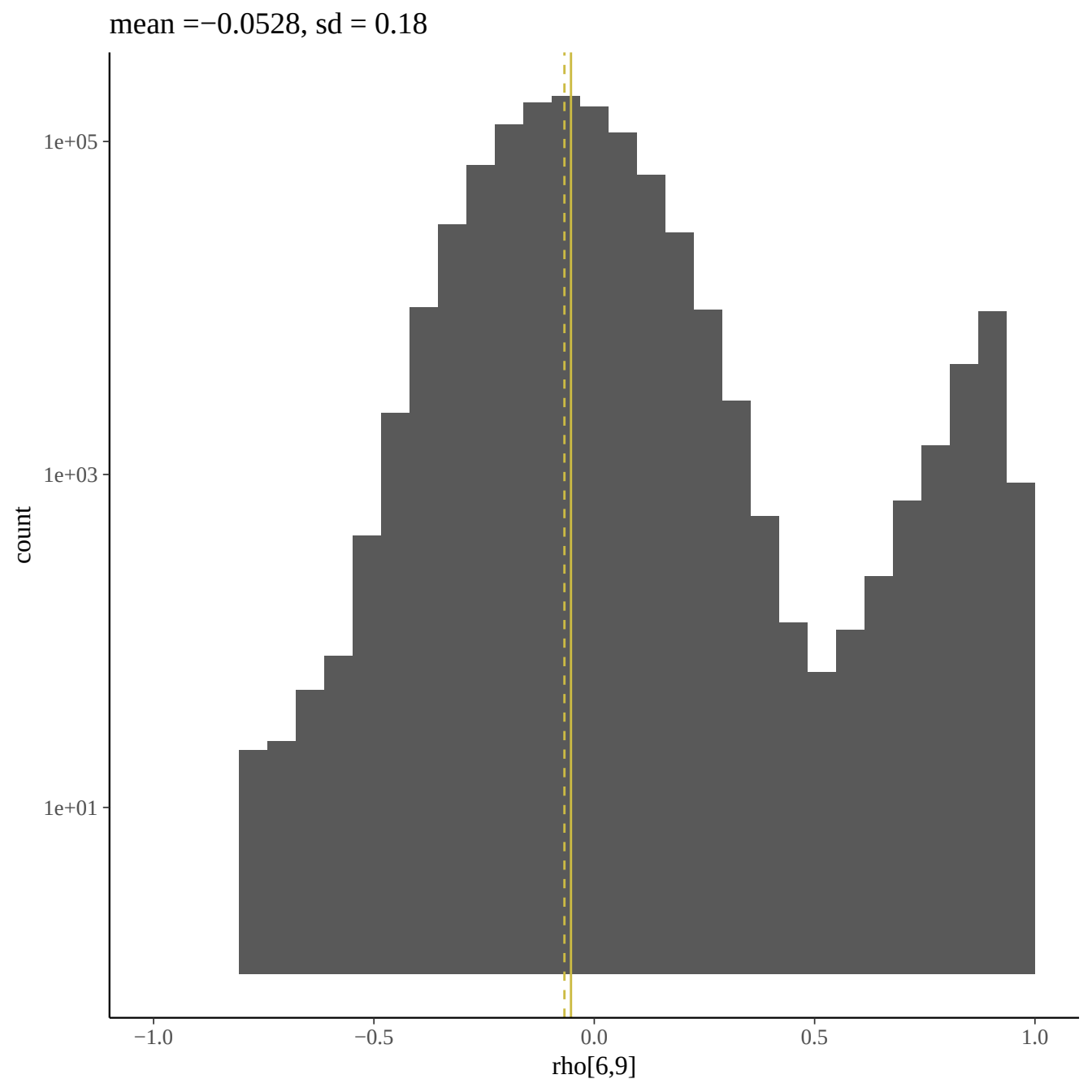

mean =0.0353, sd = 0.177
1e+05
1e+03
1e+01
count
−1.0
−0.5
0.0
0.5
1.0
rho[1,9]

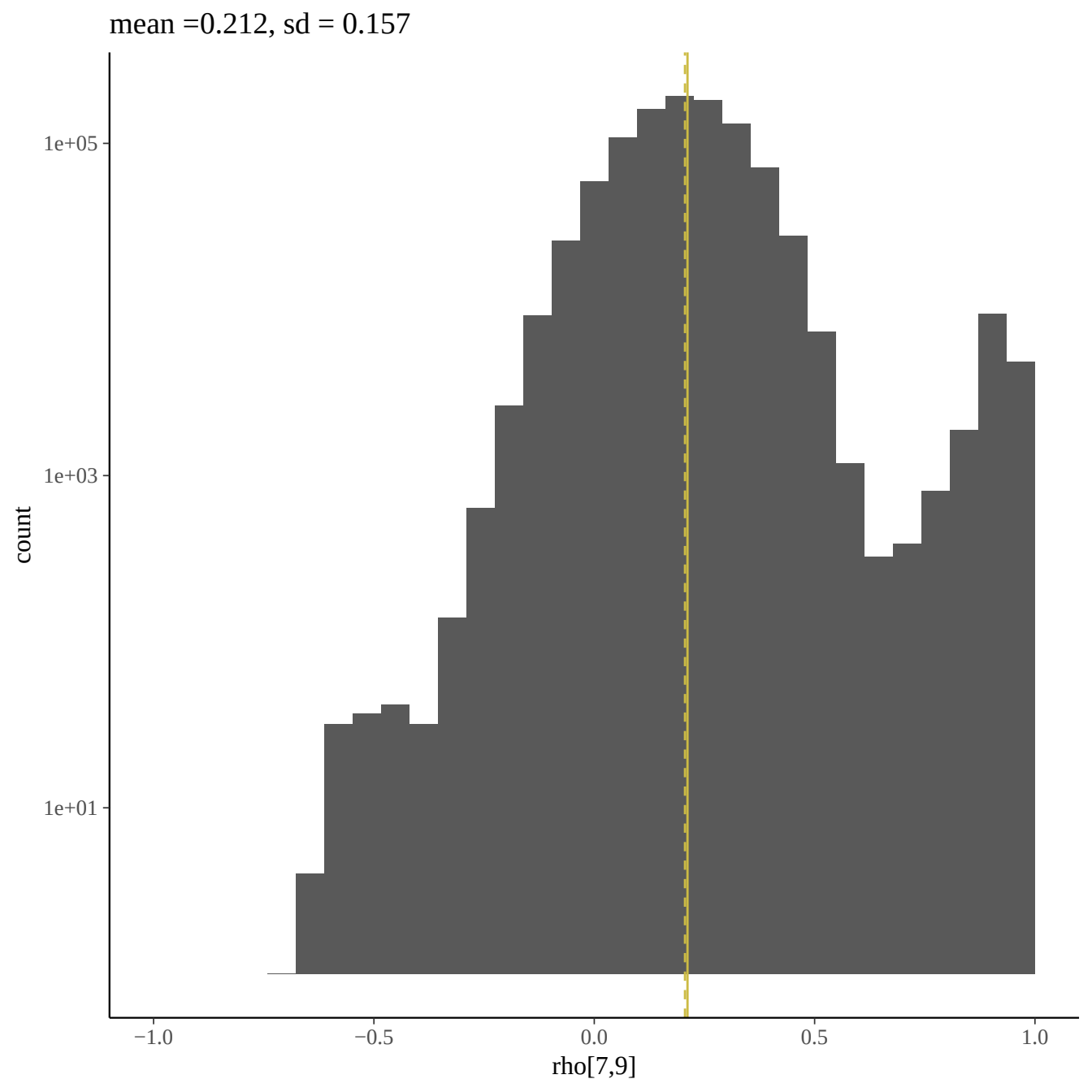

mean =−0.0187, sd = 0.177
1e+05
1e+03
1e+01
count
−1.0
−0.5
0.0
0.5
1.0
rho[2,9]

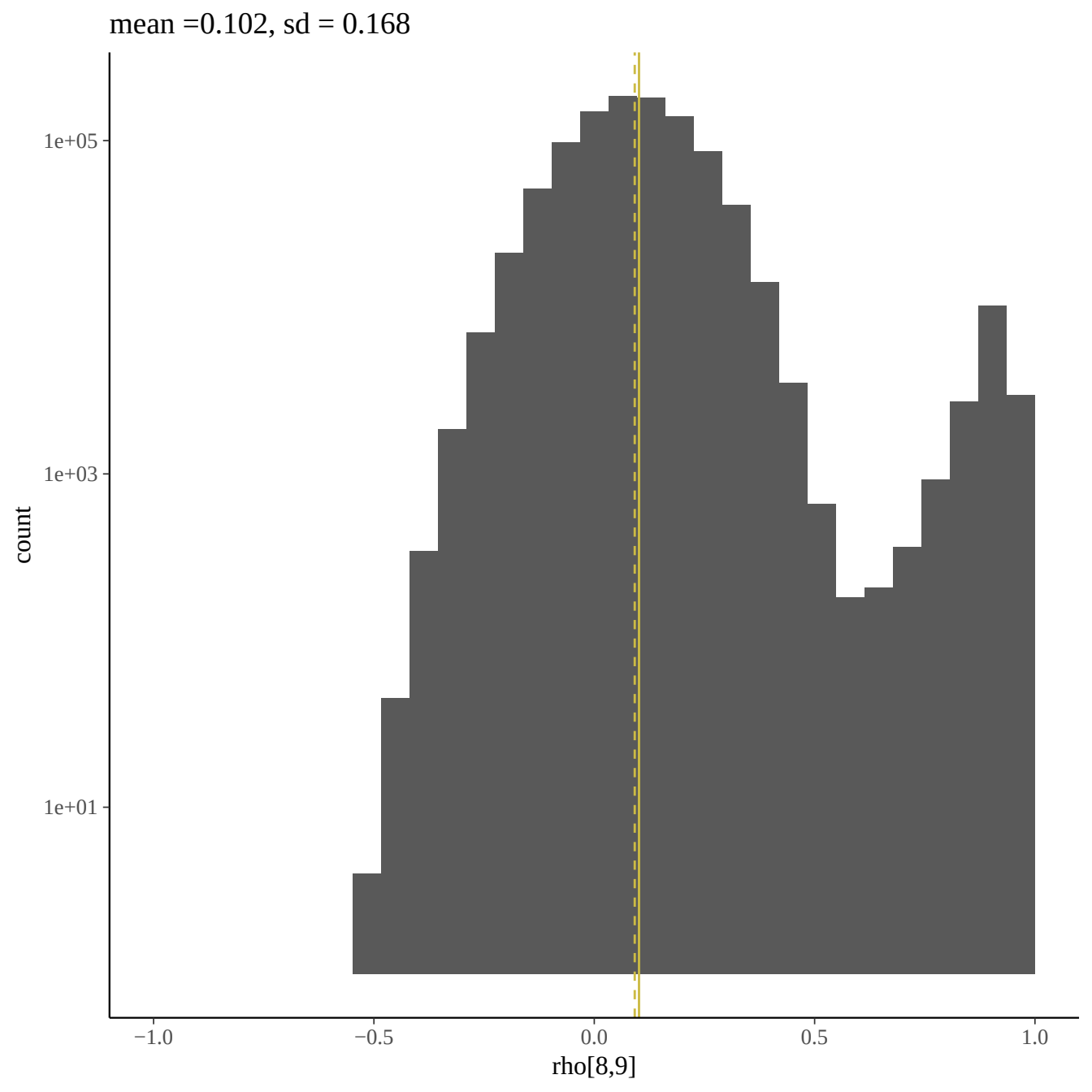

mean =0.135, sd = 0.143
1e+05
1e+03
1e+01
count
−1.0
−0.5
0.0
0.5
1.0
rho[3,9]

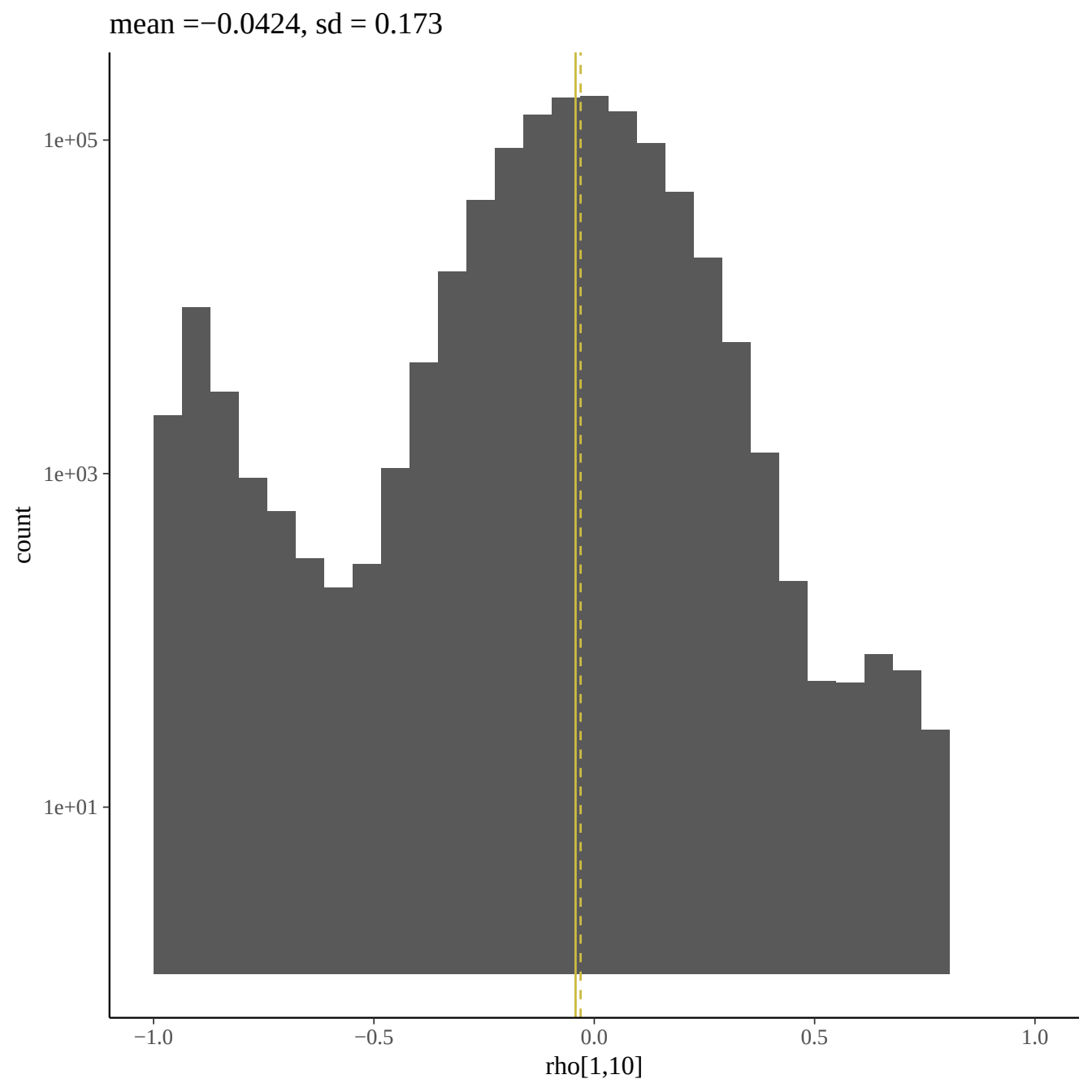

mean =0.02, sd = 0.161
1e+05
1e+03
1e+01
count
−1.0
−0.5
0.0
0.5
1.0
rho[4,9]

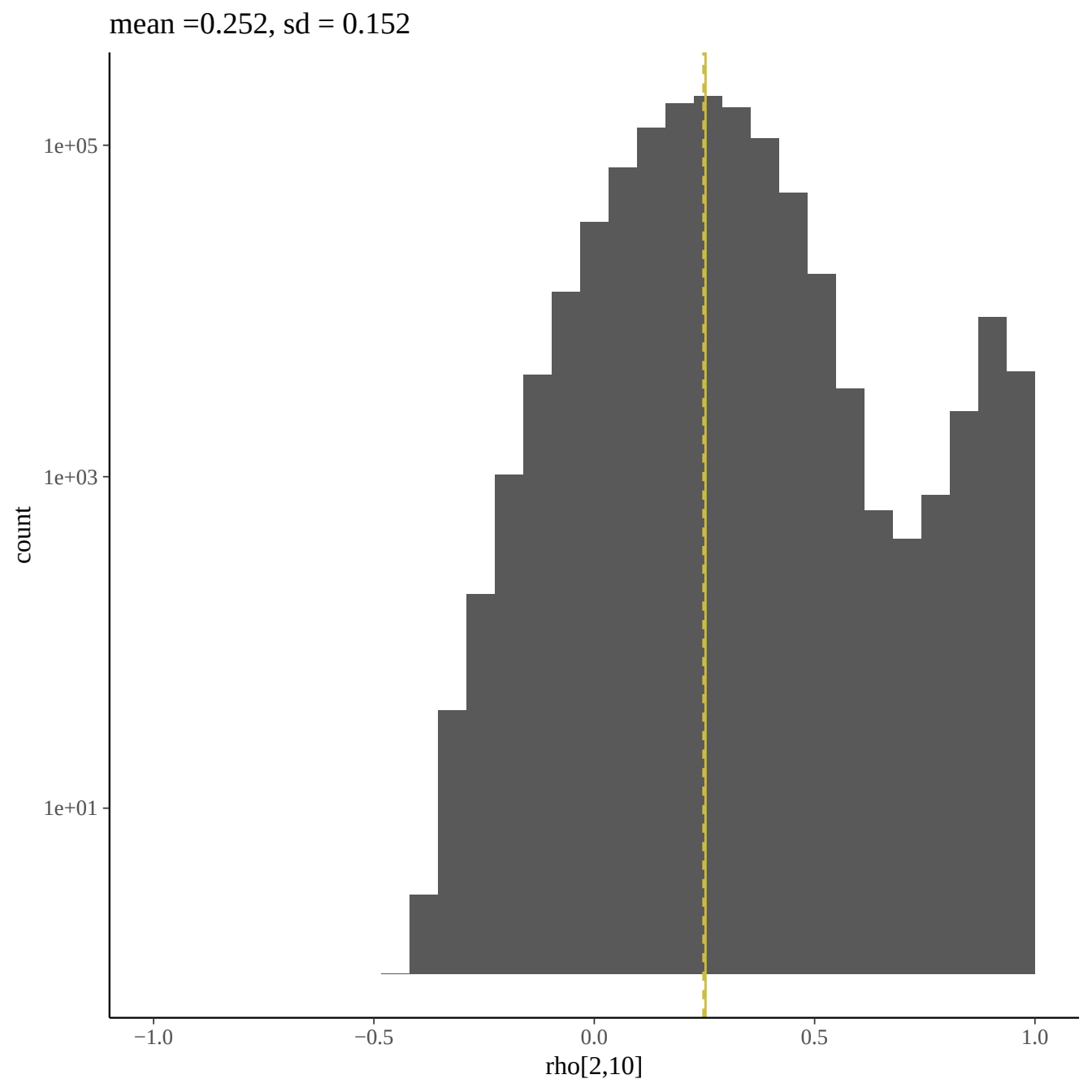

mean =0.227, sd = 0.136
1e+05
1e+03
1e+01
count
−1.0
−0.5
0.0
0.5
1.0
rho[5,9]

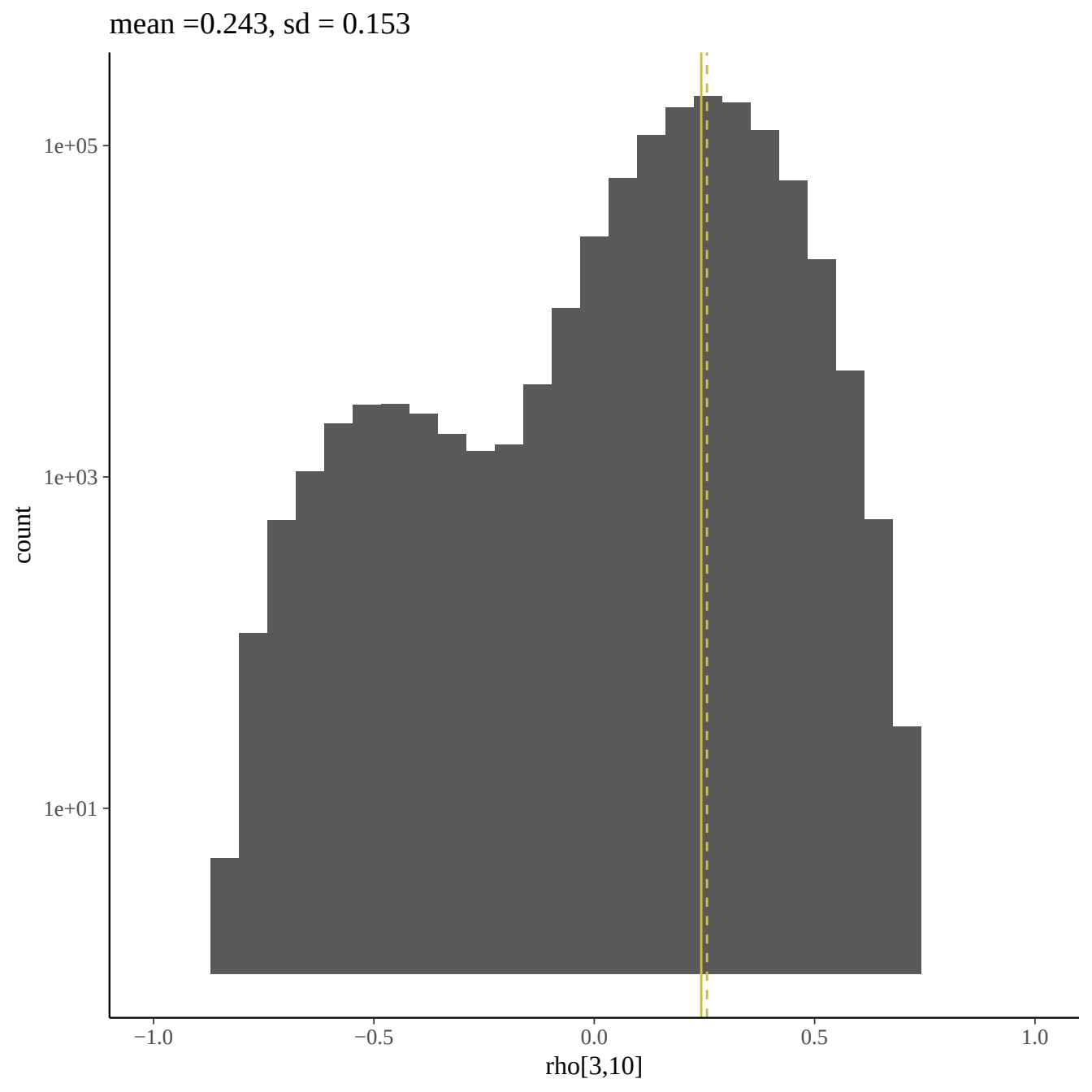

mean =−0.0528, sd = 0.18
1e+05
1e+03
1e+01
count
−1.0
−0.5
0.0
0.5
1.0
rho[6,9]

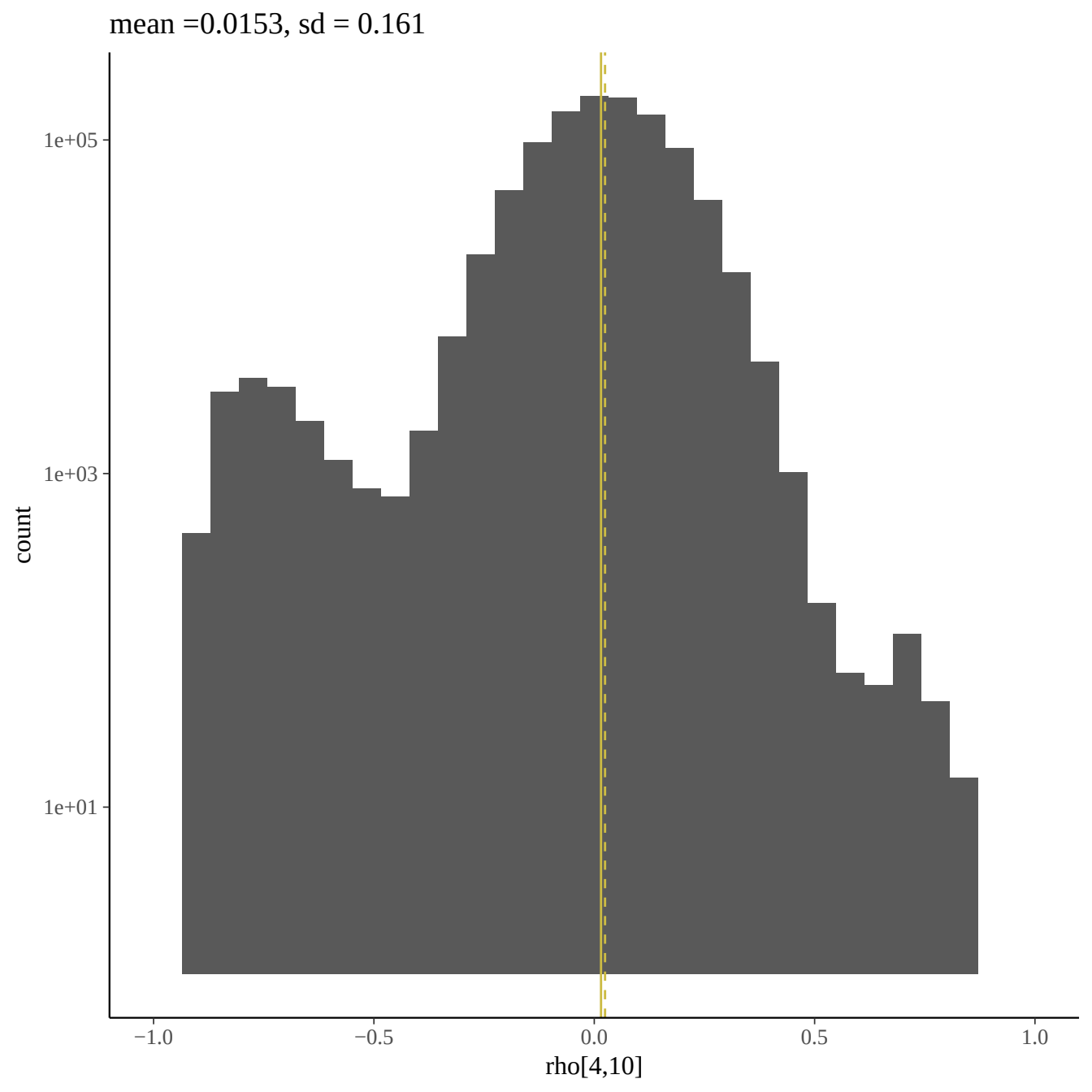

mean =0.212, sd = 0.157
1e+05
1e+03
1e+01
count
−1.0
−0.5
0.0
0.5
1.0
rho[7,9]

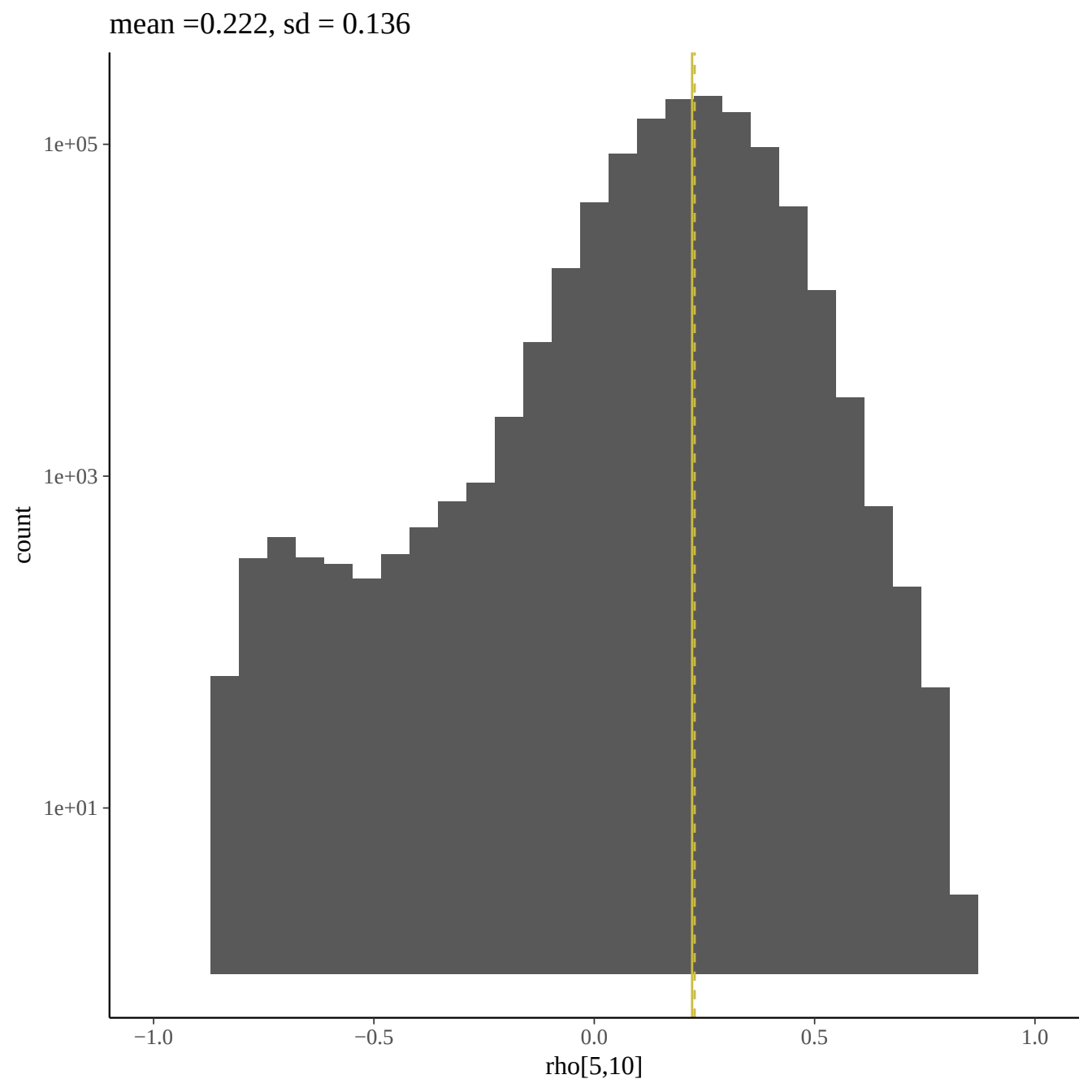

mean =0.102, sd = 0.168
1e+05
1e+03
1e+01
count
−1.0
−0.5
0.0
0.5
1.0
rho[8,9]

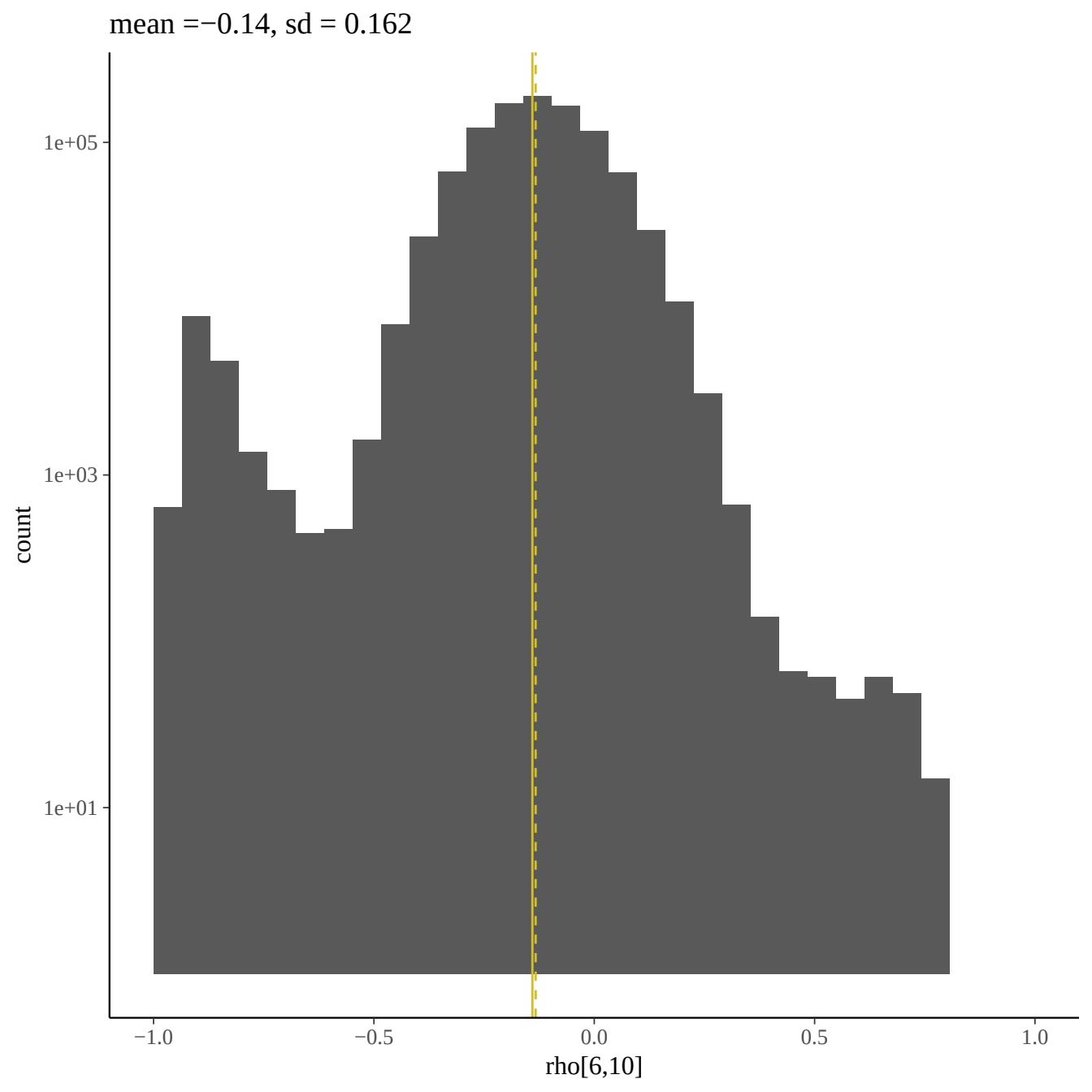

mean =−0.0424, sd = 0.173
1e+05
1e+03
1e+01
count
−1.0
−0.5
0.0
0.5
1.0
rho[1,10]

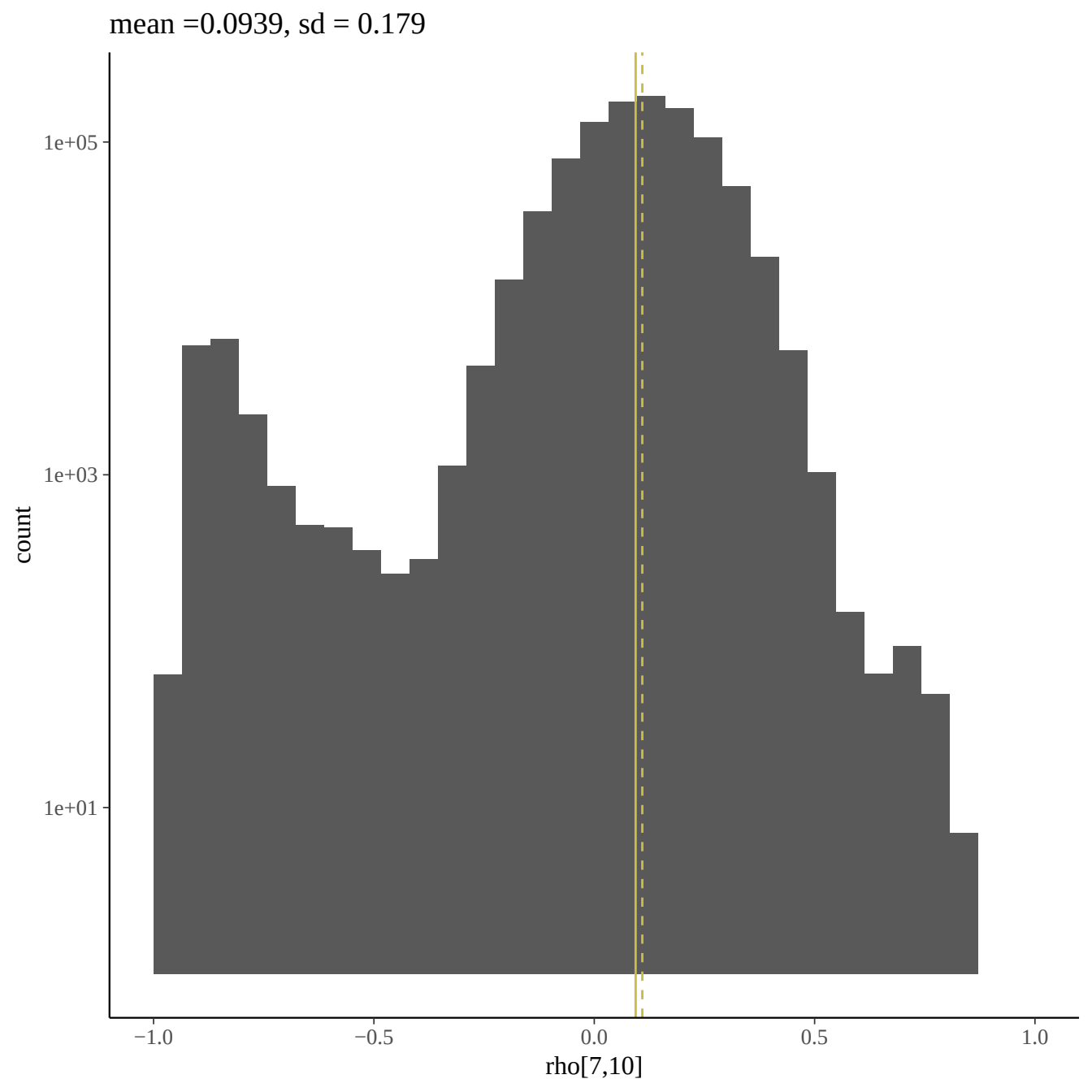

mean =0.252, sd = 0.152
1e+05
1e+03
1e+01
count
−1.0
−0.5
0.0
0.5
1.0
rho[2,10]

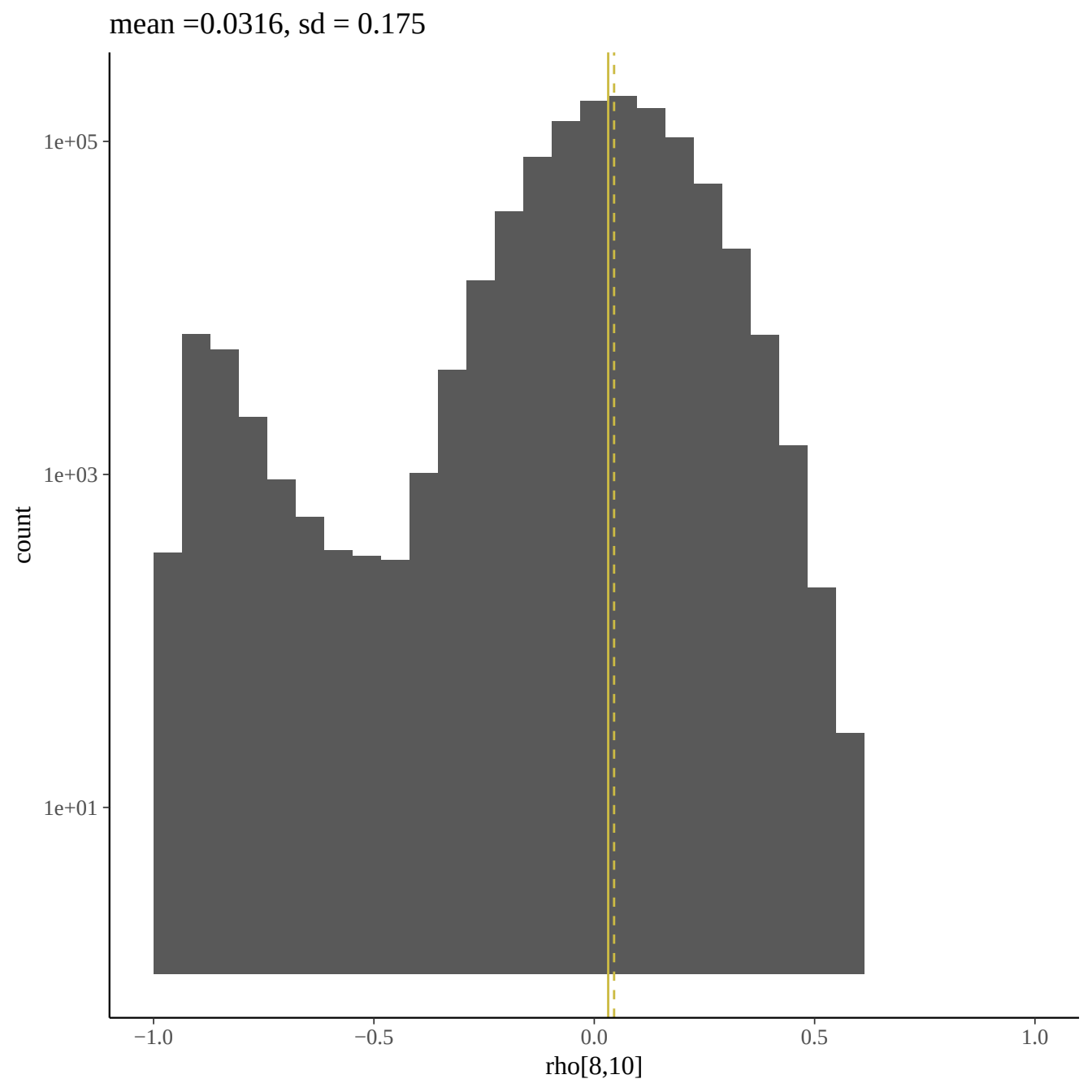

mean =0.243, sd = 0.153
1e+05
1e+03
1e+01
count
−1.0
−0.5
0.0
0.5
1.0
rho[3,10]

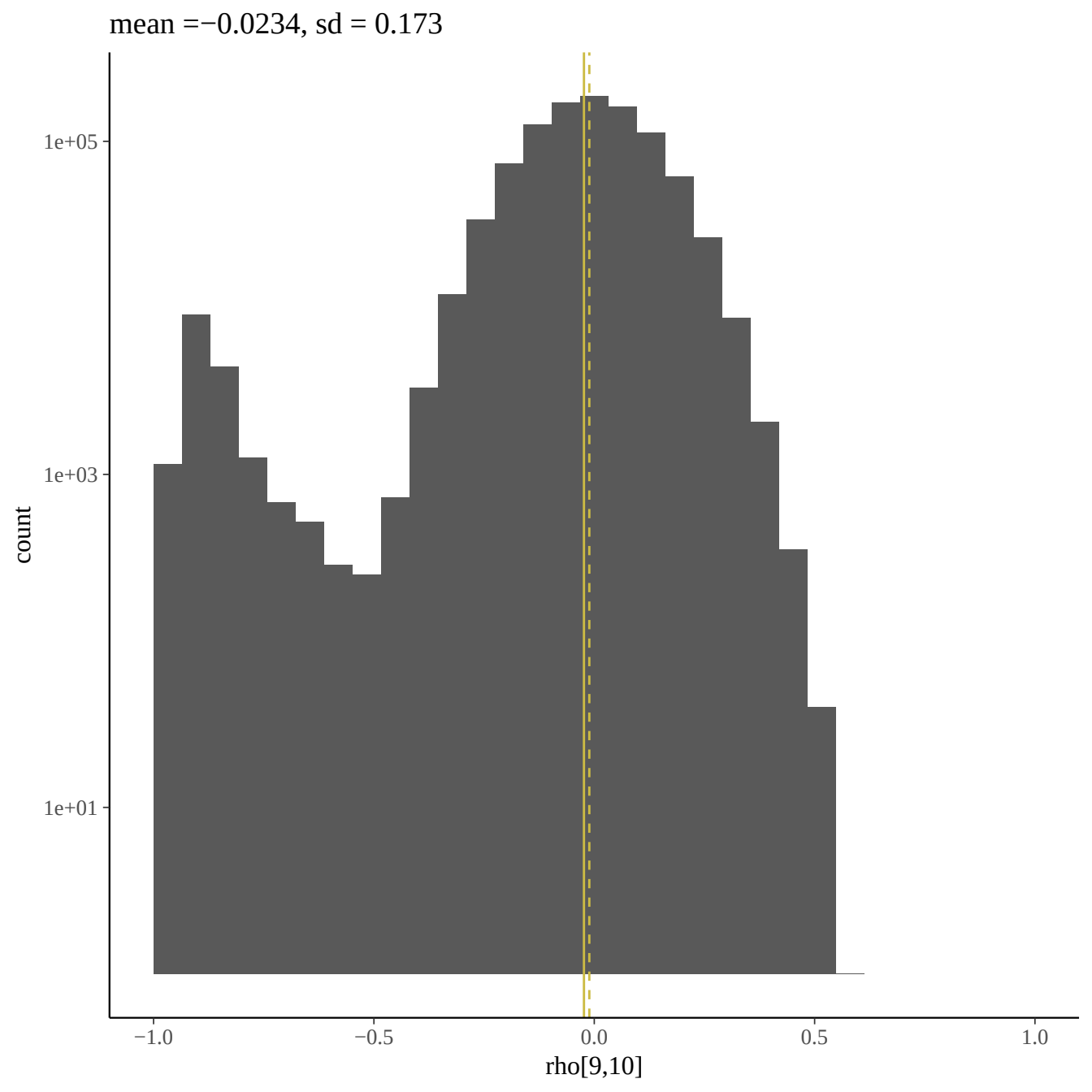

mean =0.0153, sd = 0.161
1e+05
1e+03
1e+01
count
−1.0
−0.5
0.0
0.5
1.0
rho[4,10]

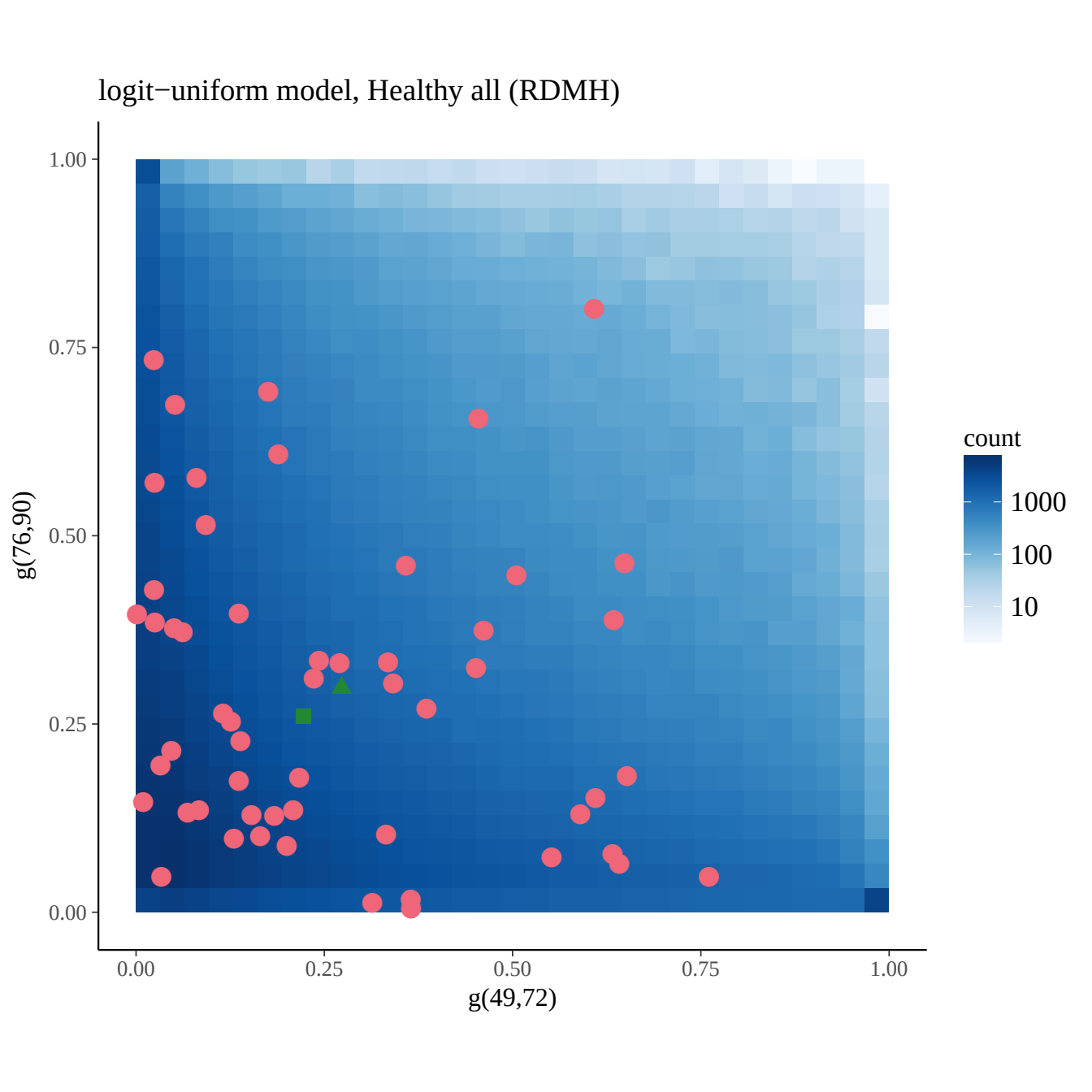

mean =0.222, sd = 0.136
1e+05
1e+03
1e+01
count
−1.0
−0.5
0.0
0.5
1.0
rho[5,10]

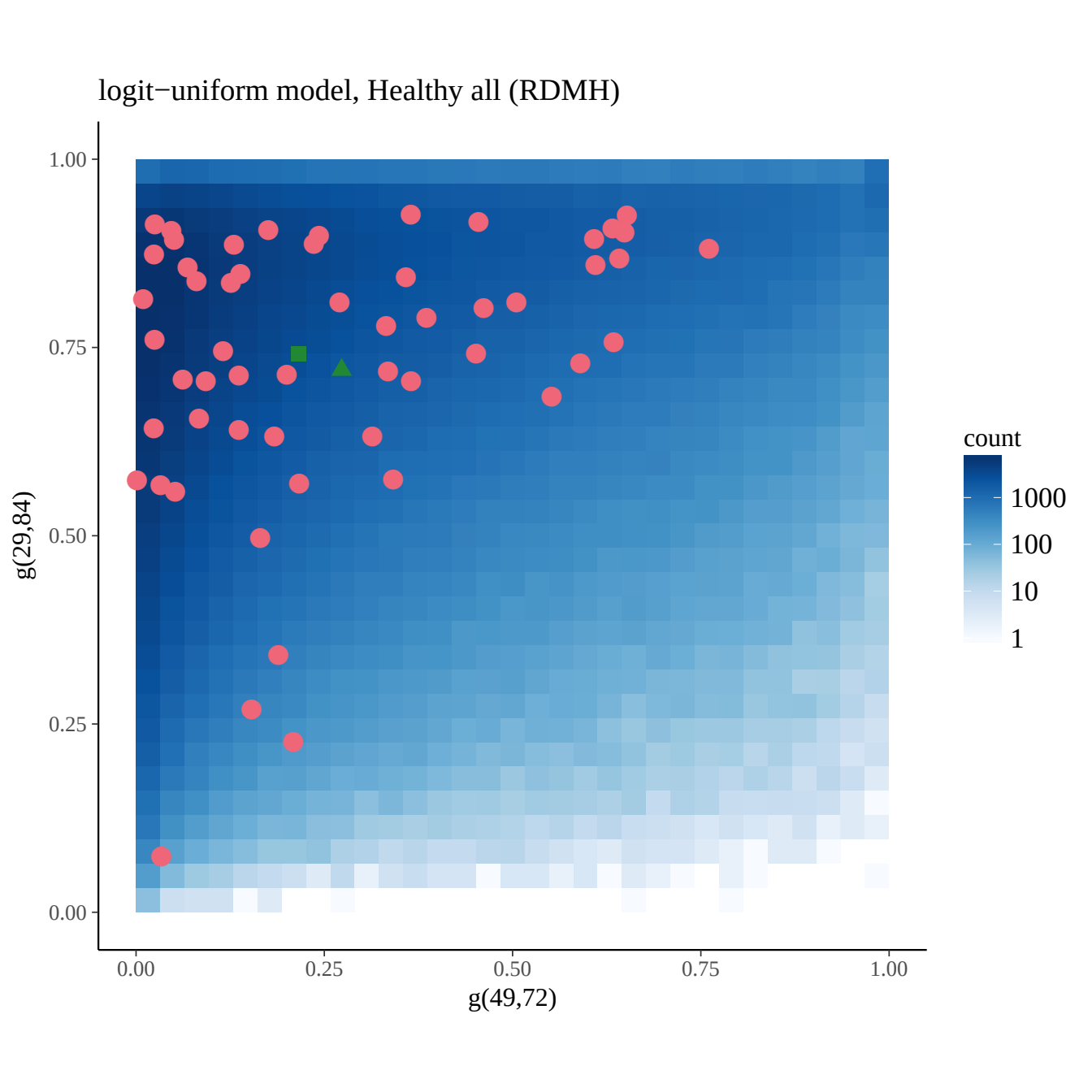

mean =−0.14, sd = 0.162
1e+05
1e+03
1e+01
count
−1.0
−0.5
0.0
0.5
1.0
rho[6,10]

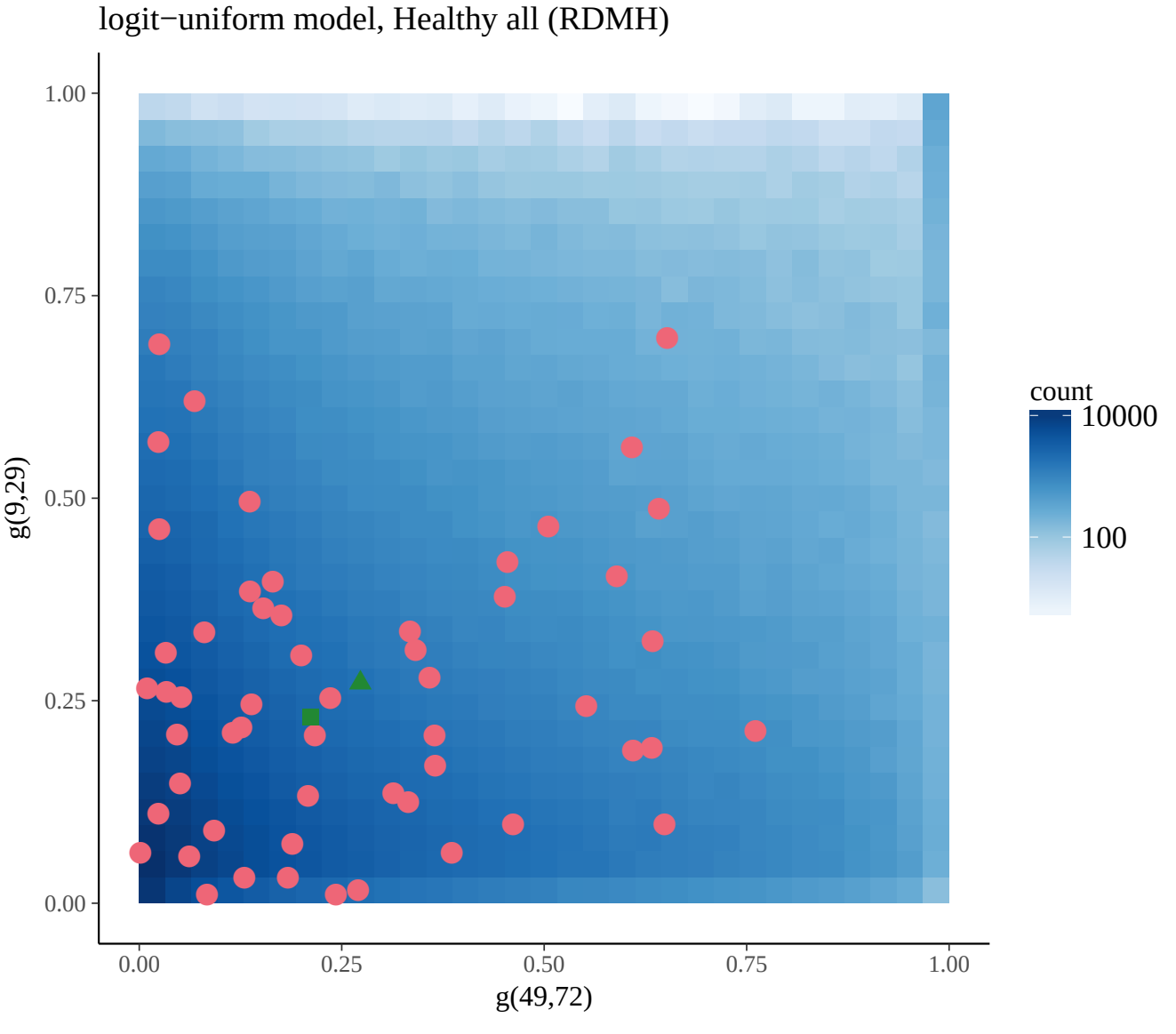

mean =0.0939, sd = 0.179
1e+05
1e+03
1e+01
count
−1.0
−0.5
0.0
0.5
1.0
rho[7,10]

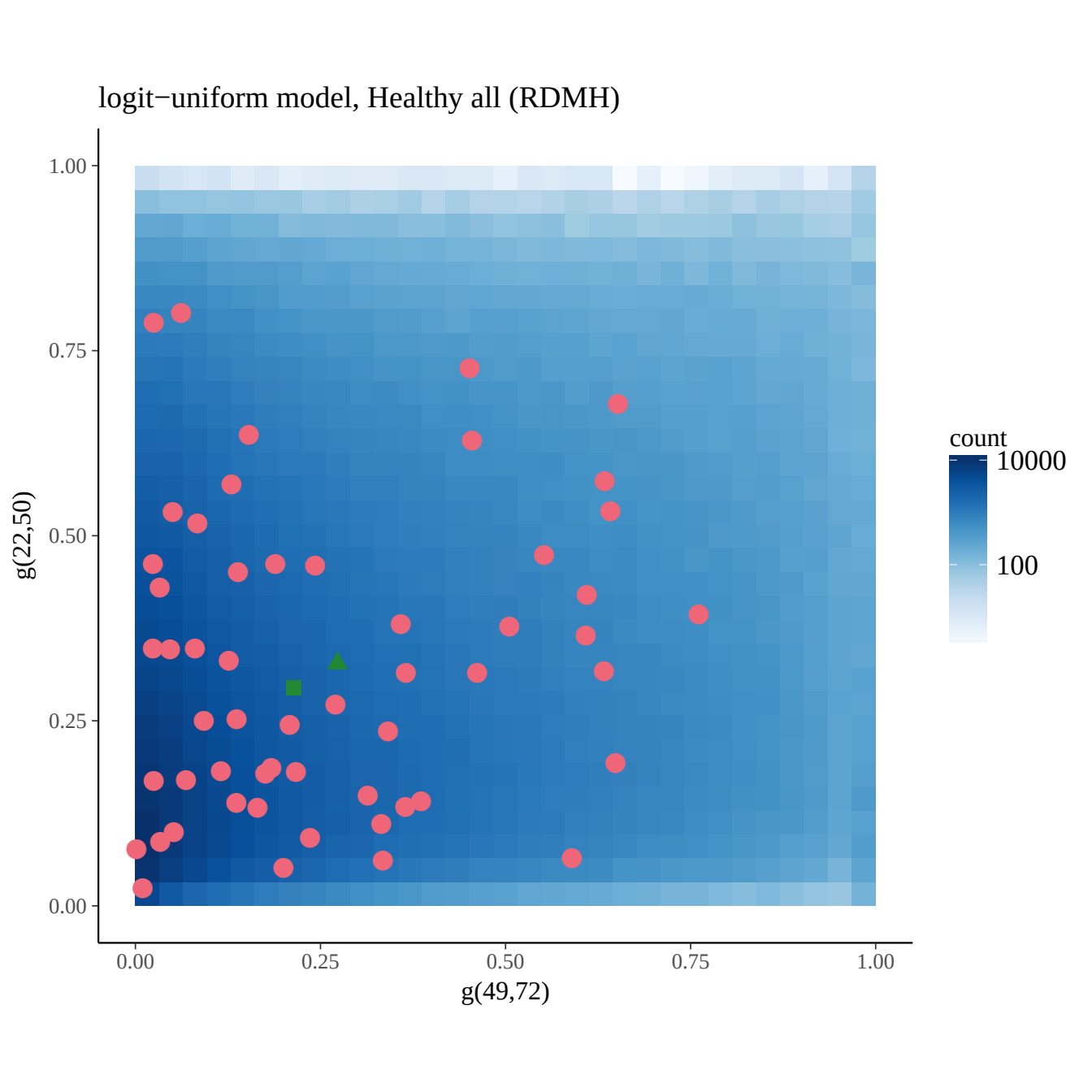

mean =0.0316, sd = 0.175
1e+05
1e+03
1e+01
count
−1.0
−0.5
0.0
0.5
1.0
rho[8,10]

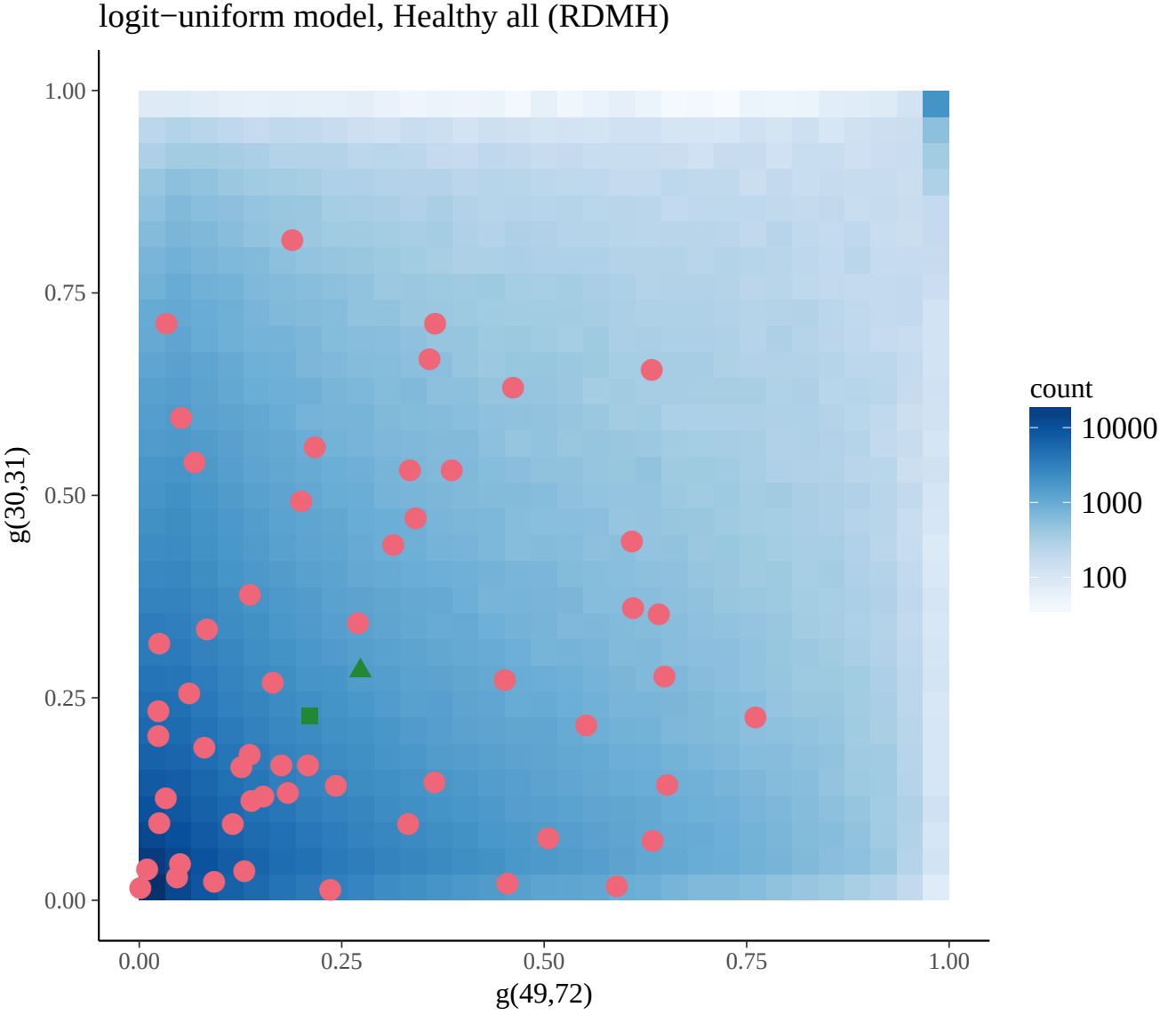

mean =−0.0234, sd = 0.173
1e+05
1e+03
1e+01
count
−1.0
−0.5
0.0
0.5
1.0
rho[9,10]

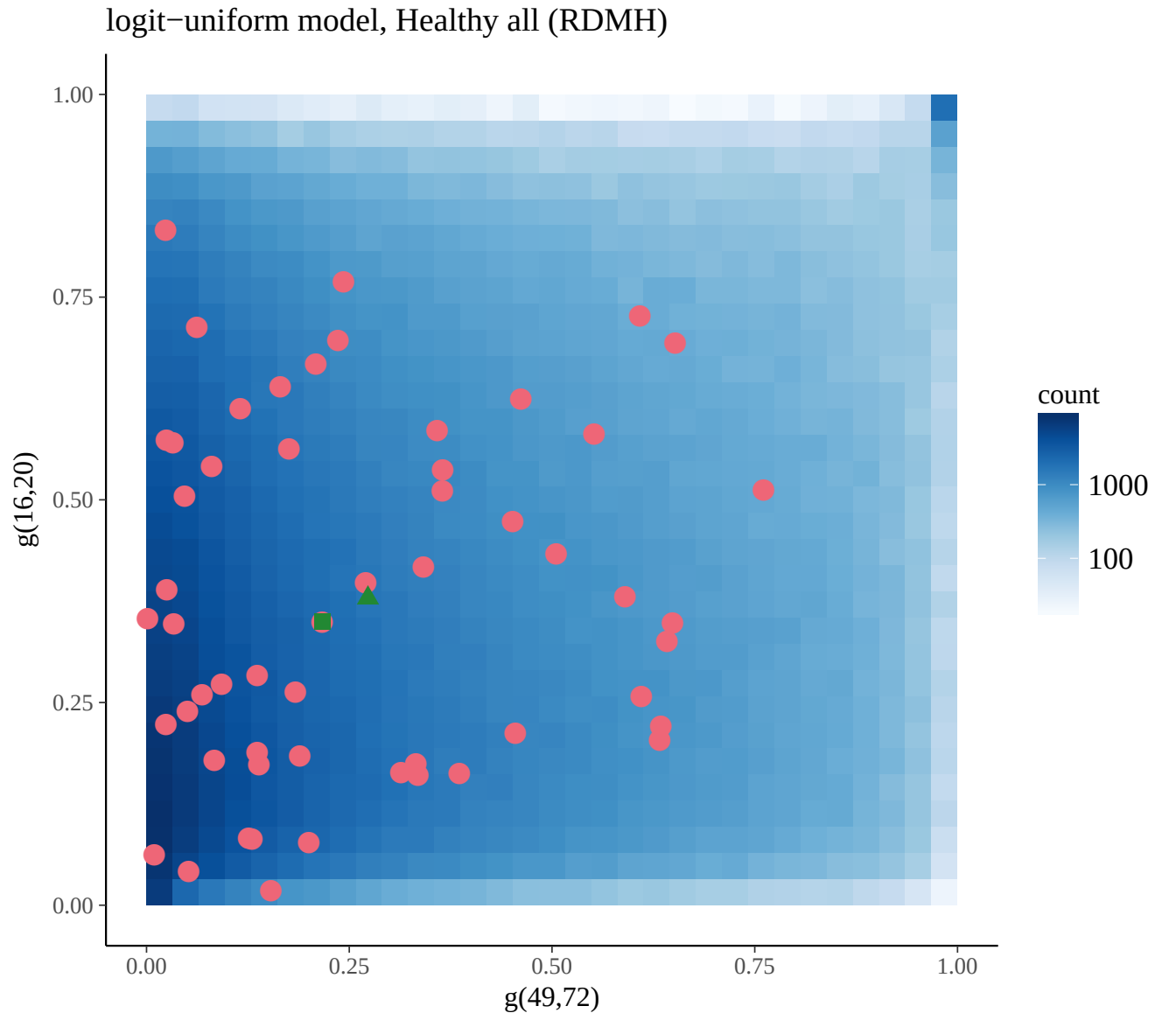

logit−uniform model, Healthy all (RDMH)
1.00
0.75
0.50
0.25
0.00
g(76,90)
0.00
0.25
0.50
0.75
1.00
g(49,72)
count
1000
100
10


logit−uniform model, Healthy all (RDMH)
1.00
0.75
0.50
0.25
0.00
g(29,84)
0.00
0.25
0.50
0.75
1.00
g(49,72)
count
1000
100
10
1


logit−uniform model, Healthy all (RDMH)
1.00
0.75
0.50
0.25
0.00
g(9,29)
0.00
0.25
0.50
0.75
1.00
g(49,72)
count
10000
100


logit−uniform model, Healthy all (RDMH)
1.00
0.75
0.50
0.25
0.00
g(22,50)
0.00
0.25
0.50
0.75
1.00
g(49,72)
count
10000
100


logit−uniform model, Healthy all (RDMH)
1.00
0.75
0.50
0.25
0.00
g(30,31)
0.00
0.25
0.50
0.75
1.00
g(49,72)
count
10000
1000
100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0.50
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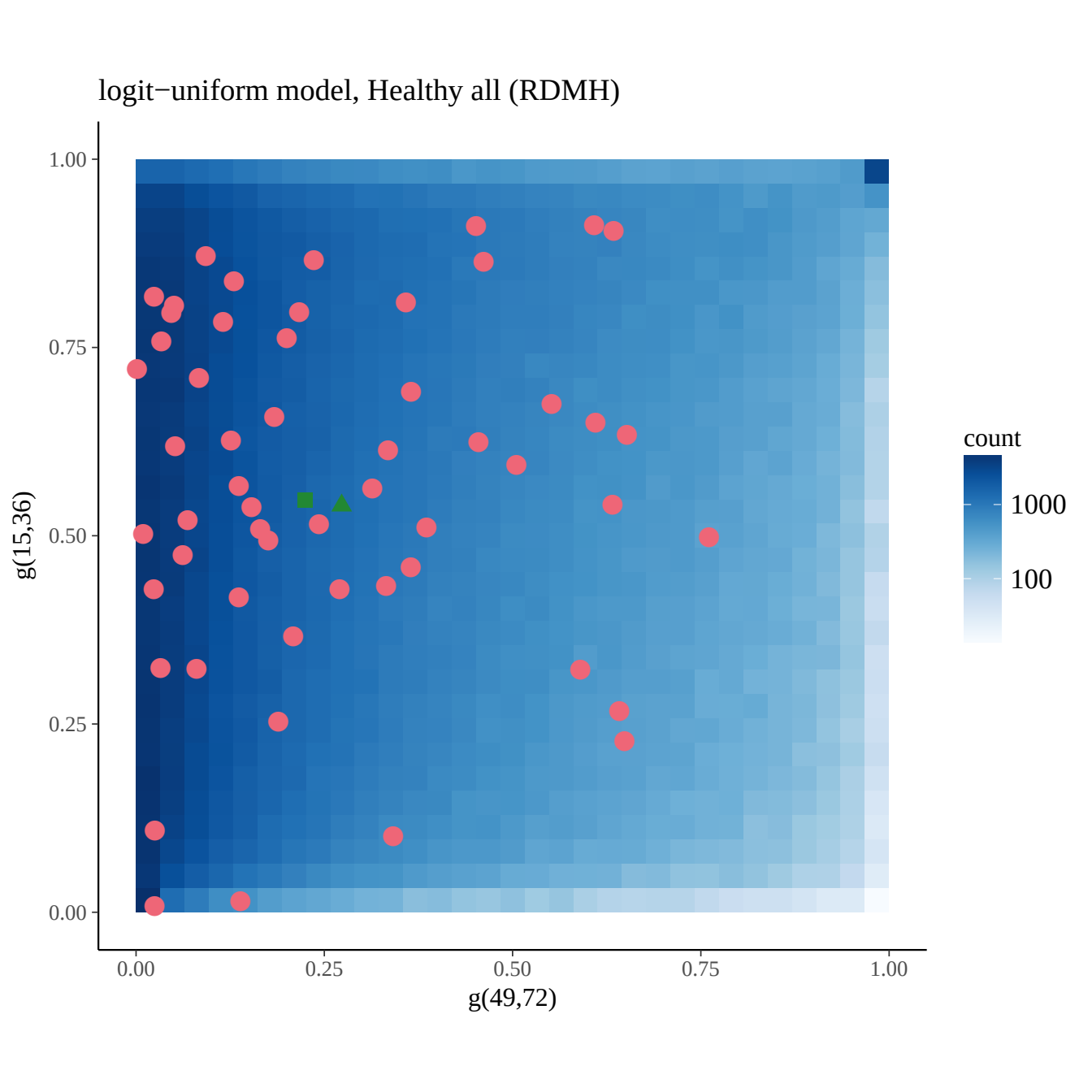
0.25
0.50
0.75
1.00
g(49,72)
count
1000
100


logit−uniform model, Healthy all (RDMH)
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0.50
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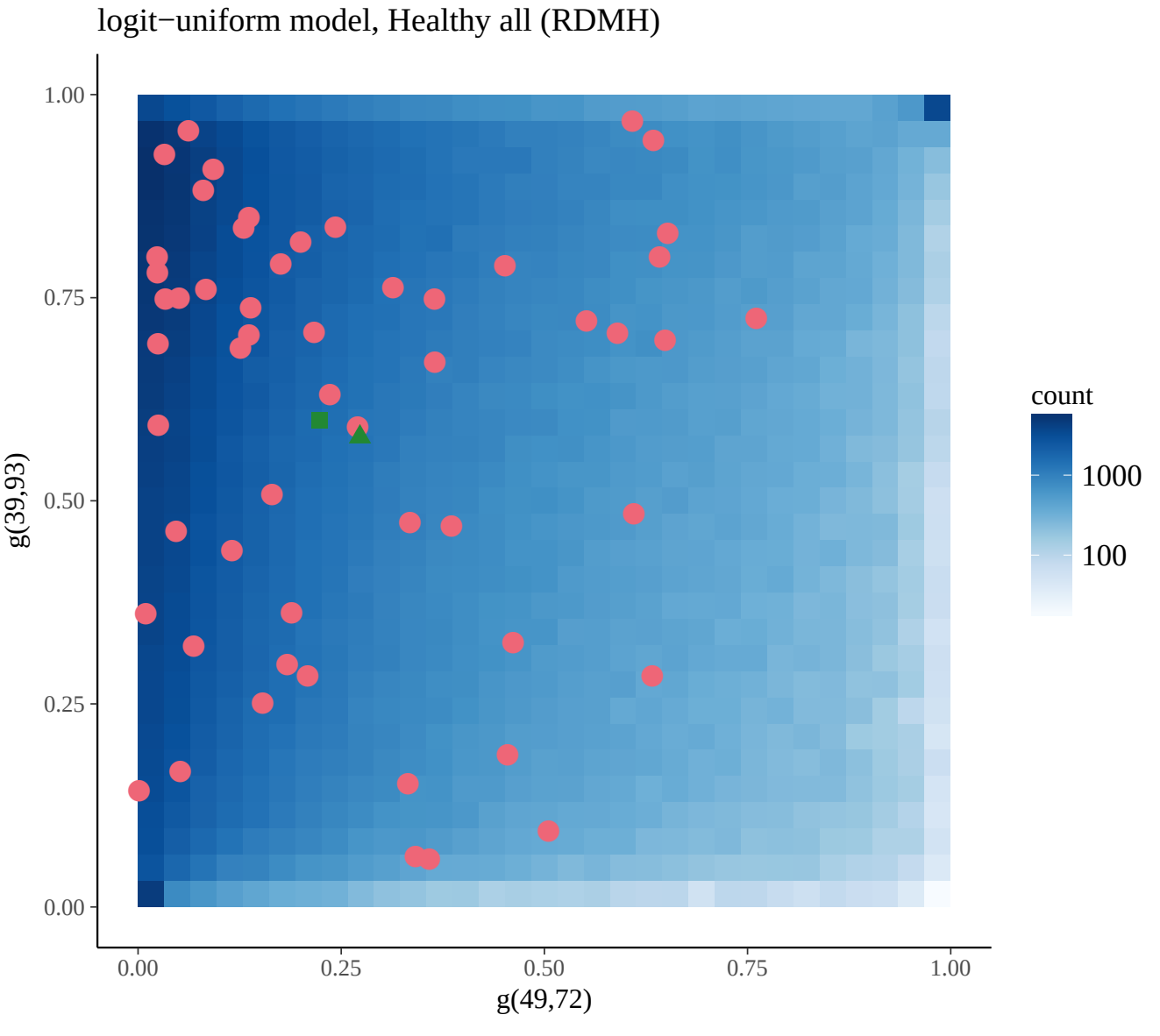
0.00
0.25
0.50
0.75
1.00
g(49,72)
count
1000
100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0.50
0.75
1.00
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count
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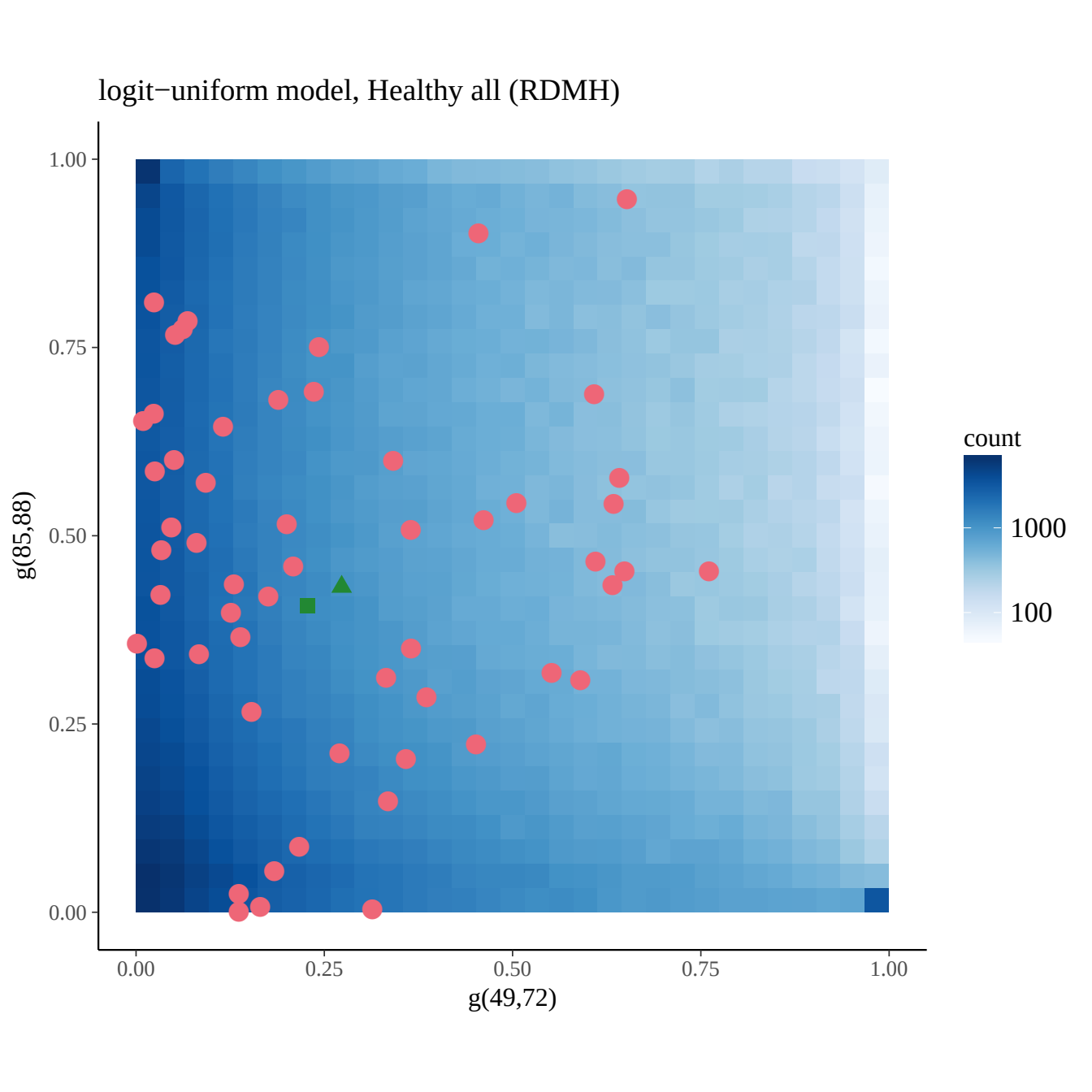
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0.75
1.00
g(49,72)
count
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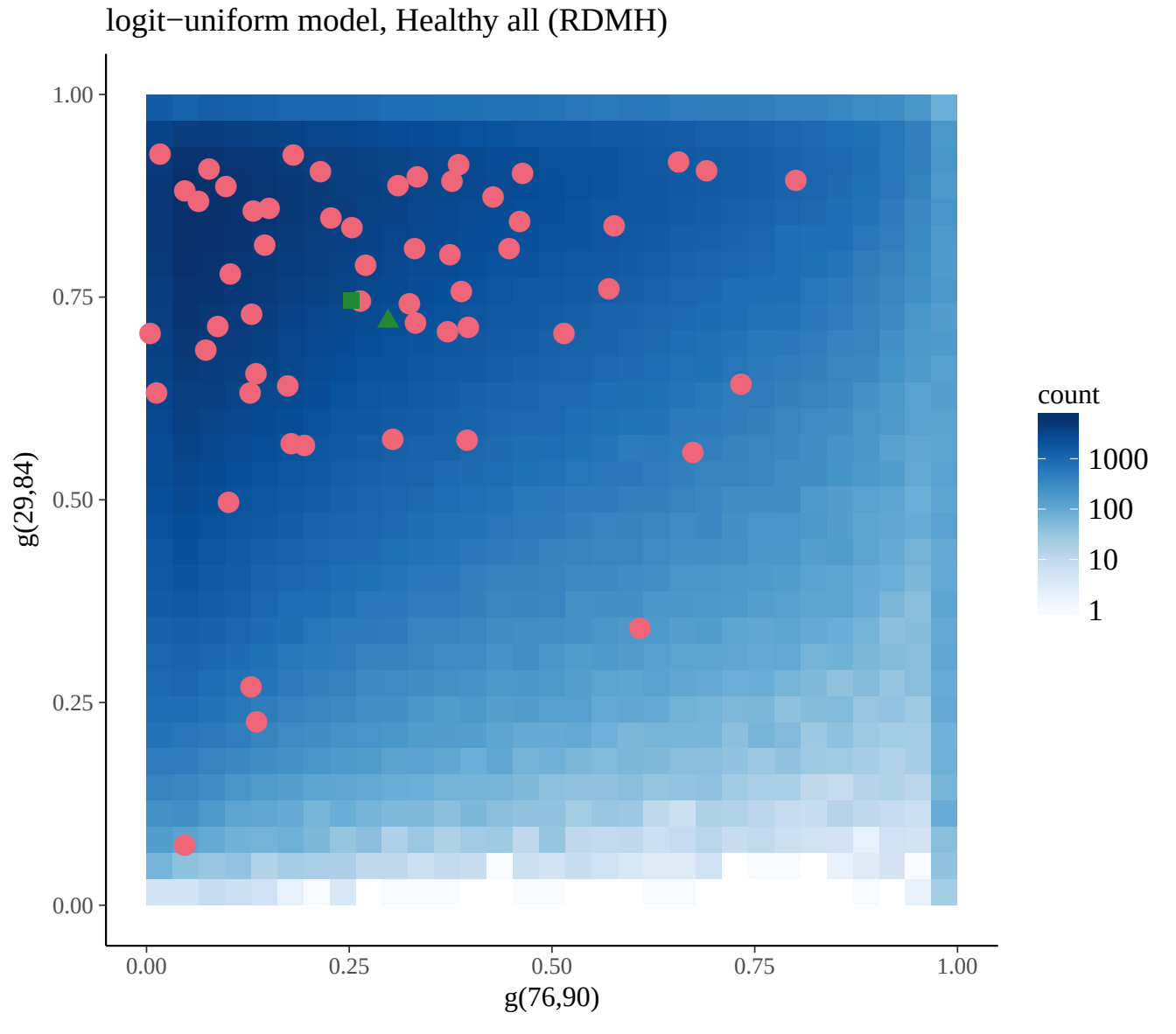
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0.50
0.25
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0.00
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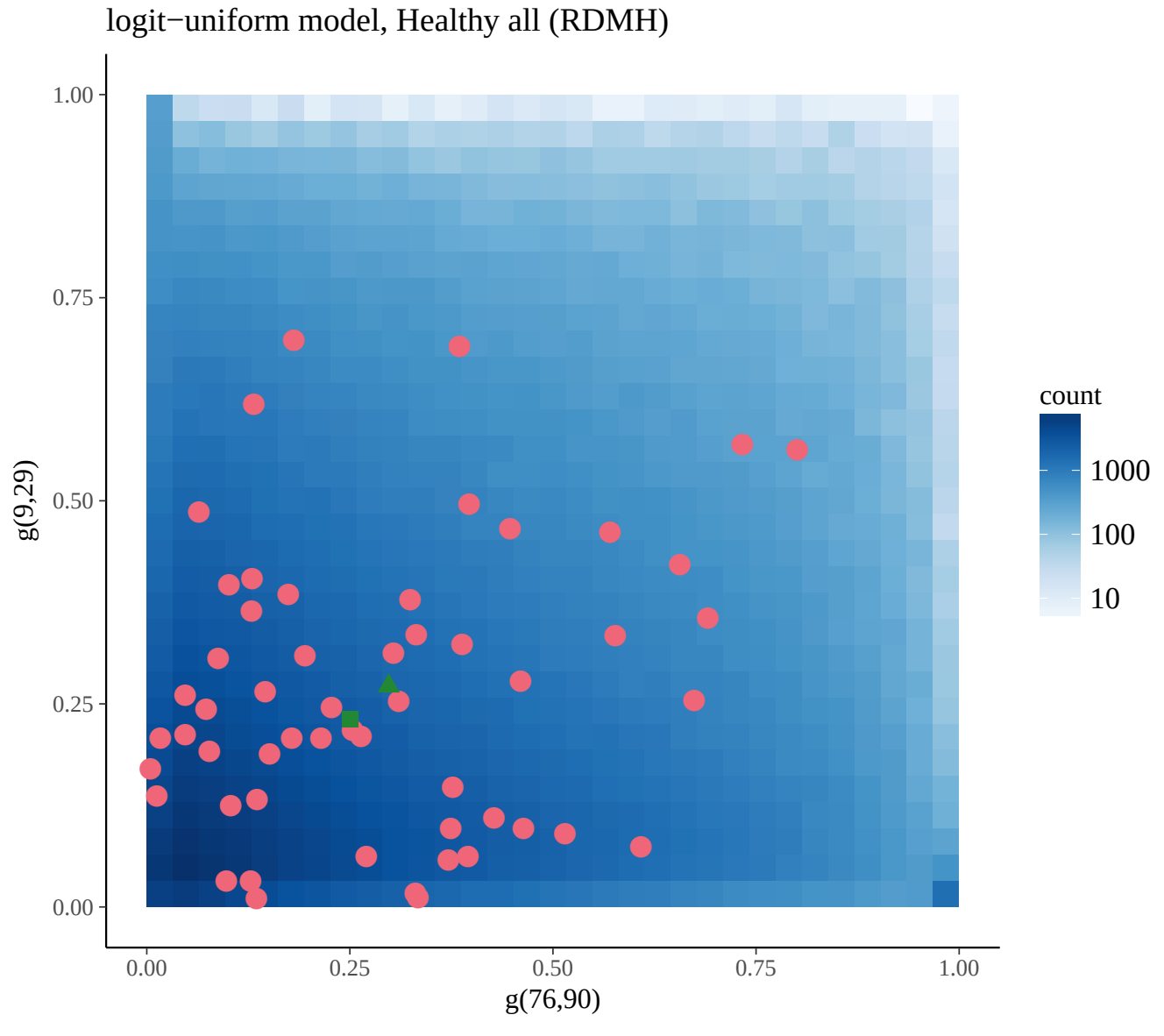
0.50
0.75
1.00
g(76,90)
count
1000
100
10
1


logit−uniform model, Healthy all (RDMH)
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0.50
0.75
1.00
g(76,90)
count
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100
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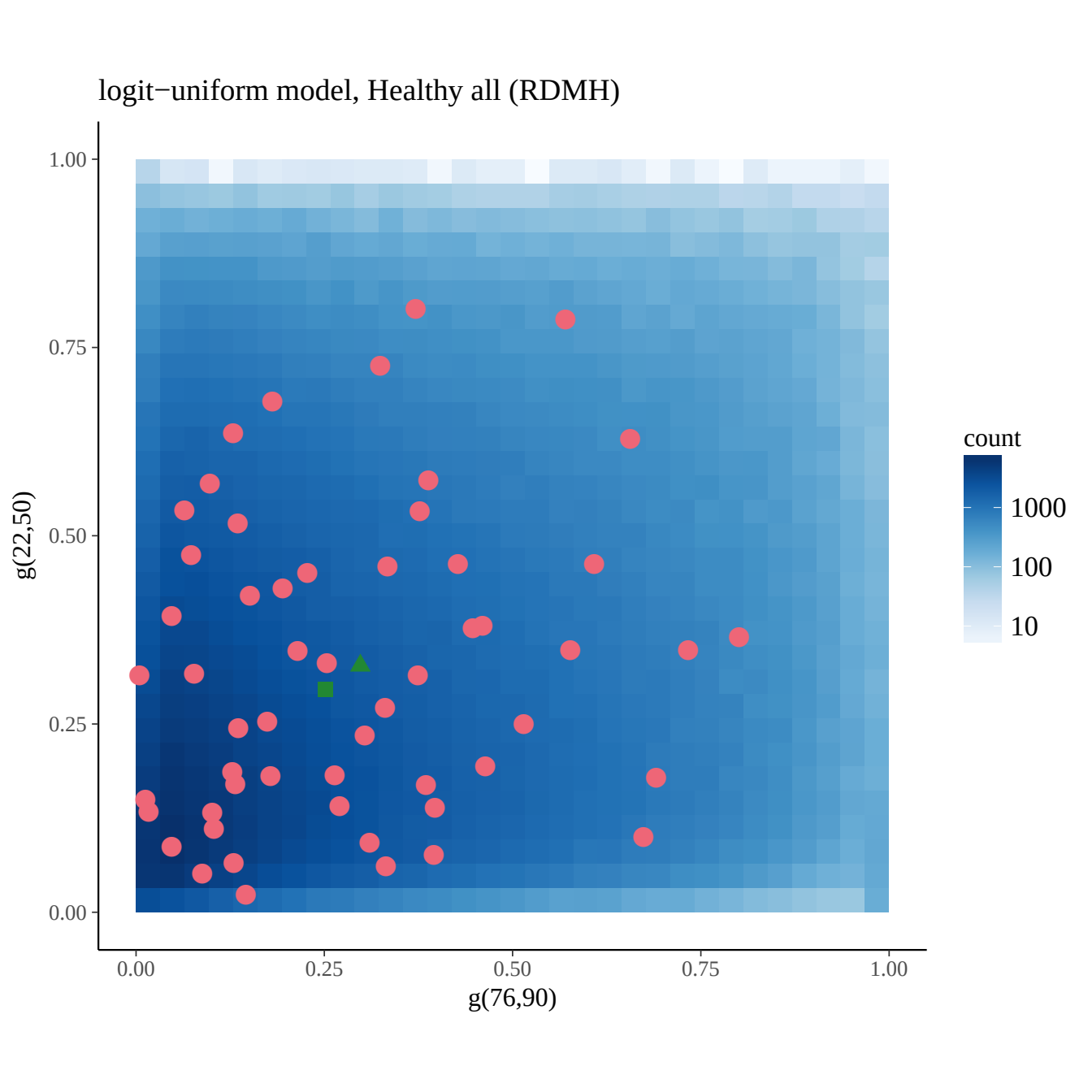
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0.25
0.50
0.75
1.00
g(76,90)
count
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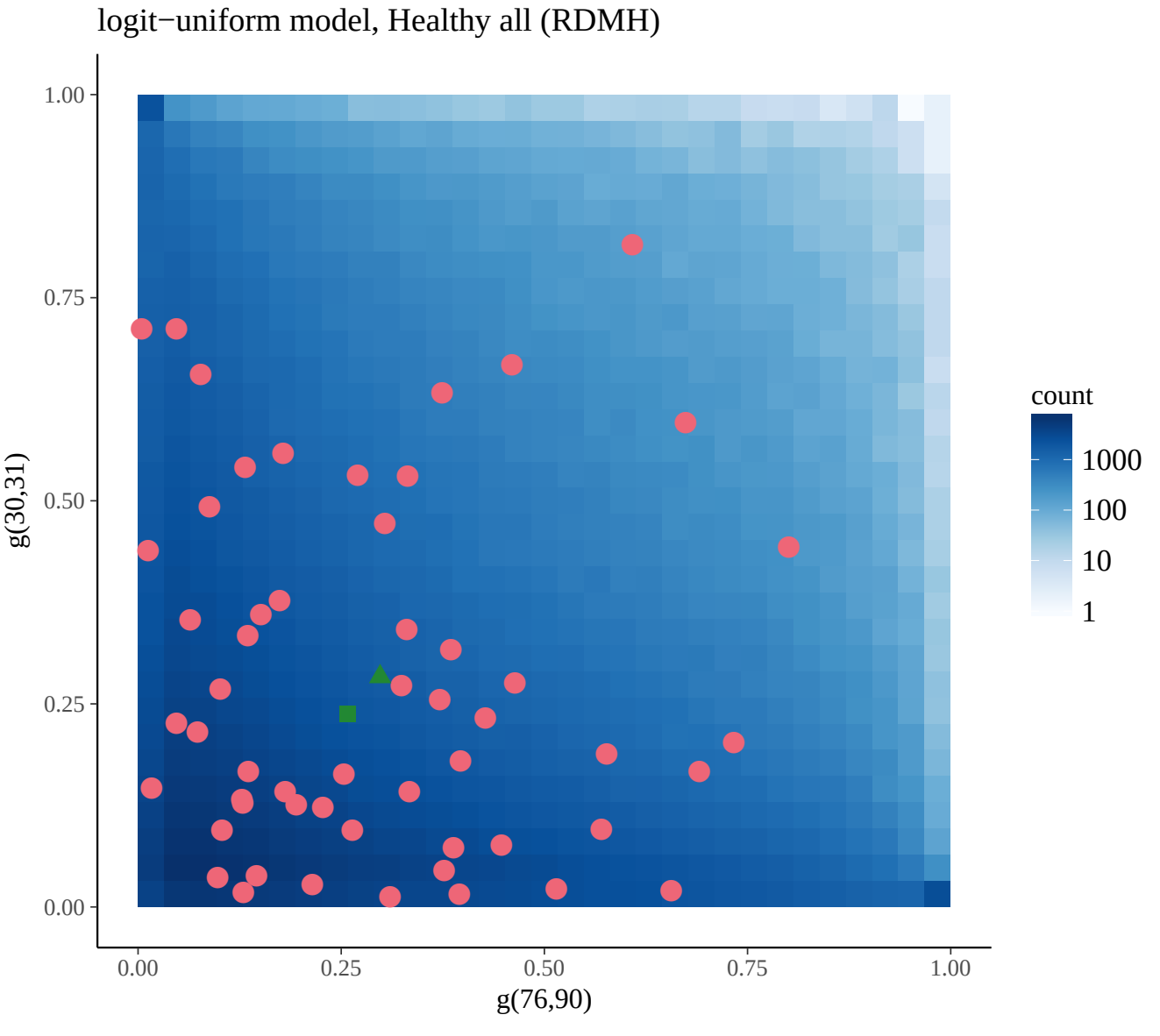
100
10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0.50
0.25
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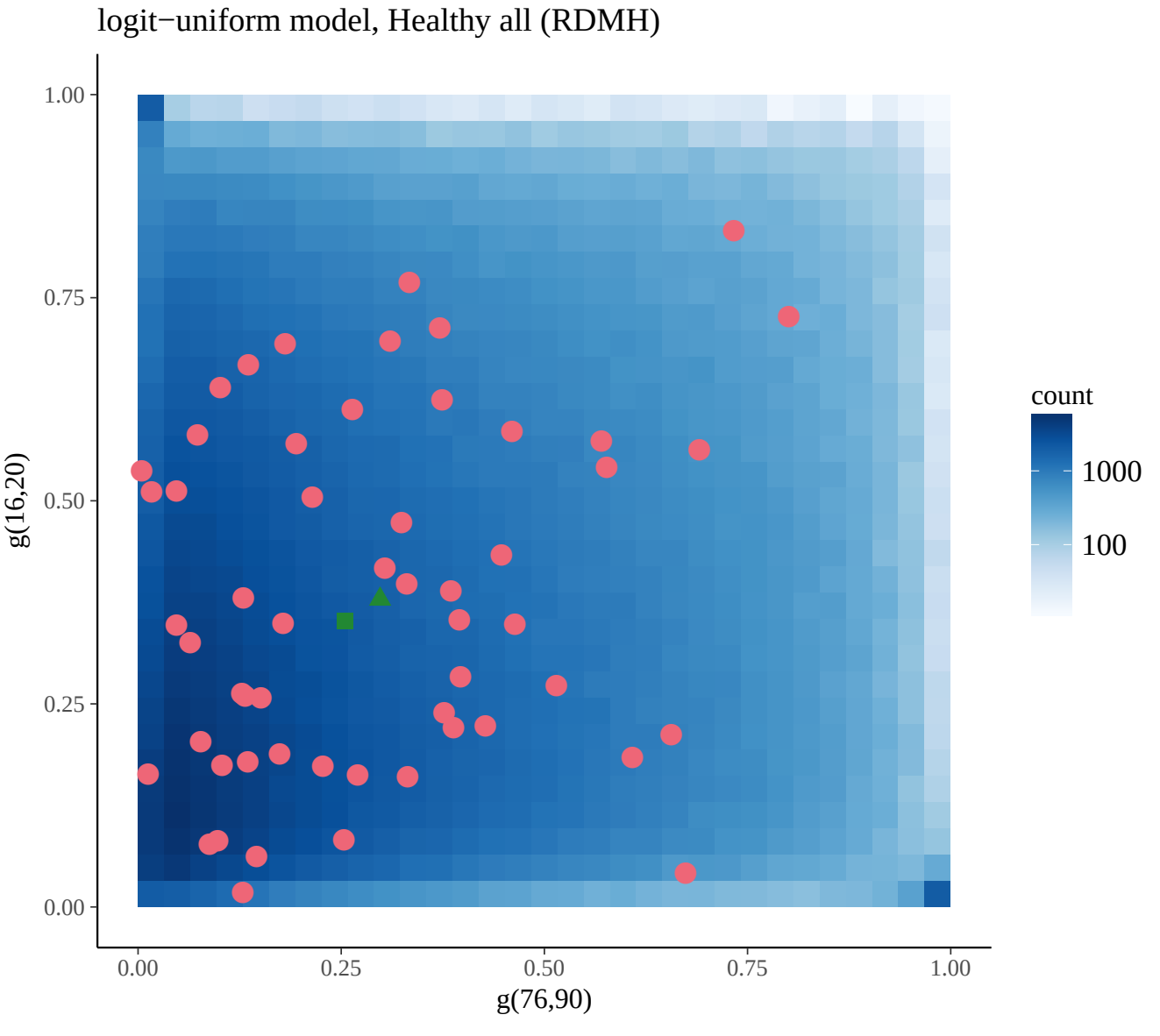
0.00
0.25
0.50
0.75
1.00
g(76,90)
count
1000
100
10
1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0.50
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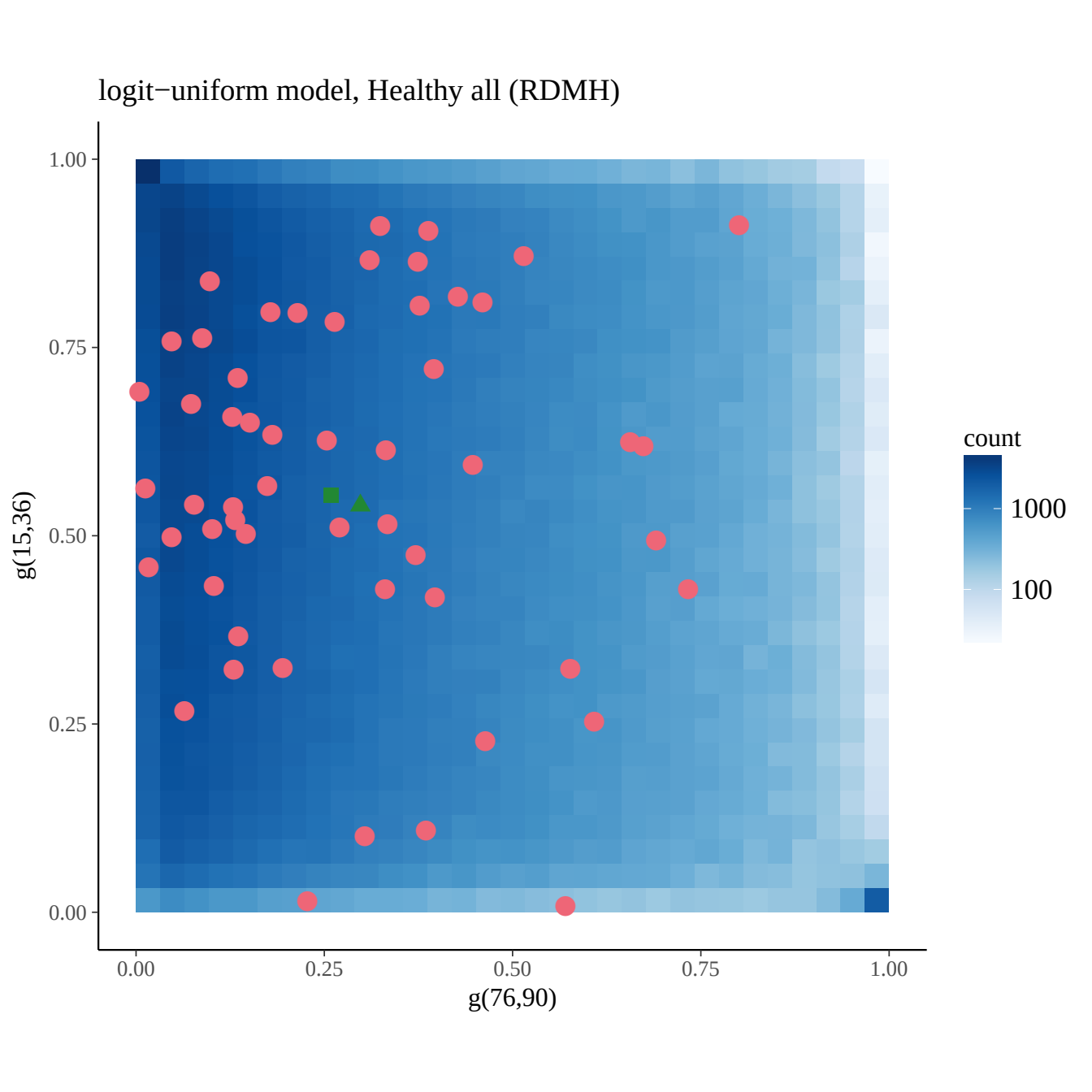
0.00
0.25
0.50
0.75
1.00
g(76,90)
count
1000
100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0.50
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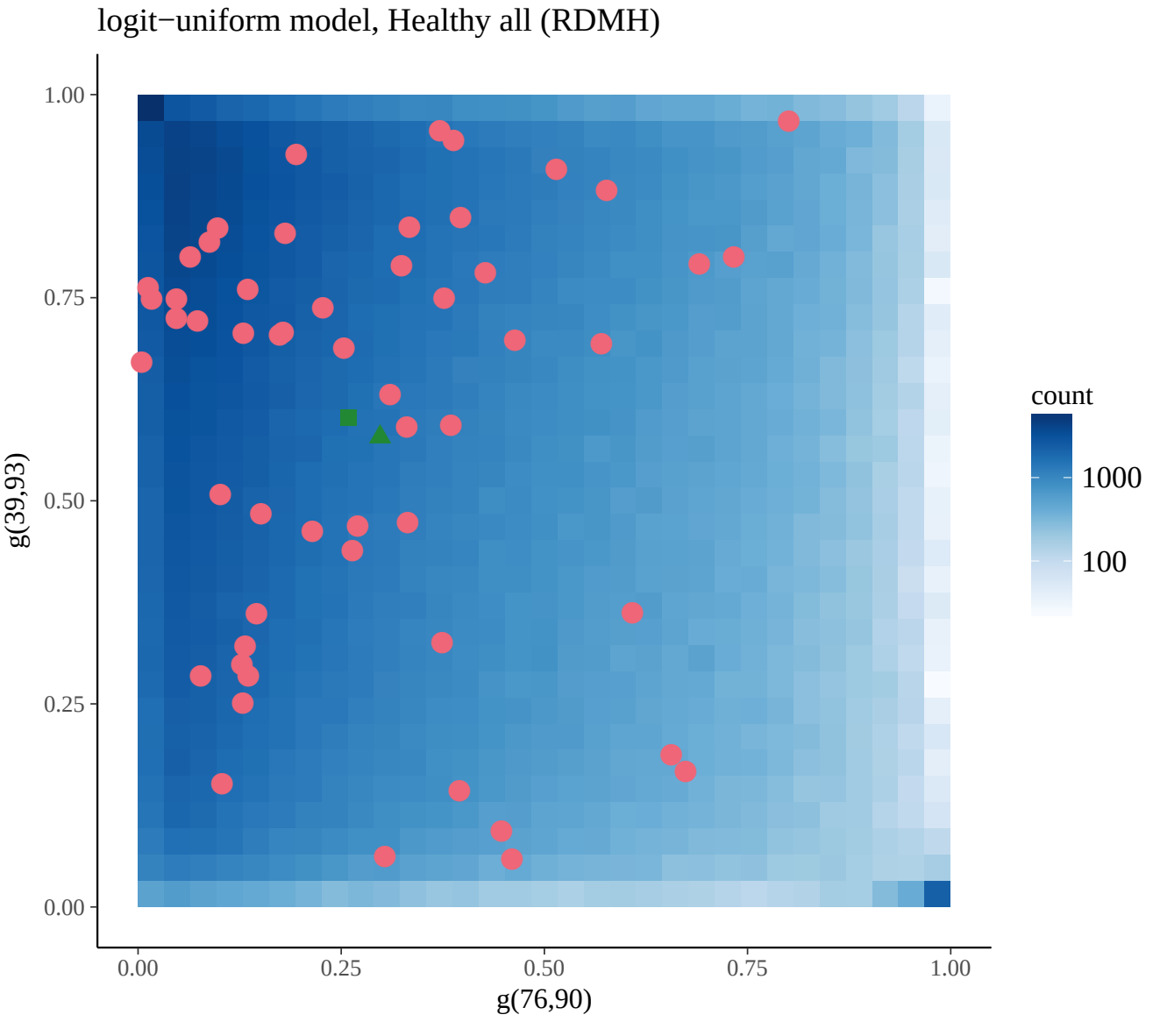
0.25
0.50
0.75
1.00
g(76,90)
count
1000
100


logit−uniform model, Healthy all (RDMH)
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0.50
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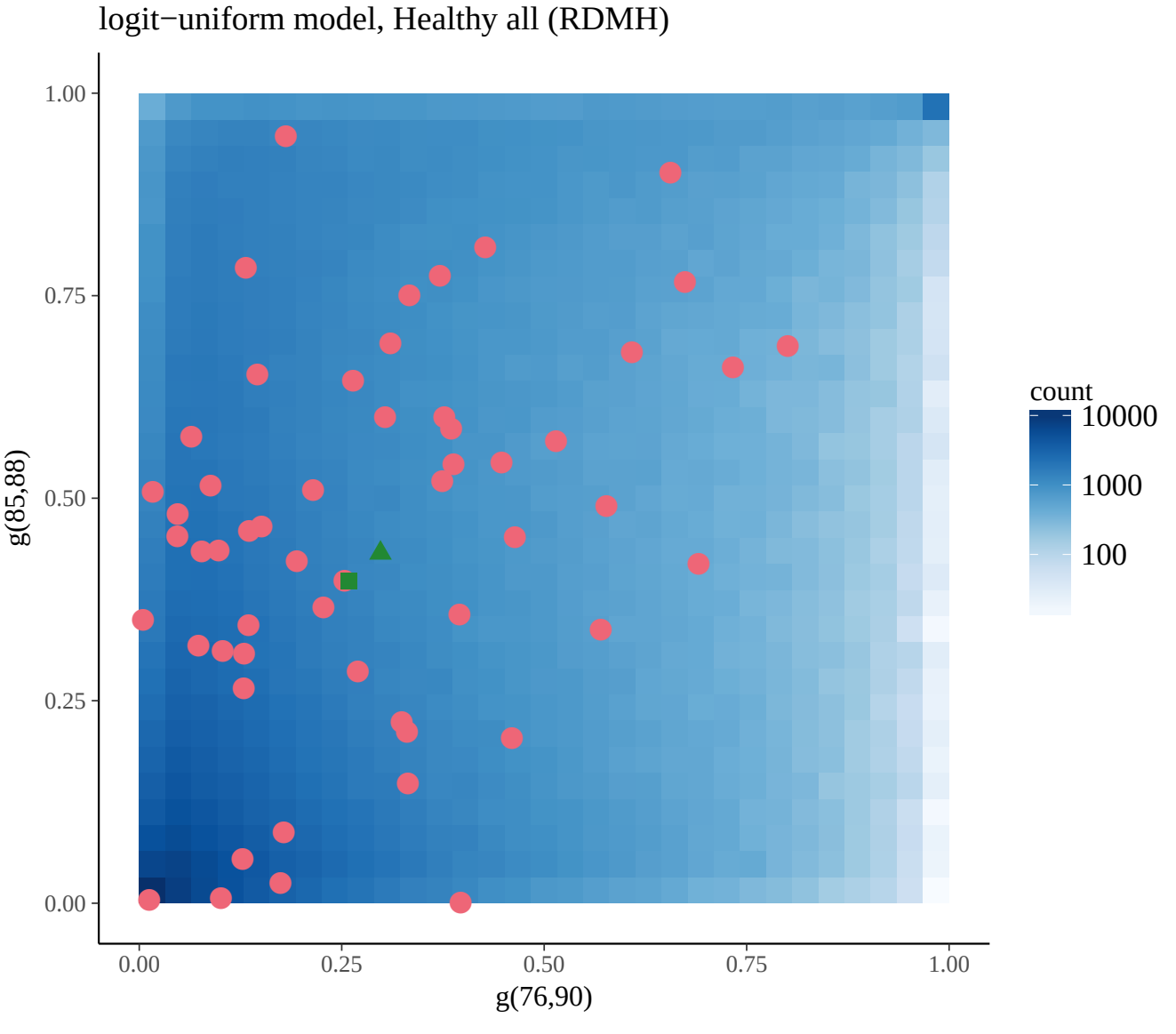
0.25
0.50
0.75
1.00
g(76,90)
count
1000
100


logit−uniform model, Healthy all (RDMH)
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0.75
0.50
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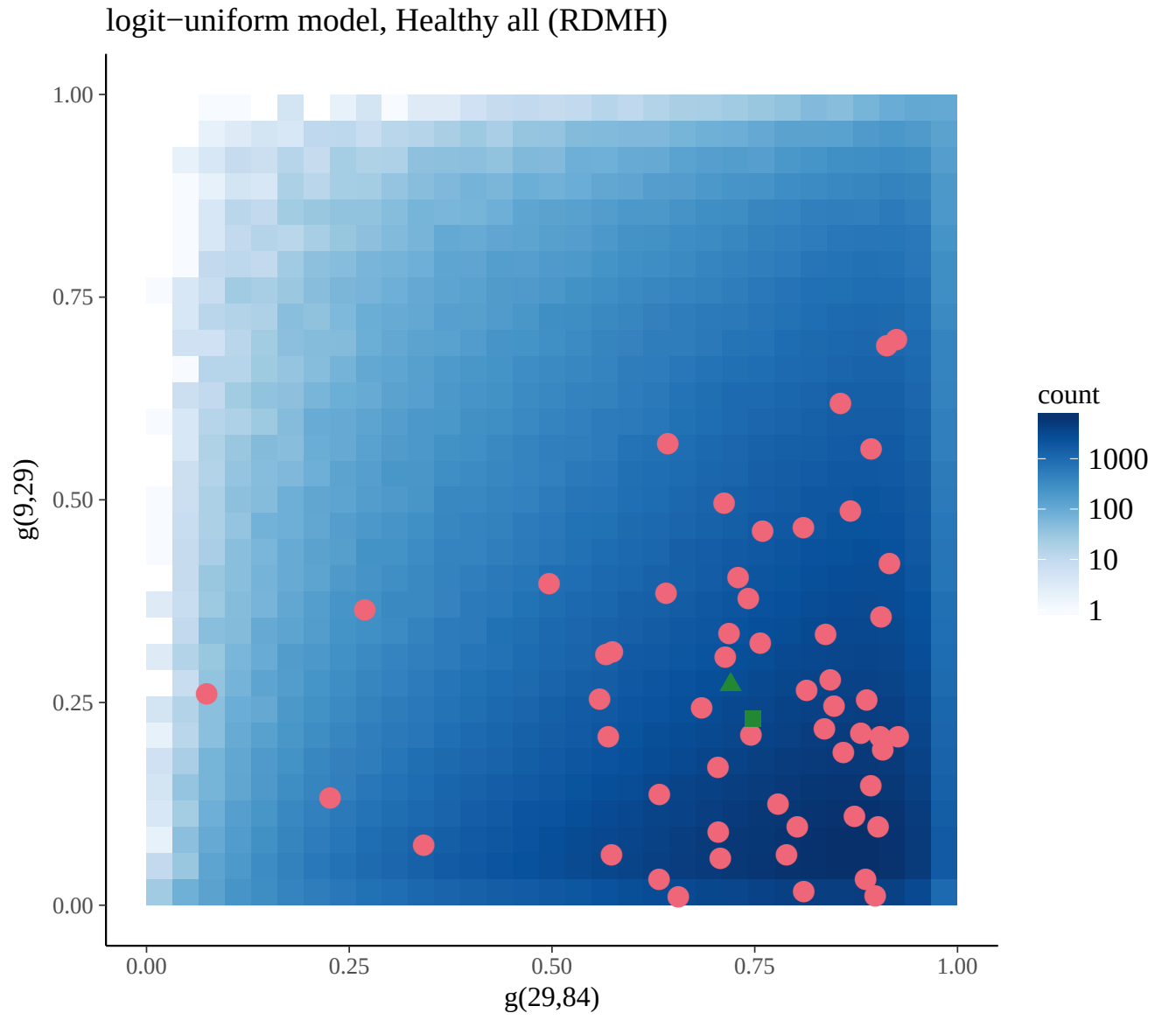
0.25
0.50
0.75
1.00
g(76,90)
count
10000
1000
100


logit−uniform model, Healthy all (RDMH)
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0.50
0.25
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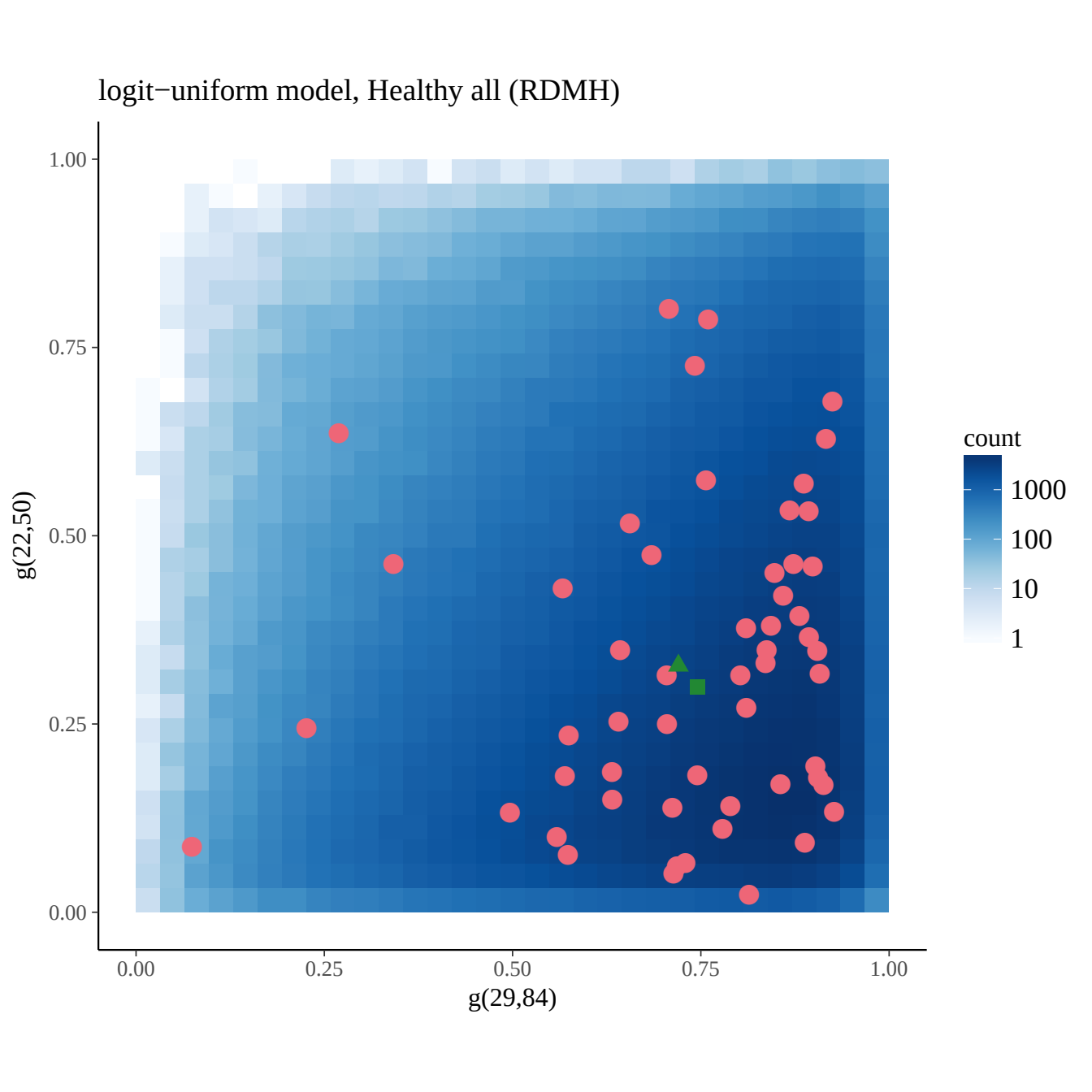
0.00
0.25
0.50
0.75
1.00
g(29,84)
count
1000
100
10
1


logit−uniform model, Healthy all (RDMH)
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0.75
0.50
0.25
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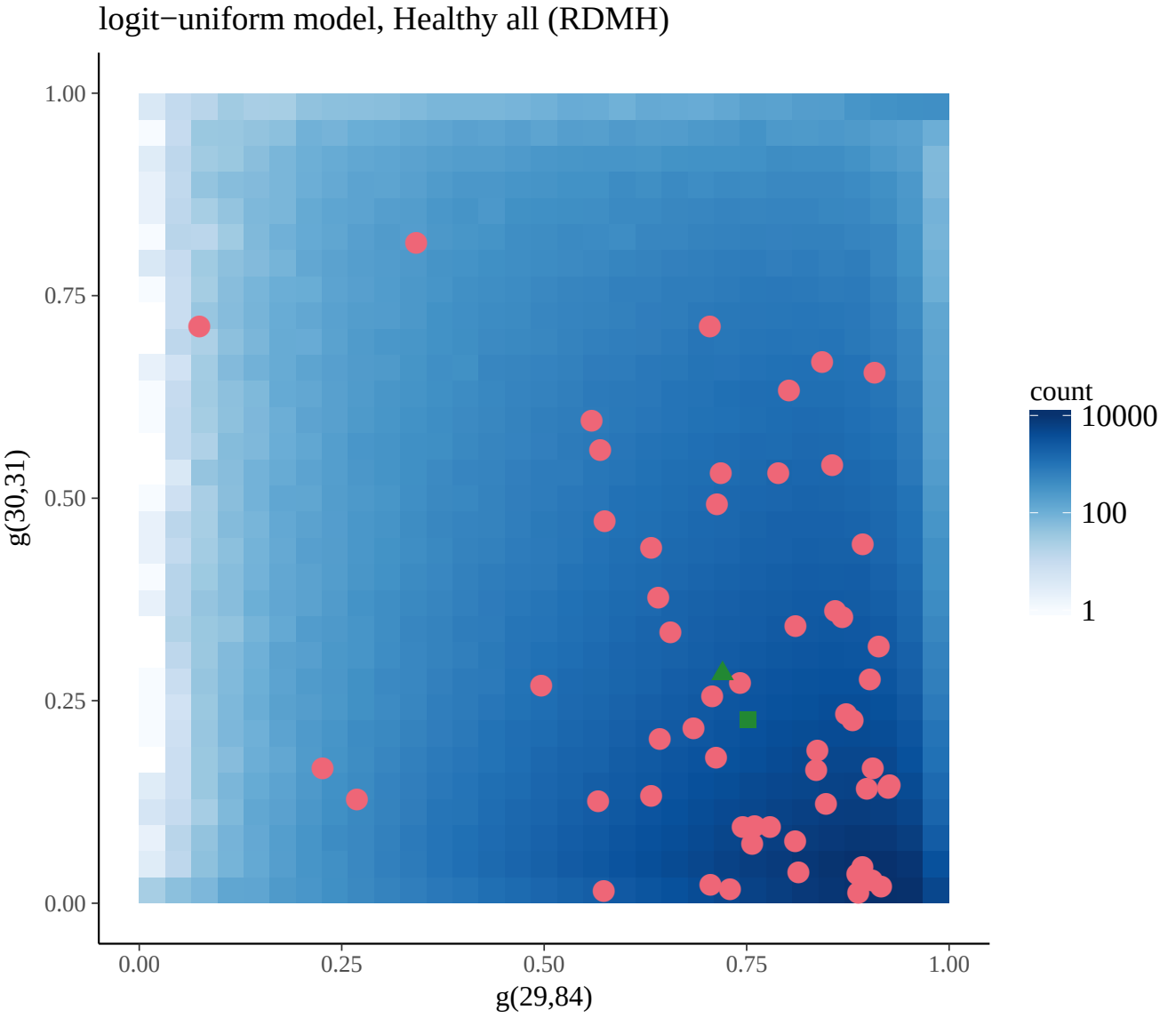
0.00
0.25
0.50
0.75
1.00
g(29,84)
count
1000
100
10
1


logit−uniform model, Healthy all (RDMH)
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0.75
0.50
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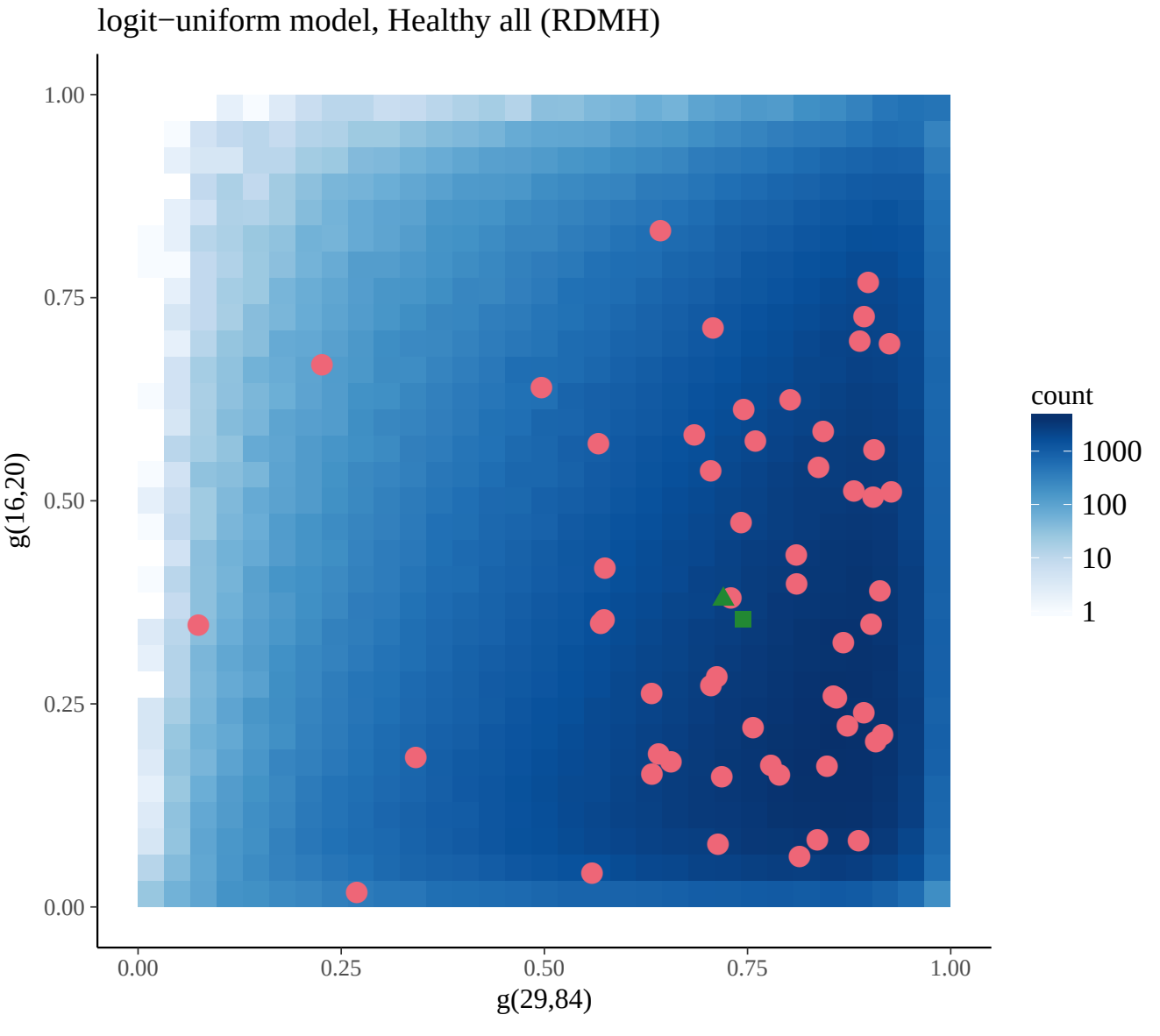
0.00
0.25
0.50
0.75
1.00
g(29,84)
count
10000
100
1


logit−uniform model, Healthy all (RDMH)
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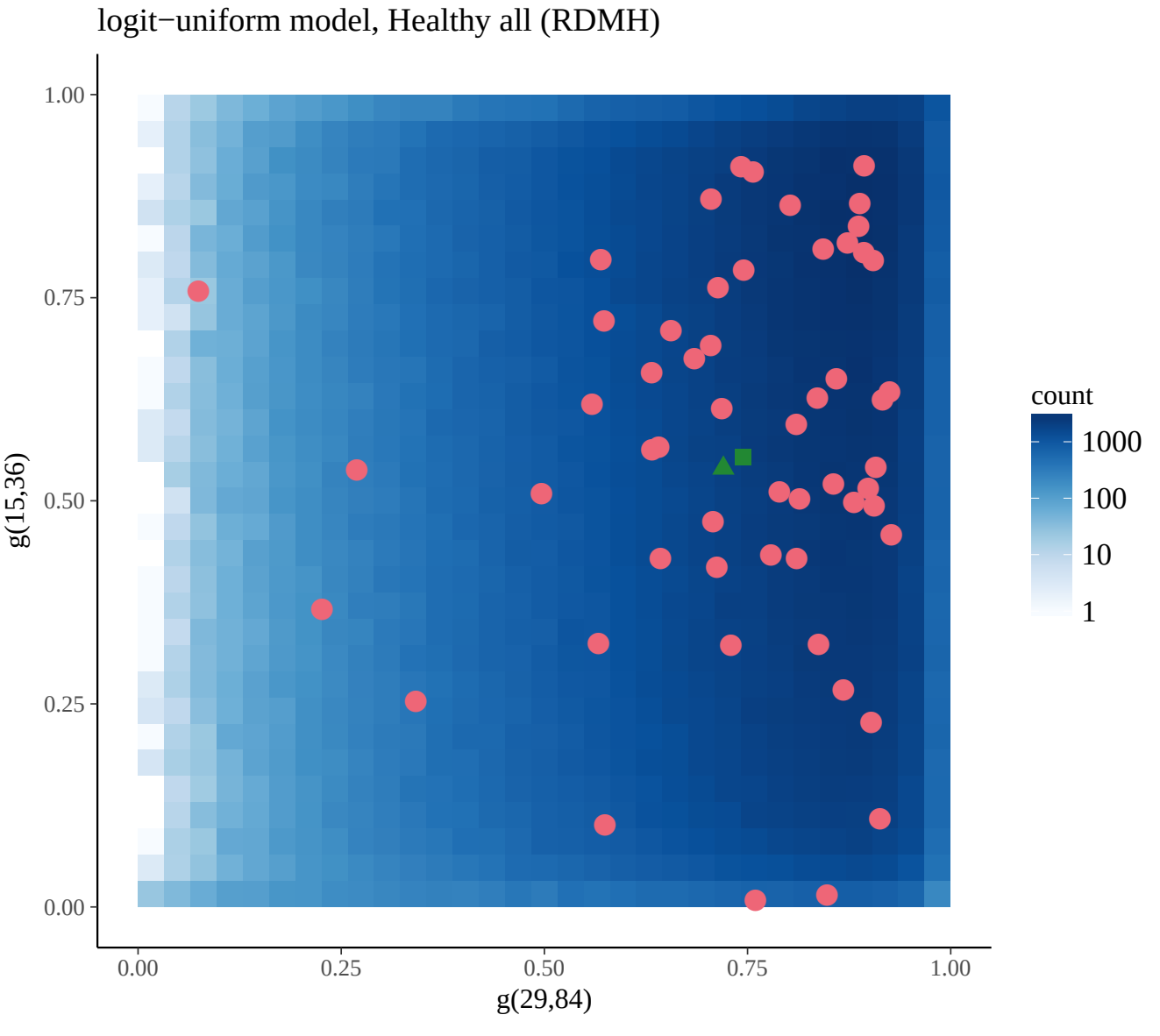
0.75
0.50
0.25
0.00
g(16,20)
0.00
0.25
0.50
0.75
1.00
g(29,84)
count
1000
100
10
1


logit−uniform model, Healthy all (RDMH)
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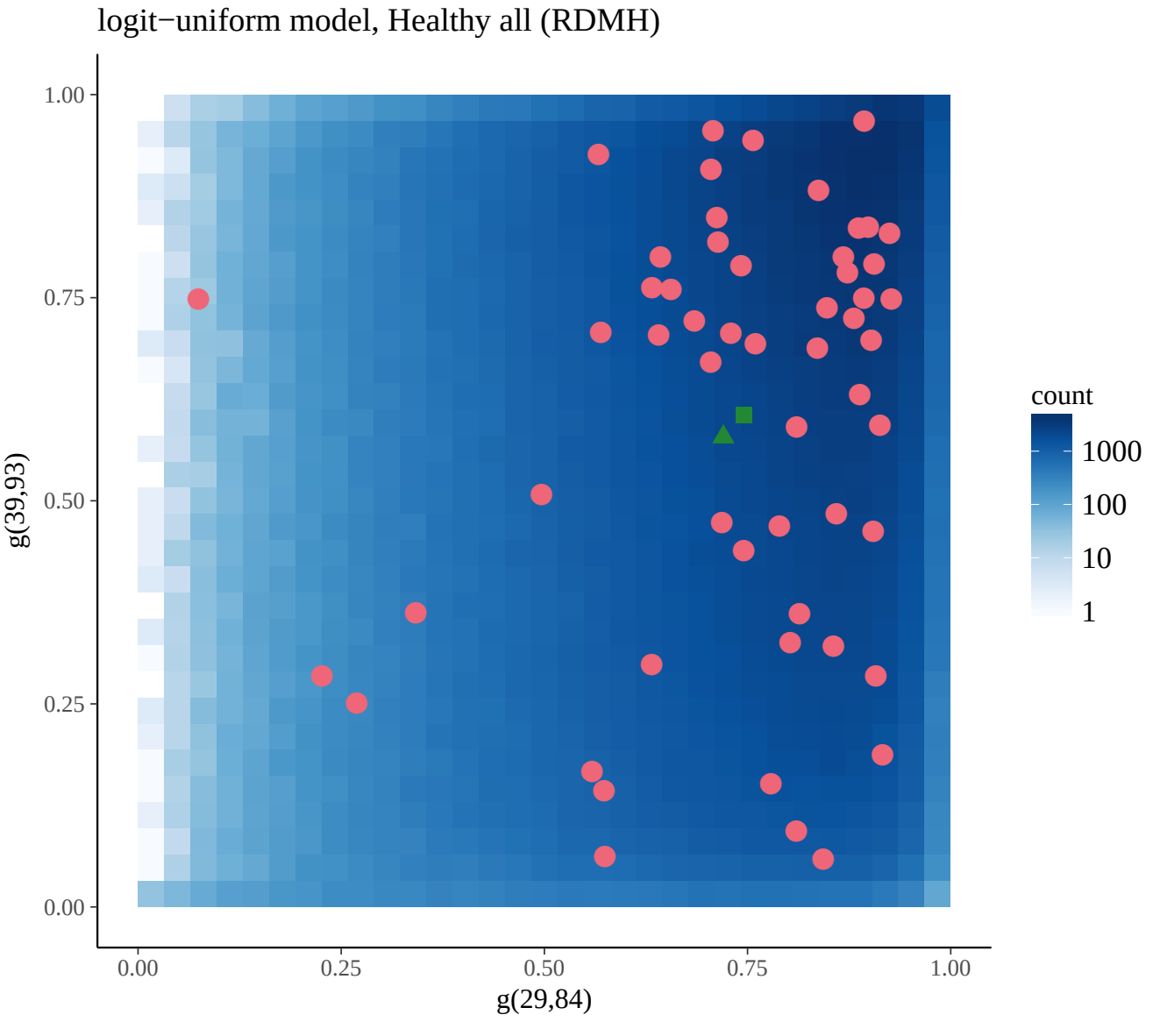
0.75
0.50
0.25
0.00
g(15,36)
0.00
0.25
0.50
0.75
1.00
g(29,84)
count
1000
100
10
1


logit−uniform model, Healthy all (RDMH)
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0.75
0.50
0.25
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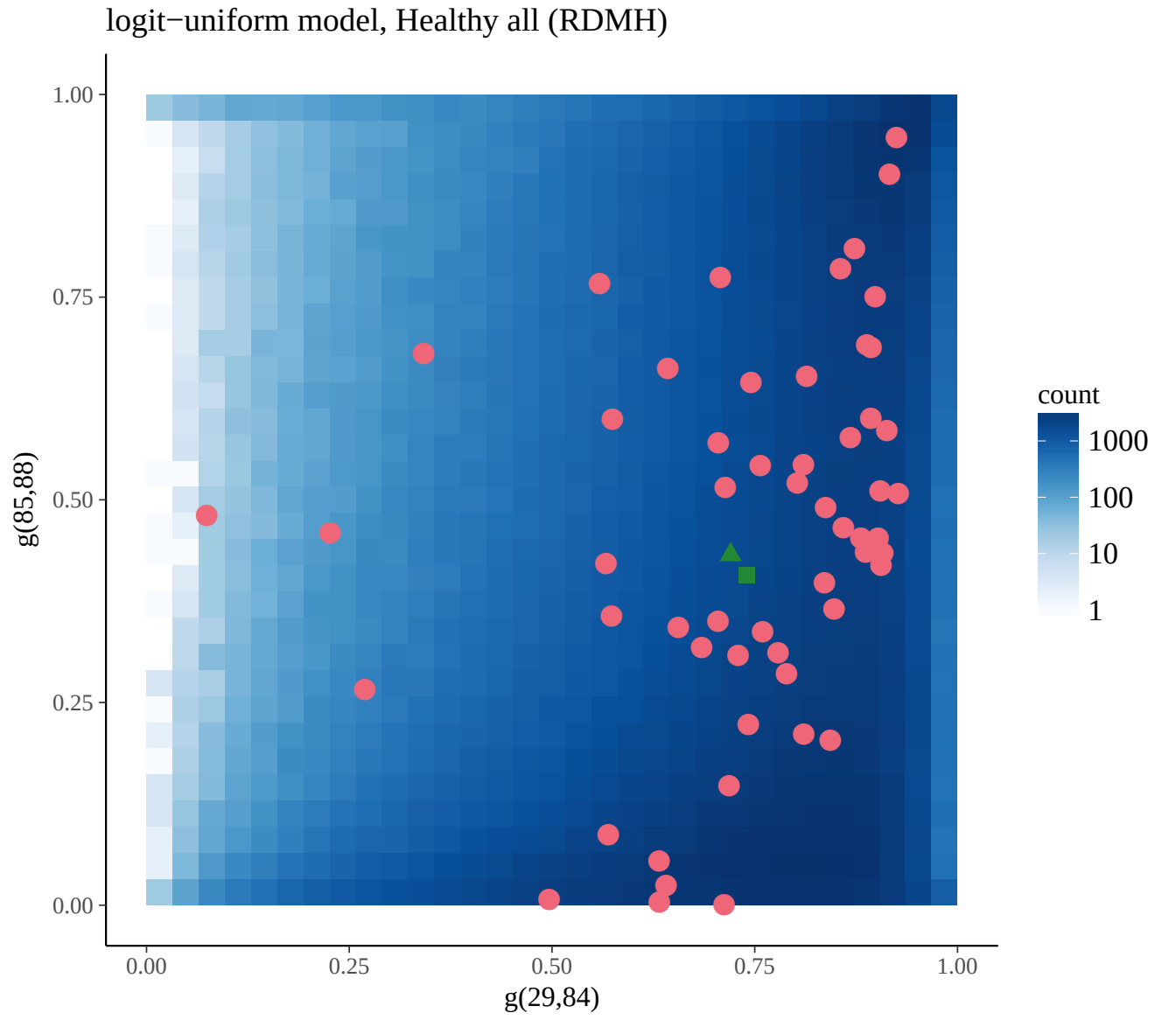
0.00
0.25
0.50
0.75
1.00
g(29,84)
count
1000
100
10
1


logit−uniform model, Healthy all (RDMH)
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0.75
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0.25
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0.00
0.25
0.50
0.75
1.00
g(29,84)
count
1000
100
10
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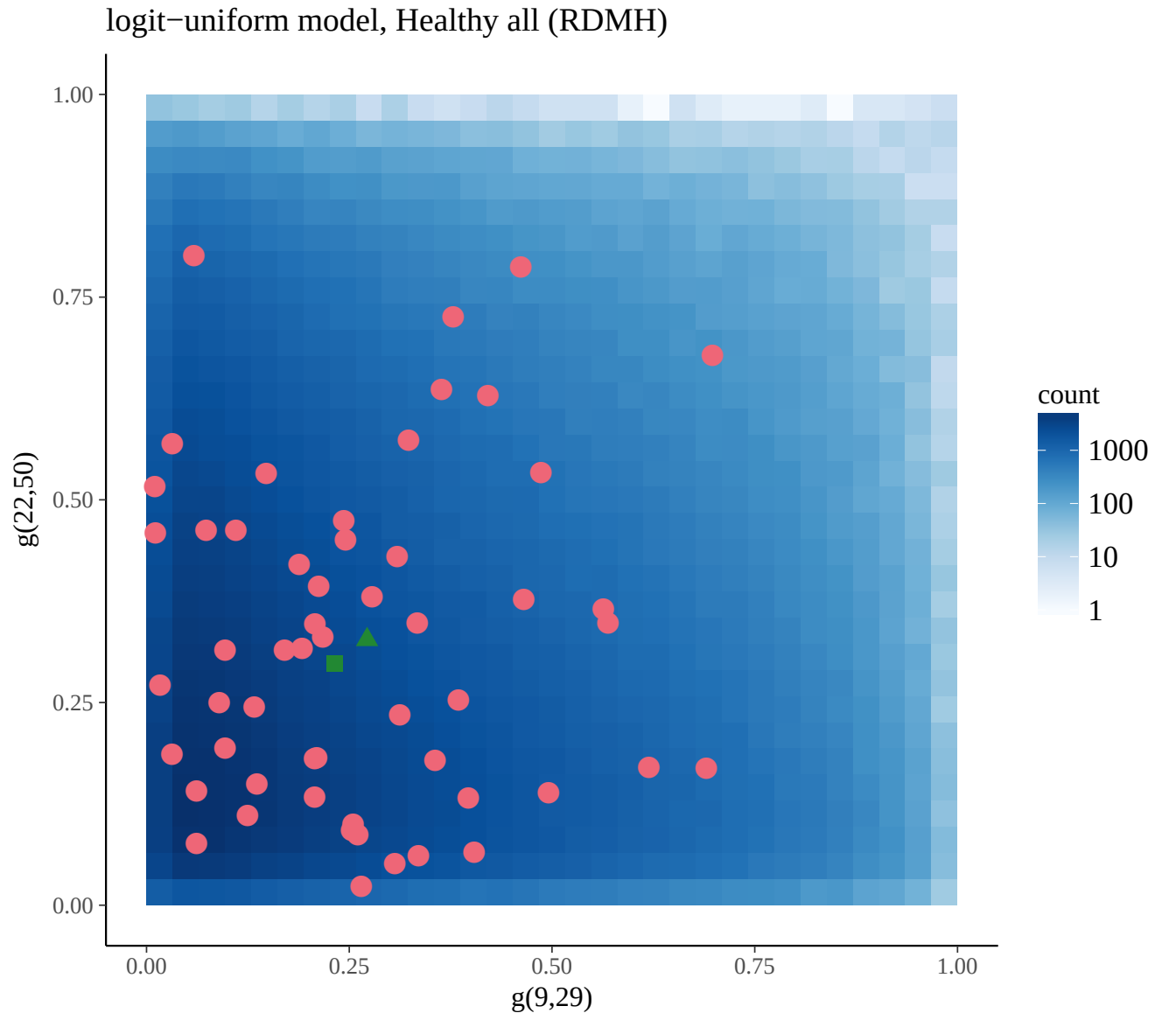
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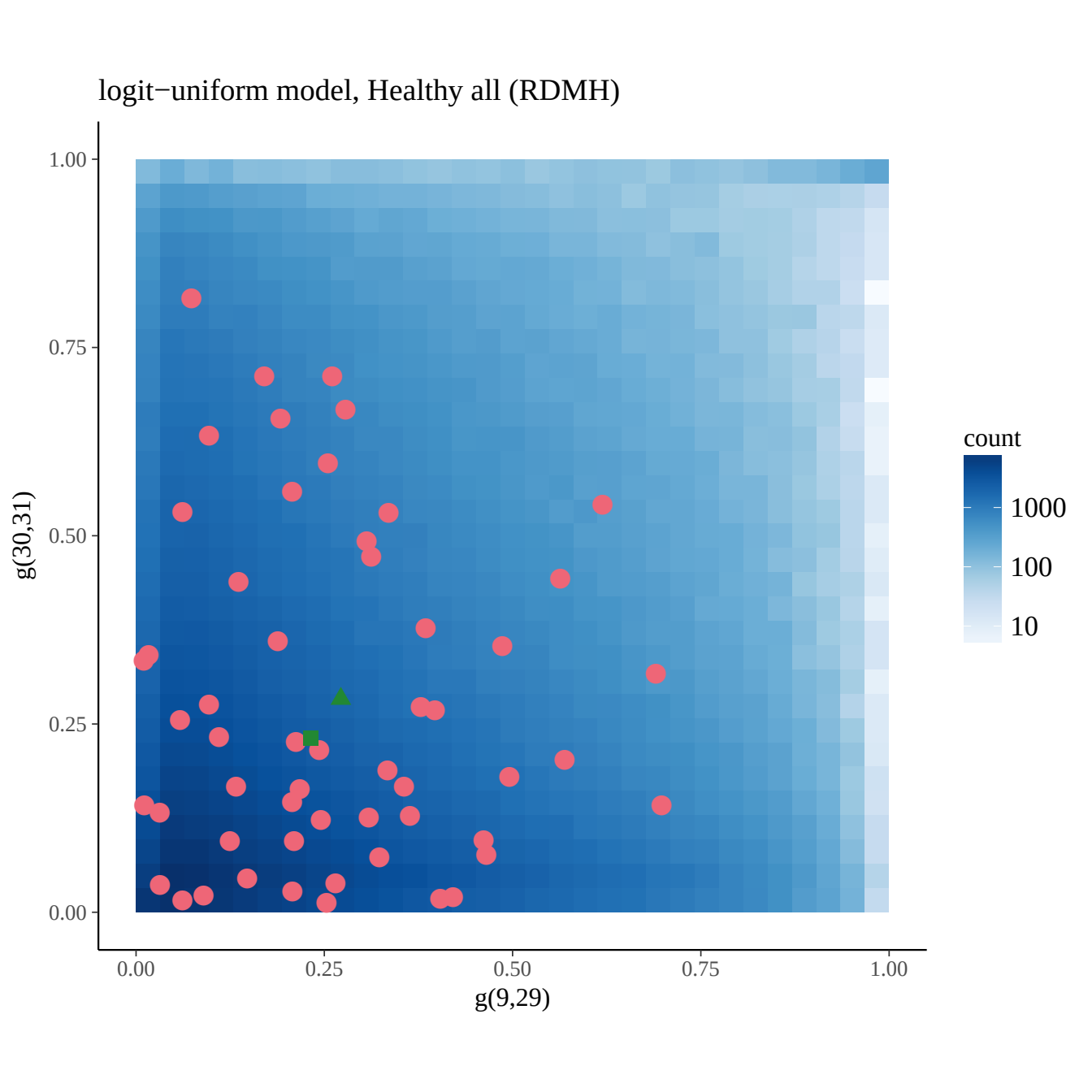
0.25
0.50
0.75
1.00
g(9,29)
count
1000
100
10
1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0.50
0.75
1.00
g(9,29)
count
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100
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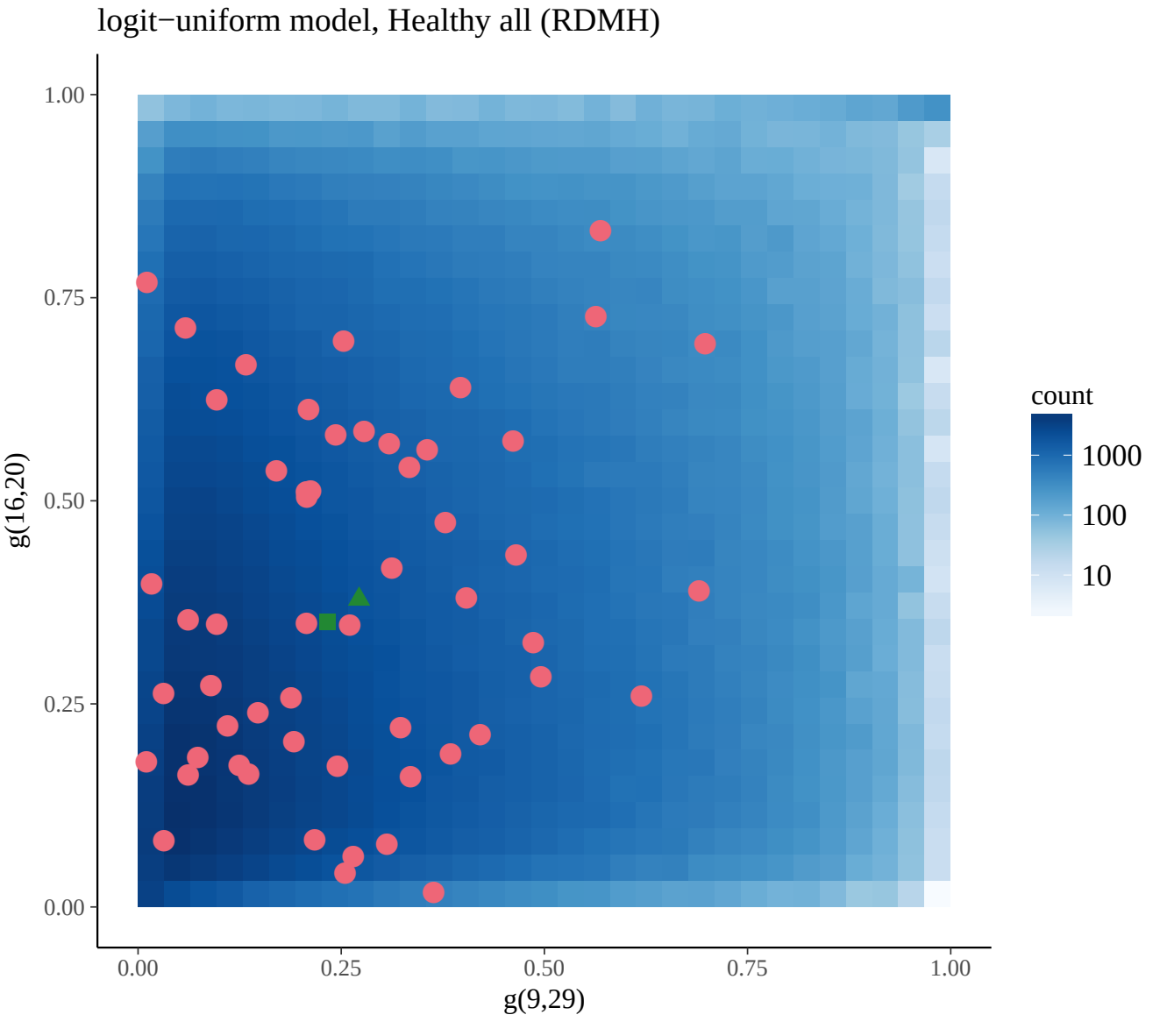
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0.00
0.25
0.50
0.75
1.00
g(9,29)
count
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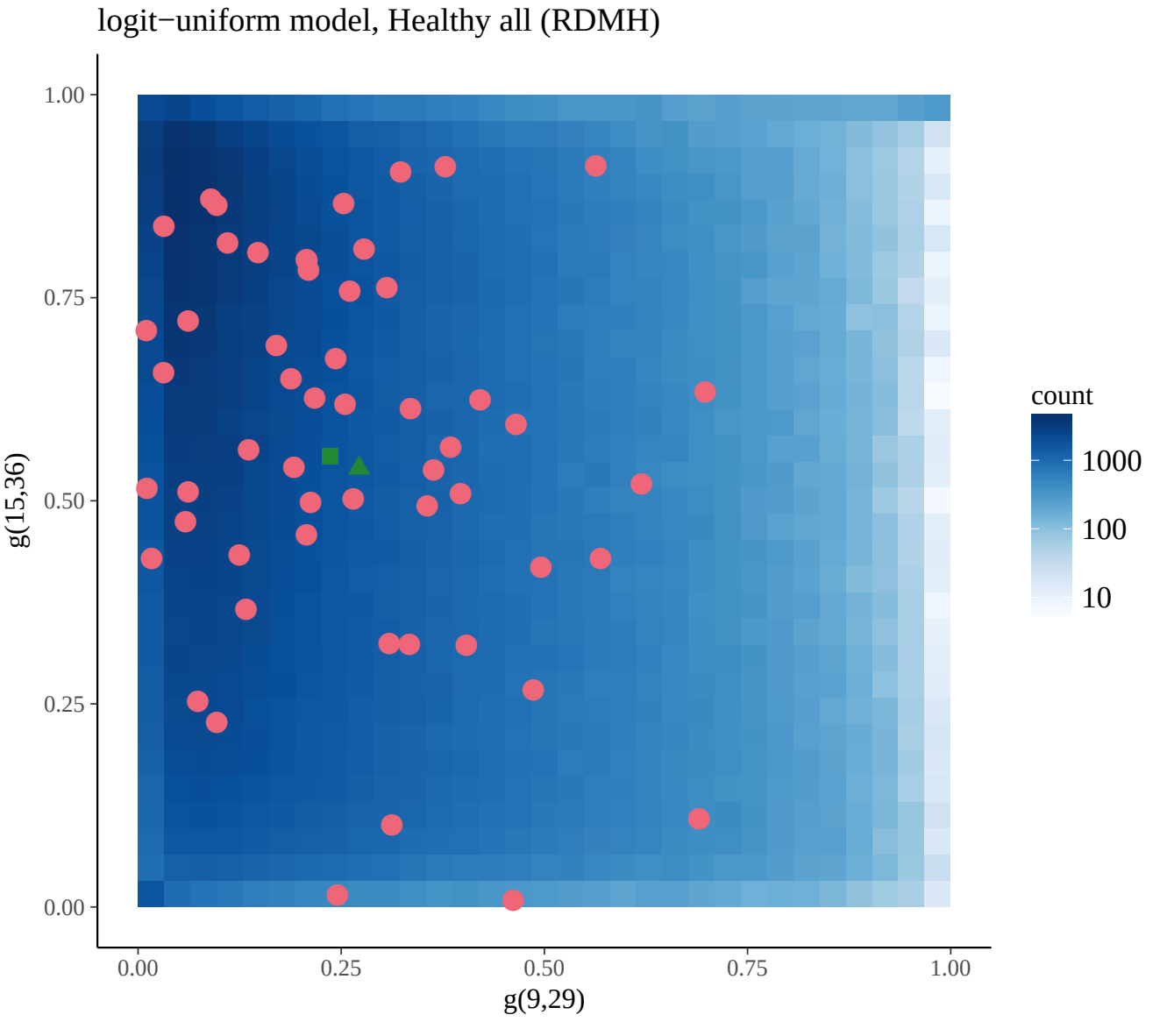
100
10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0.00
0.25
0.50
0.75
1.00
g(9,29)
count
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100
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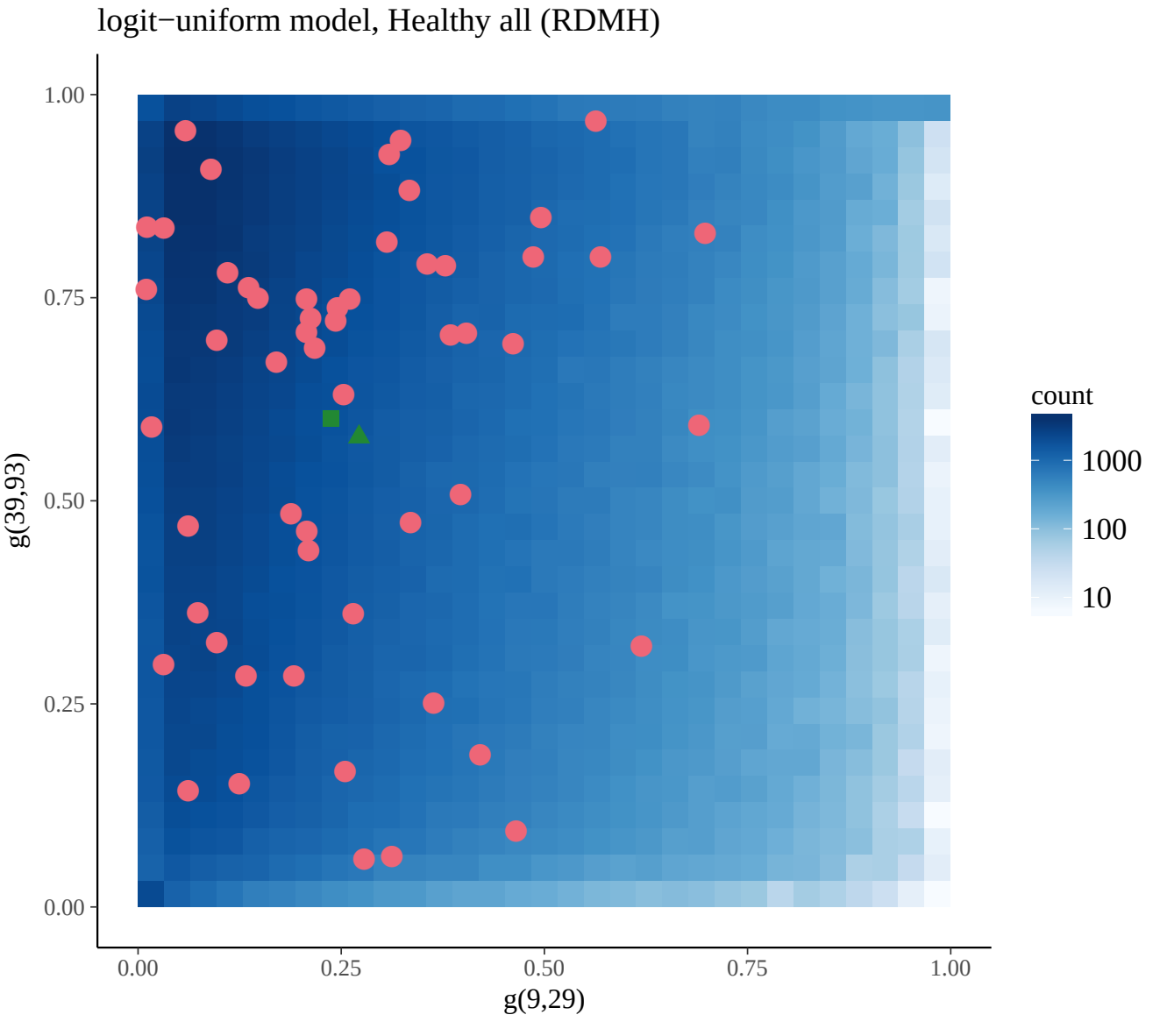
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0.50
0.75
1.00
g(9,29)
count
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100
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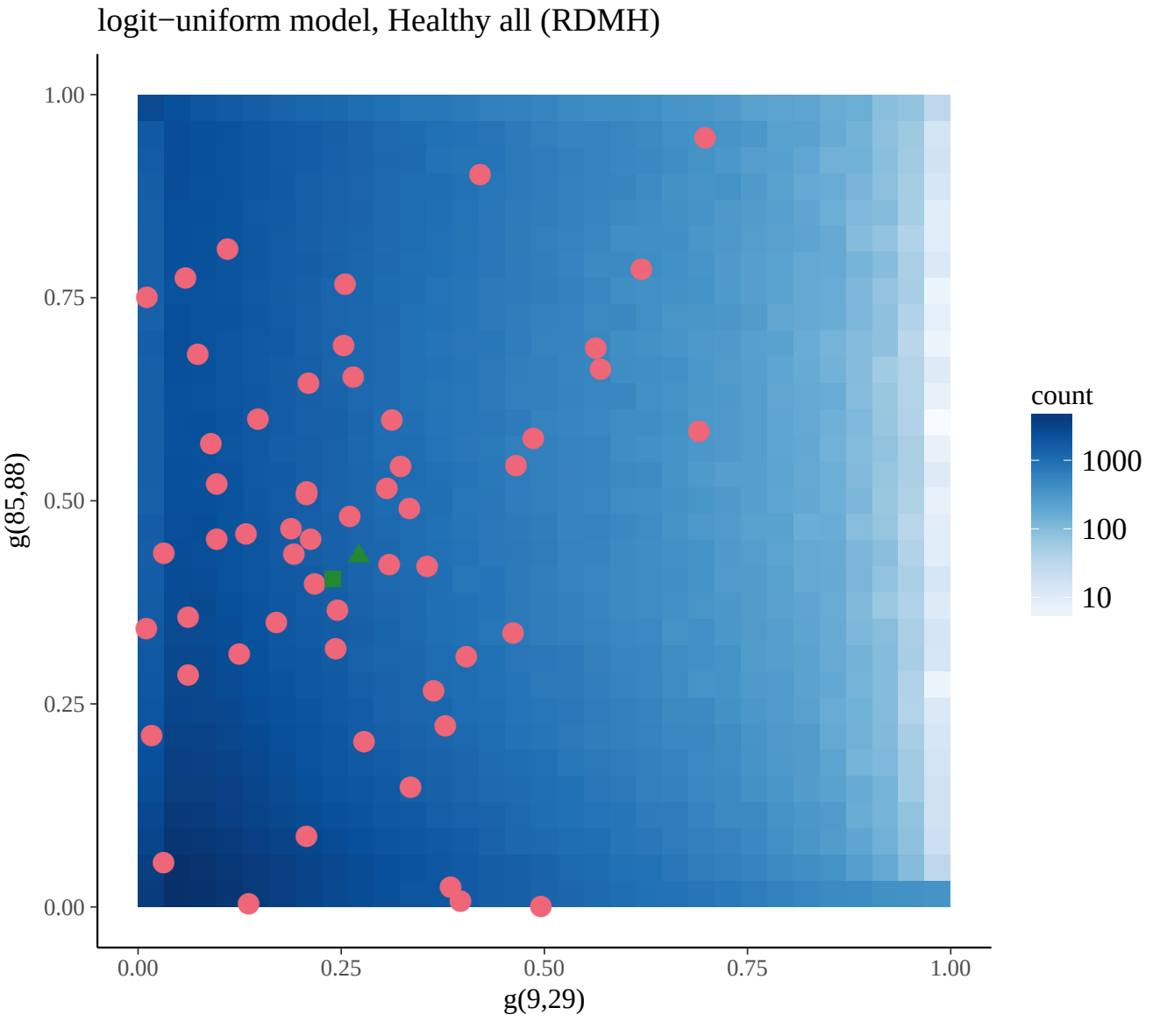
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0.50
0.75
1.00
g(9,29)
count
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100
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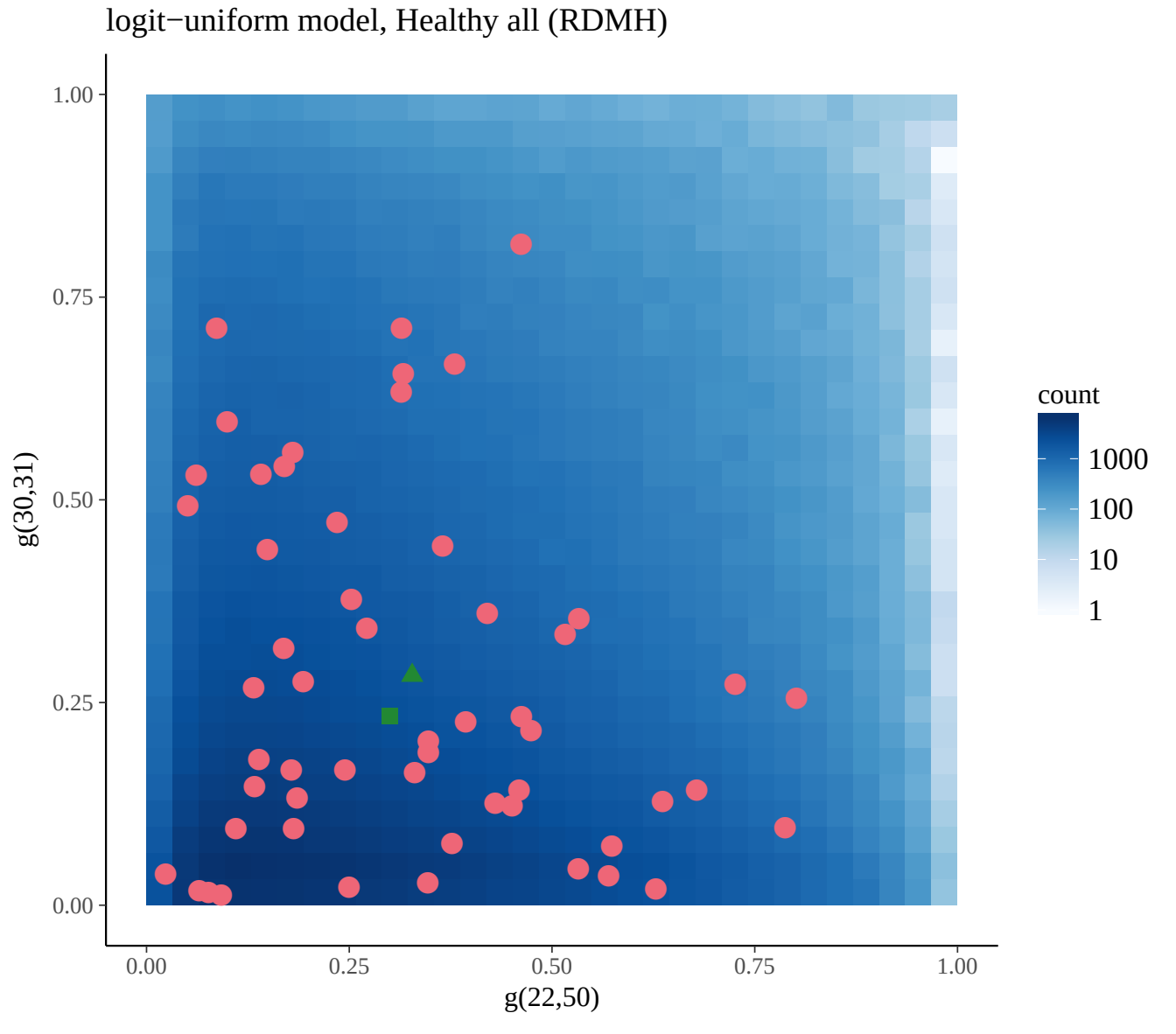
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0.75
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g(22,50)
count
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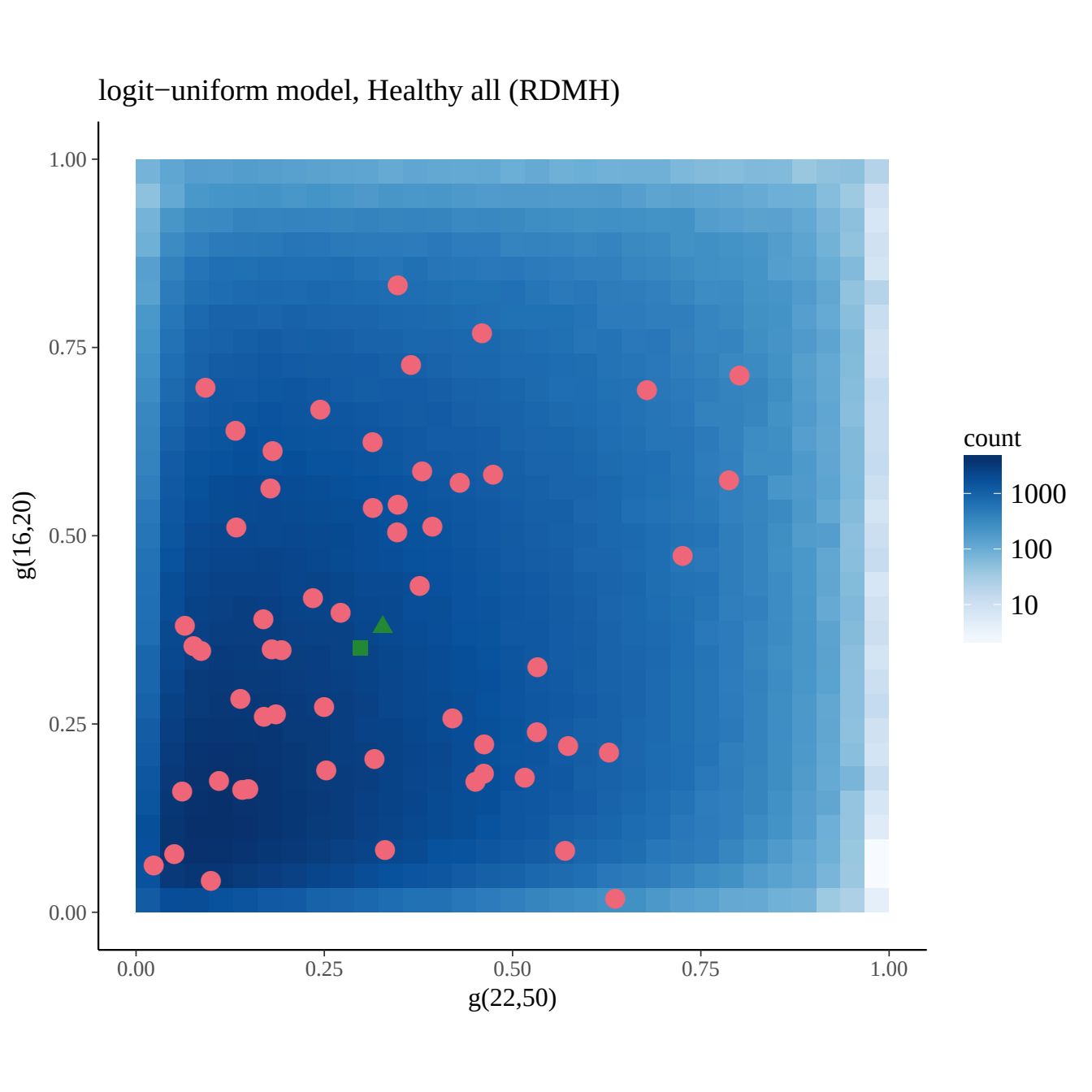
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0.75
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count
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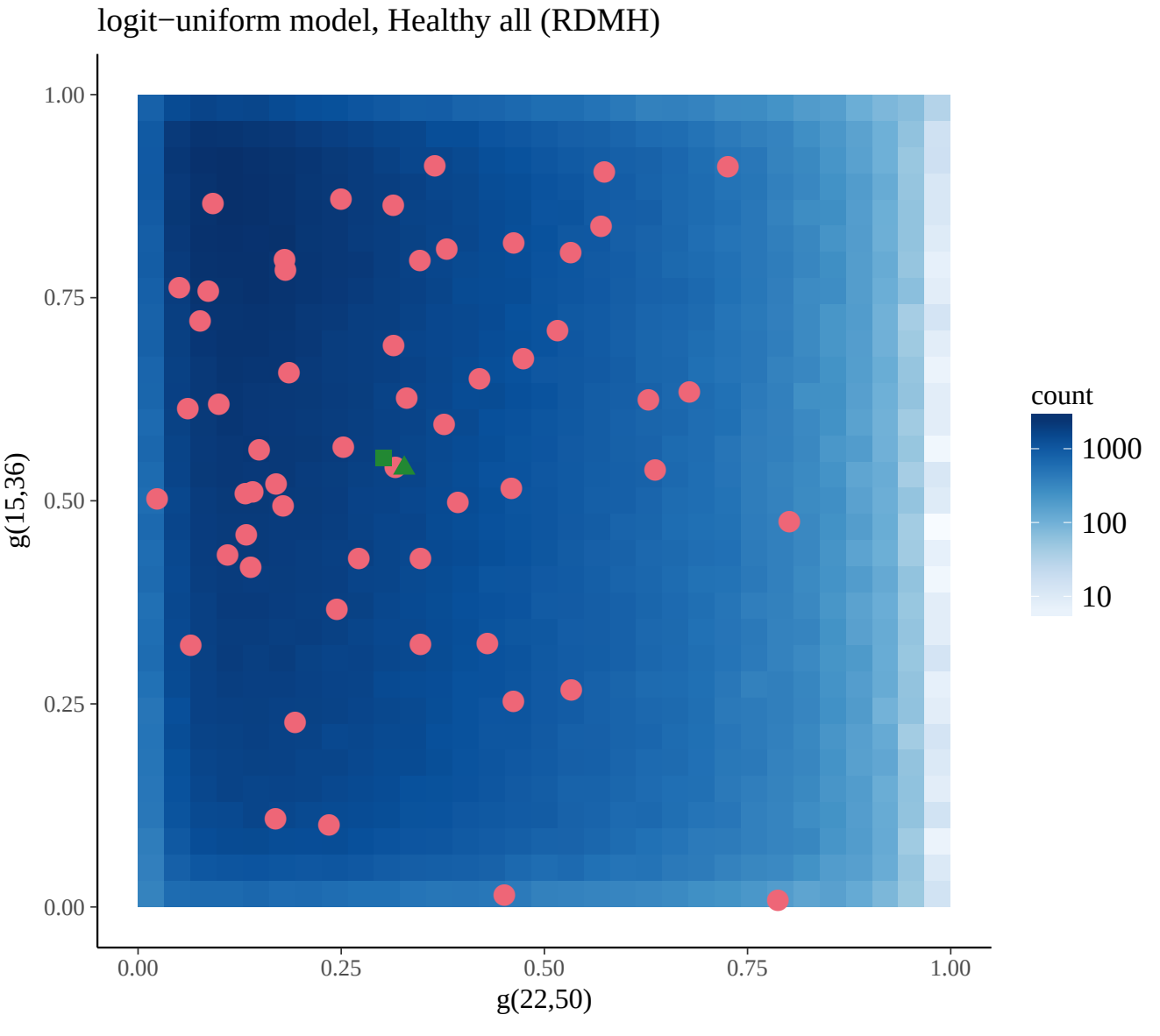
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1.00
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count
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100
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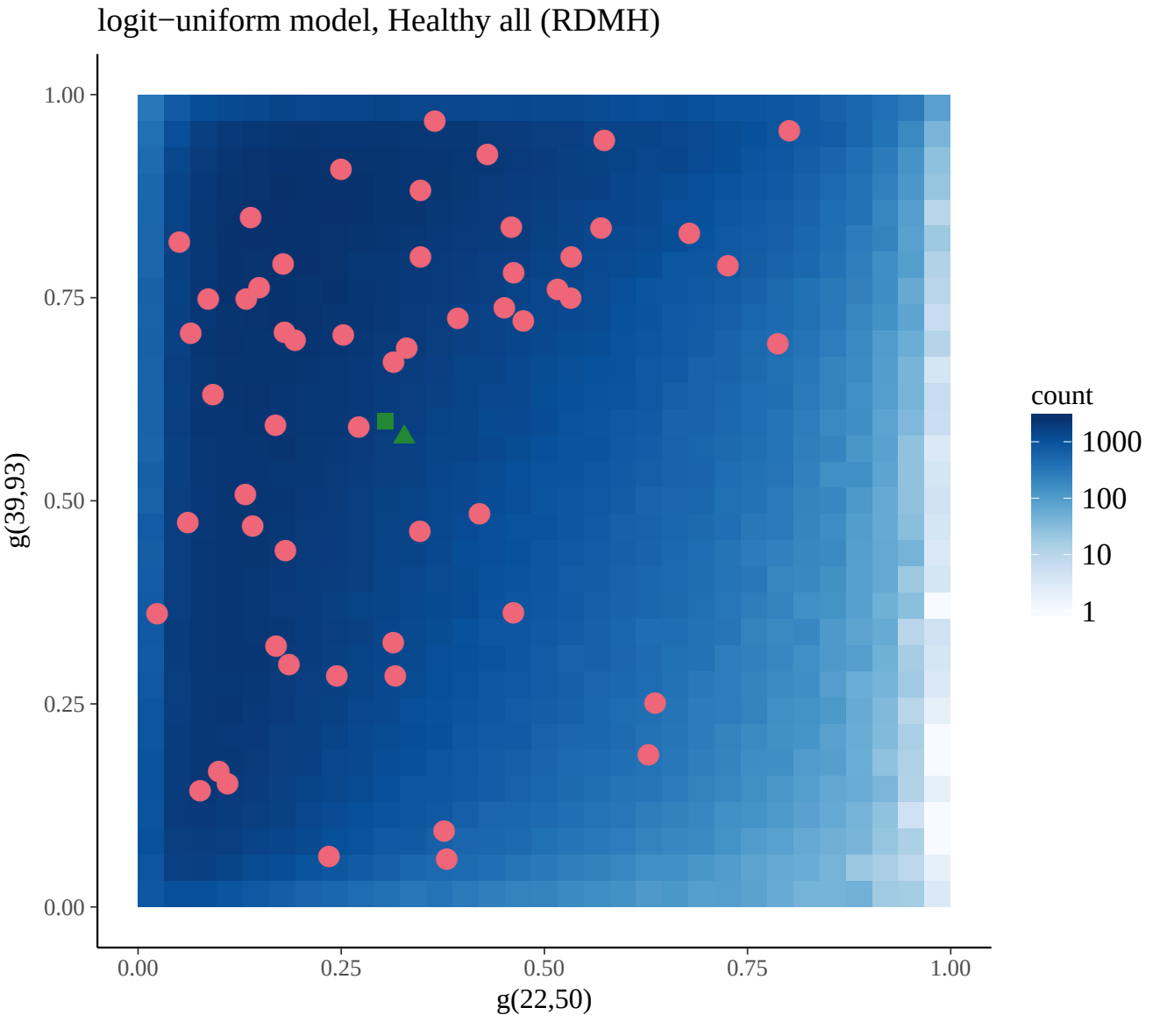
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0.75
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g(22,50)
count
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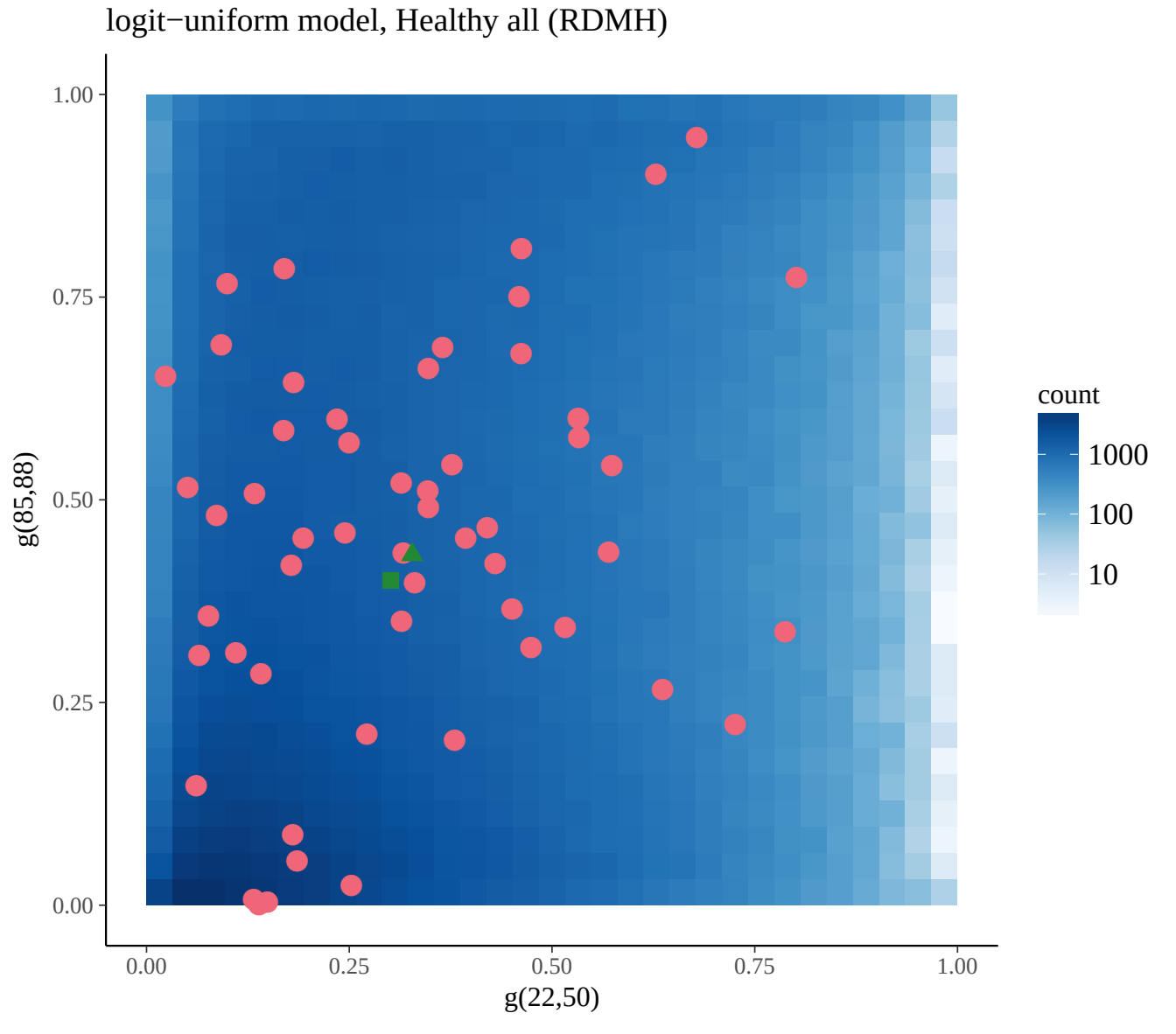
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1.00
g(22,50)
count
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100
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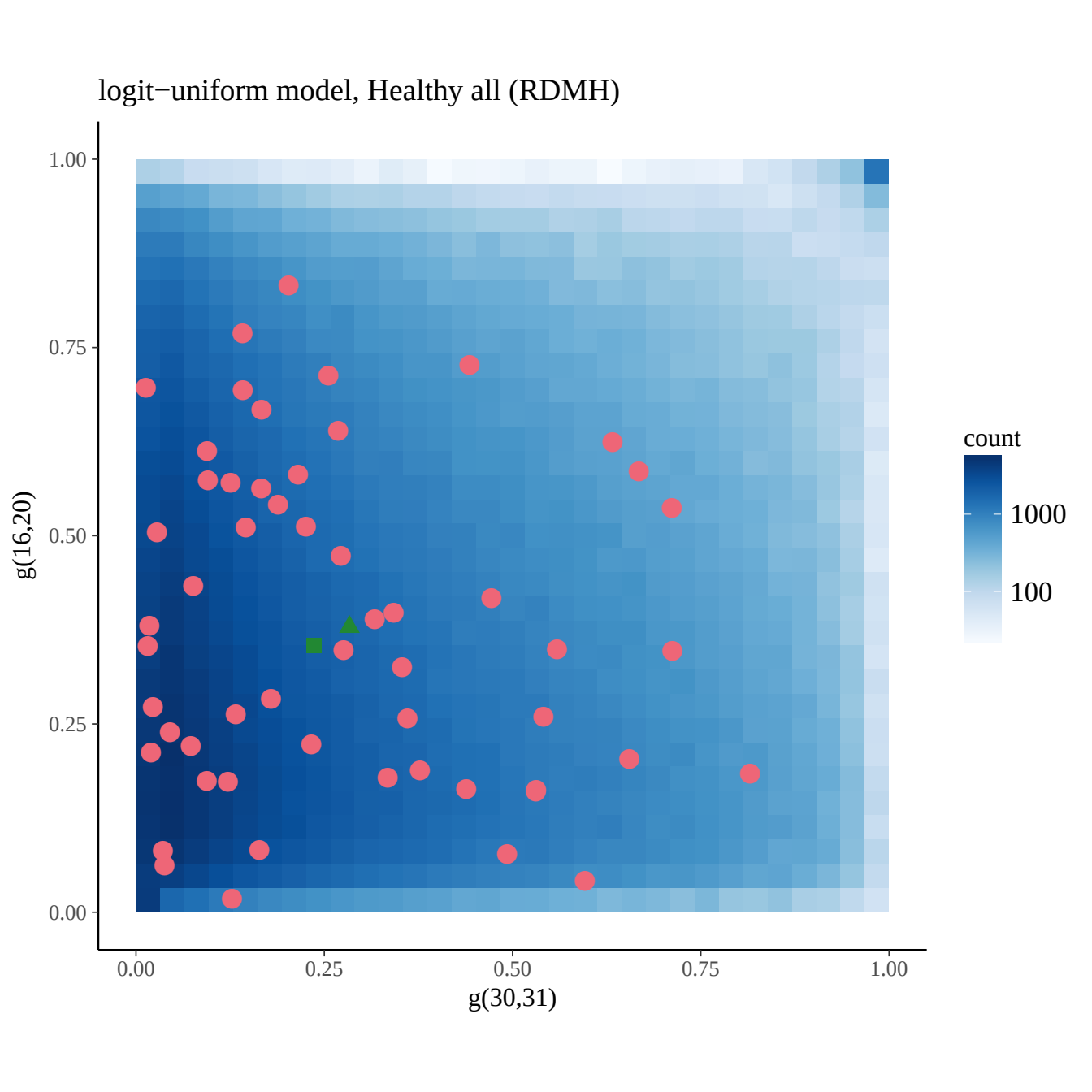
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1.00
g(30,31)
count
1000
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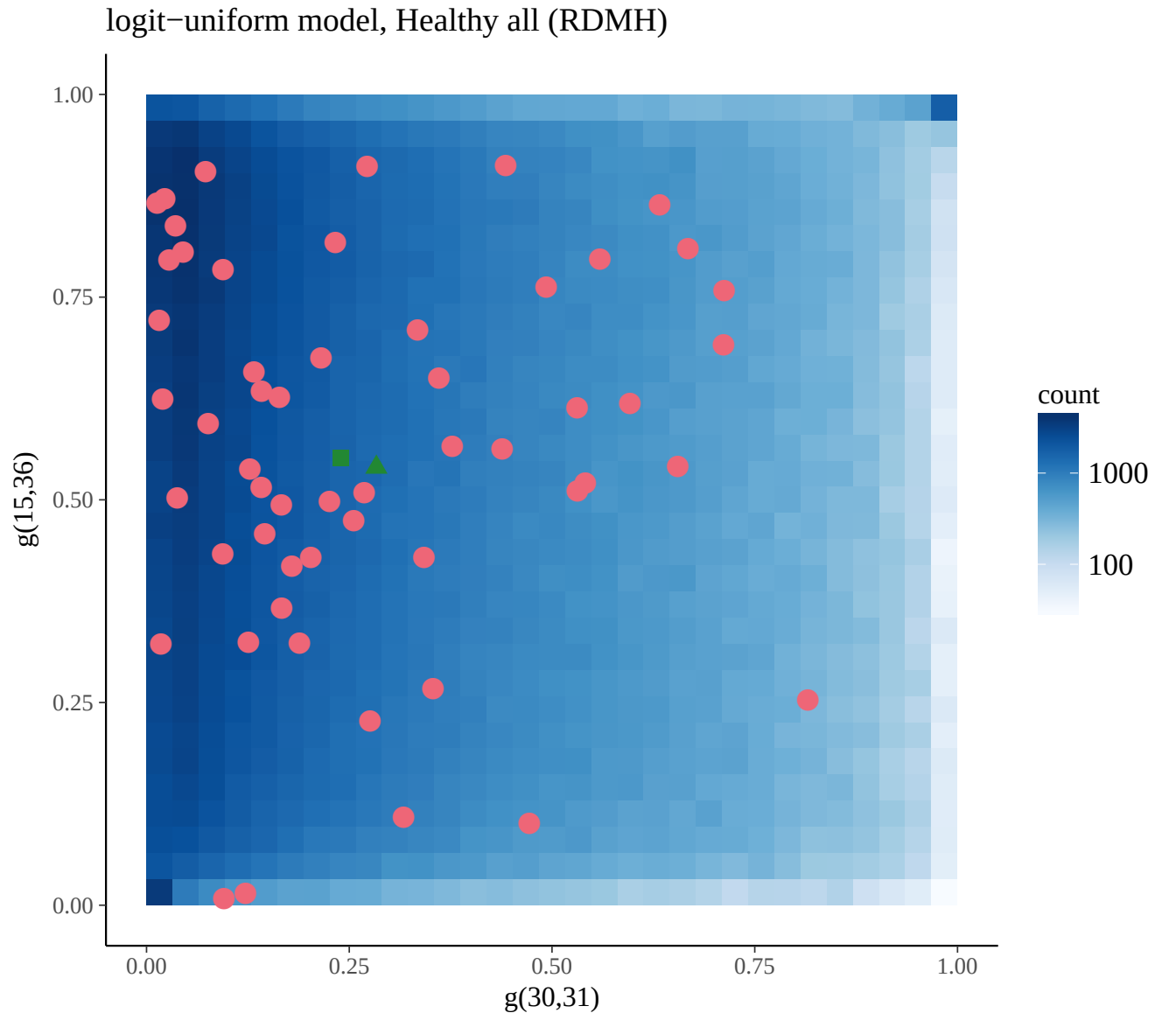
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0.75
1.00
g(30,31)
count
1000
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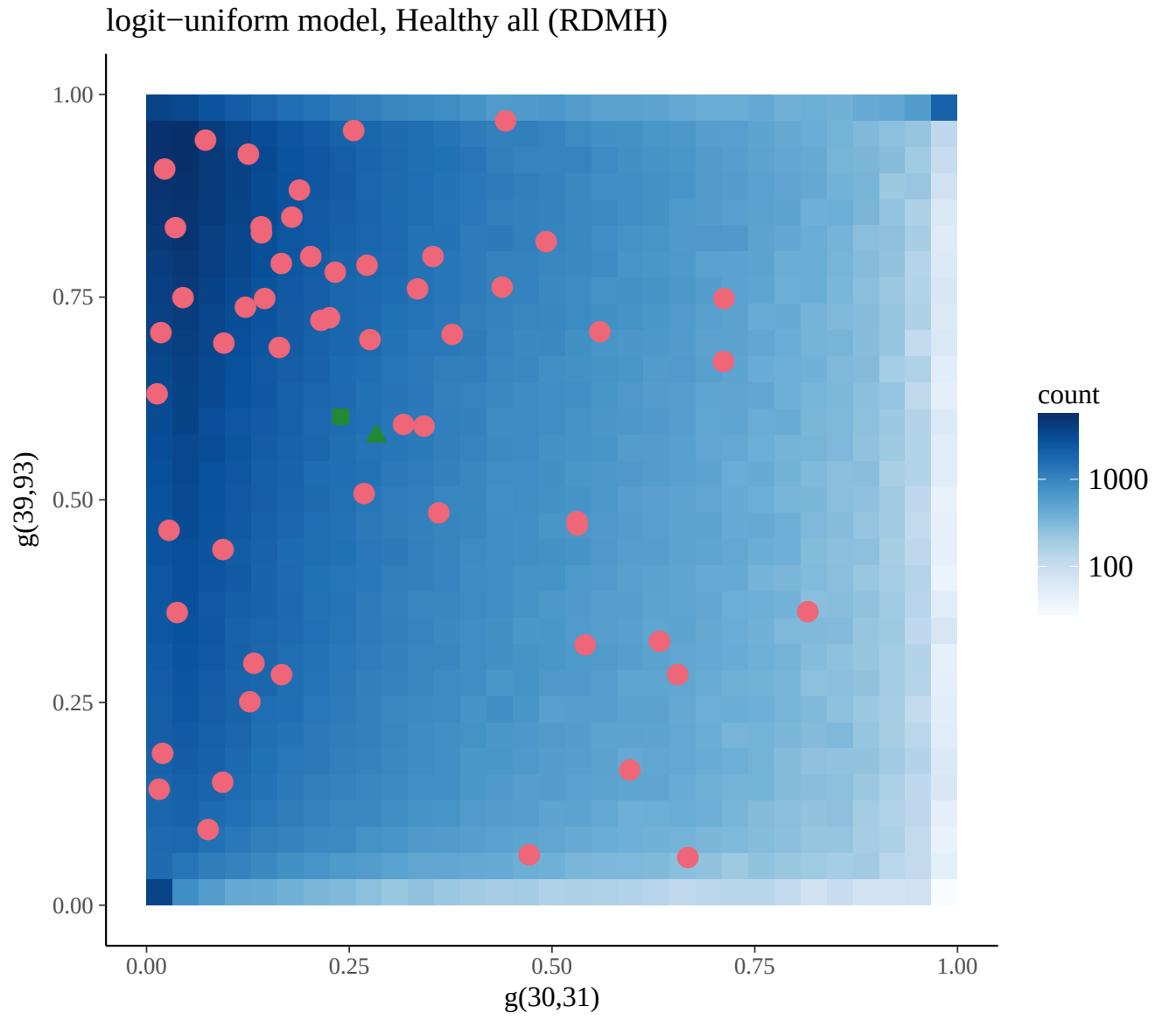
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0.50
0.75
1.00
g(30,31)
count
1000
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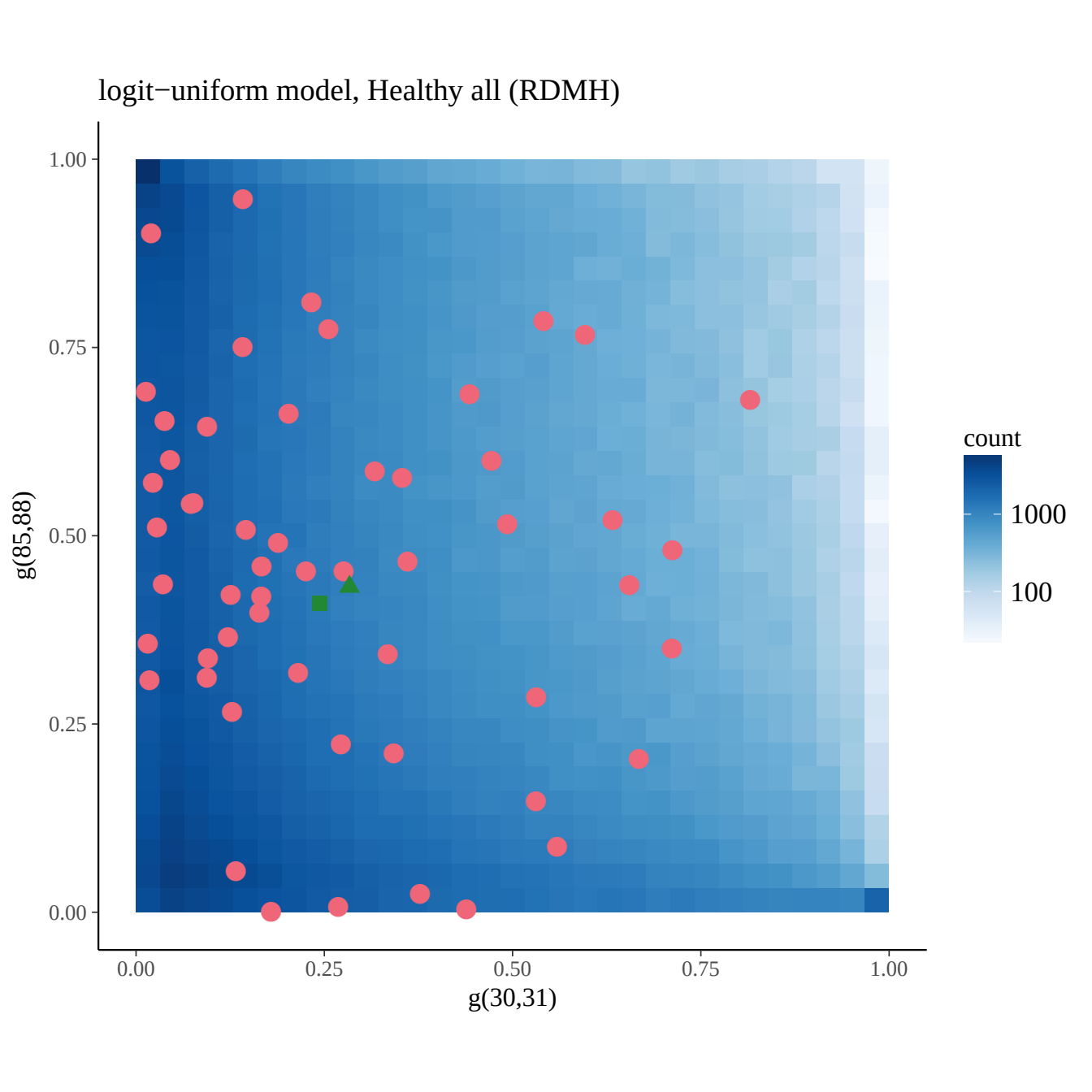
logit−uniform model, Healthy all (RDMH)
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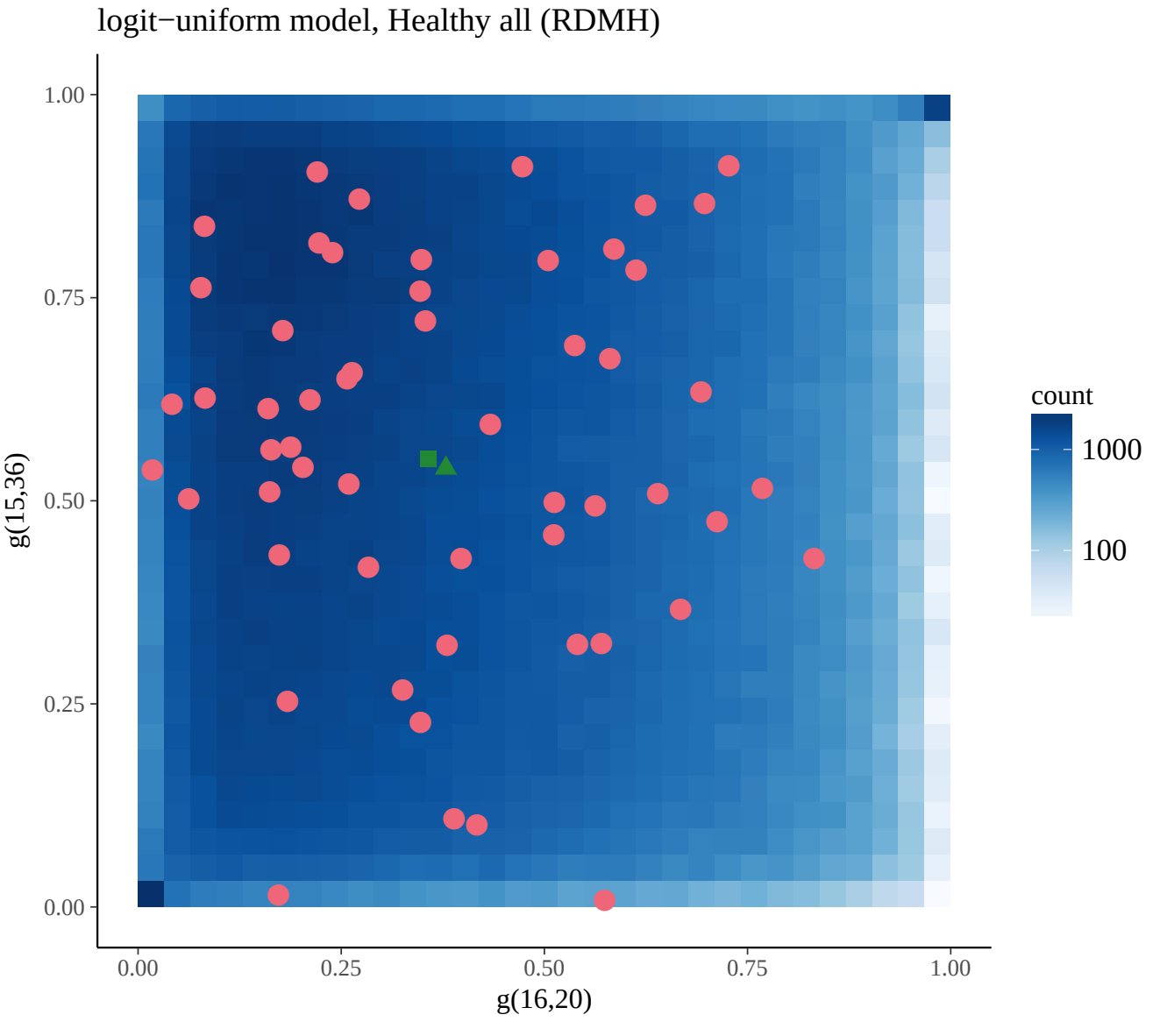
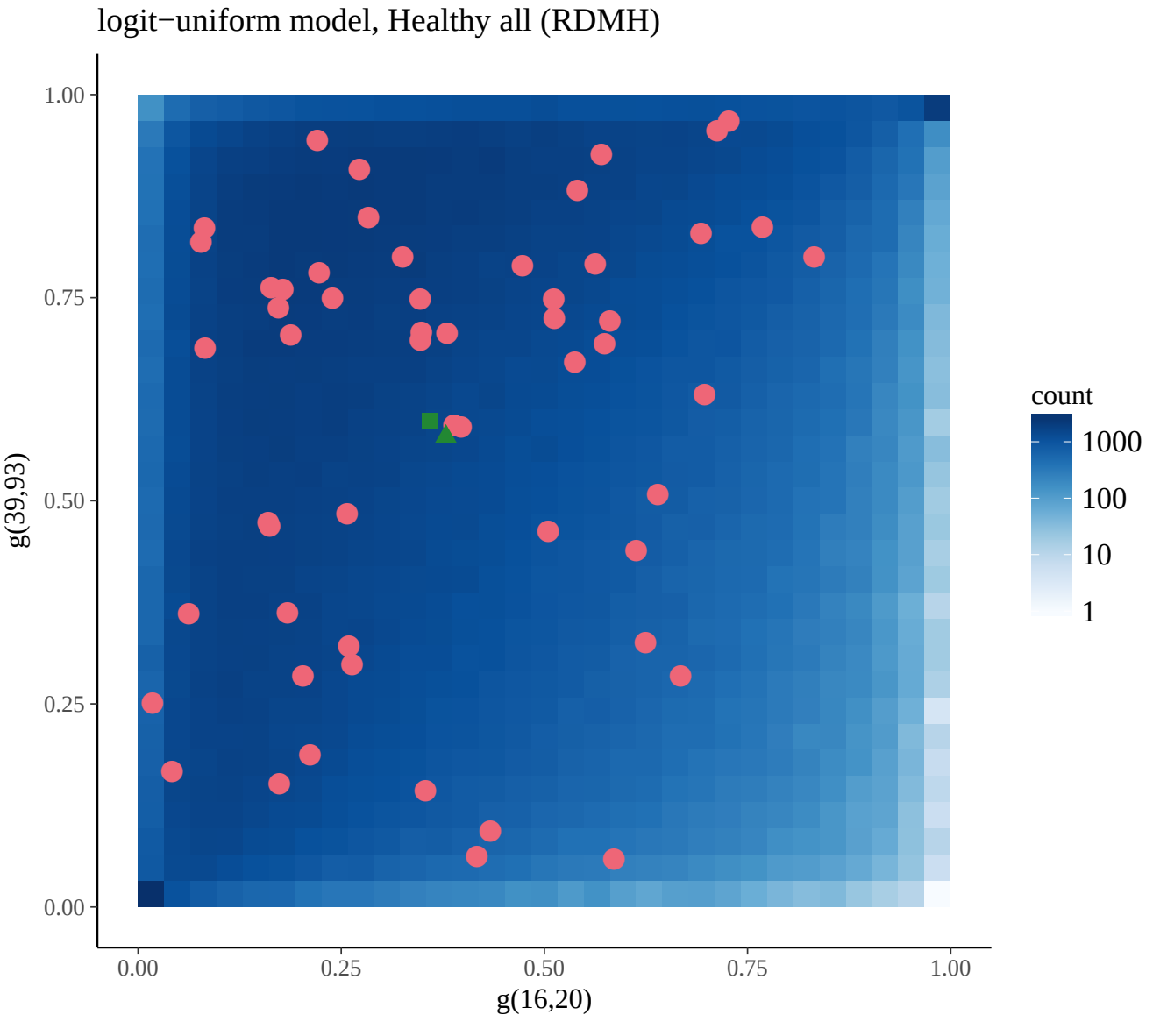
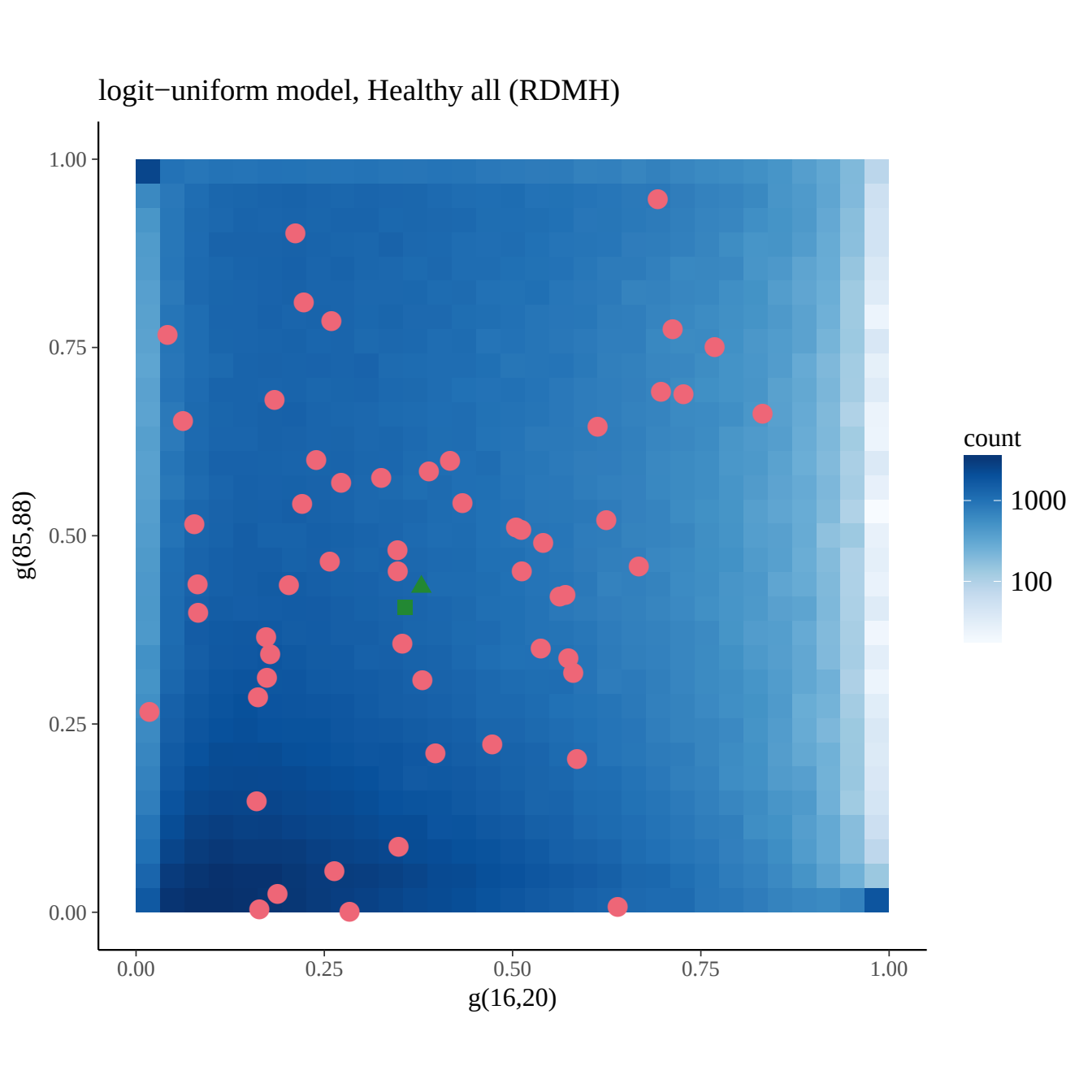
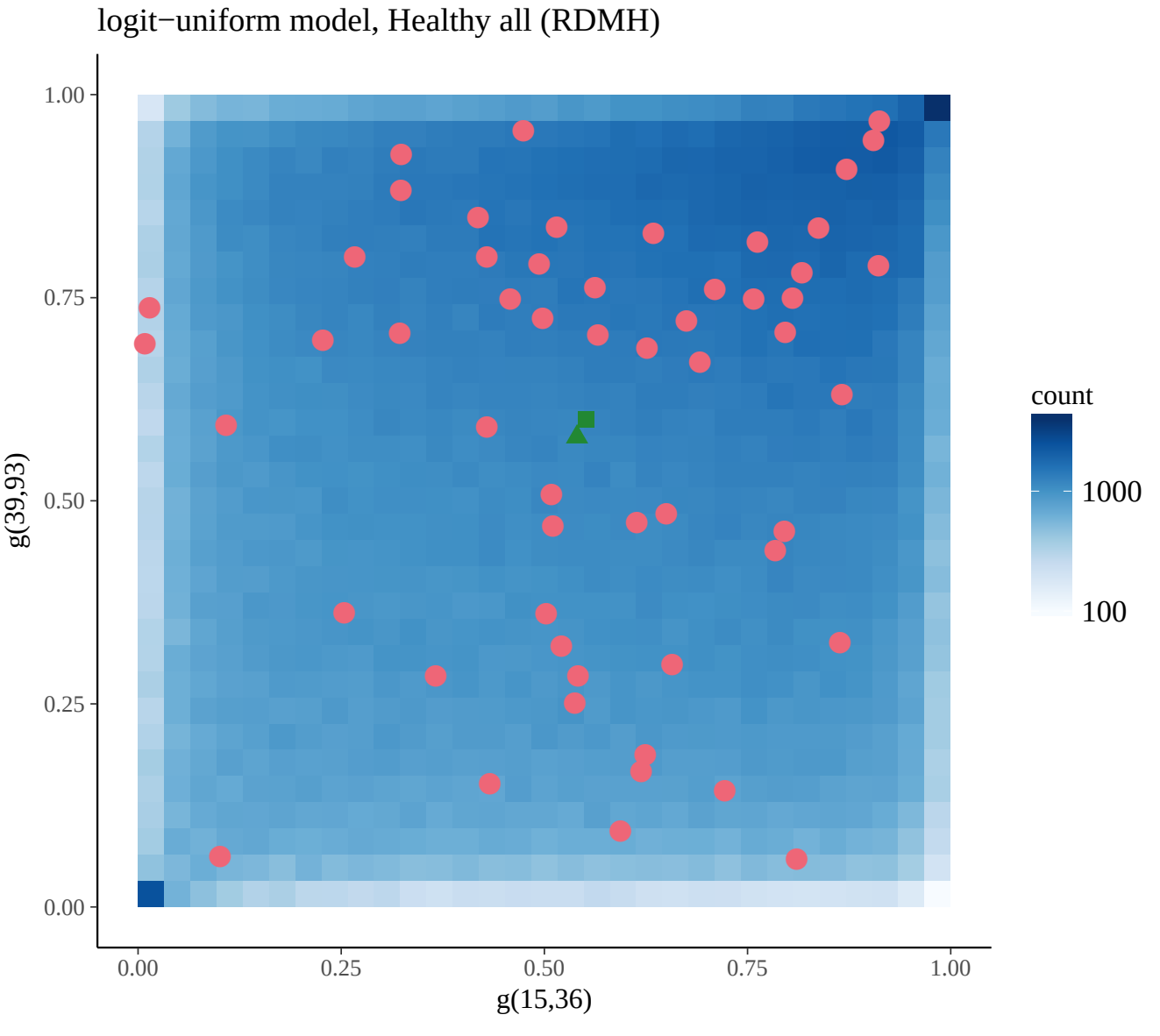
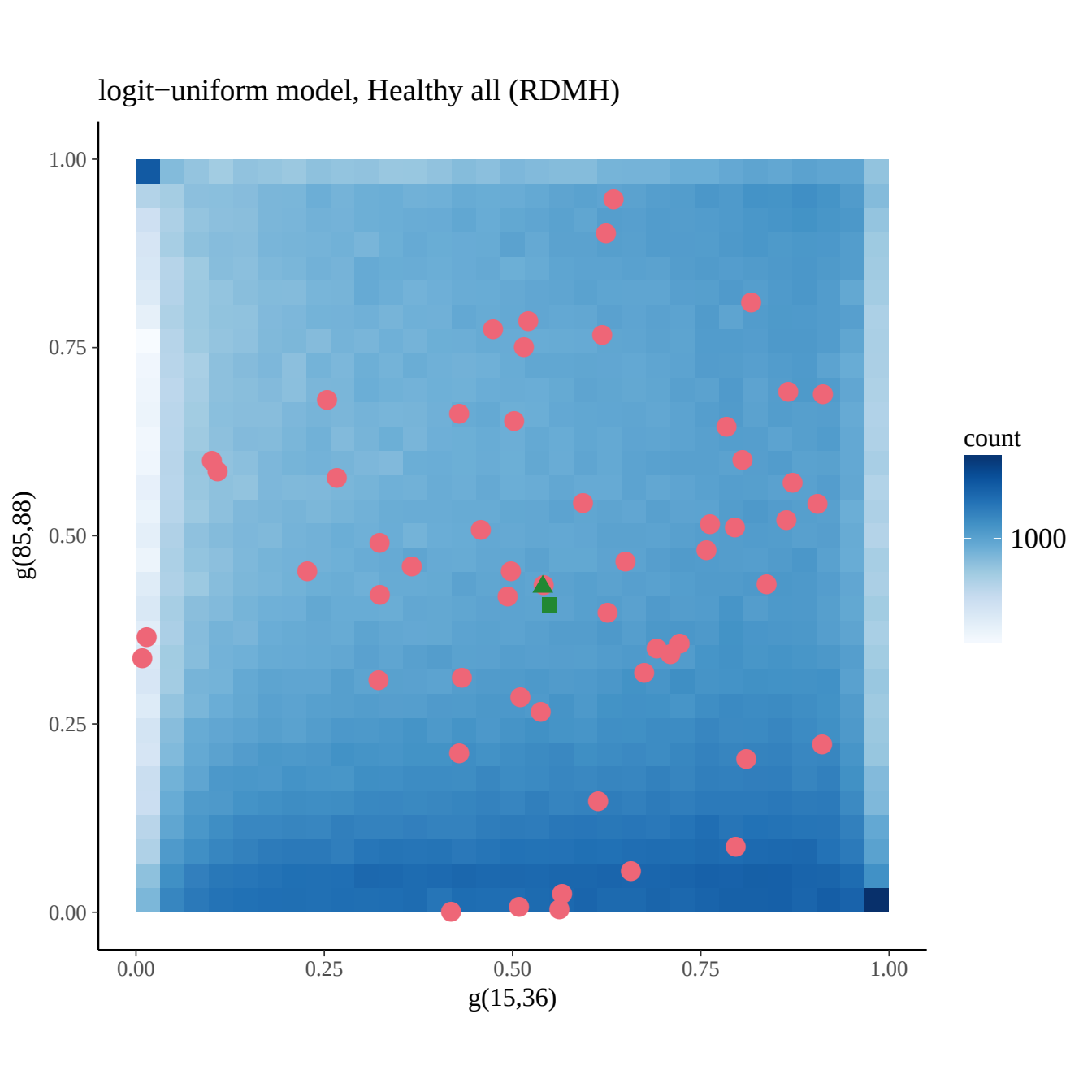
0.75
1.00
g(30,31)
count
1000
100

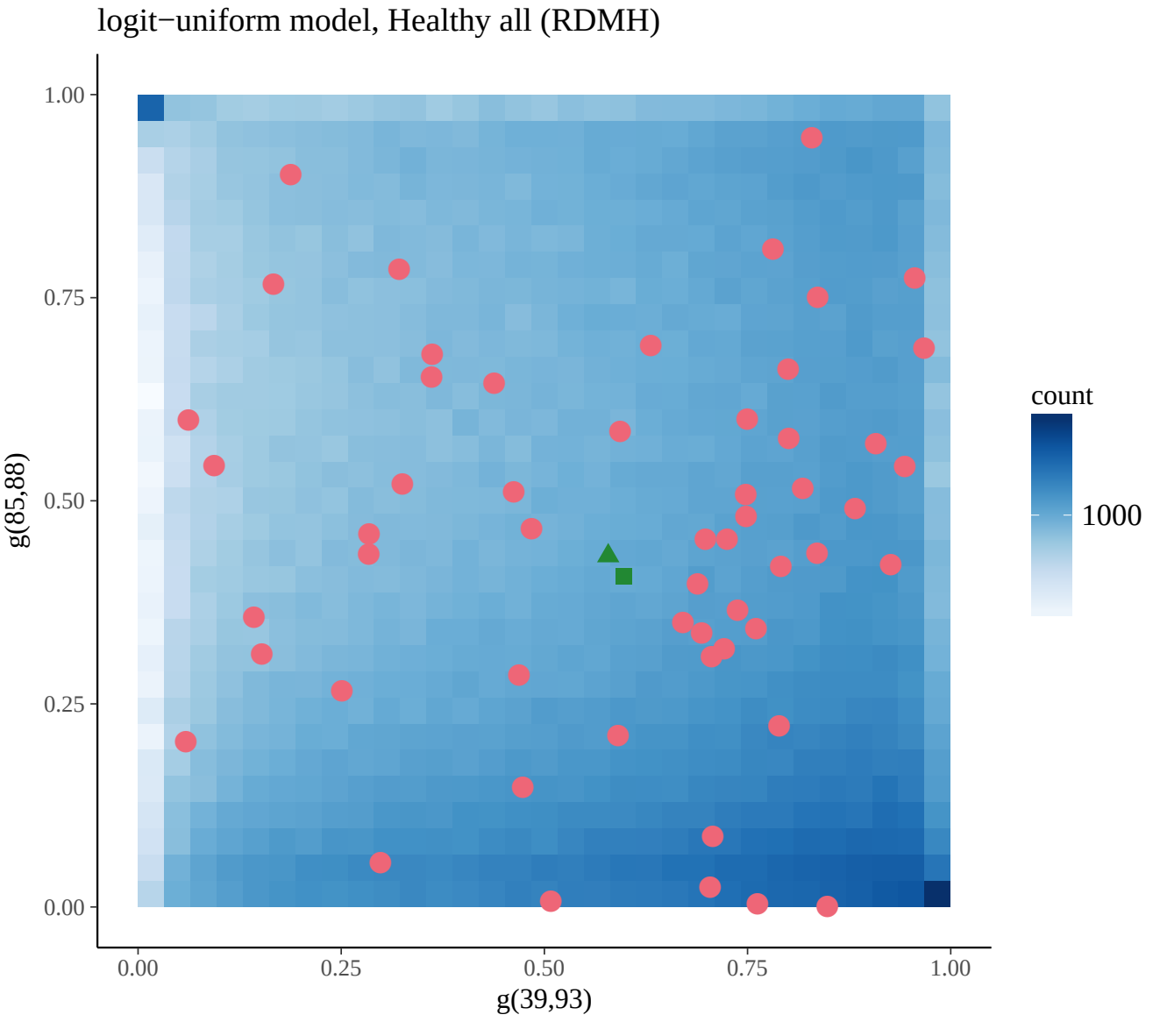

logit−uniform model, Healthy all (RDMH)
1.00
0.75
0.50
0.25
0.00
g(15,36)
0.00
0.25
0.50
0.75
1.00
g(16,20)
count
1000
100


logit−uniform model, Healthy all (RDMH)
1.00
0.75
0.50
0.25
0.00
g(39,93)
0.00
0.25
0.50
0.75
1.00
g(16,20)
count
1000
100
10
1


logit−uniform model, Healthy all (RDMH)
1.00
0.75
0.50
0.25
0.00
g(85,88)
0.00
0.25
0.50
0.75
1.00
g(16,20)
count
1000
100


logit−uniform model, Healthy all (RDMH)
1.00
0.75
0.50
0.25
0.00
g(39,93)
0.00
0.25
0.50
0.75
1.00
g(15,36)
count
1000
100


logit−uniform model, Healthy all (RDMH)
1.00
0.75
0.50
0.25
0.00
g(85,88)
0.00
0.25
0.50
0.75
1.00
g(15,36)
count
1000


logit−uniform model, Healthy all (RDMH)
1.00
0.75
0.50
0.25
0.00
g(85,88)
0.00
0.25
0.50
0.75
1.00
g(39,93)
count
1000