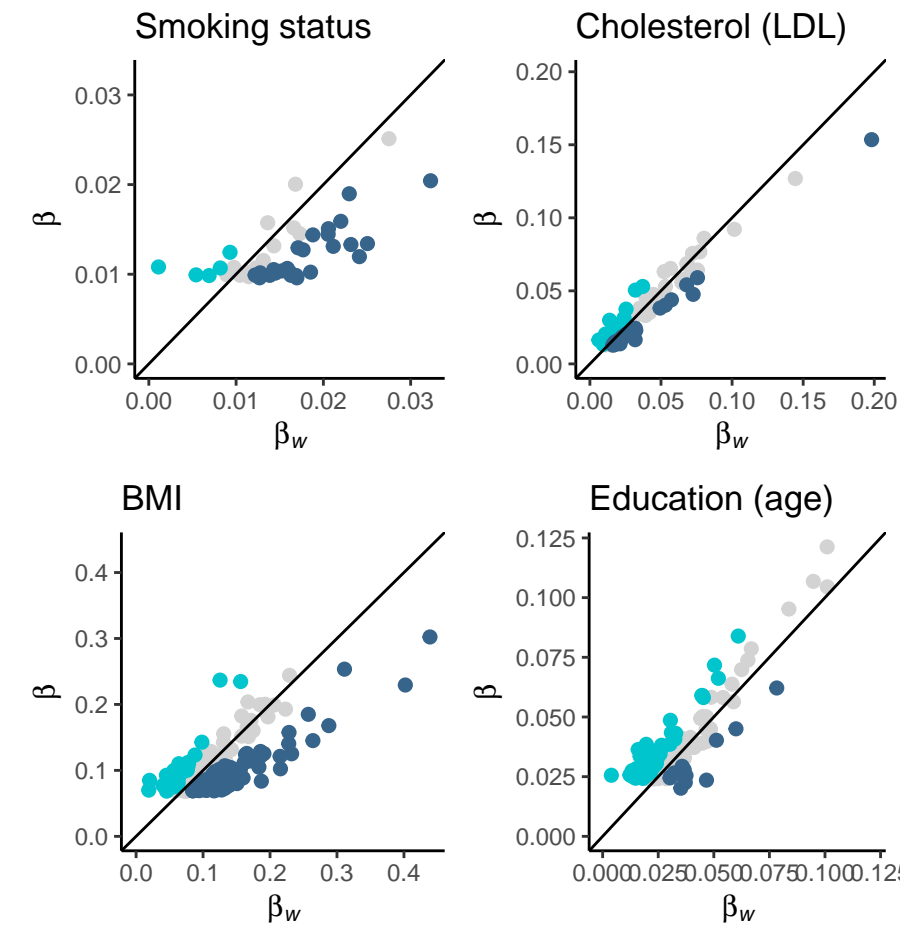
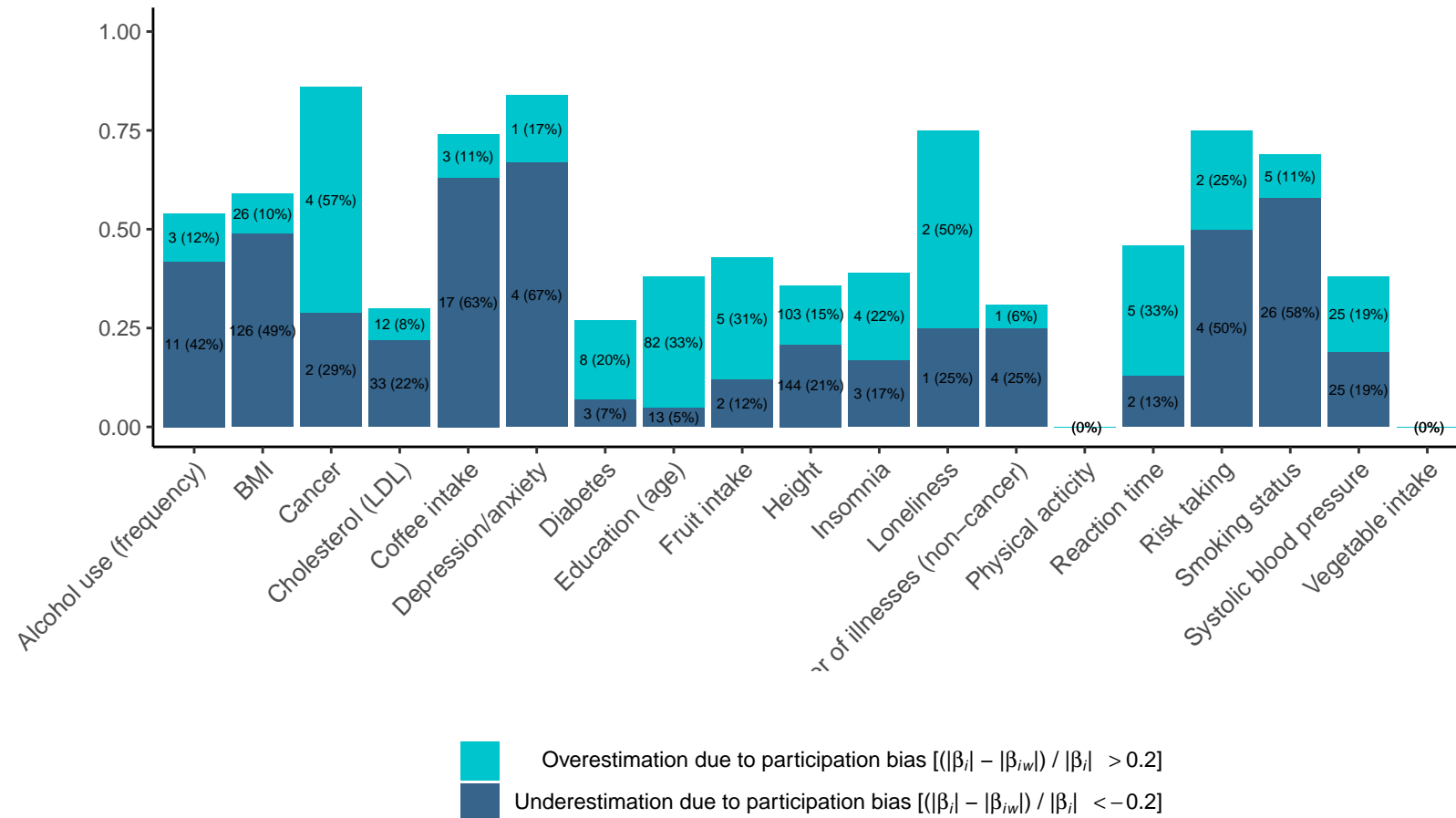


A

Change in effect estimates for genome-wide variants

**B**

Number of genome-wide variants

