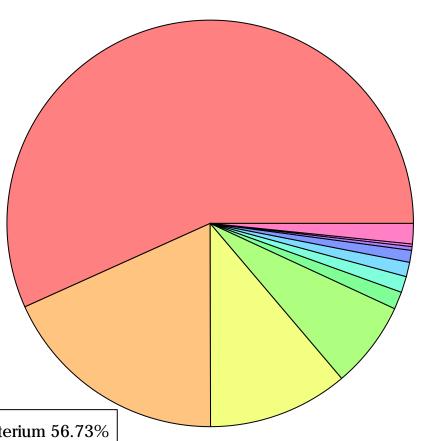
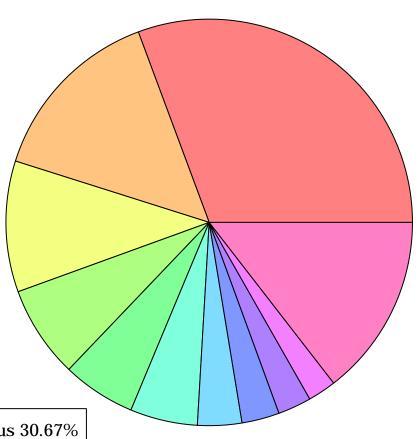
# This sample's microbiome, broken down by genus:



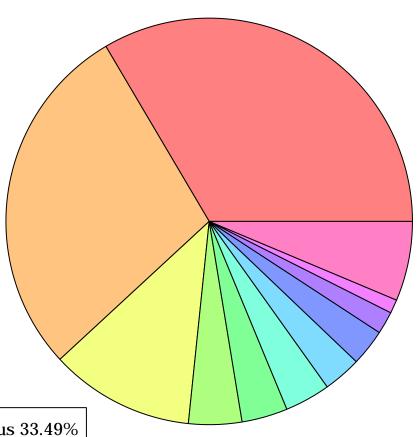
- Propionibacterium 56.73%
- Anaerococcus 18.29%
- □ Peptoniphilus 11.17%
- Staphylococcus 6.91%
- □ Corynebacterium 1.39%
- □ Campylobacter 1.28%
- Negativicoccus 1.15%
- Finegoldia 0.95%
- Streptococcus 0.3%
- Dietzia 0.2%
- Other 1.62%

# This sample's microbiome, broken down by genus:



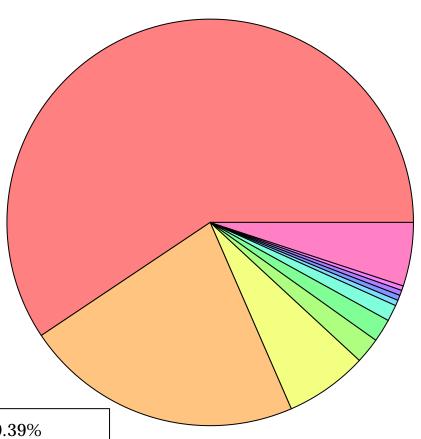
- Streptococcus 30.67%
- Haemophilus 14.45%
- ☐ Granulicatella 10.42%
- Prevotella 7.33%
- Veillonella 5.84%
- Rothia 5.36%
- **■** Kingella 3.52%
- **■** Gemella 2.99%
- Fusobacterium 2.66%
- Actinomyces 2.25%
- Other 14.51%

# This sample's microbiome, broken down by genus:



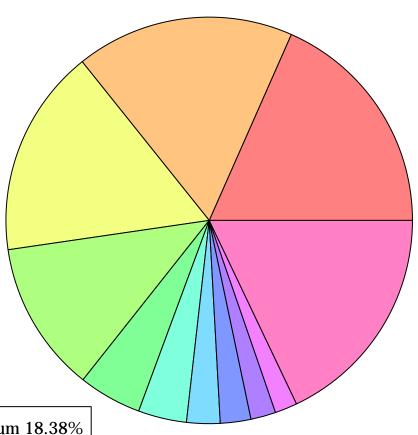
- Streptococcus 33.49%
- Veillonella 28.39%
- **■** Atopobium 11.49%
- Actinomyces 4.23%
- **■** Granulicatella 3.65%
- Haemophilus 3.62%
- Rothia 3.02%
- Prevotella 2.91%
- Megasphaera 1.79%
- □ Campylobacter 1.09%
- Other 6.33%

# This sample's microbiome, broken down by genus:



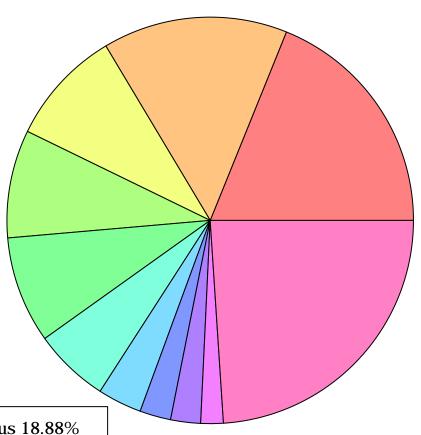
- Neisseria 59.39%
- □ Corynebacterium 22.15%
- ☐ Staphylococcus 6.58%
- Stenoxybacter 2.04%
- Anaerococcus 1.83%
- Streptococcus 1.3%
- Micrococcus 0.45%
- Propionibacterium 0.42%
- Vitreoscilla 0.41%
- Enhydrobacter 0.37%
- Other 5.06%

# This sample's microbiome, broken down by genus:



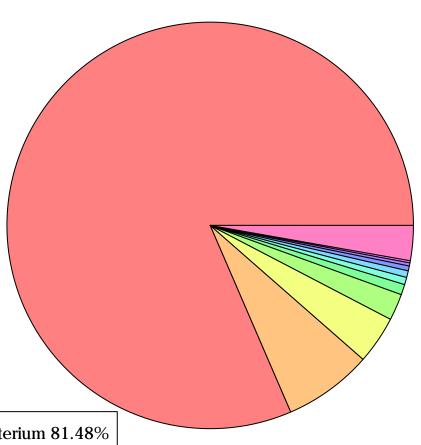
- Fusobacterium 18.38%
- Veillonella 17.36%
- □ Streptococcus 16.57%
- Haemophilus 12.03%
- Prevotella 5.02%
- Actinomyces 3.85%
- □ Campylobacter 2.66%
- Porphyromonas 2.43%
- **■** Atopobium 1.99%
- Gemella 1.78%
- Other 17.93%

# This sample's microbiome, broken down by genus:



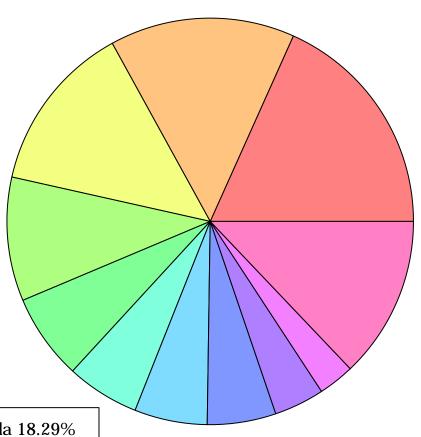
- Streptococcus 18.88%
- Staphylococcus 14.7%
- ☐ Mycoplasma 9.21%
- □ Propionibacterium 8.59%
- **■** Corynebacterium 8.49%
- Veillonella 6.02%
- Prevotella 3.51%
- Lactobacillus 2.47%
- Anaerococcus 2.38%
- Peptoniphilus 1.79%
- Other 23.96%

# This sample's microbiome, broken down by genus:



- Propionibacterium 81.48%
- Mycoplasma 7.06%
- ☐ Staphylococcus 3.81%
- □ Corynebacterium 2.11%
- Calothrix 0.83%
- Streptococcus 0.58%
- Dietzia 0.53%
- Anaerococcus 0.39%
- Enhydrobacter 0.25%
- Lactobacillus 0.17%
- Other 2.78%

# This sample's microbiome, broken down by genus:



- Granulicatella 18.29%
- Streptococcus 14.71%
- ☐ Haemophilus 13.49%
- Rothia 9.85%
- Veillonella 6.82%
- **■** Porphyromonas 5.81%
- Neisseria 5.79%
- Actinomyces 5.46%
- Abiotrophia 4%
- □ Capnocytophaga 2.87%
- Other 12.9%