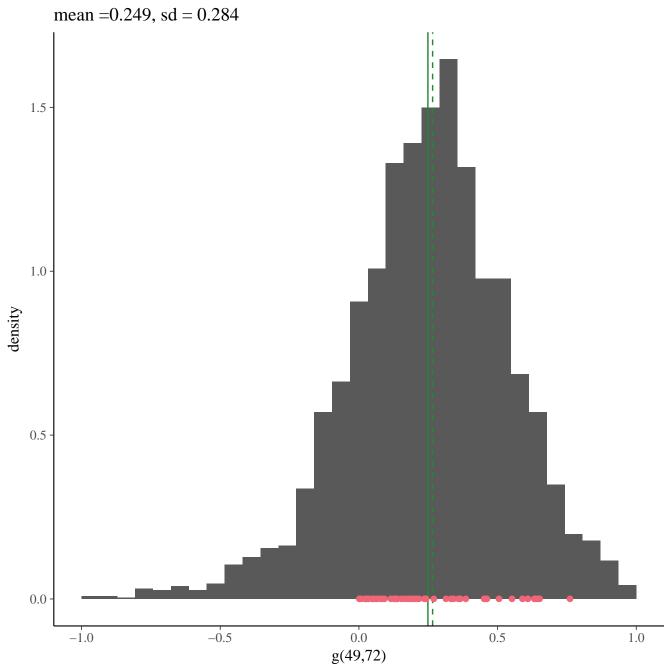
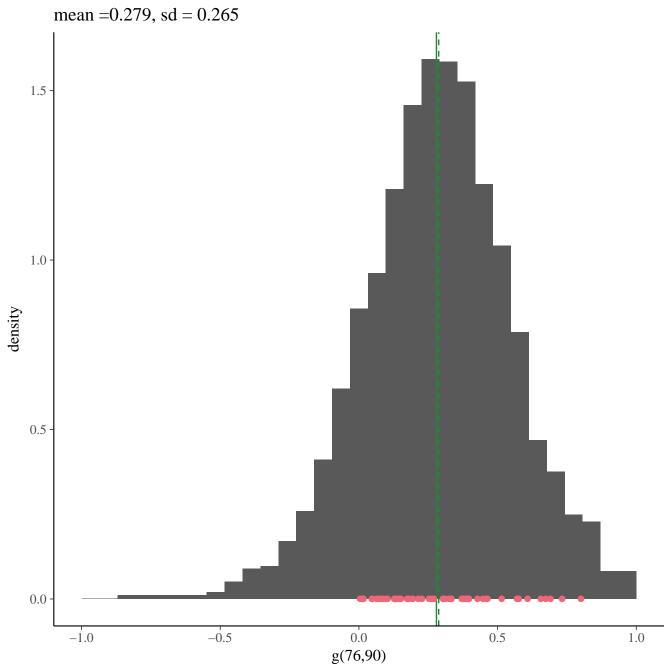
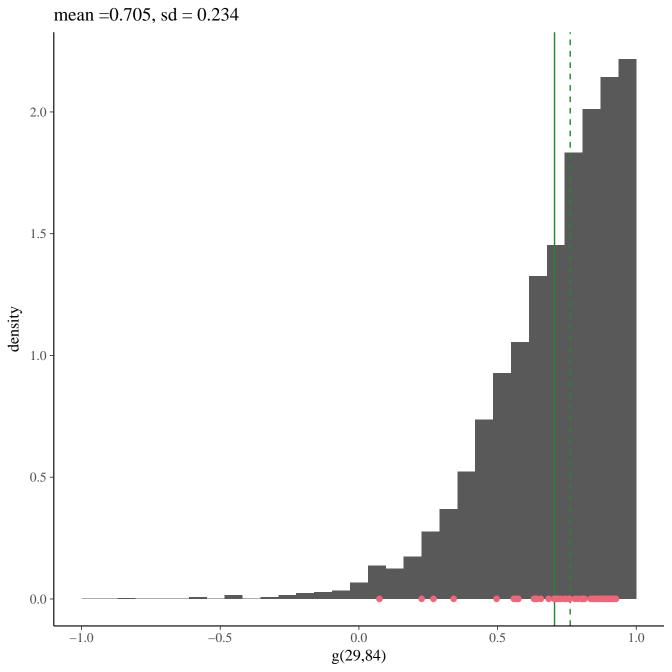
10 graph prop, 55 indiv

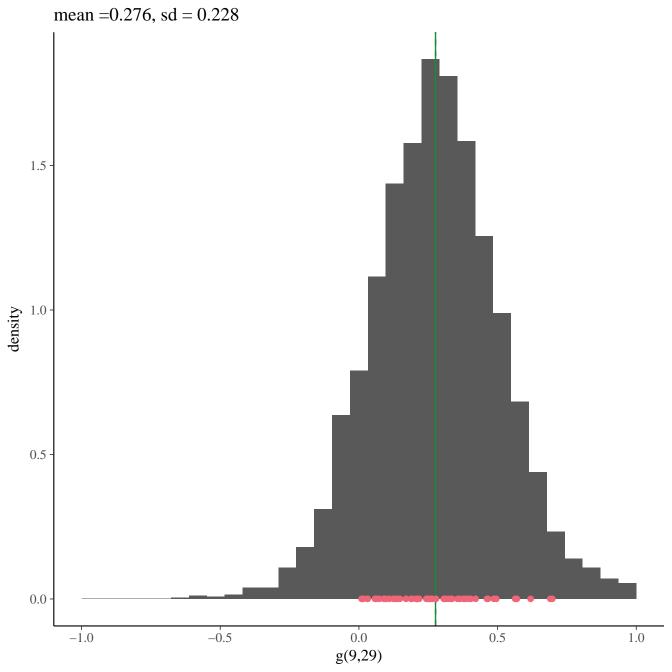
LML = 12.4358093345952

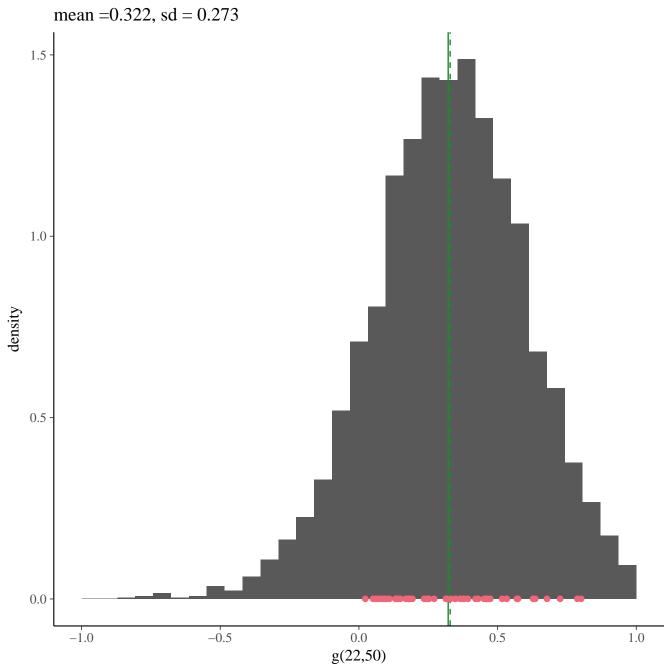
time = 391.06

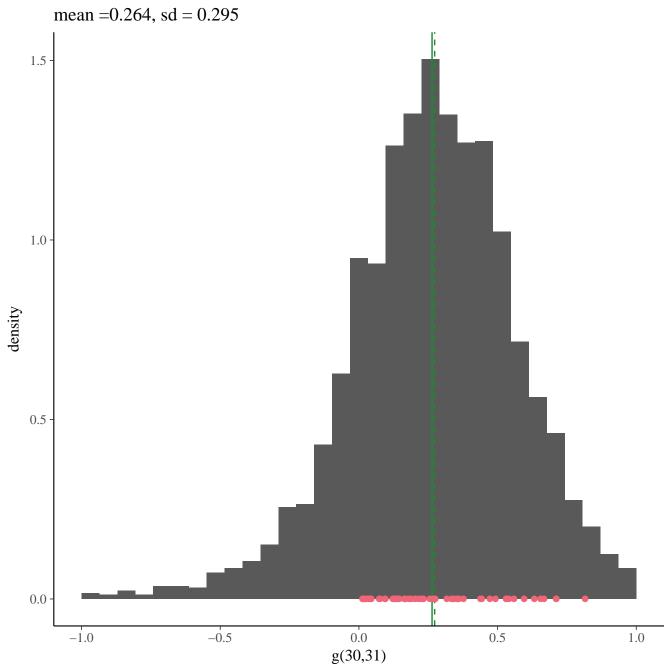


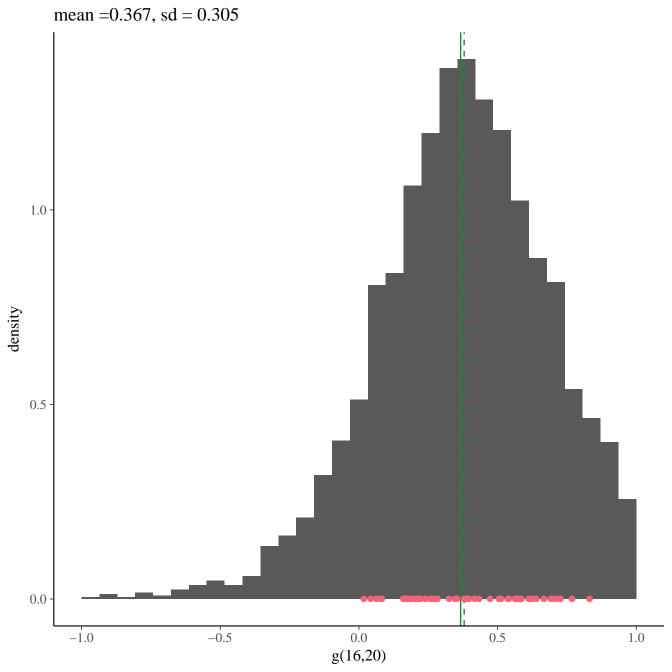


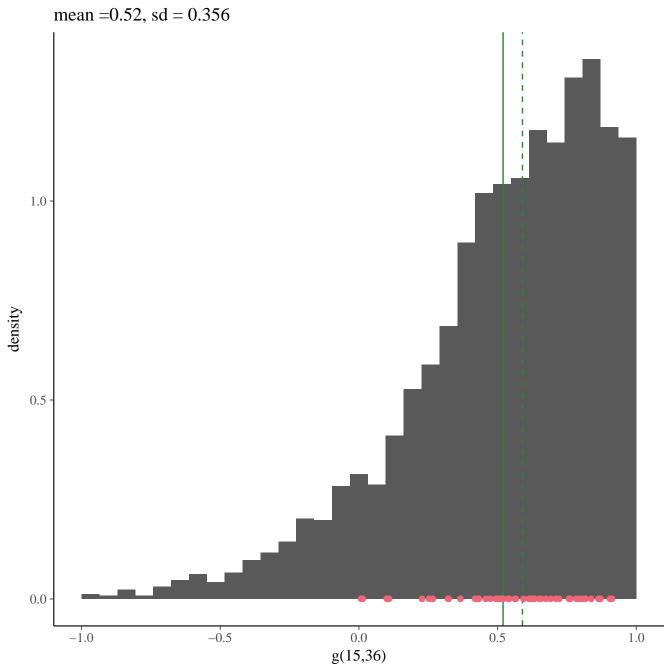


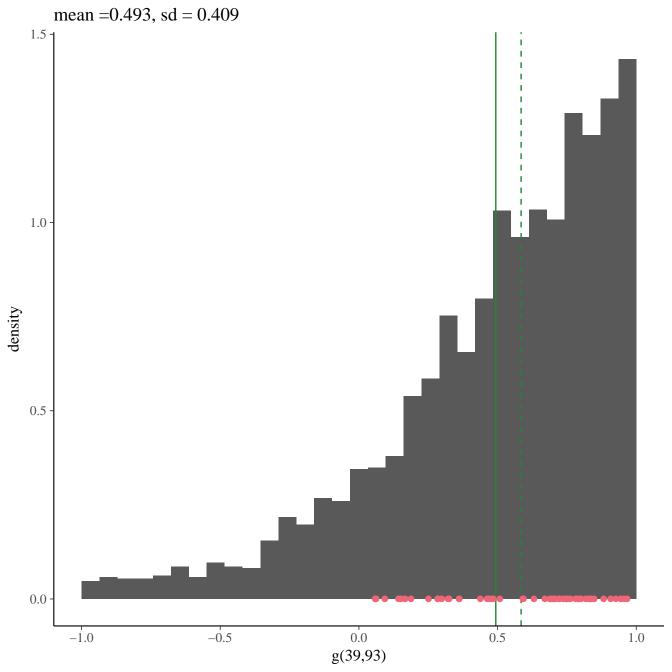


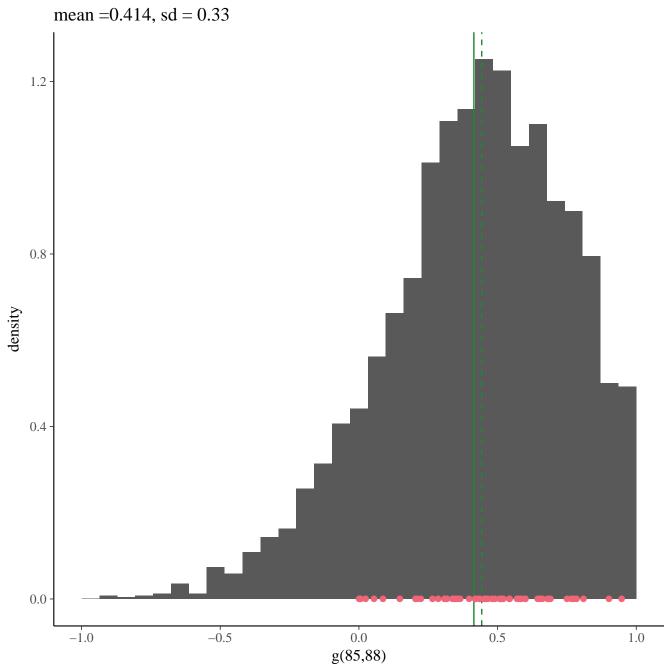


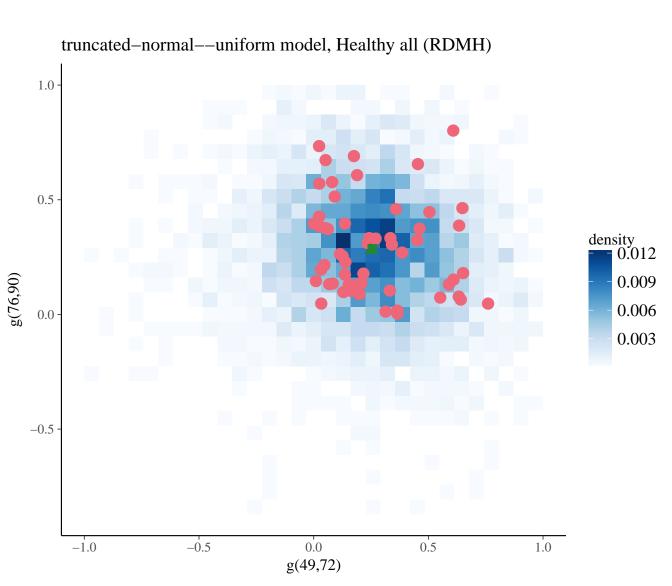


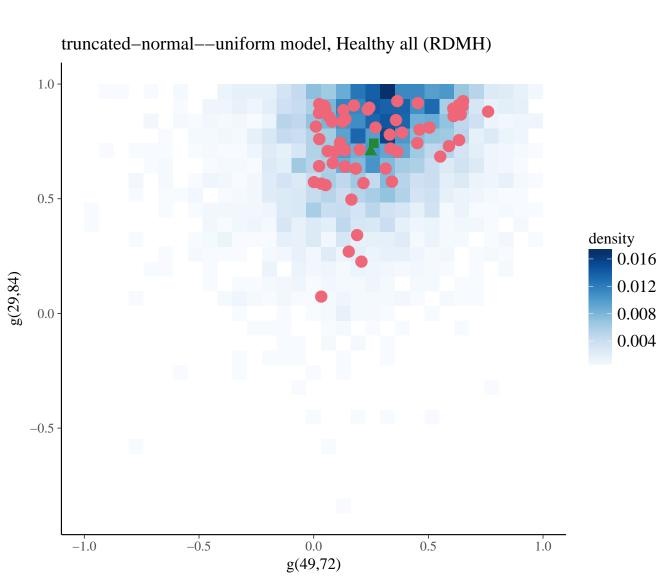


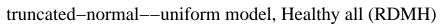


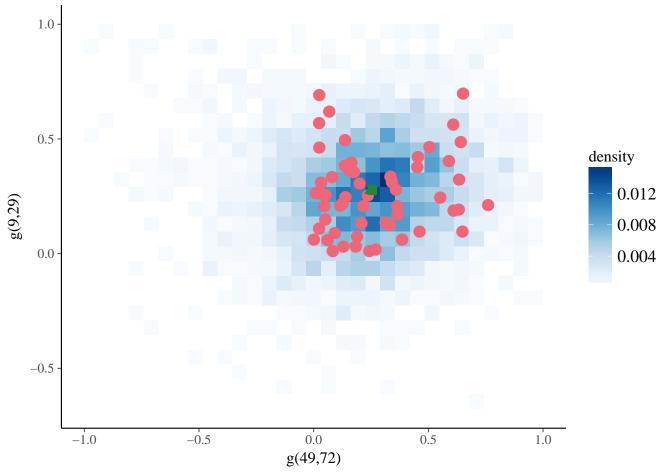


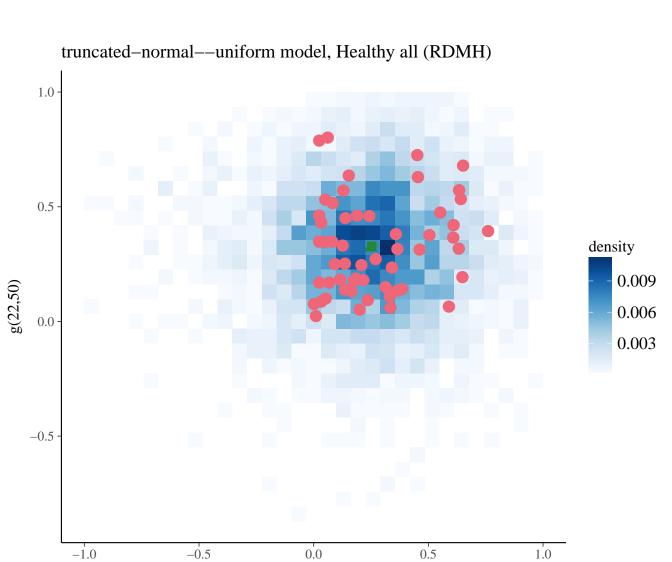




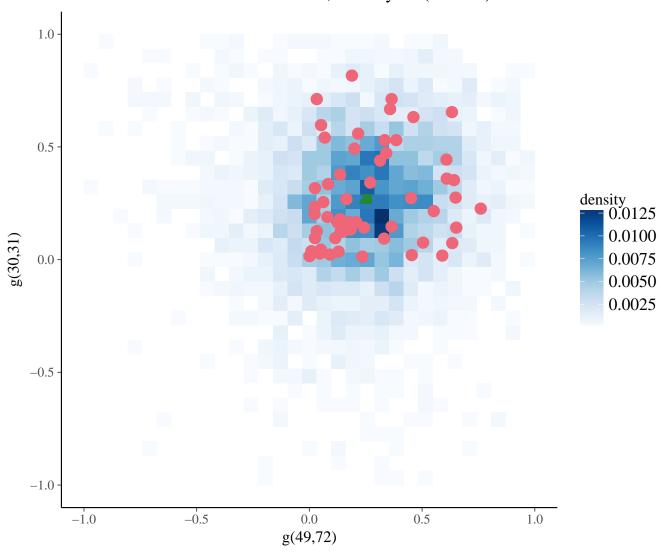


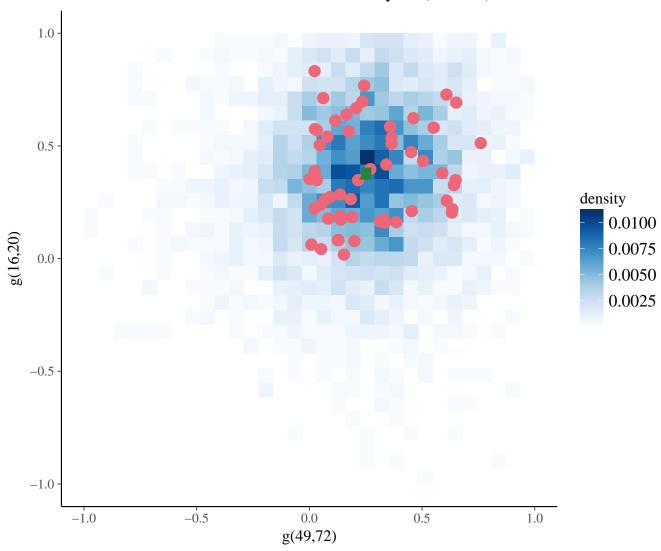


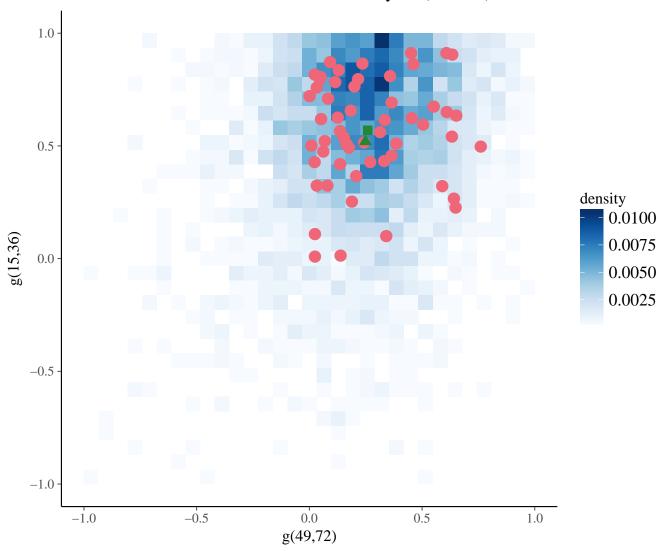


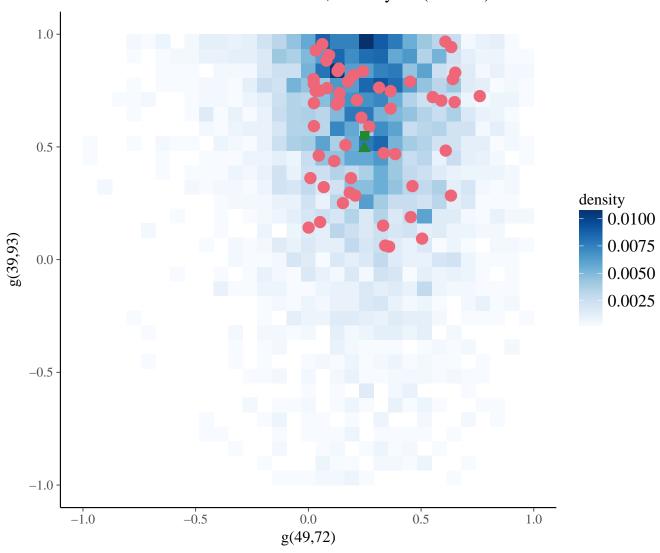


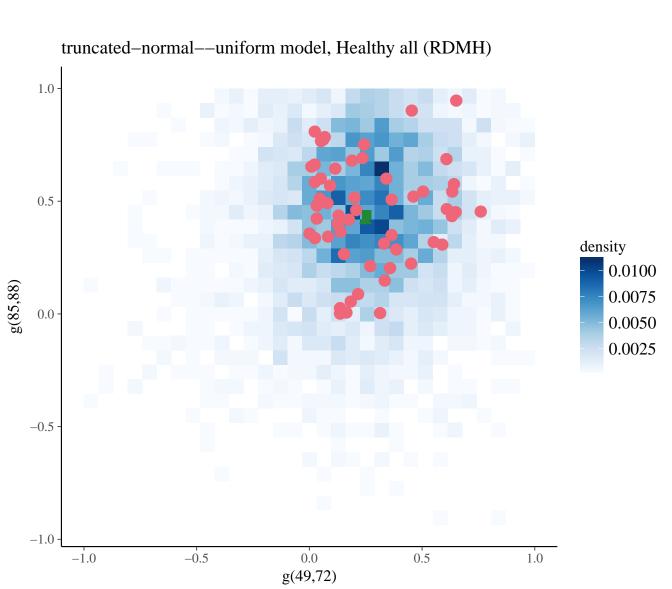
g(49,72)











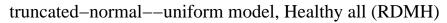
truncated-normal--uniform model, Healthy all (RDMH) 1.0 0.5 density 0.016 0.012 0.008 0.0 0.004 -0.5

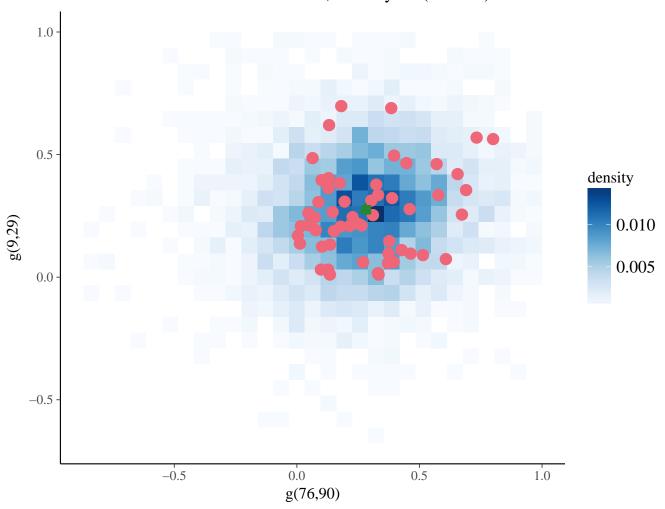
> 0.0 g(76,90)

-0.5

0.5

1.0





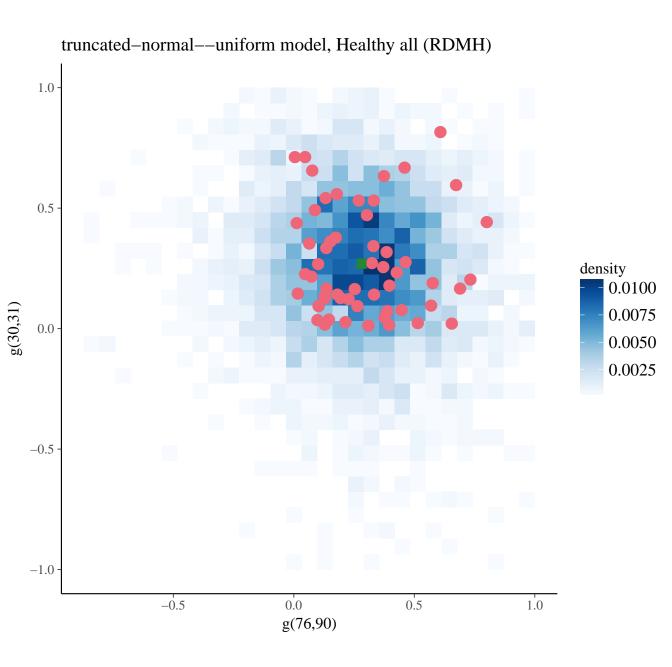
truncated-normal--uniform model, Healthy all (RDMH) 1.0 0.5 density 0.009 0.006 0.0 0.003 -0.5 -

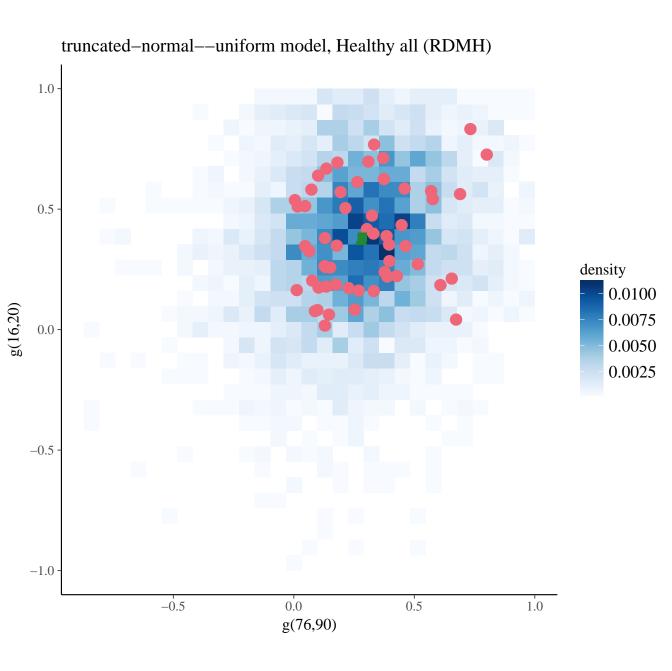
> 0.0 g(76,90)

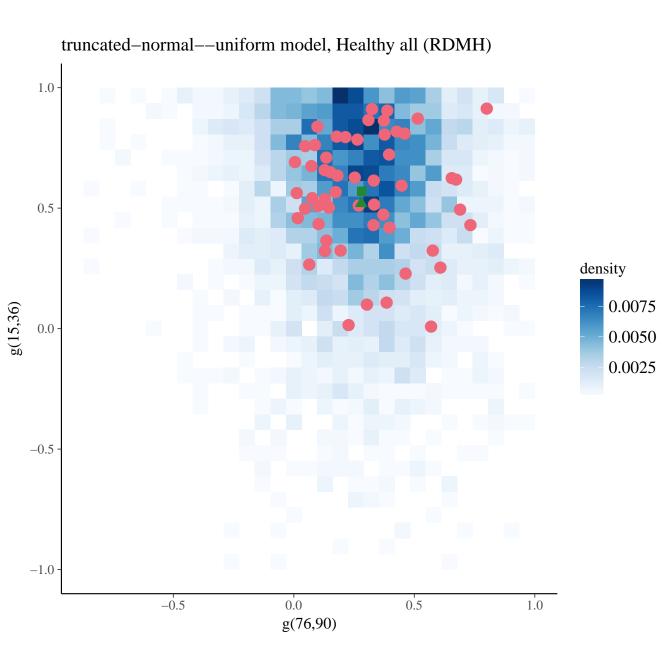
0.5

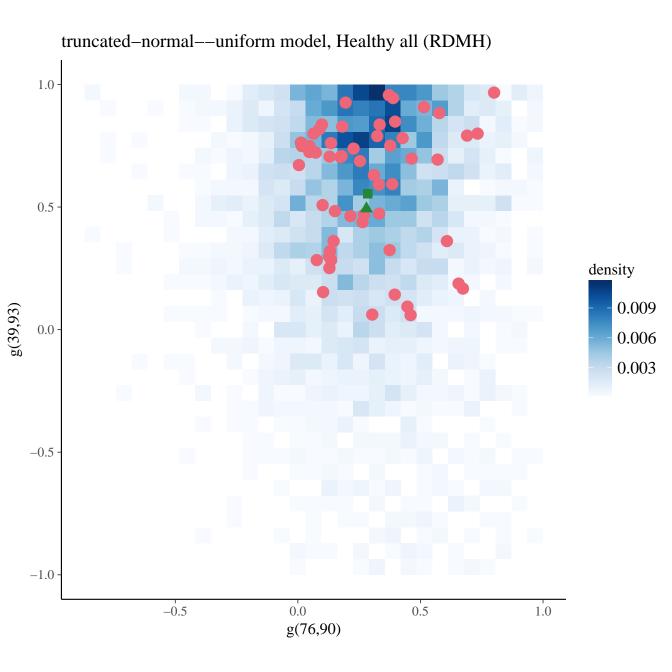
1.0

g(22,50)









truncated-normal--uniform model, Healthy all (RDMH) 1.0 0.5 density 0.009 0.0 0.006 0.003 -0.5

0.0

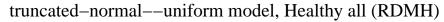
g(76,90)

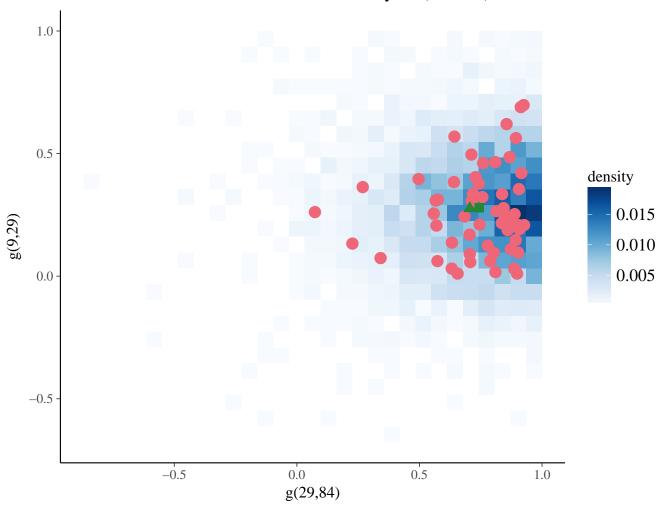
0.5

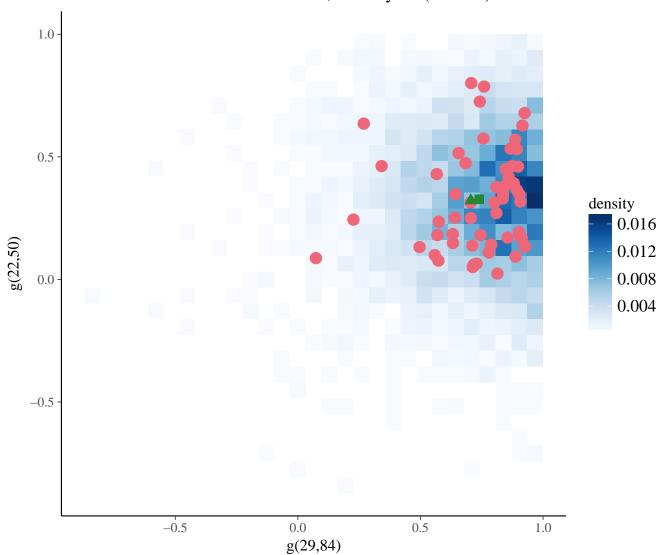
1.0

g(85,88)

-1.0







truncated-normal--uniform model, Healthy all (RDMH) 1.0 0.5 density 0.010 0.0 0.005 -0.5 -1.0

0.0

g(29,84)

0.5

1.0

g(30,31)

truncated-normal--uniform model, Healthy all (RDMH) 1.0 0.5 density 0.010 0.0 0.005 -0.5-1.0

0.0

g(29,84)

0.5

1.0

g(16,20)

truncated-normal--uniform model, Healthy all (RDMH) 1.0 0.5 density g(15,36)0.010 0.0 0.005 -0.5-1.00.5 -0.5 0.0 1.0 g(29,84)

