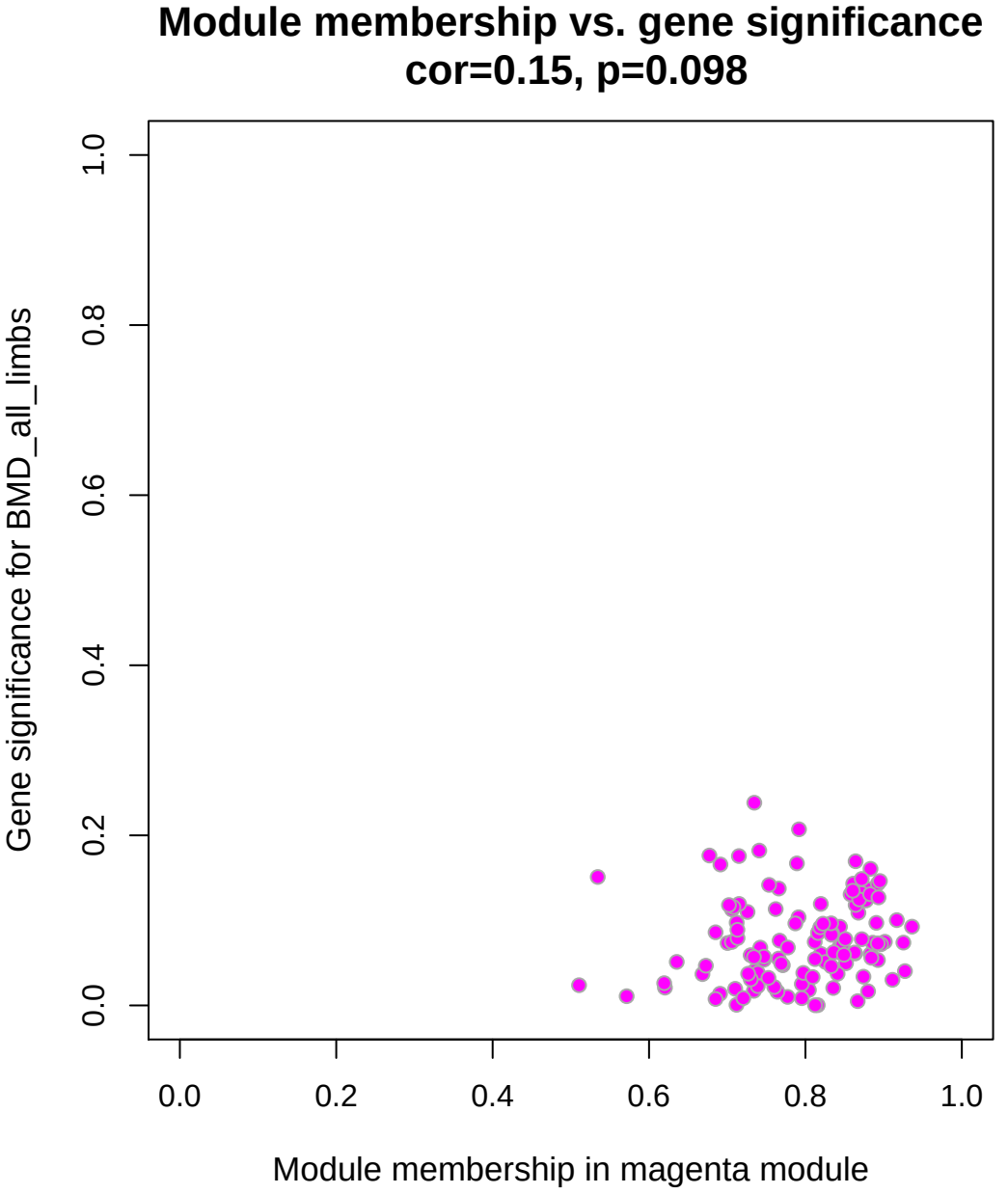
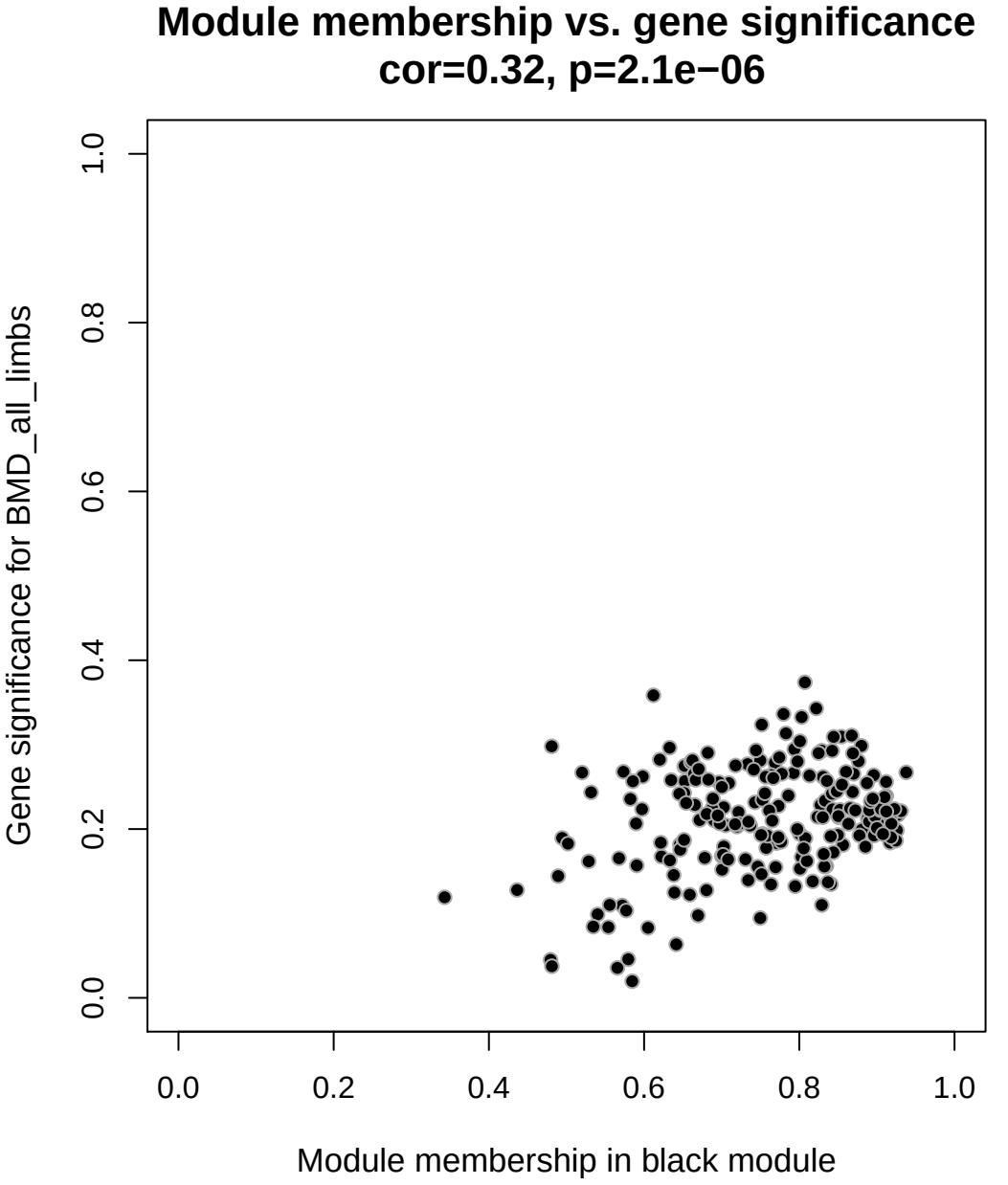
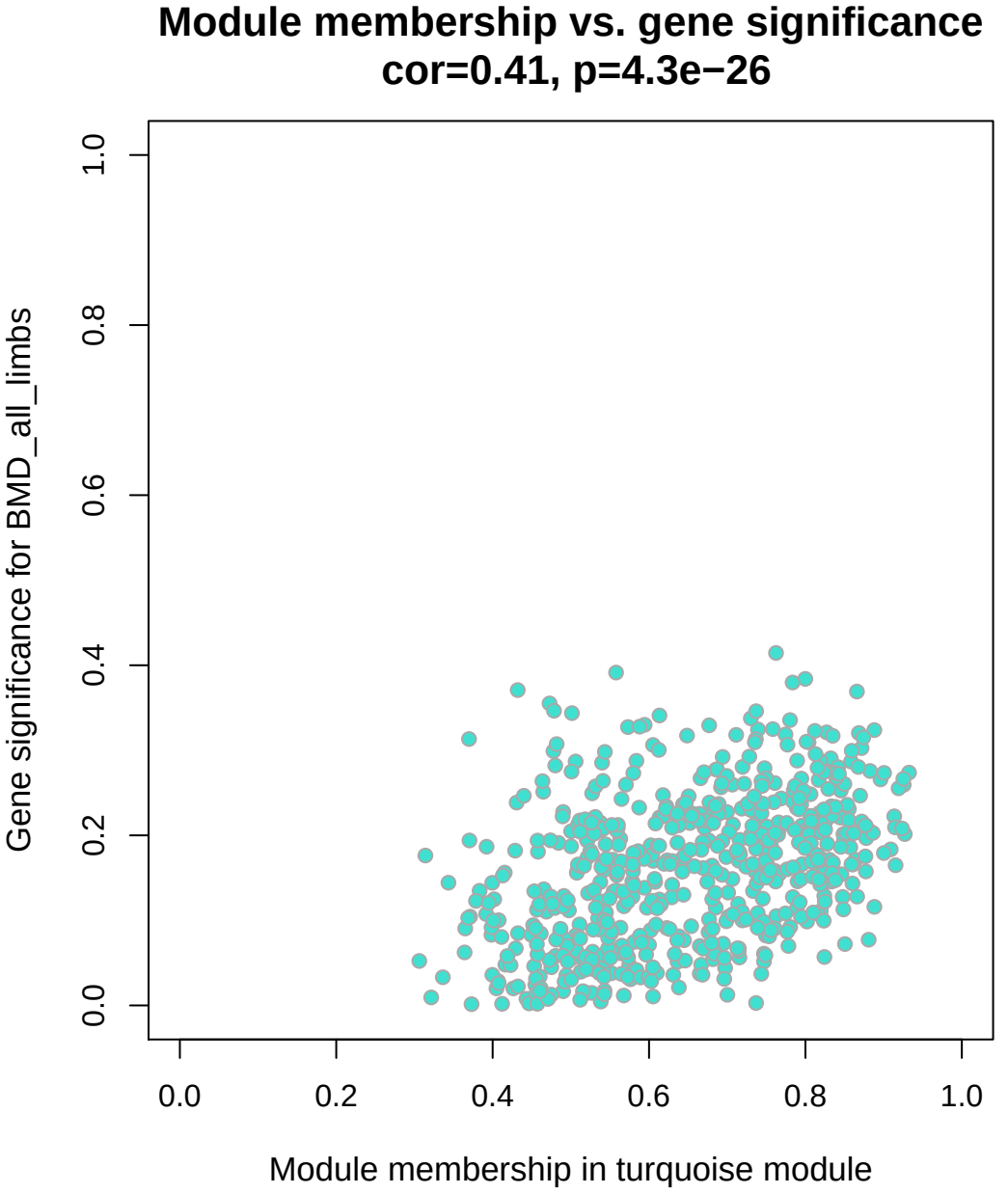
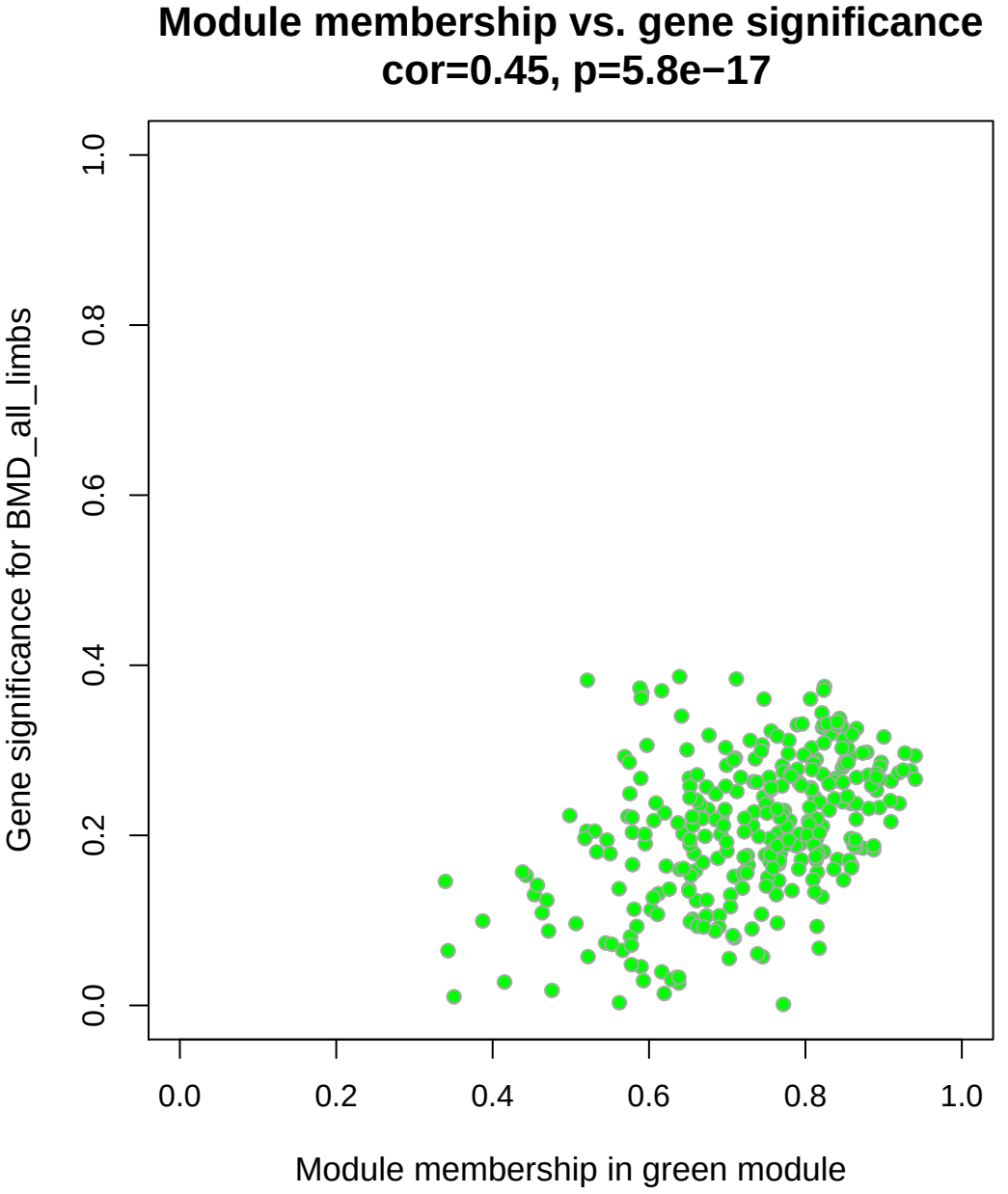
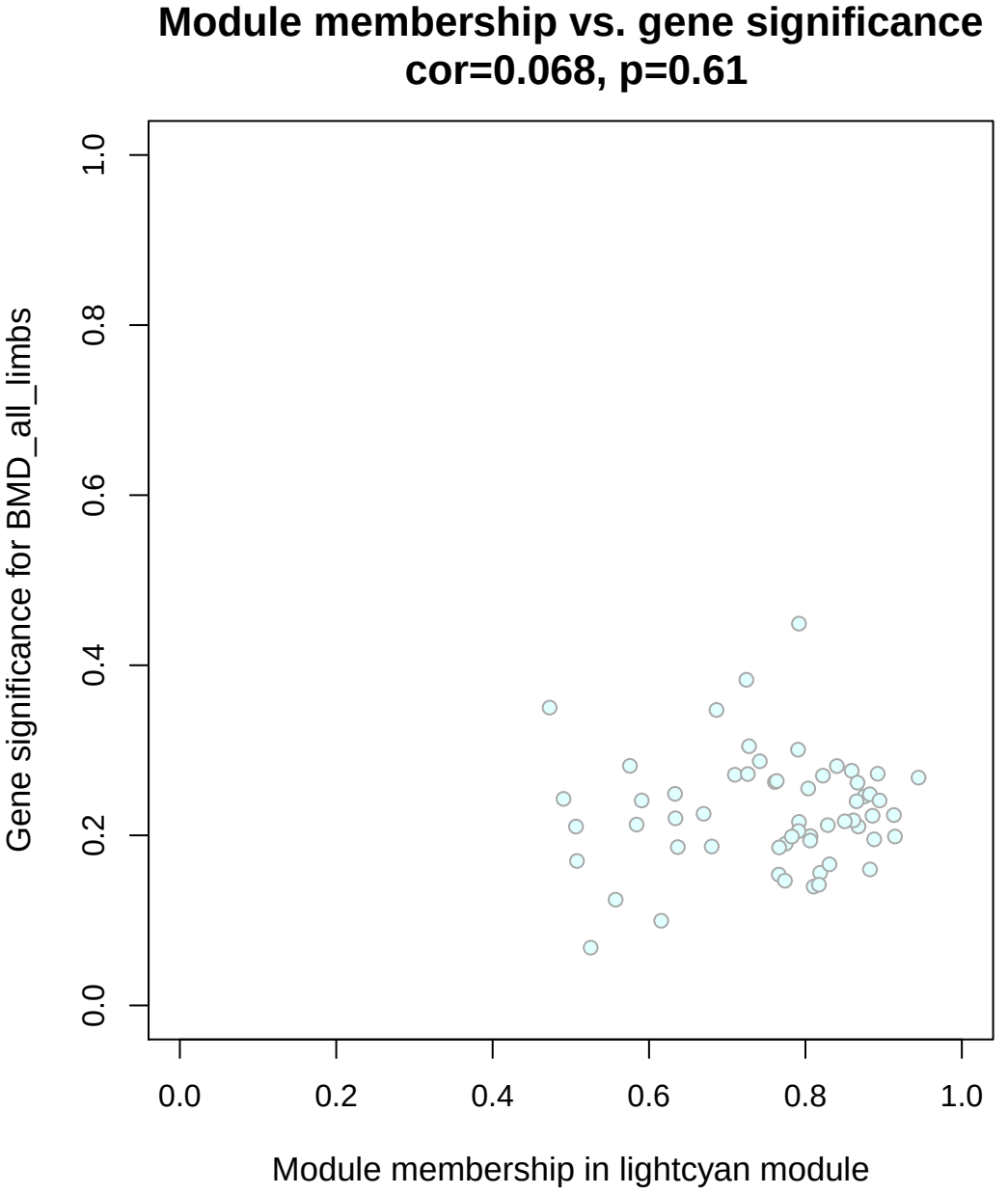
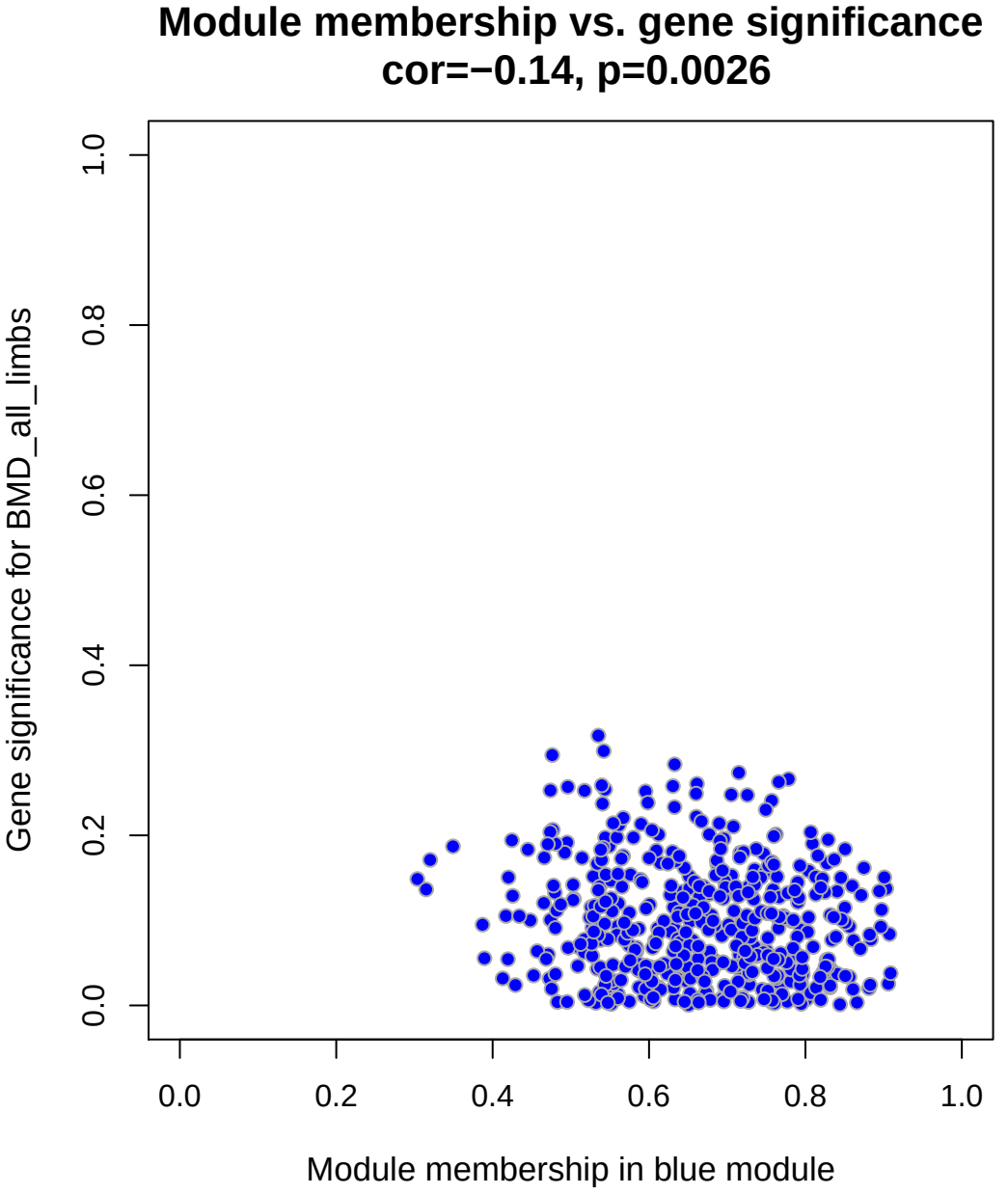

Module membership vs. gene significance
cor=0.15, p=0.098
Gene significance for BMD_all_limbs
0.0
0.2
0.4
0.6
0.8
1.0
Module membership in magenta module

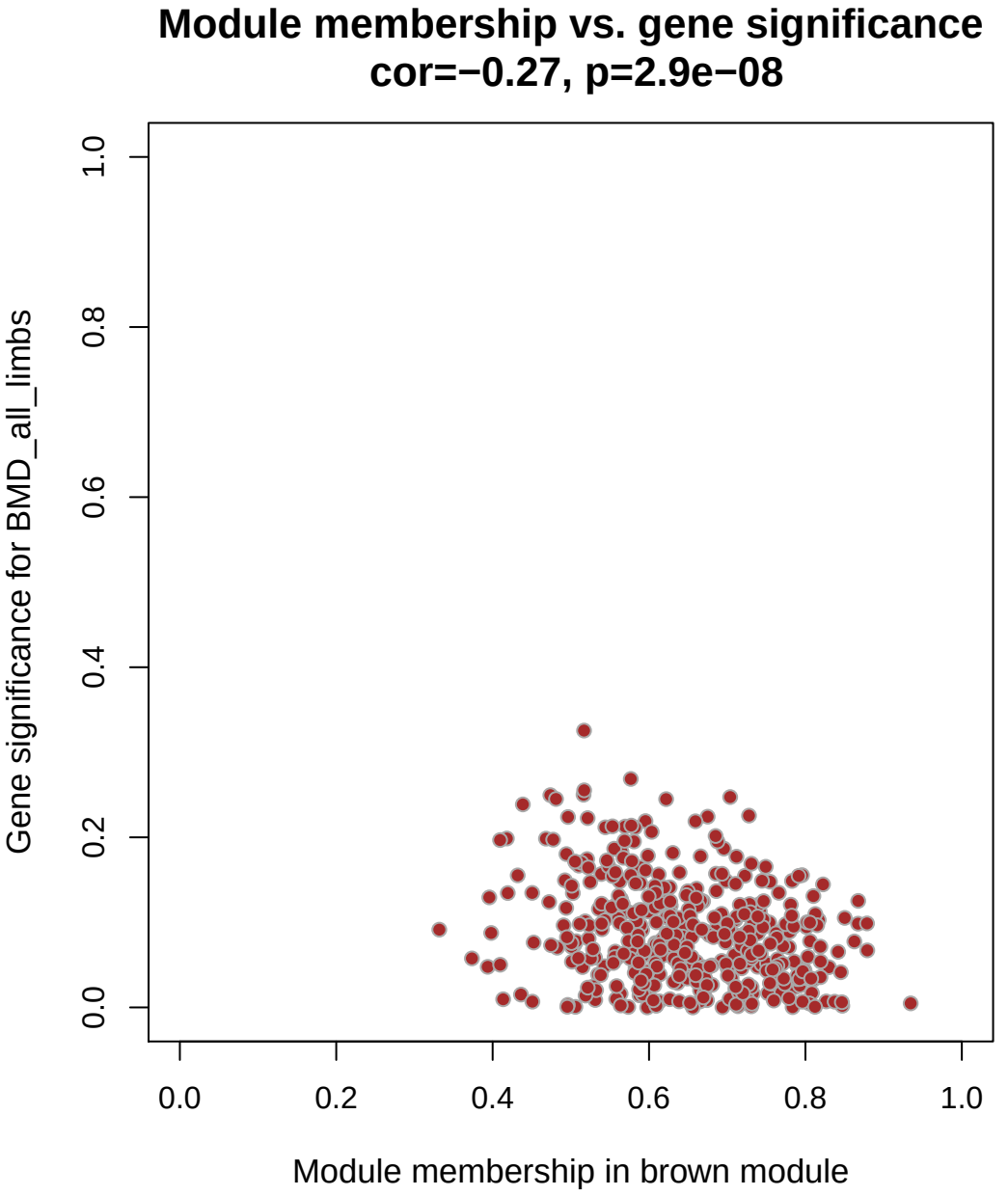

Module membership vs. gene significance
cor=0.32, p=2.1e−06
Gene significance for BMD_all_limbs
0.0
0.2
0.4
0.6
0.8
1.0
Module membership in black module
0.0
0.2
0.4
0.6
0.8
1.0

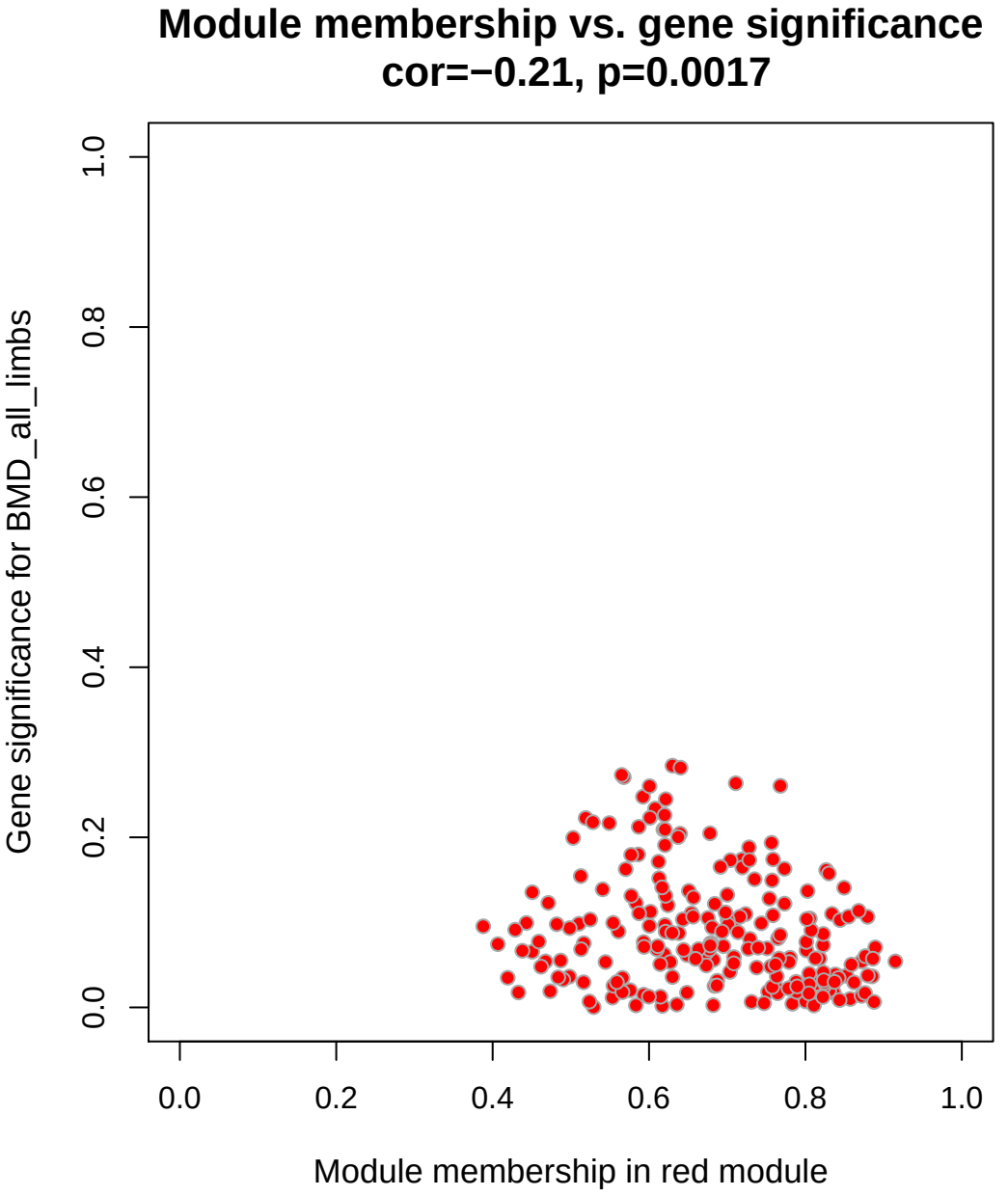

Module membership vs. gene significance
cor=0.41, p=4.3e−26
Gene significance for BMD_all_limbs
0.0
0.2
0.4
0.6
0.8
1.0
Module membership in turquoise module

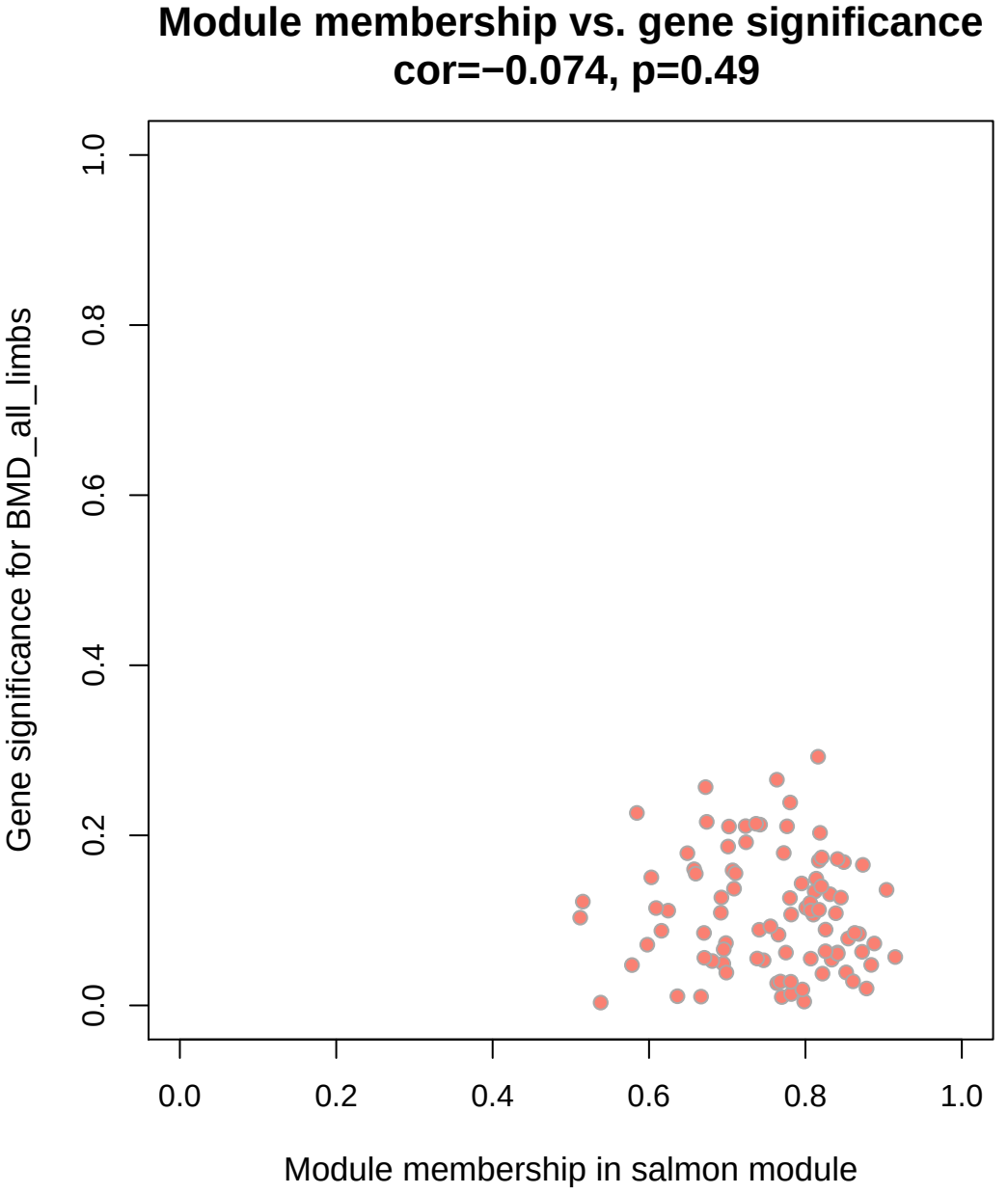

Module membership vs. gene significance
cor=0.45, p=5.8e−17
Gene significance for BMD_all_limbs
0.0
0.2
0.4
0.6
0.8
1.0
Module membership in green module
0.0
0.2
0.4
0.6
0.8
1.0

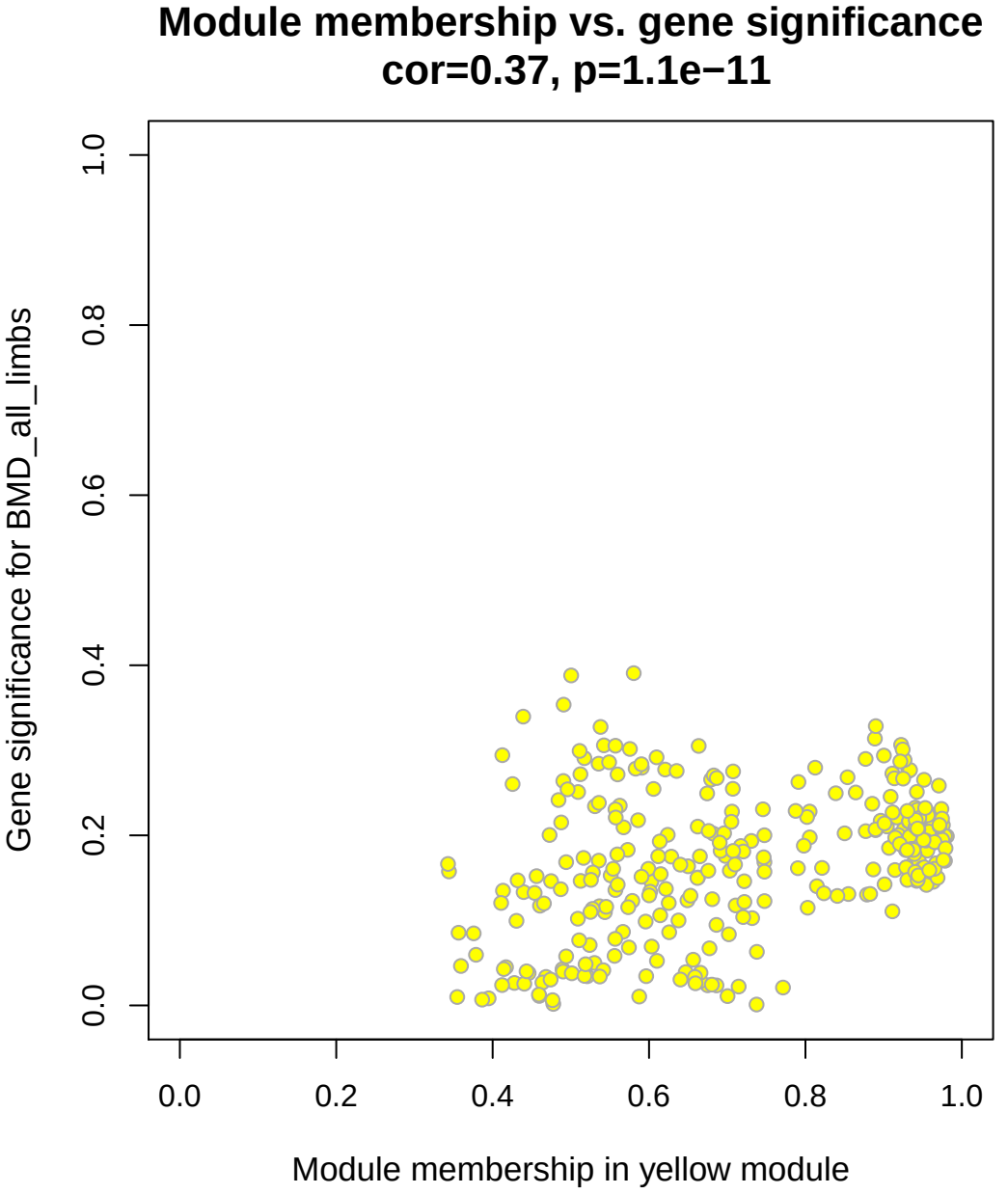

Module membership vs. gene significance
cor=0.068, p=0.61
Gene significance for BMD_all_limbs
0.0
0.2
0.4
0.6
0.8
1.0
Module membership in lightcyan module
0.0
0.2
0.4
0.6
0.8
1.0

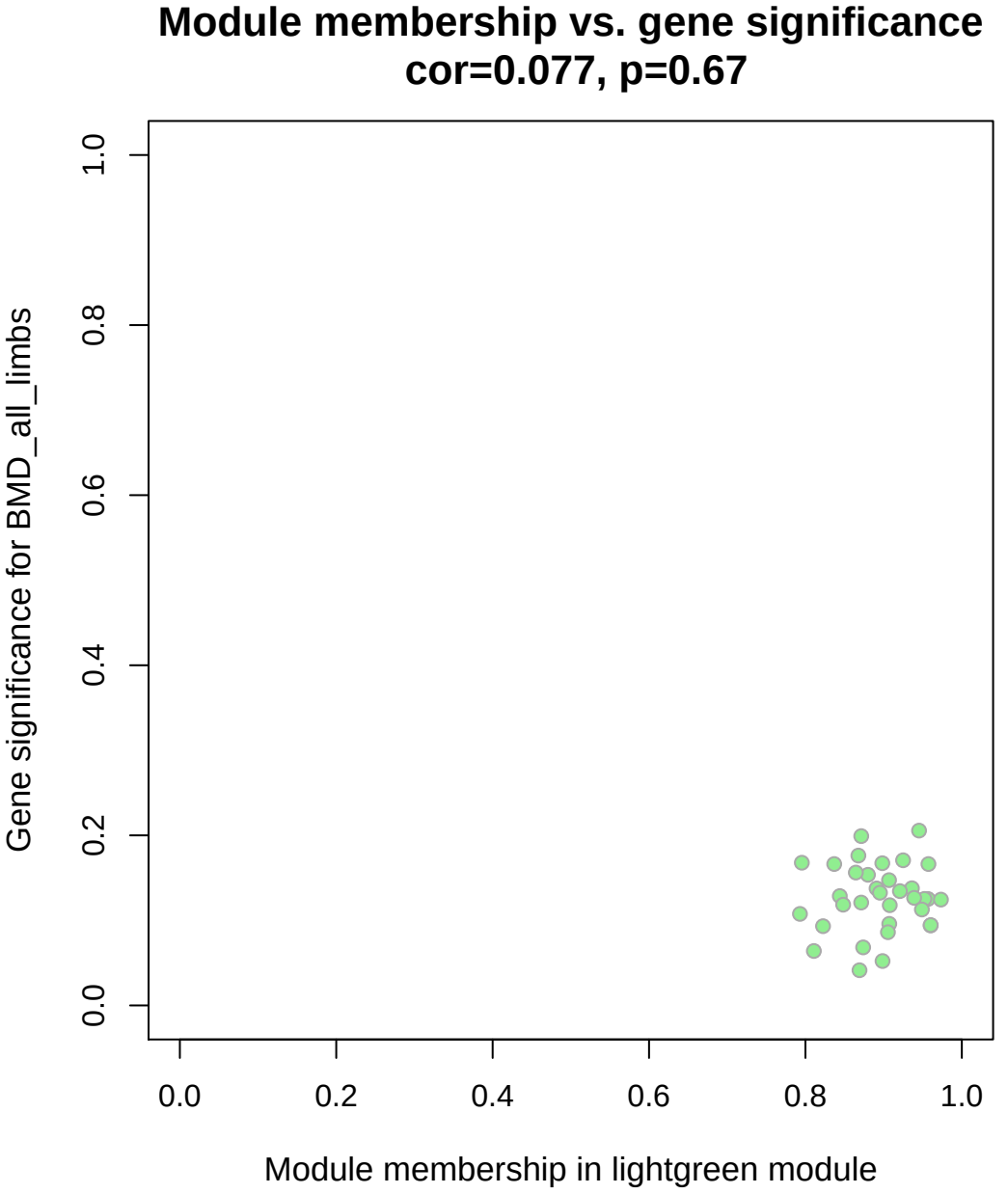

Module membership vs. gene significance
cor=−0.14, p=0.0026
Gene significance for BMD_all_limbs
0.0
0.2
0.4
0.6
0.8
1.0
Module membership in blue module

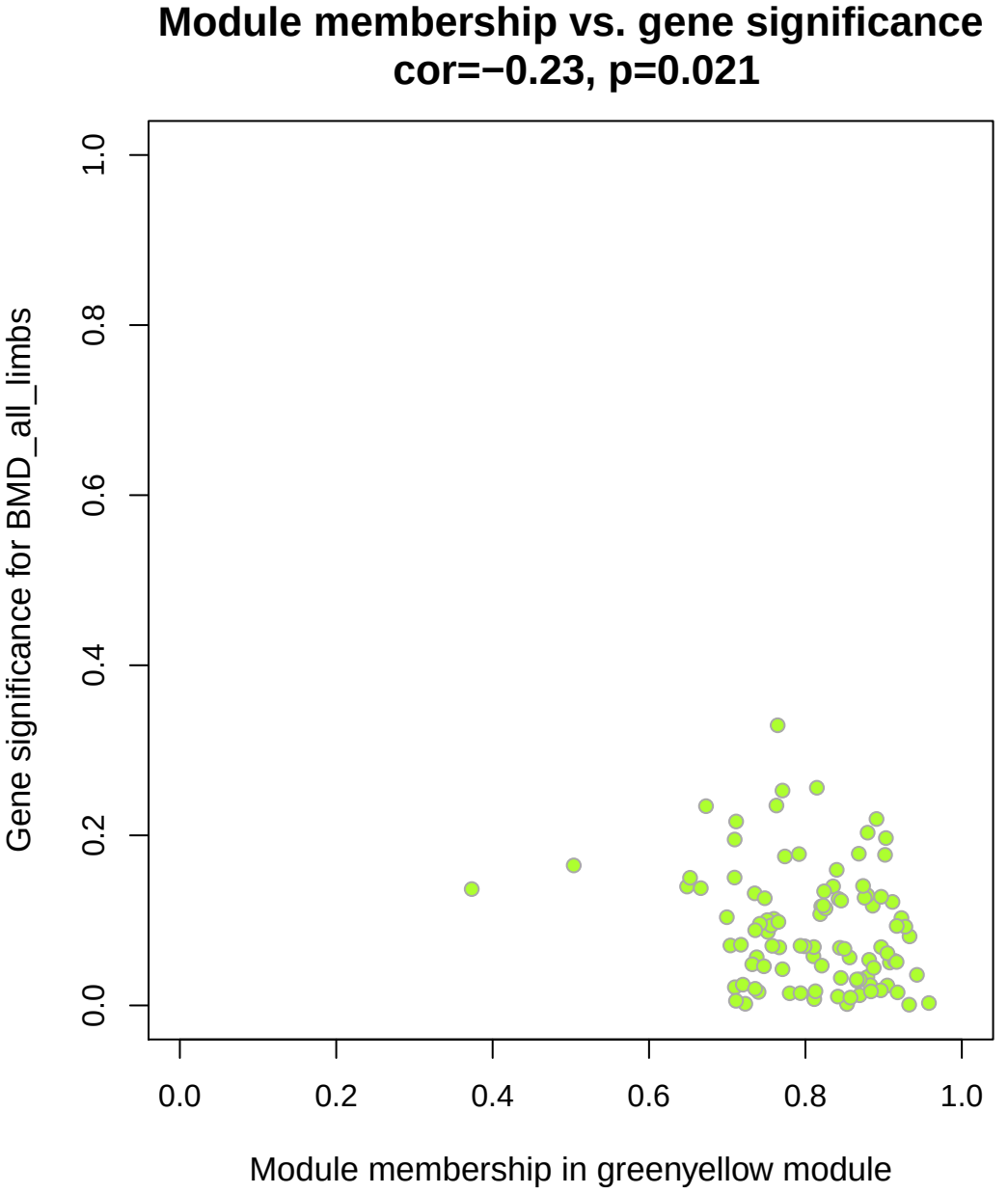

Module membership vs. gene significance
cor=−0.27, p=2.9e−08
Gene significance for BMD_all_limbs
0.0
0.2
0.4
0.6
0.8
1.0
Module membership in brown module

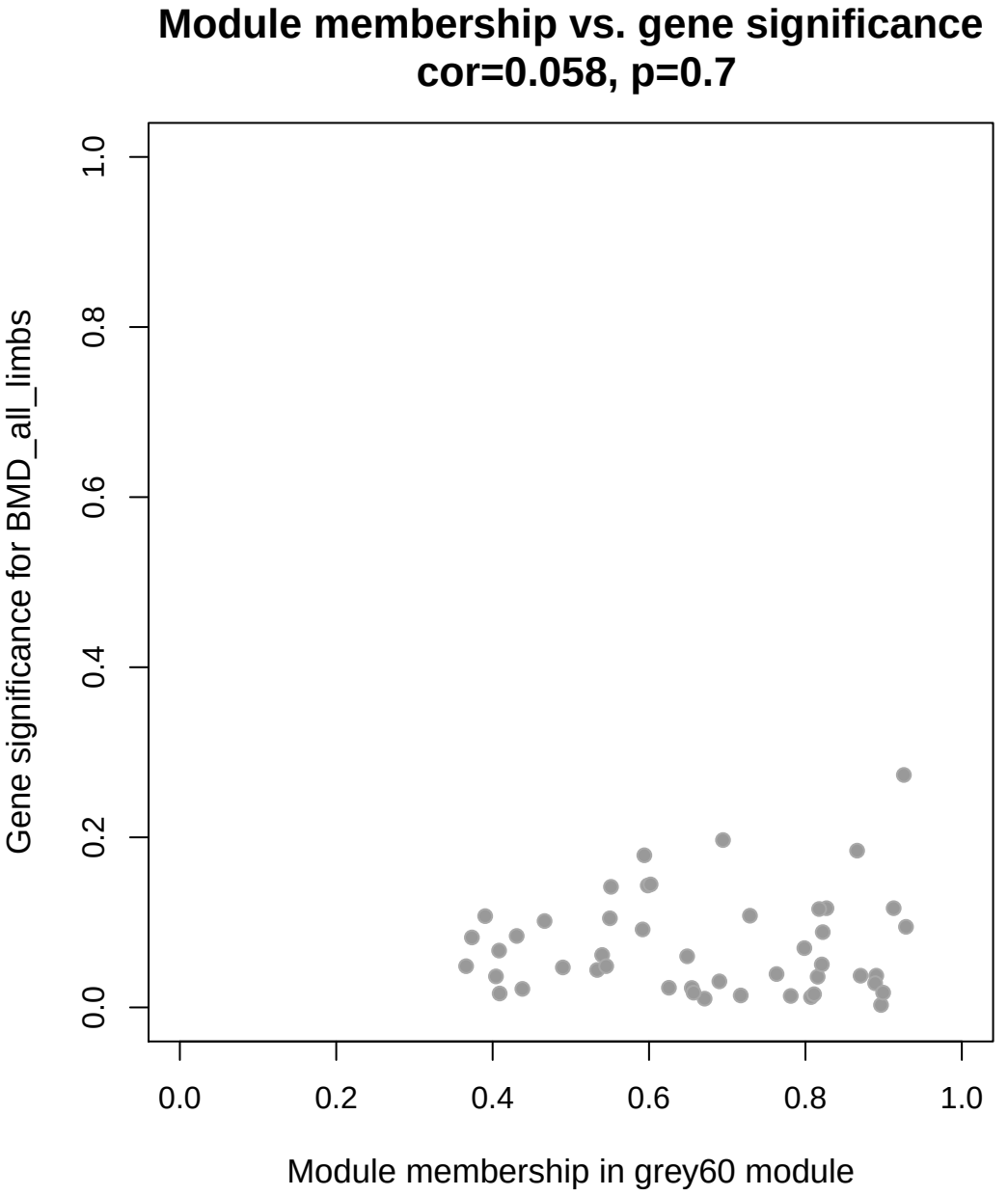

Module membership vs. gene significance
cor=−0.21, p=0.0017
Gene significance for BMD_all_limbs
0.0
0.2
0.4
0.6
0.8
1.0
Module membership in red module

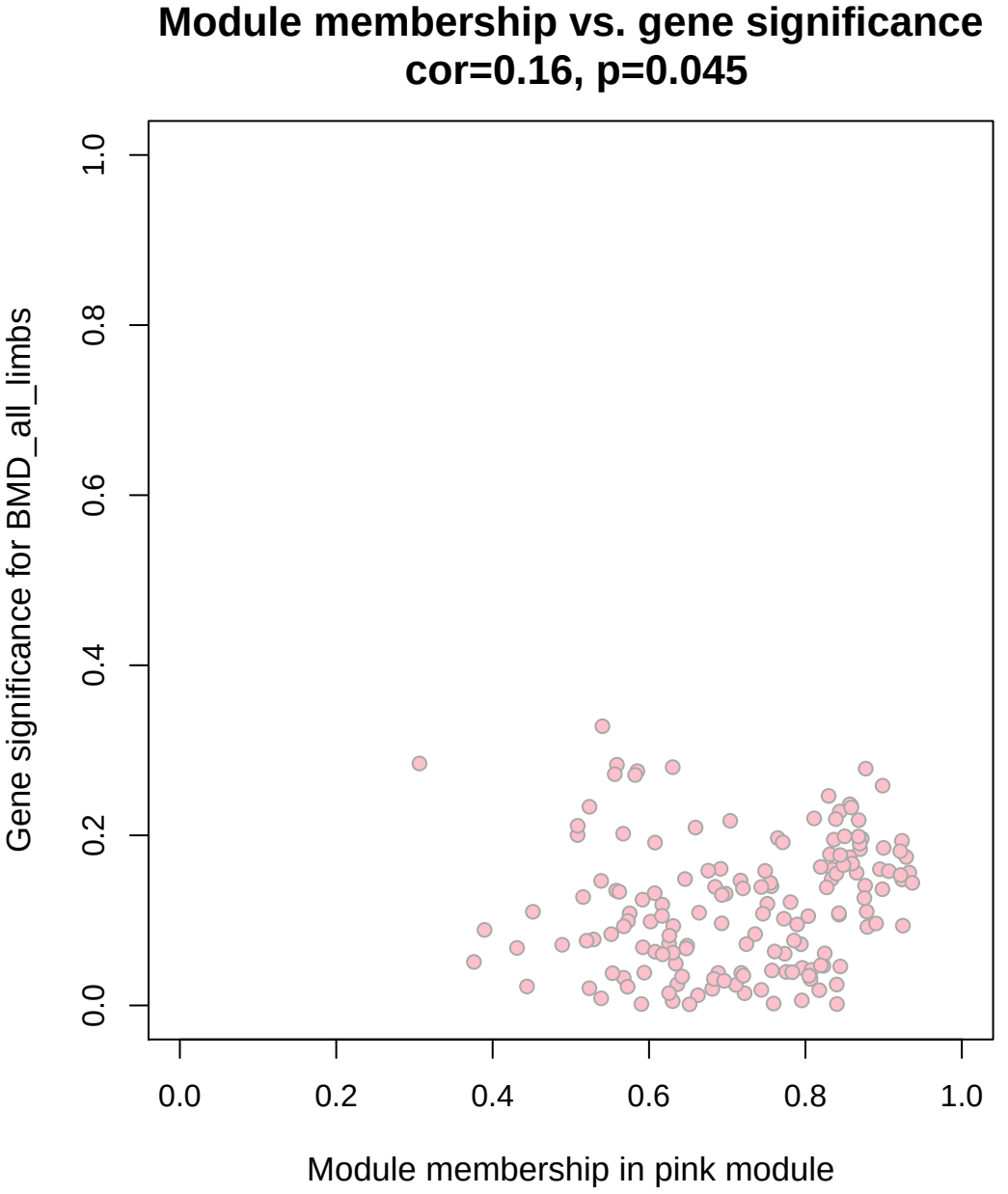

Module membership vs. gene significance
cor=−0.074, p=0.49
Gene significance for BMD_all_limbs
0.0
0.2
0.4
0.6
0.8
1.0
Module membership in salmon module

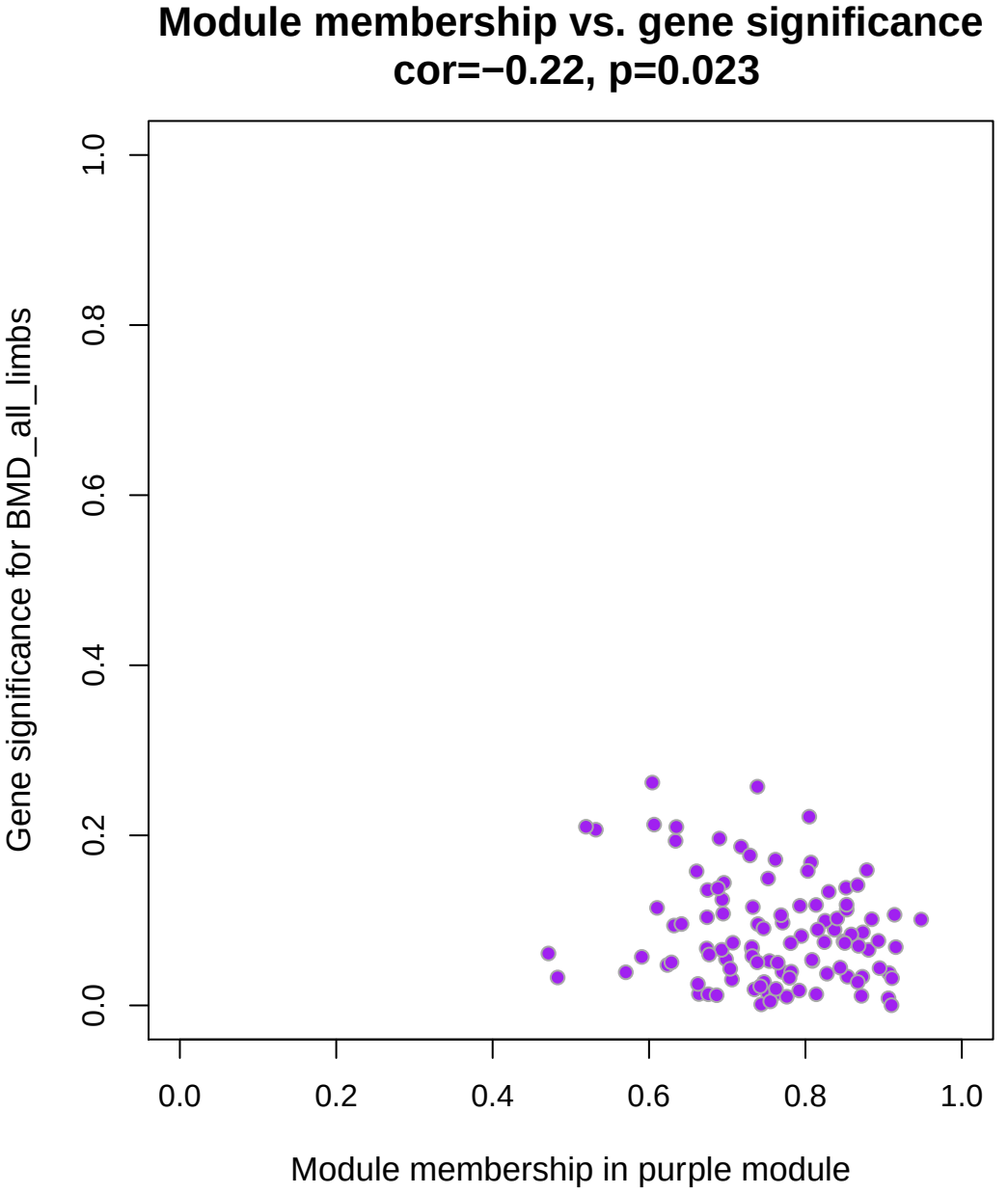

Module membership vs. gene significance
cor=0.37, p=1.1e−11
Gene significance for BMD_all_limbs
0.0
0.2
0.4
0.6
0.8
1.0
Module membership in yellow module
0.0
0.2
0.4
0.6
0.8
1.0

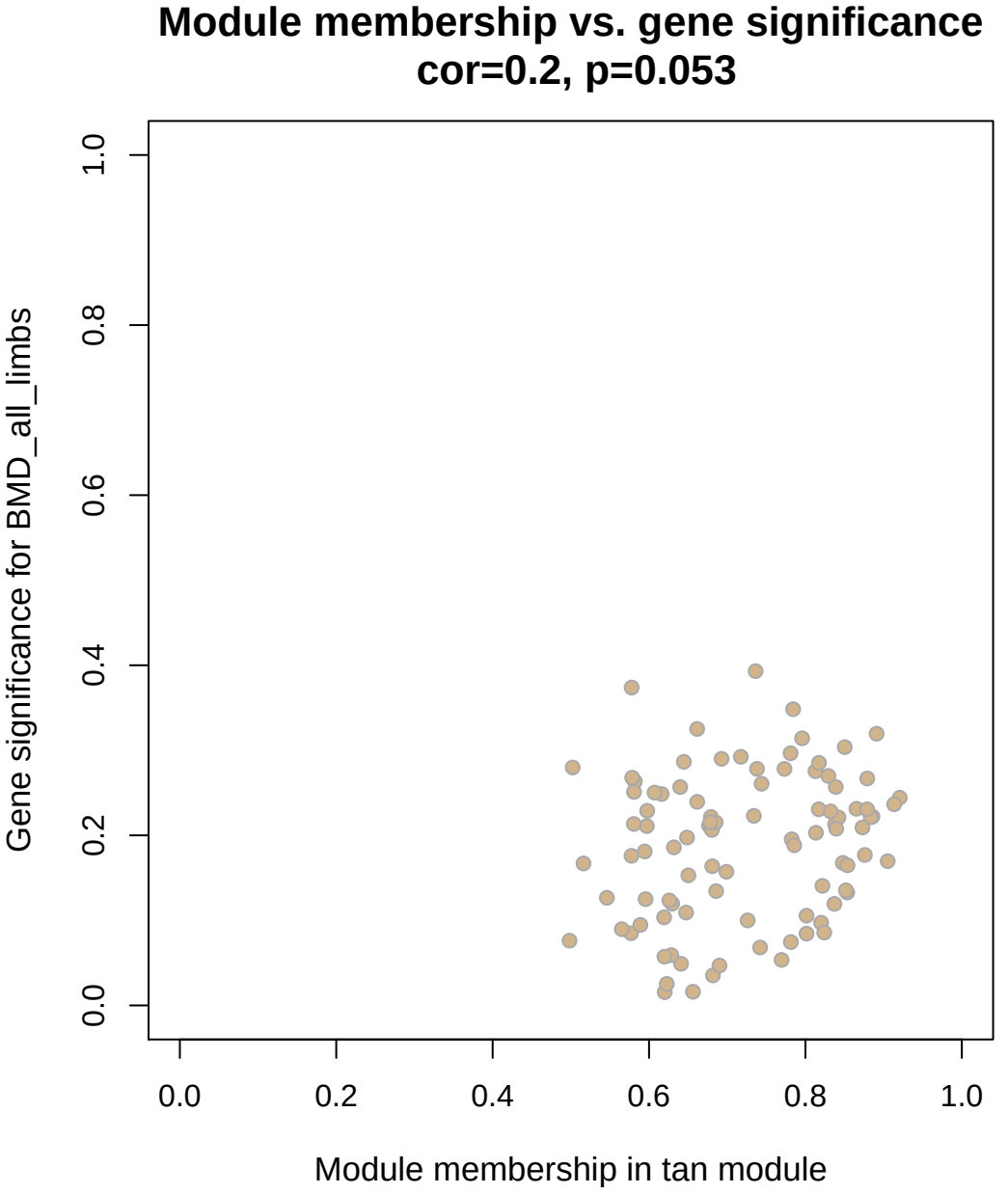

Module membership vs. gene significance
cor=0.077, p=0.67
Gene significance for BMD_all_limbs
0.0
0.2
0.4
0.6
0.8
1.0
Module membership in lightgreen module

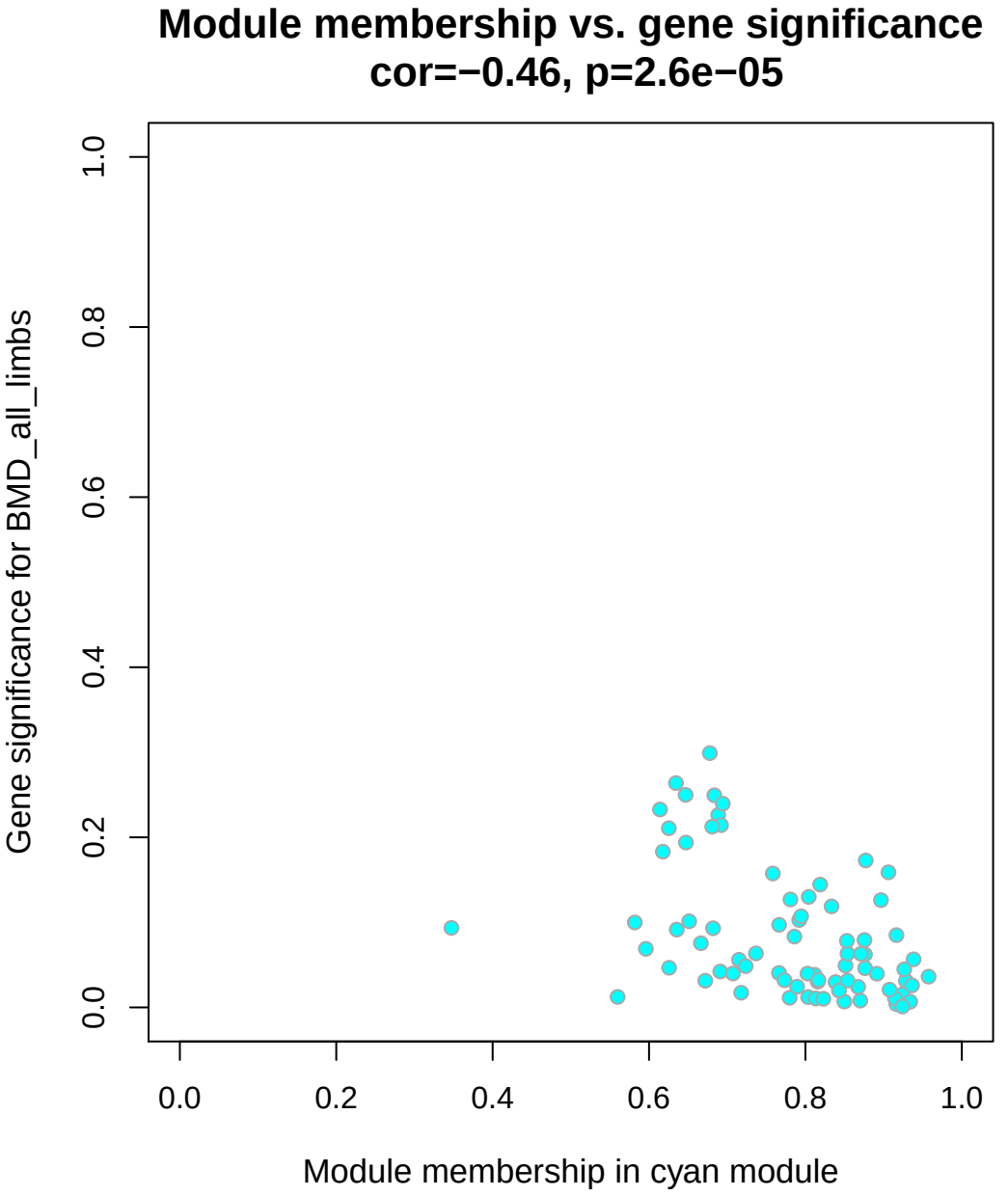

Module membership vs. gene significance
cor=−0.23, p=0.021
Gene significance for BMD_all_limbs
0.0
0.2
0.4
0.6
0.8
1.0
Module membership in greenyellow module

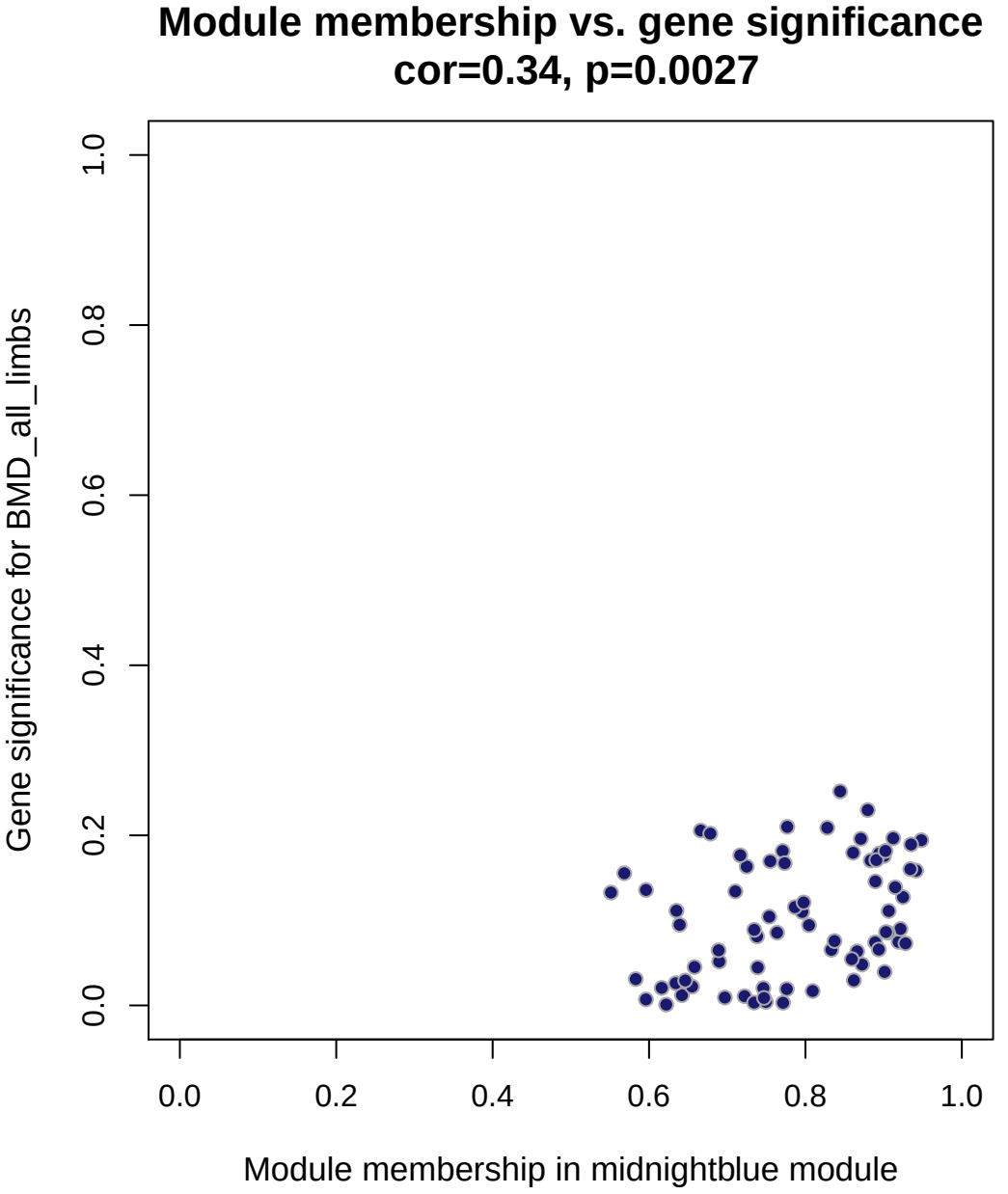

Module membership vs. gene significance
cor=0.058, p=0.7
Gene significance for BMD_all_limbs
0.0
0.2
0.4
0.6
0.8
1.0
Module membership in grey60 module
0.0
0.2
0.4
0.6
0.8
1.0

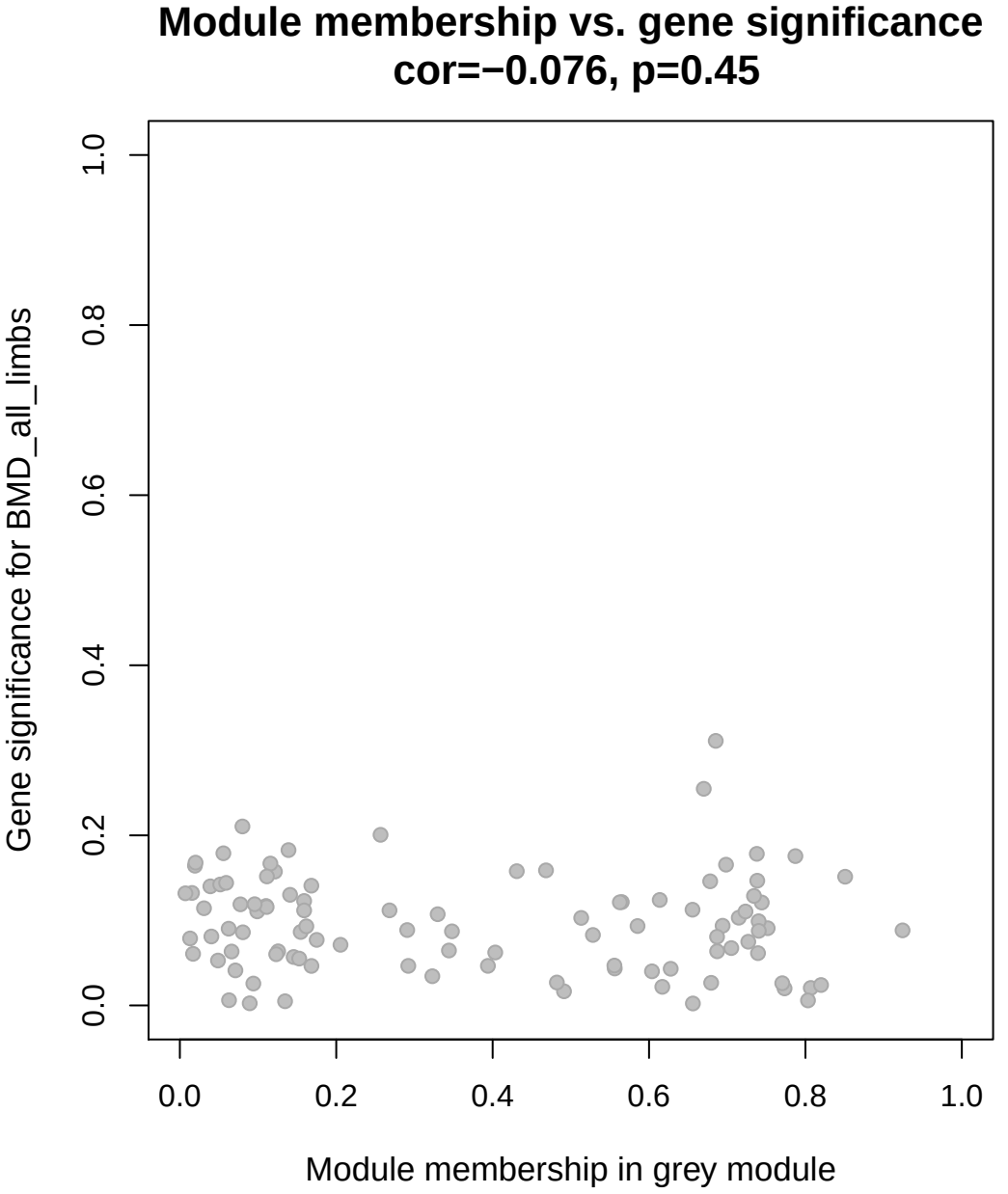

Module membership vs. gene significance
cor=0.16, p=0.045
Gene significance for BMD_all_limbs
0.0
0.2
0.4
0.6
0.8
1.0
Module membership in pink module


Module membership vs. gene significance
cor=−0.22, p=0.023
Gene significance for BMD_all_limbs
0.0
0.2
0.4
0.6
0.8
1.0
Module membership in purple module
0.0
0.2
0.4
0.6
0.8
1.0


Module membership vs. gene significance
cor=0.2, p=0.053
Gene significance for BMD_all_limbs
0.0
0.2
0.4
0.6
0.8
1.0
Module membership in tan module
0.0
0.2
0.4
0.6
0.8
1.0


Module membership vs. gene significance
cor=−0.46, p=2.6e−05
Gene significance for BMD_all_limbs
0.0
0.2
0.4
0.6
0.8
1.0
Module membership in cyan module


Module membership vs. gene significance
cor=0.34, p=0.0027
Gene significance for BMD_all_limbs
0.0
0.2
0.4
0.6
0.8
1.0
Module membership in midnightblue module


Module membership vs. gene significance
cor=−0.076, p=0.45
Gene significance for BMD_all_limbs
0.0
0.2
0.4
0.6
0.8
1.0
Module membership in grey module