Module membership vs. gene significance
cor=−0.11, p=0.23

Gene significance for BMD_femurs_only vs. Module membership in magenta module

Module membership vs. gene significance
cor=0.34, p=4.2e−07

Gene significance for BMD_femurs_only

Module membership in black module

Module membership vs. gene significance
cor=0.27, p=1.2e−11

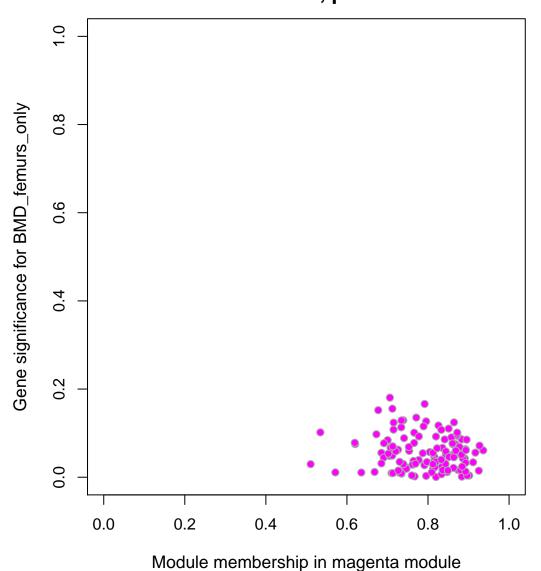**Module membership vs. gene significance**
cor=0.38, p=3.7e−12
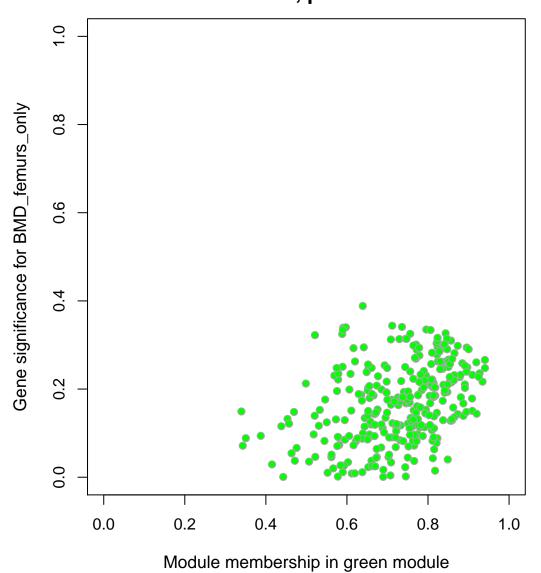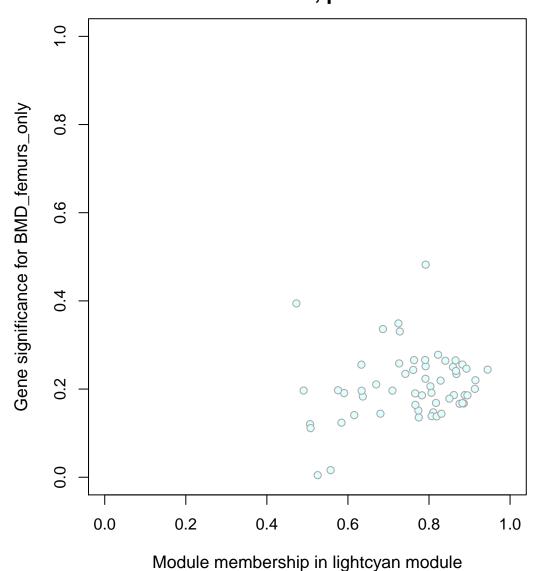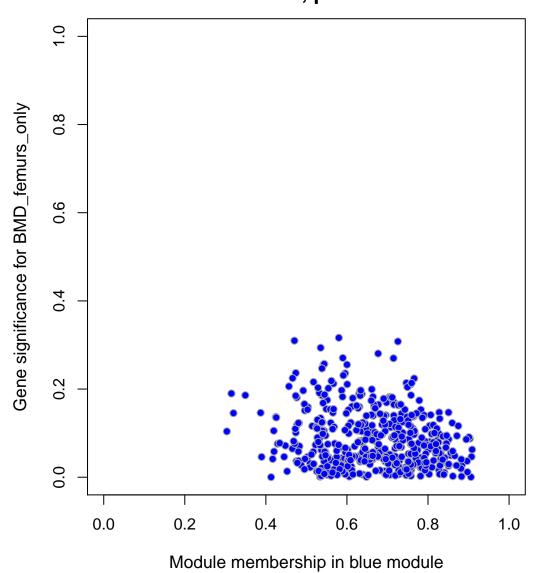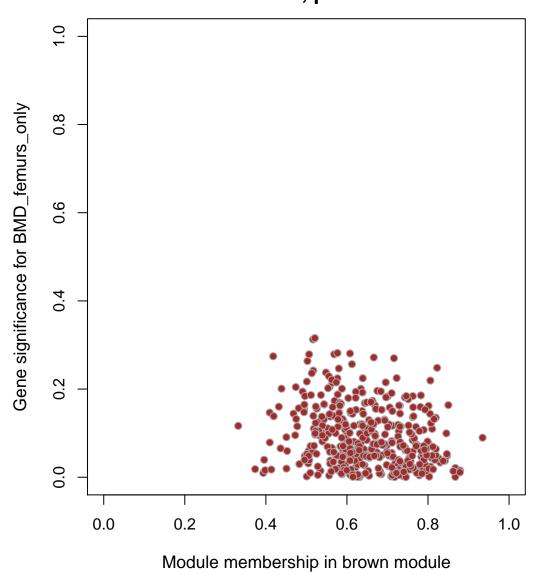
Module membership vs. gene significance
cor=0.19, p=0.15

Module membership vs. gene significance
cor=−0.19, p=4.1e−05

Gene significance for BMD_femurs_only

Module membership in blue module

Module membership vs. gene significance
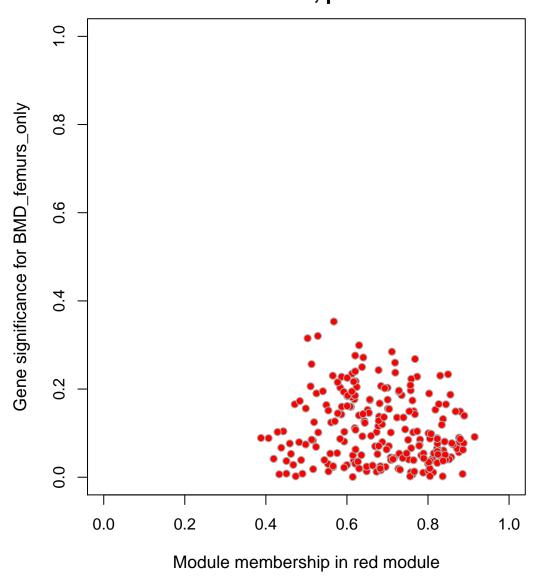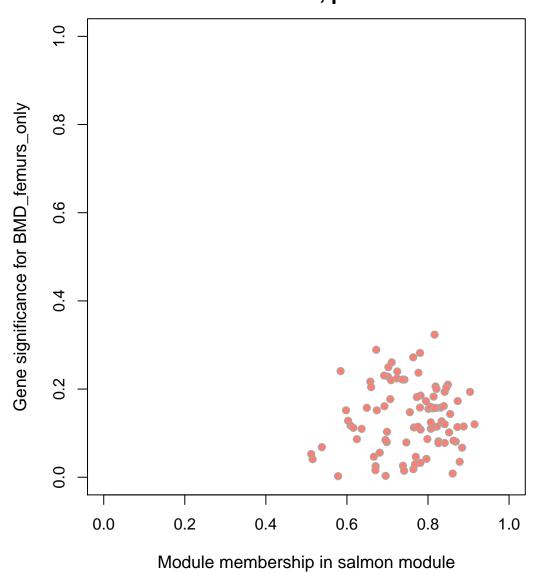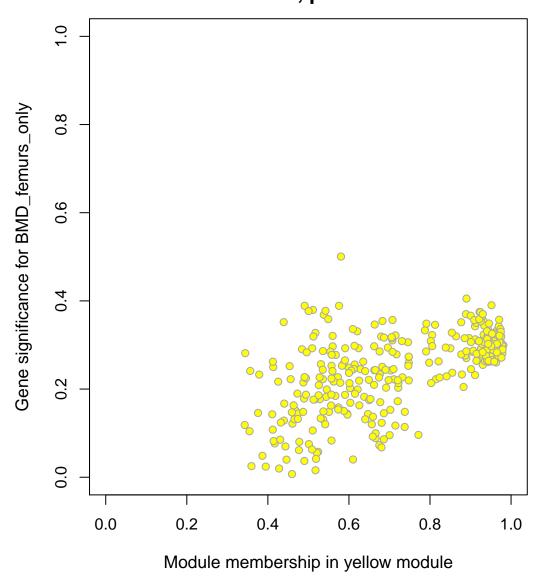**cor=−0.22, p=7.1e−06**

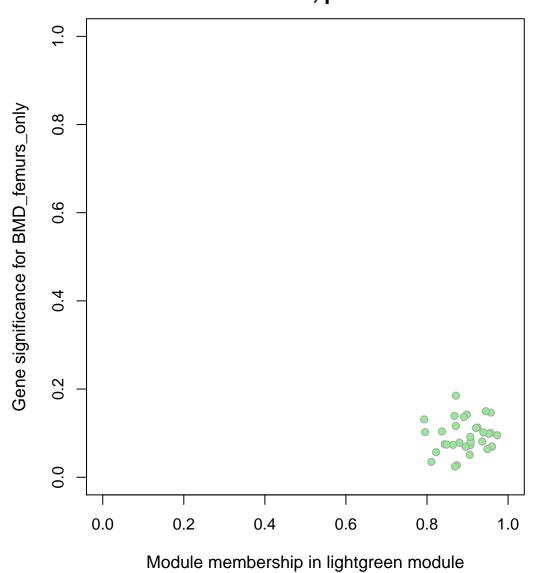**Module membership vs. gene significance**
cor=−0.13, p=0.054

Module membership vs. gene significance
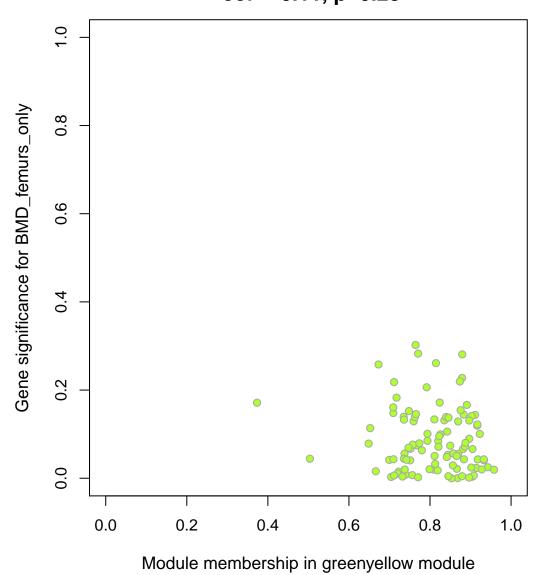**cor=0.076, p=0.47**

Module membership vs. gene significance
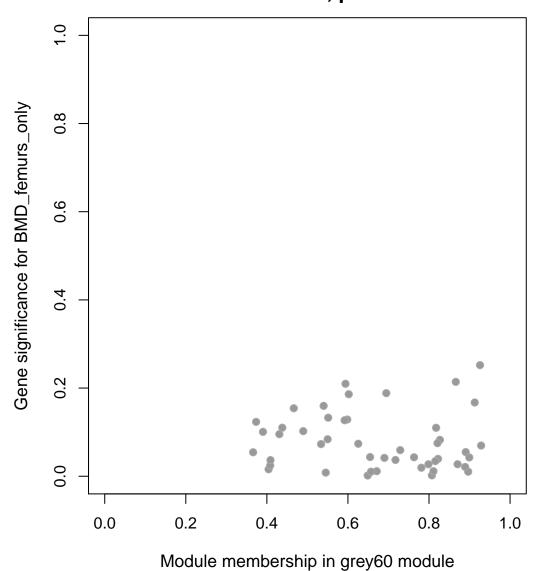**cor=0.59, p=5.1e−31**

Module membership vs. gene significance
cor=0.11, p=0.54
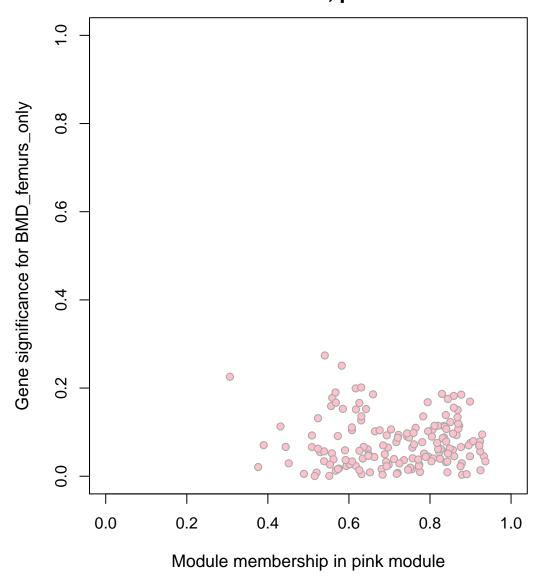
Gene significance for BMD_femurs_only

Module membership in lightgreen module

Module membership vs. gene significance
cor=−0.11, p=0.28

Gene significance for BMD_femurs_only

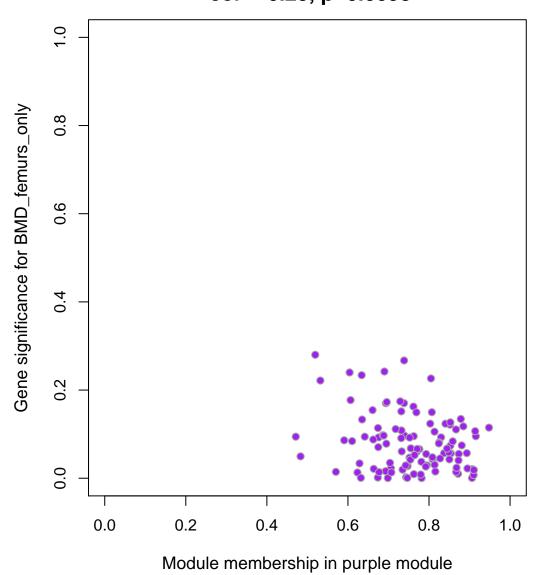Module membership in greenyellow module

Module membership vs. gene significance
cor=−0.073, p=0.63
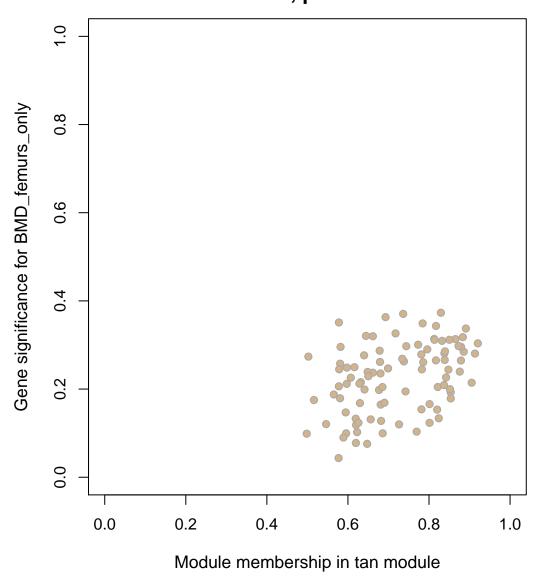
Module membership vs. gene significance
cor=−0.025, p=0.76
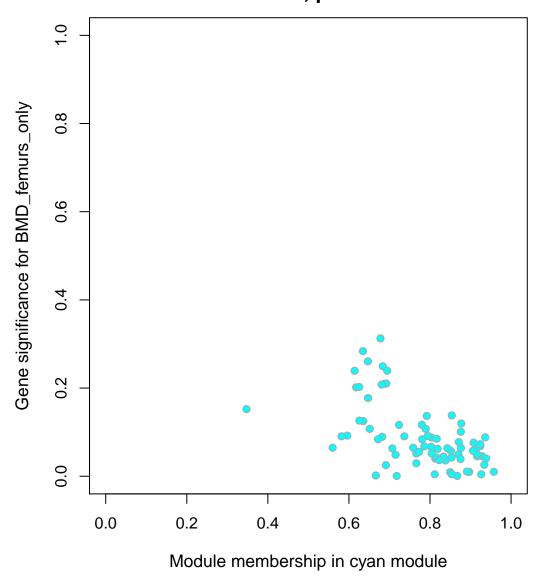
Module membership vs. gene significance
cor=−0.25, p=0.0098

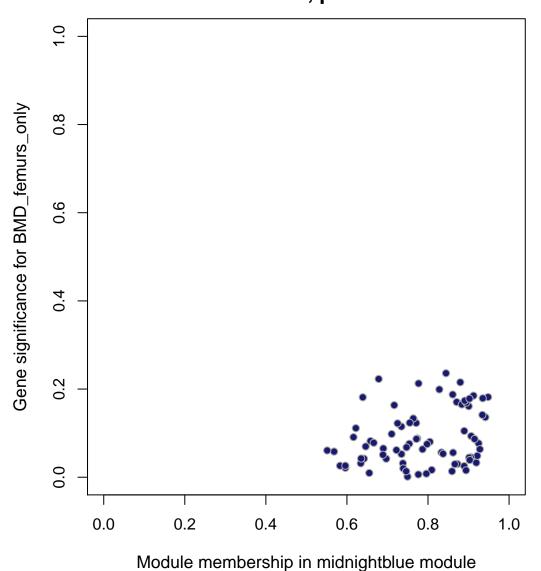Gene significance for BMD_femurs_only (y-axis), Module membership in purple module (x-axis)

Module membership vs. gene significance
cor=0.4, p=6.5e−05

Module membership vs. gene significance
cor=−0.56, p=1.2e−07

Gene significance for BMD_femurs_only

Module membership in cyan module

Module membership vs. gene significance
cor=0.27, p=0.018

Module membership vs. gene significance
cor=−0.18, p=0.075

Gene significance for BMD_femurs_only

Module membership in grey module