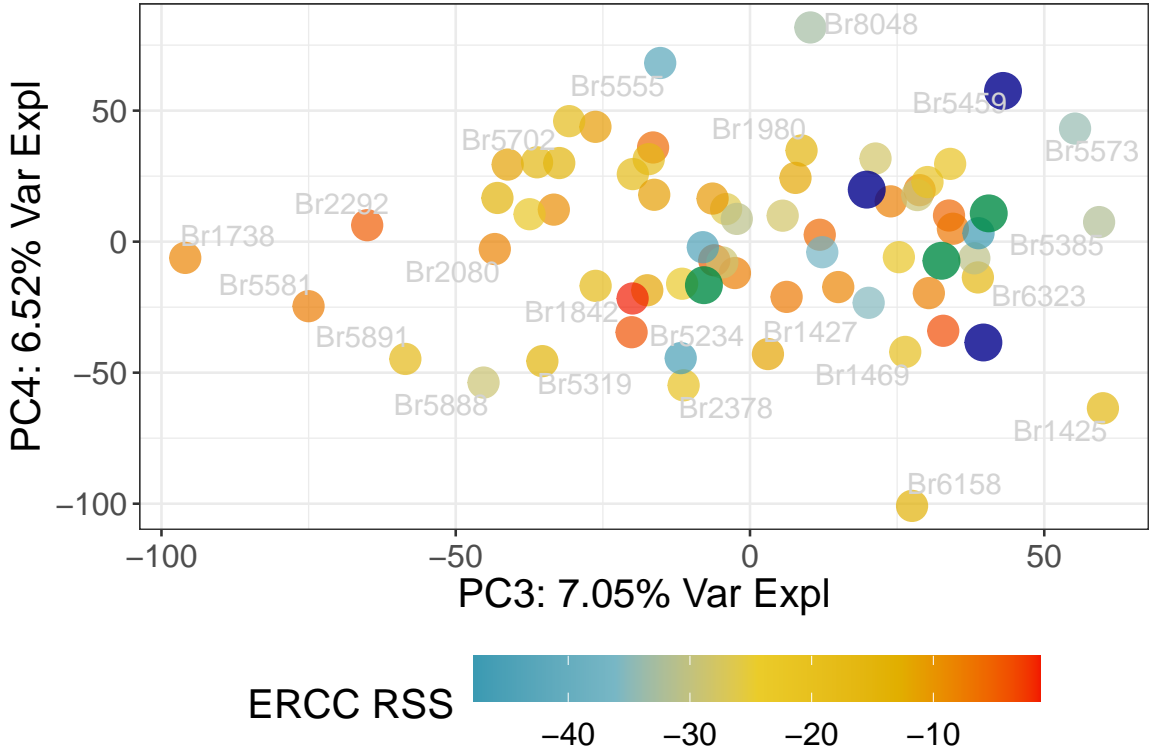


**PC3 vs PC4 by ERCC RSS**

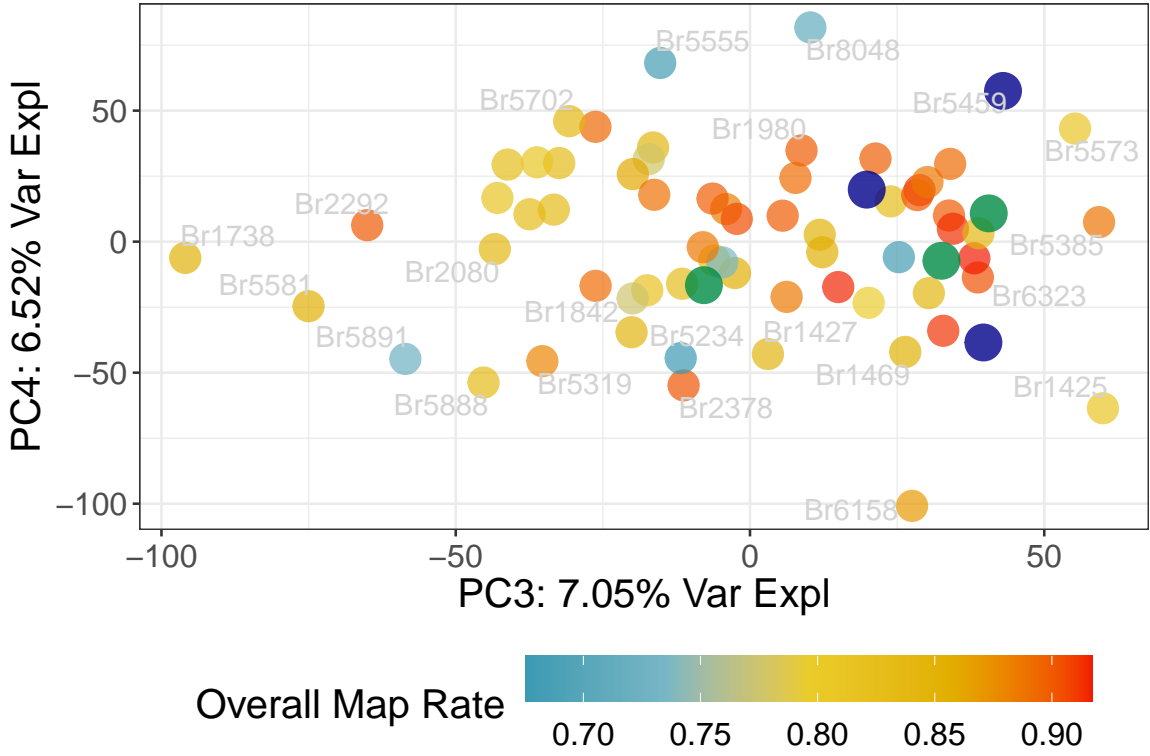
*Br5572, Br1682, and Br5446*



*Br1676, Br5459, and Br6323*

**PC3 vs PC4 by Overall Map Rate**

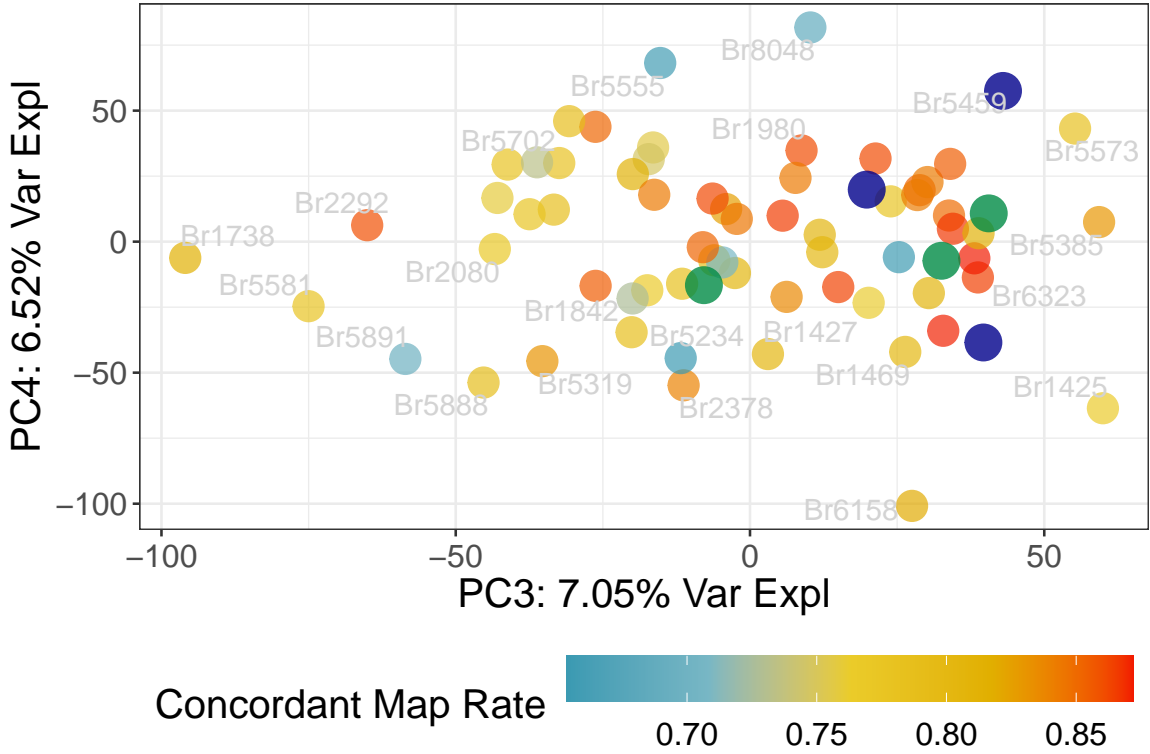
*Br5572, Br1682, and Br5446*



*Br1676, Br5459, and Br6323*

**PC3 vs PC4 by Concordant Map Rate**

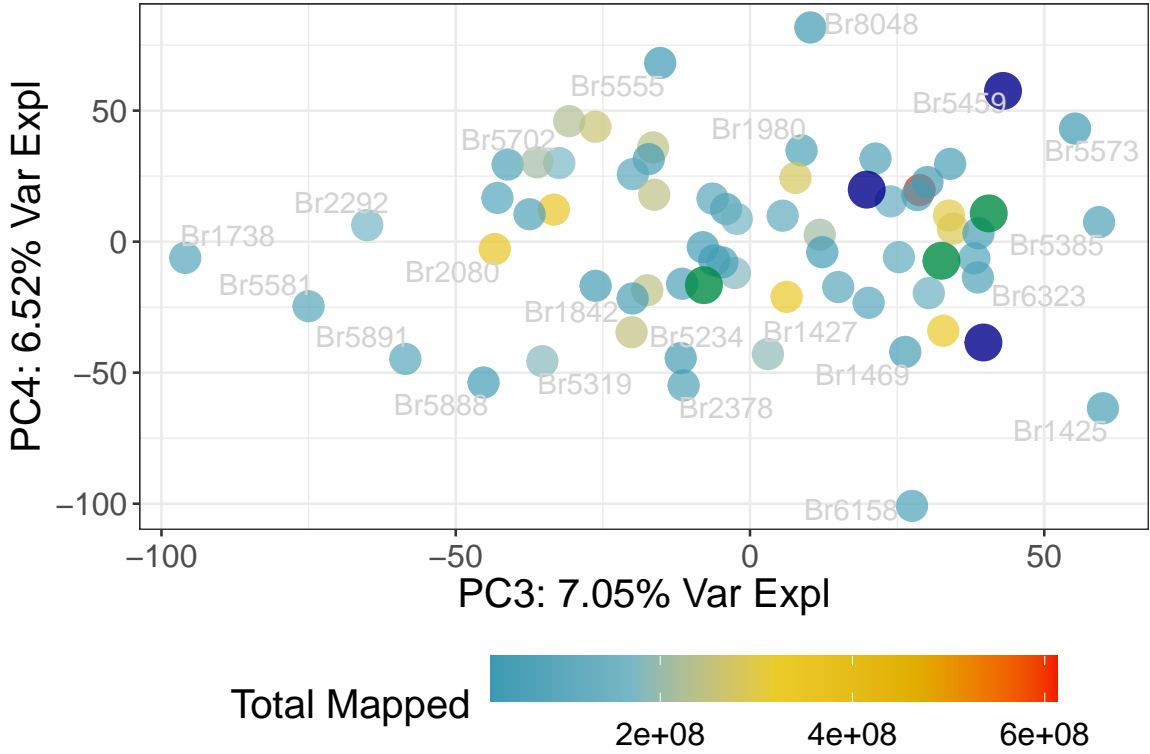
*Br5572, Br1682, and Br5446*



*Br1676, Br5459, and Br6323*

**PC3 vs PC4 by Total Mapped**

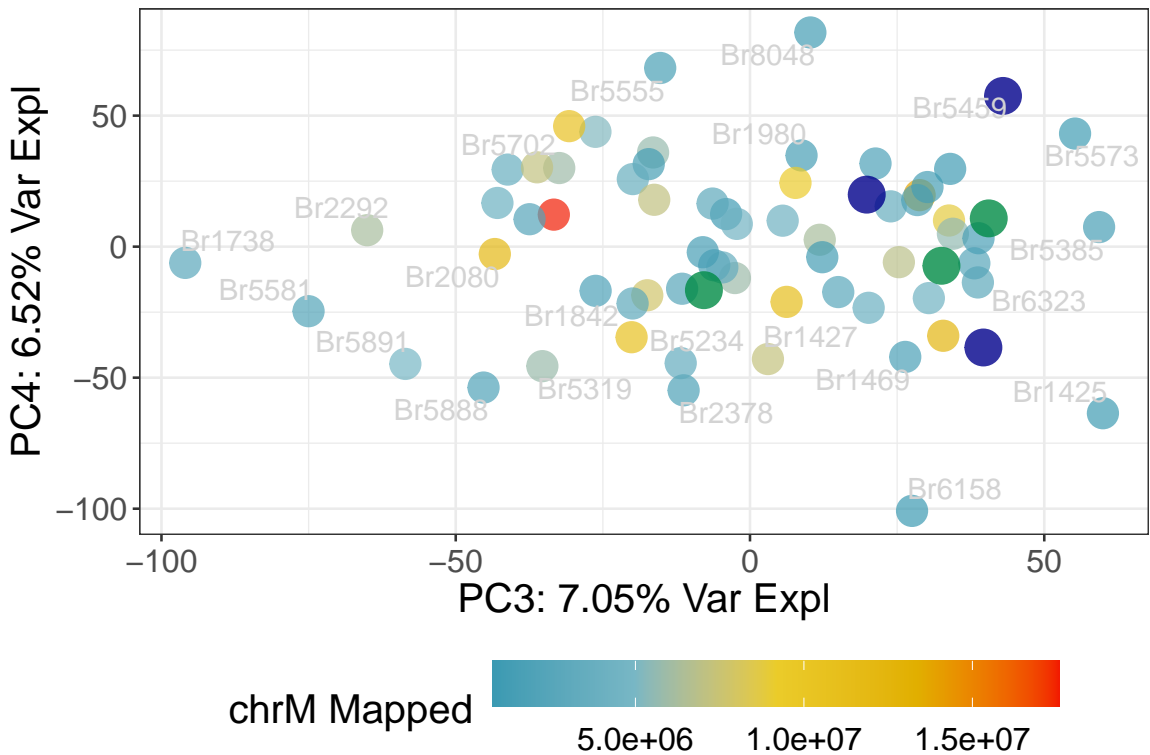
*Br5572, Br1682, and Br5446*



*Br1676, Br5459, and Br6323*

**PC3 vs PC4 by chrM Mapped**

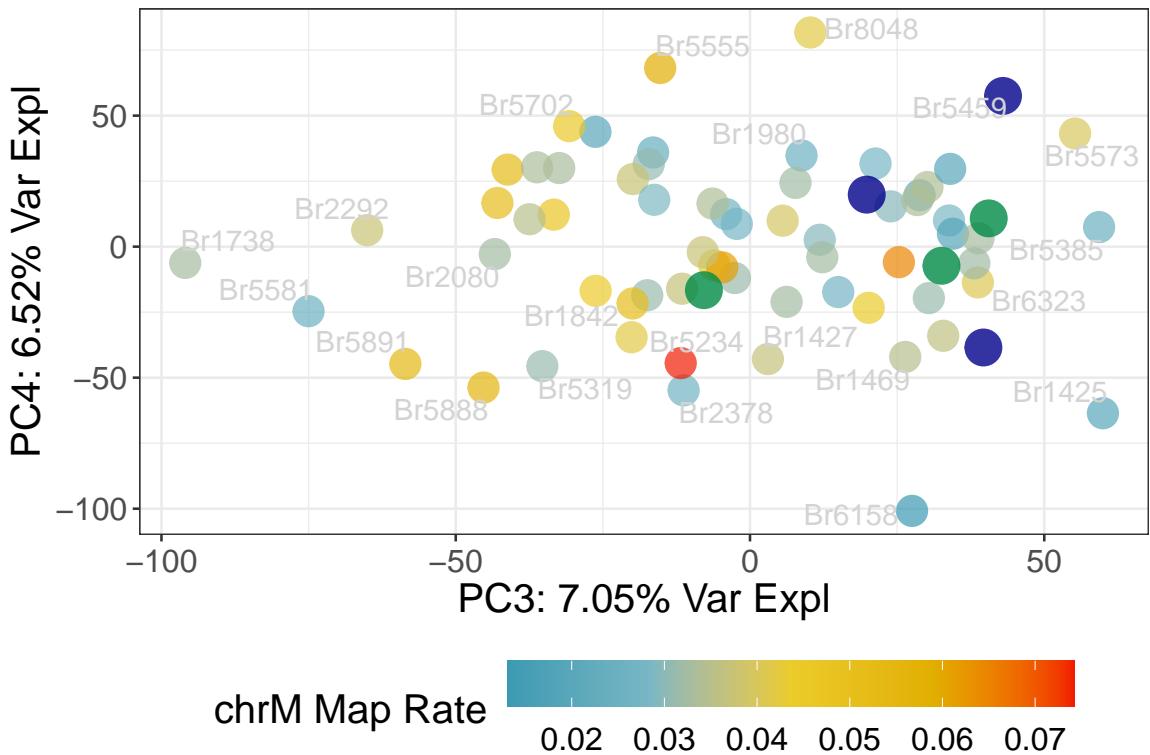
*Br5572, Br1682, and Br5446*



*Br1676, Br5459, and Br6323*

**PC3 vs PC4 by chrM Map Rate**

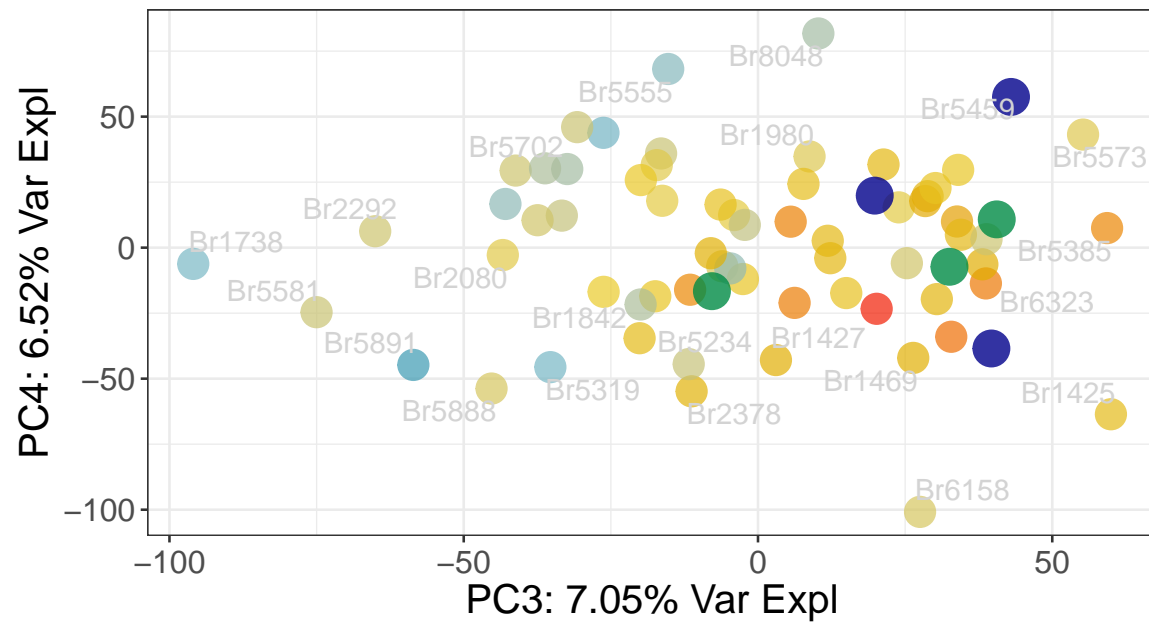
*Br5572, Br1682, and Br5446*



*Br1676, Br5459, and Br6323*

**PC3 vs PC4 by Total Assigned Genes**

*Br5572, Br1682, and Br5446*

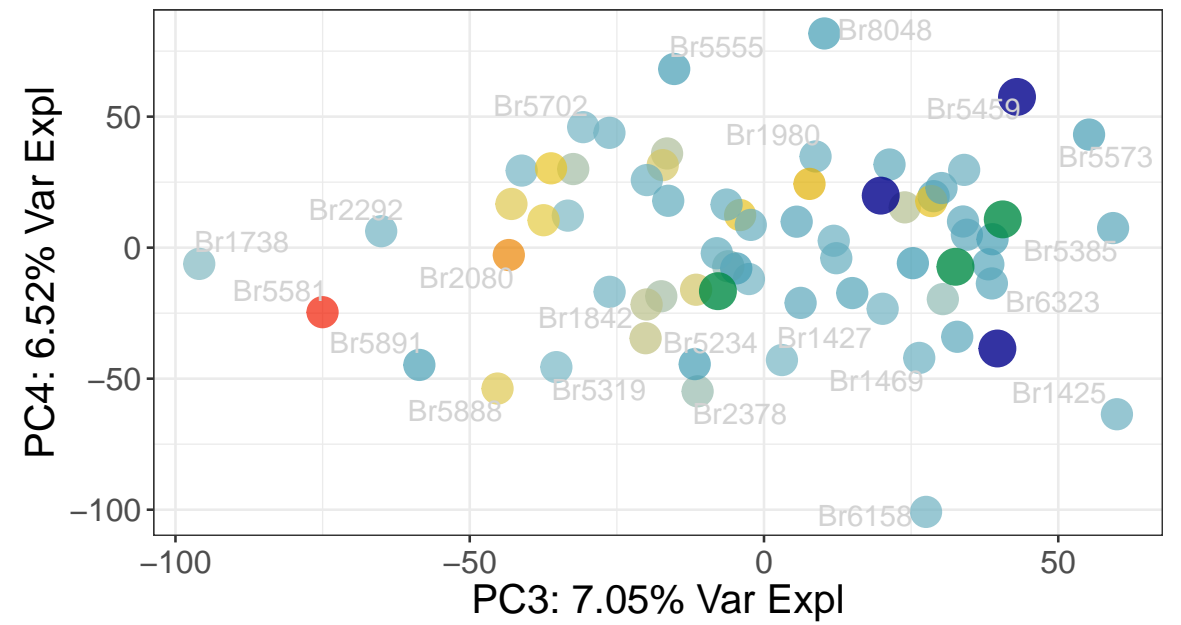


Total Assigned Genes

0.4 0.5 0.6  
*Br1676, Br5459, and Br6323*

**PC3 vs PC4 by Gene rRNA Rate**

*Br5572, Br1682, and Br5446*

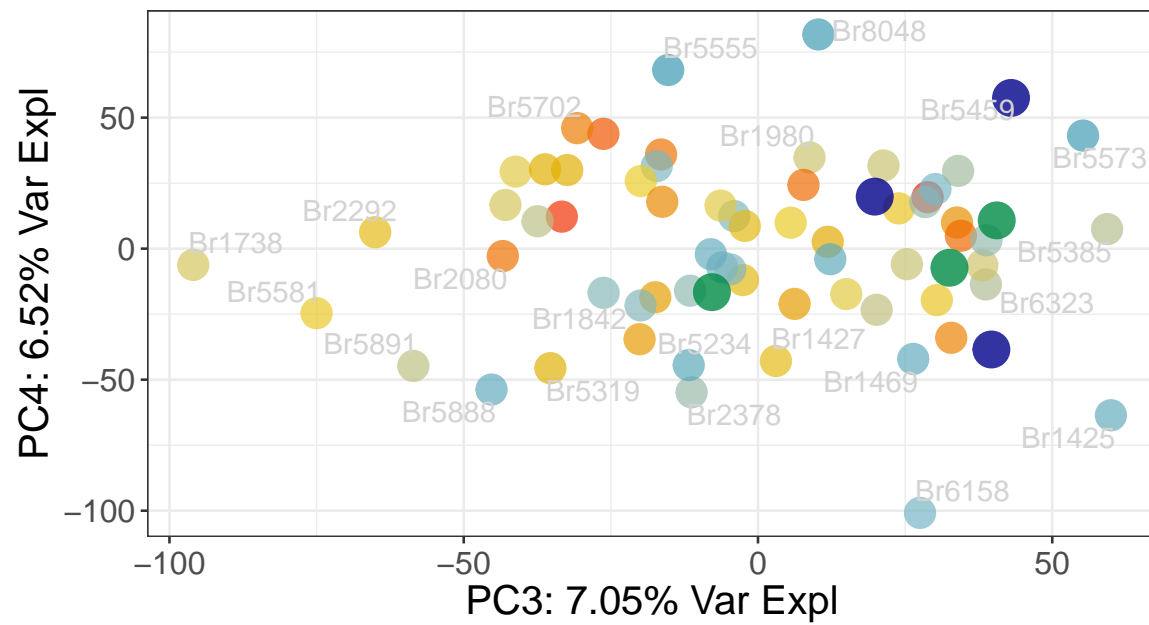


Gene rRNA Rate

0.00005 0.00010 0.00015  
*Br1676, Br5459, and Br6323*

**PC3 vs PC4 by Detected**

*Br5572, Br1682, and Br5446*

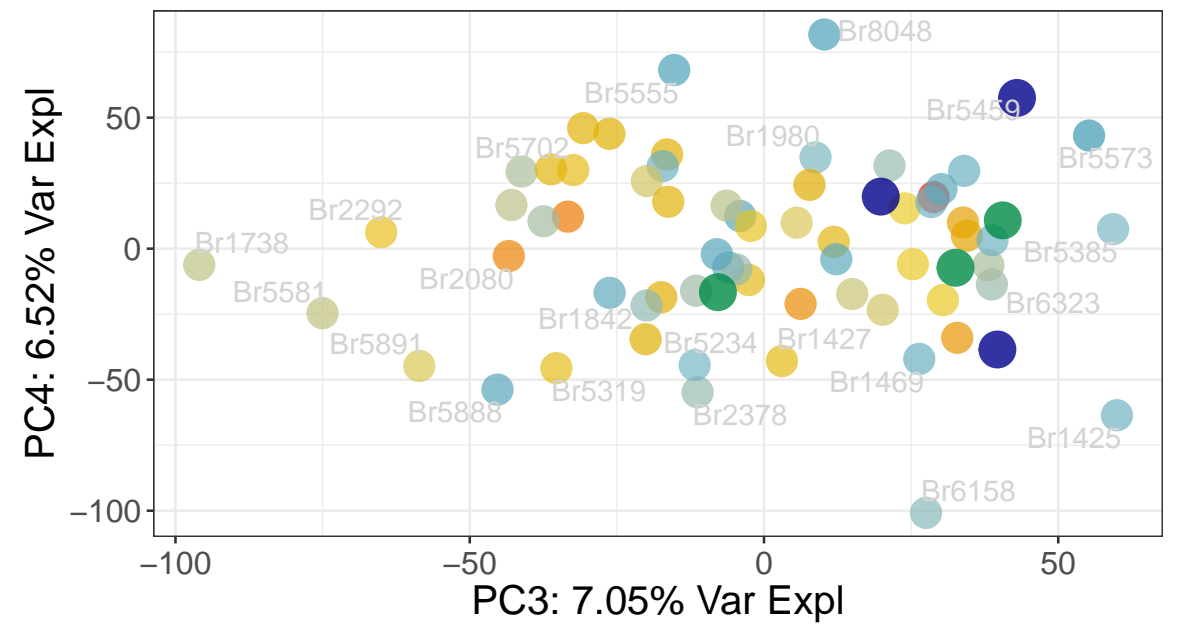


Detected

27500 30000 32500 35000  
*Br1676, Br5459, and Br6323*

**PC3 vs PC4 by Num Reads (log 10)**

*Br5572, Br1682, and Br5446*

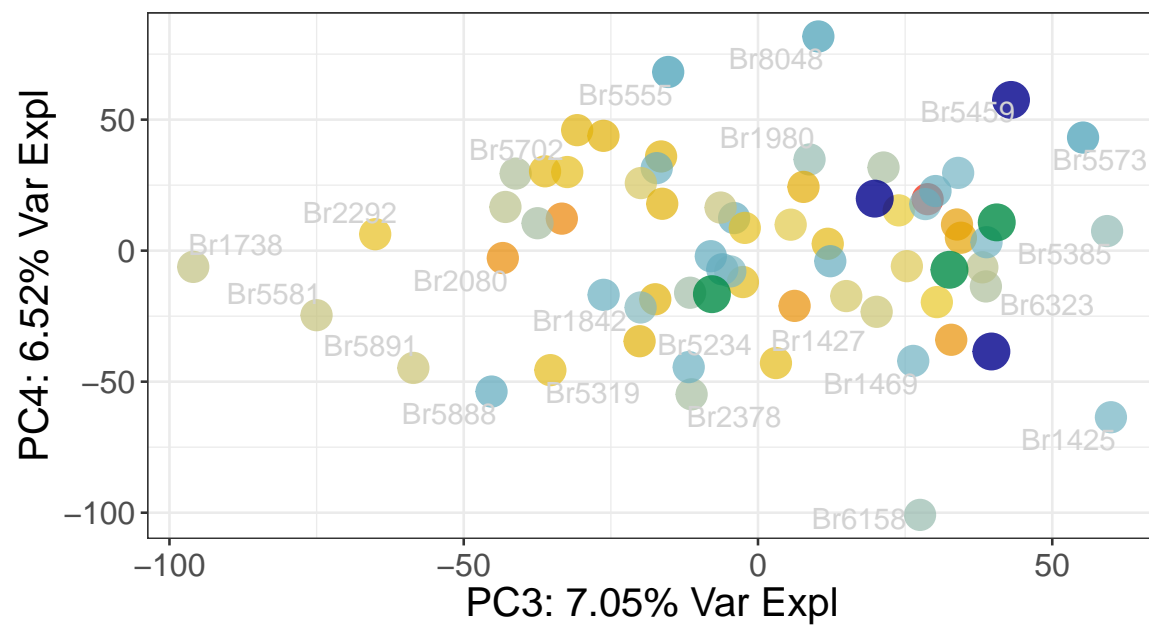


Num Reads (log 10)

0.88 0.90 0.92 0.94  
*Br1676, Br5459, and Br6323*

**PC3 vs PC4 by Num Mapped (log 10)**

*Br5572, Br1682, and Br5446*

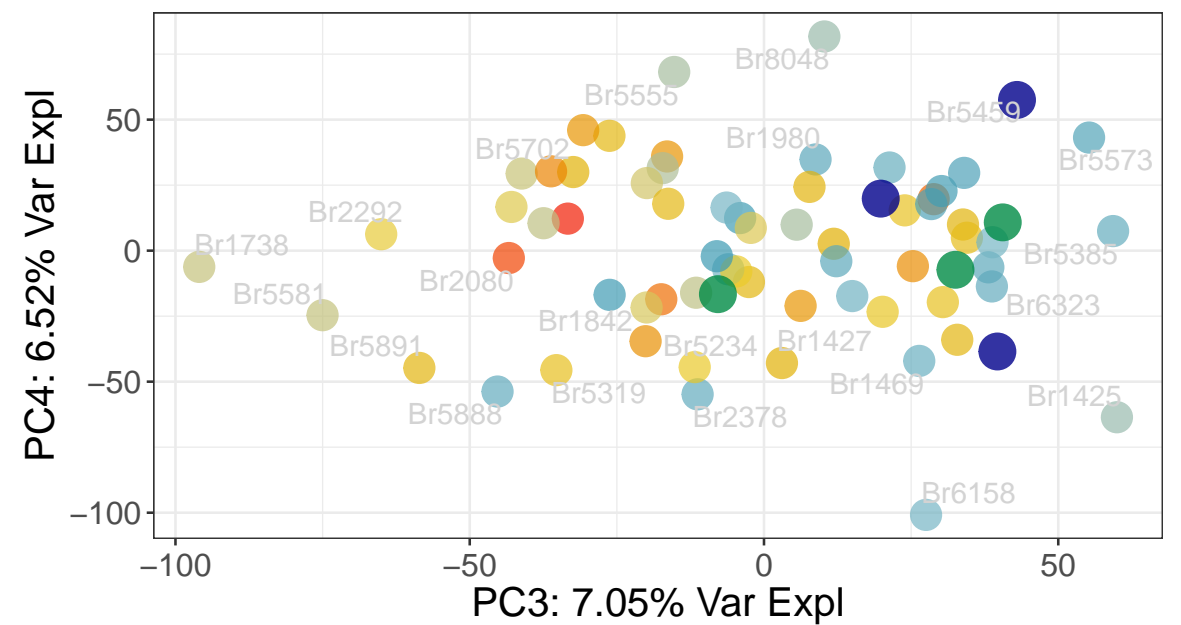


Num Mapped (log 10)

0.88 0.90 0.92 0.94  
*Br1676, Br5459, and Br6323*

**PC3 vs PC4 by Num Unmapped (log 10)**

*Br5572, Br1682, and Br5446*



Num Unmapped (log 10)

6.9 7.2 7.5 7.8  
*Br1676, Br5459, and Br6323*