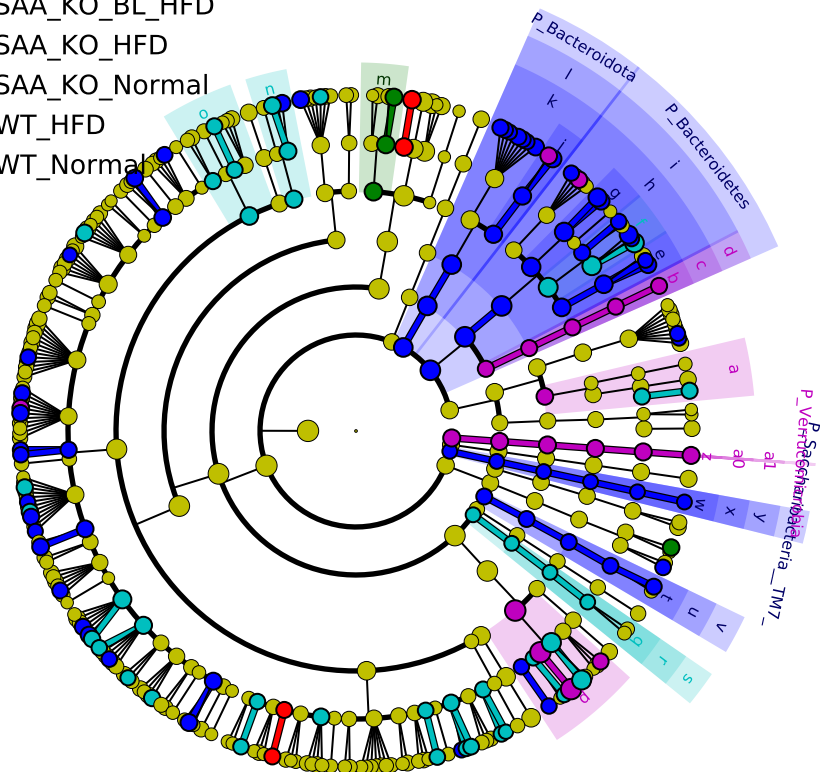


# Significantly Represented Taxa

- SAA\_KO\_BL\_HFD
- SAA\_KO\_HFD
- SAA\_KO\_Normal
- WT\_HFD
- WT\_Normal



- a: O\_Coriobacteriales
- b: F\_Bacteroidetes\_F\_1
- c: O\_Bacteroidetes\_O\_1
- d: C\_Bacteroidetes\_C\_1
- e: F\_Bacteroidaceae
- f: F\_Muribaculaceae
- g: F\_Prevotellaceae
- h: O\_Bacteroidales
- i: C\_Bacteroidia
- j: F\_Bacteroidaceae
- k: O\_Bacteroidales
- l: C\_Bacteroidia
- m: F\_Streptococcaceae
- n: F\_Clostridiaceae
- o: F\_Eubacteriales\_F\_1
- p: F\_Erysipelotrichaceae
- q: F\_Mollicutes\_F\_2
- r: O\_Mollicutes\_O\_2
- s: C\_Mollicutes
- t: F\_Sutterellaceae
- u: O\_Burkholderiales
- v: C\_Betaproteobacteria
- w: F\_Saccharibacteria\_TM7\_
- x: O\_Saccharibacteria\_TM7\_
- y: C\_Saccharibacteria\_TM7\_
- z: F\_Akkermansiaceae
- a0: O\_Verrucomicrobiales
- a1: C\_Verrucomicrobiales