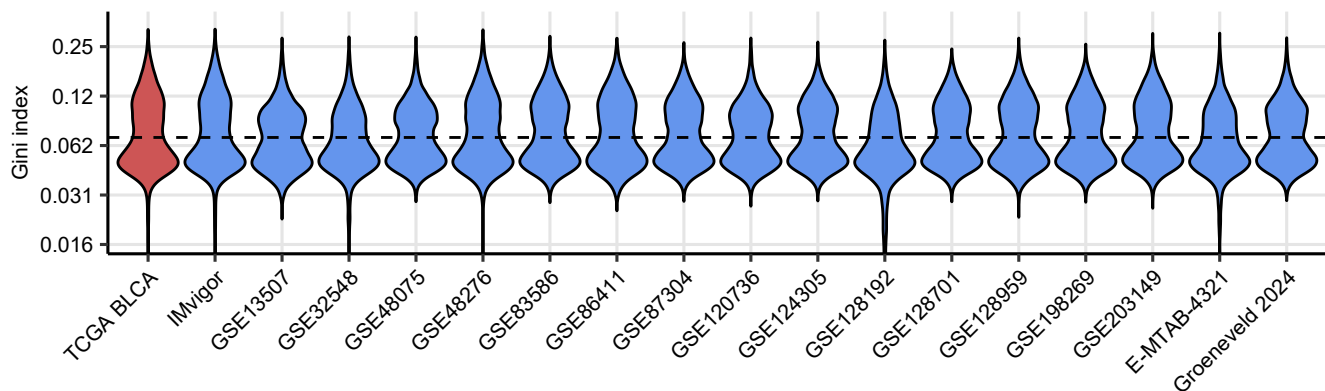
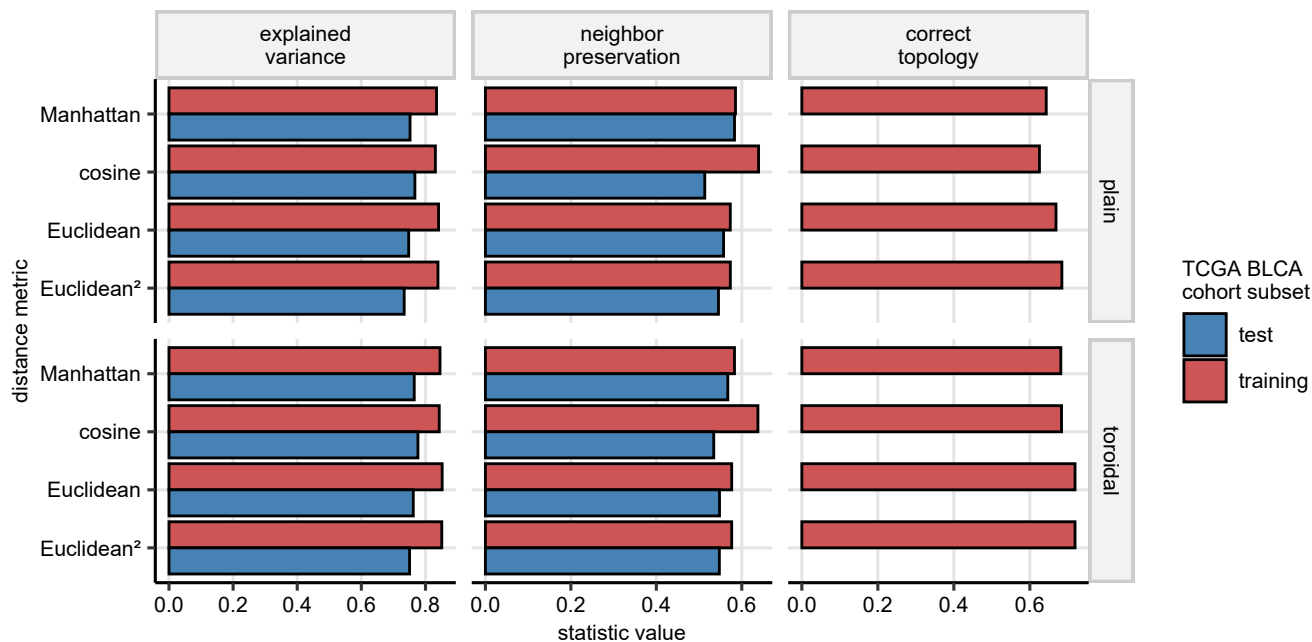


A**Choice of variably expressed genes**

total variables: n = 12394

**B****Choice of metagene-defining SOM algorithm: model performance**

total genes: n = 3234, total metagenes: n = 256

**C****Choice of metagene-defining SOM algorithm: metagene size distribution**

total genes: n = 3234, total metagenes: n = 256

