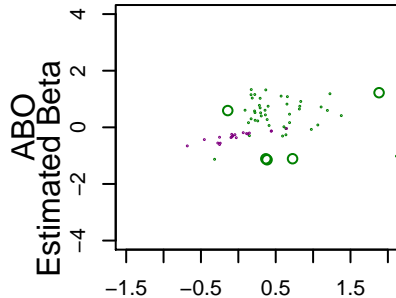
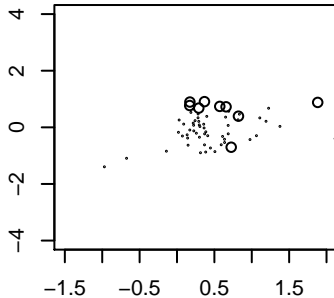


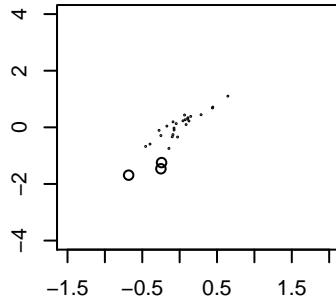
Holo-estimated



Plant-estimated



Microbe-estimated

ABA

This scatter plot shows the relationship between the known effect and the Holo-estimated effect for the ABA trait. The y-axis ranges from -4 to 4, and the x-axis ranges from -1.5 to 1.5. Data points are colored green and purple. Most points are clustered between -0.5 and 1.0 on the x-axis and -1 and 2 on the y-axis. There are several large green circles representing outliers, notably one at approximately (1.8, 2.5) and another at (0.4, -1.2).

This scatter plot shows the relationship between the known effect and the Plant-estimated effect for the ABA trait. The y-axis ranges from -4 to 4, and the x-axis ranges from -1.5 to 1.5. Data points are black dots, with several large black circles representing outliers. The points are more spread out than in the Holo-estimated plot, with outliers reaching up to x=1.8 and y=2.5.

This scatter plot shows the relationship between the known effect and the Microbe-estimated effect for the ABA trait. The y-axis ranges from -4 to 4, and the x-axis ranges from -1.5 to 1.5. Data points are black dots, with several large black circles representing outliers. The points are tightly clustered along a diagonal line from approximately (-0.5, -0.5) to (0.5, 0.5).

Known effect