Overlapping fungal genera between TCGA WGS and RNA–Seq and Weizmann datasets
(note: intersection among features found in rep200 database)

Legend: Found in Weizmann and TCGA | Weizmann only

Y-axis: Genus count

X-axis: Cancer/tissue type (tumor and NAT combined when available)

Bone: 11, 3
Breast: 38, 0
Colorectal: 10, 0
Glioblastoma: 10, 0
Lung: 28, 0
Melanoma: 13, 2
Ovarian: 24, 0
Pancreatic: 8, 5