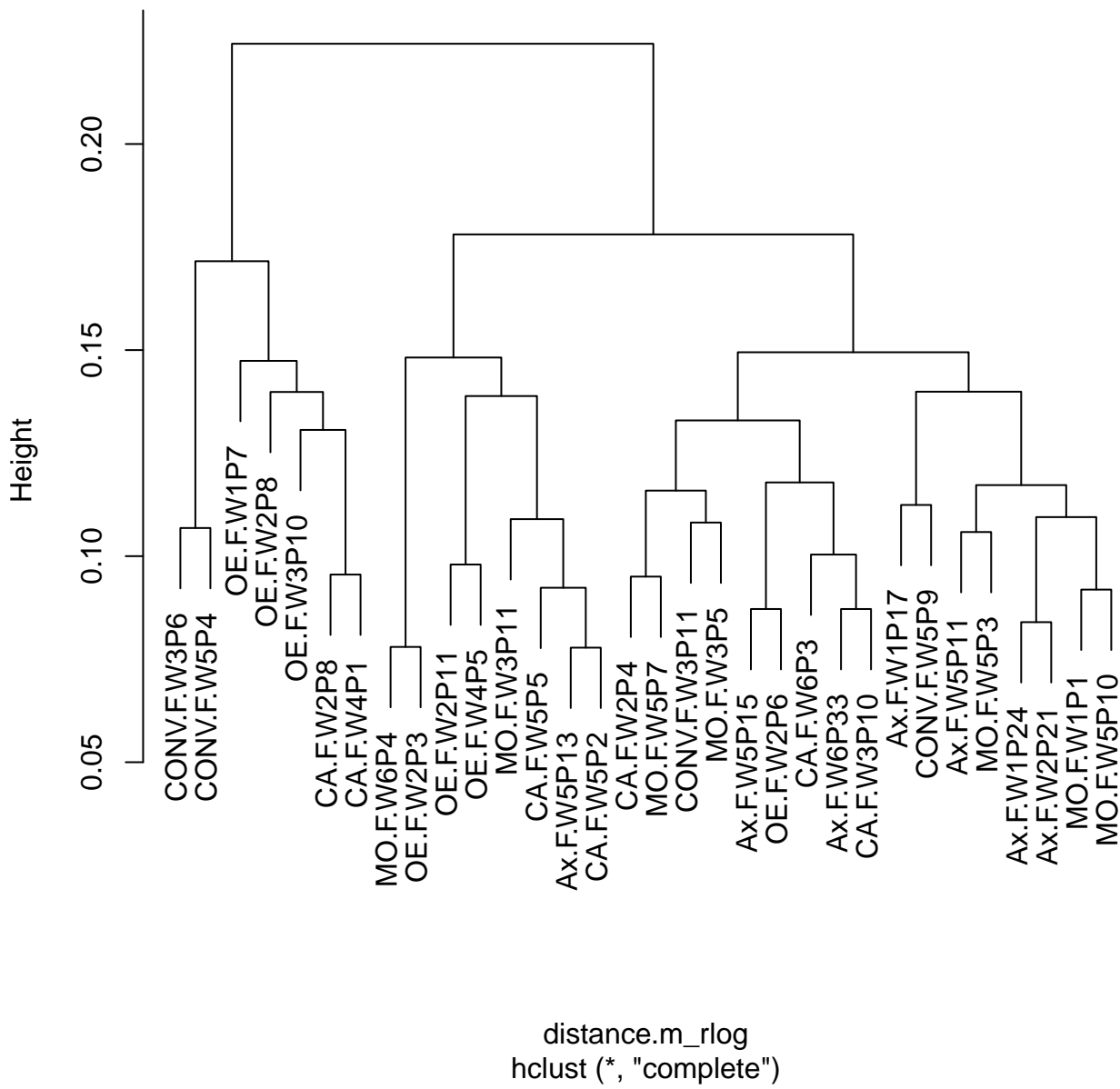
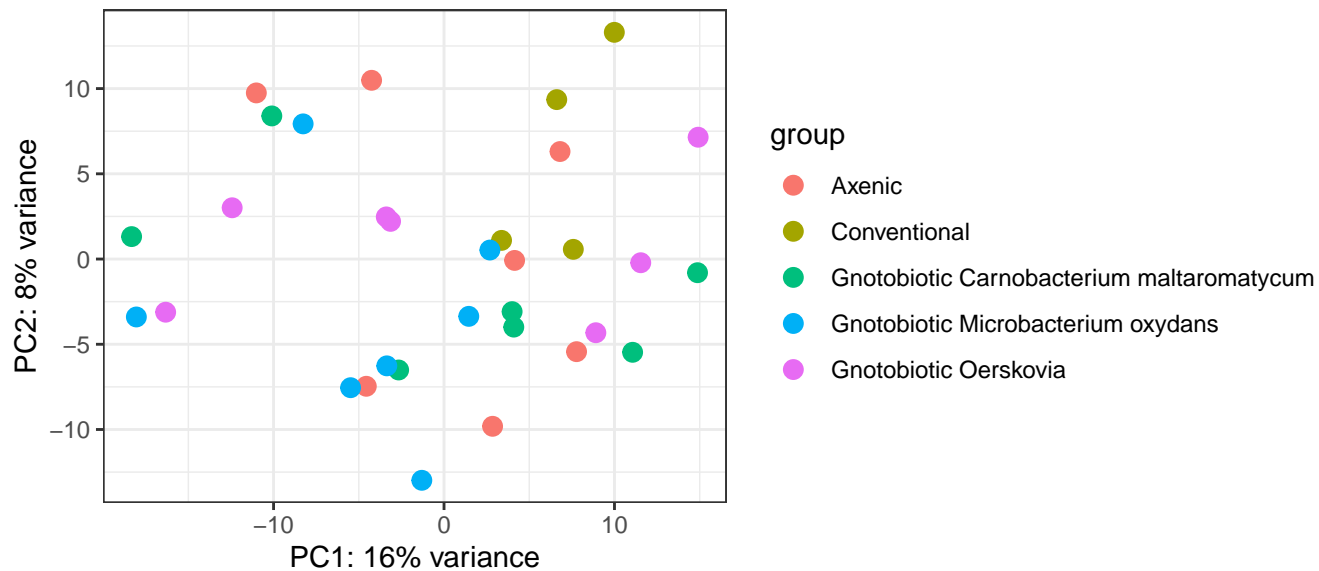


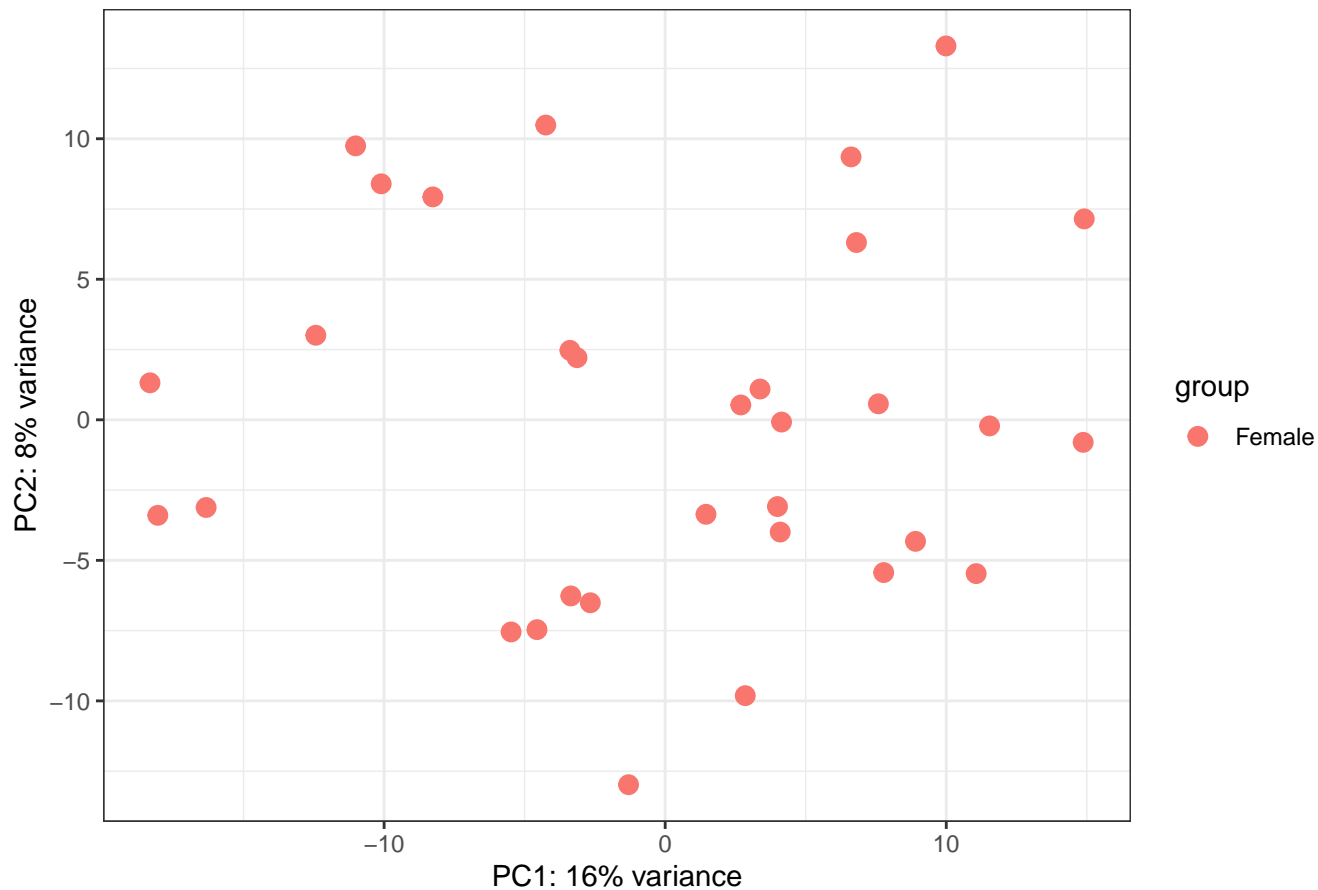
**rlog transformed read counts**  
**distance: Pearson correlation**



Rlog transformed counts



# Rlog transformed counts



**By modality**

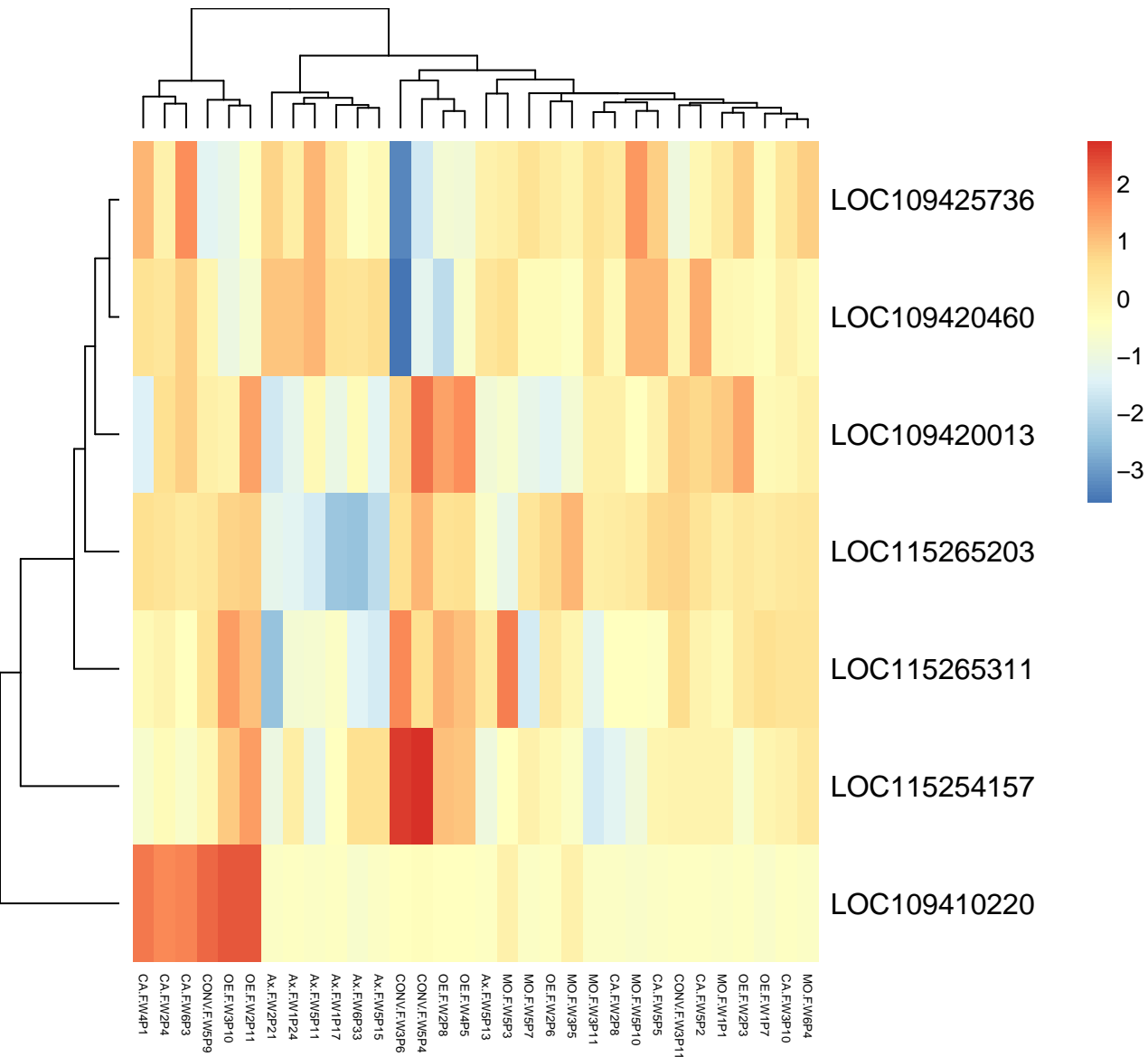
Heatmap showing the expression of seven genes across 28 samples, clustered by modality. The color scale ranges from -3 (blue) to 2 (red).

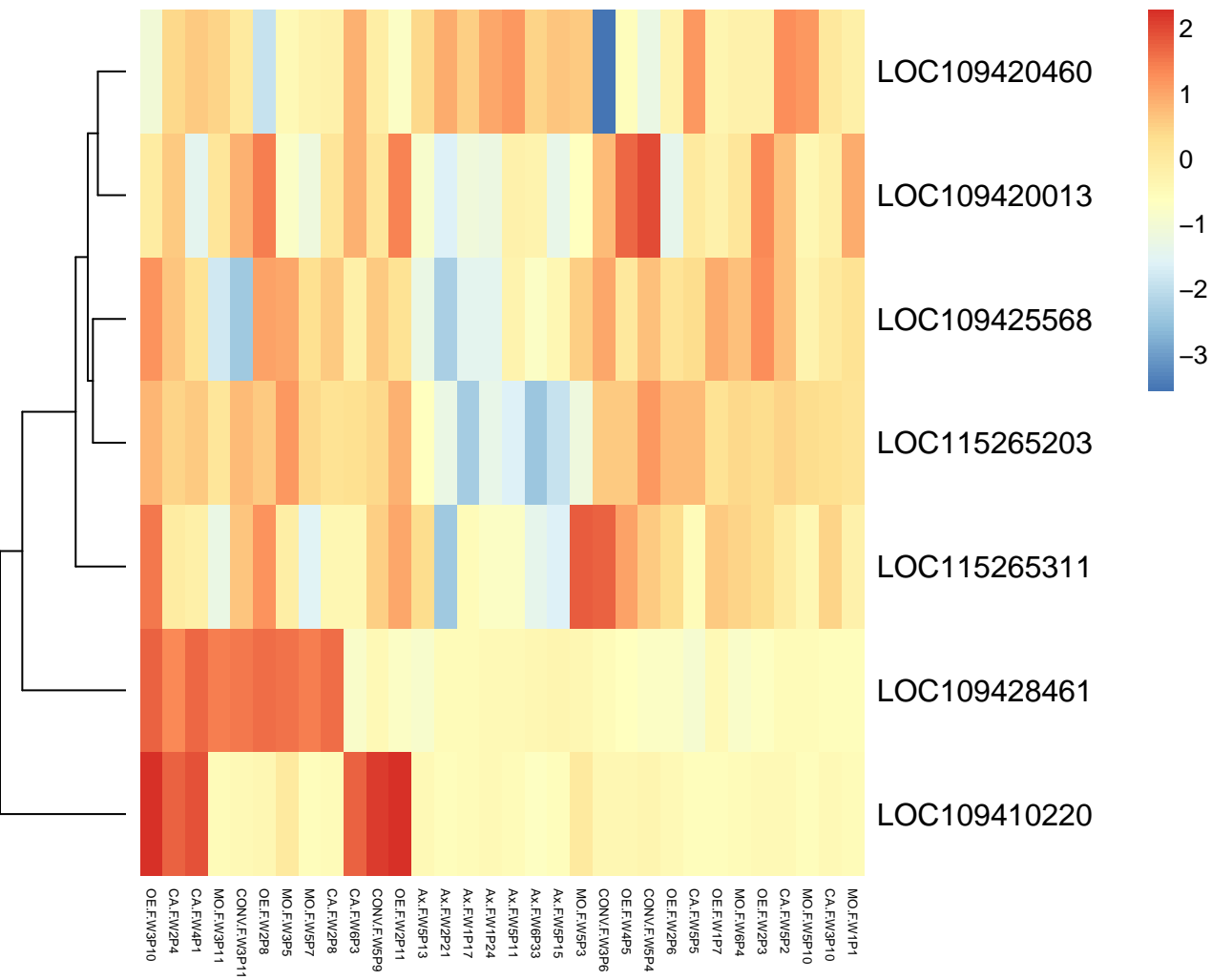
Genes (rows):

- LOC109425736
- LOC109420460
- LOC109420013
- LOC115265203
- LOC115265311
- LOC115254157
- LOC109410220

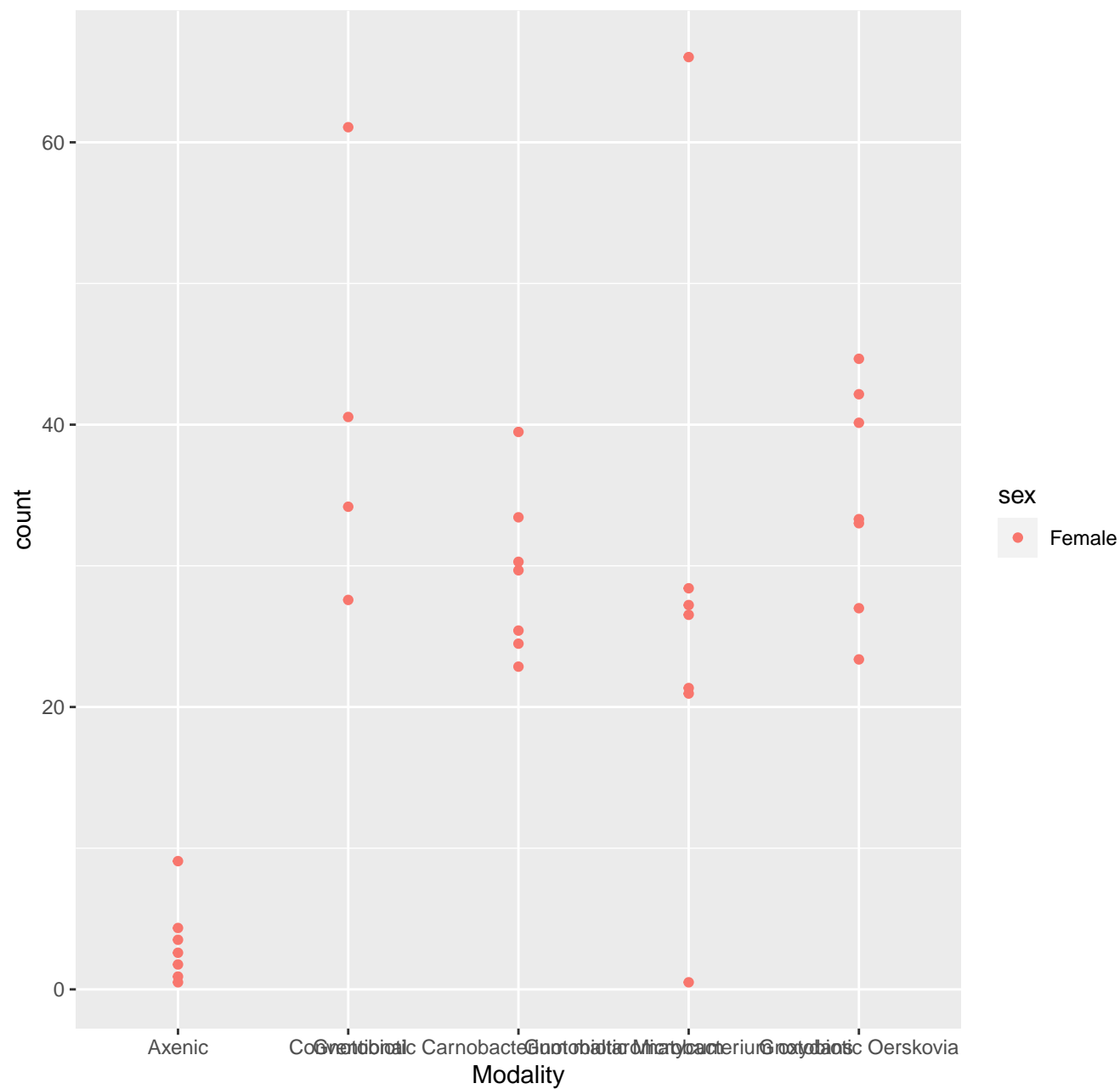
Samples (columns):

- CA.FW4P1
- CA.FW2P4
- CA.FW6P3
- CONV.FW6P9
- OE.FW3P10
- OE.FW2P11
- AX.FW2P21
- AX.FW1P24
- AX.FW5P11
- AX.FW6P33
- AX.FW1P17
- AX.FW5P15
- CONV.FW3P6
- CONV.FW5P4
- OE.FW2P8
- OE.FW4P5
- AX.FW5P13
- MO.FW5P3
- MO.FW5P7
- OE.FW2P6
- MO.FW3P5
- MO.FW3P11
- CA.FW2P8
- MO.FW5P10
- CA.FW5P5
- CONV.FW3P11
- CA.FW5P2
- MO.FW1P1
- OE.FW2P3
- OE.FW1P7
- CA.FW3P10
- MO.FW6P4





LOC115265203



**MA plot: group1 vs group2 (padj < 0.05)**

