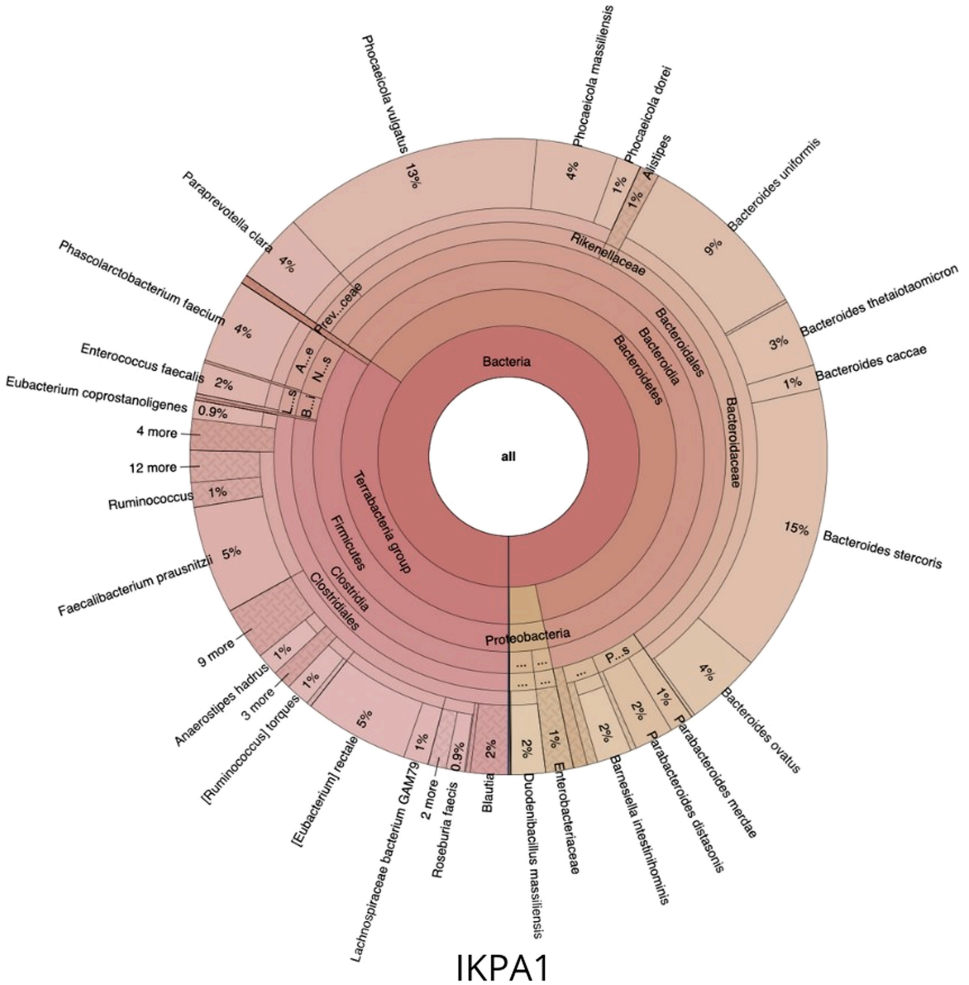
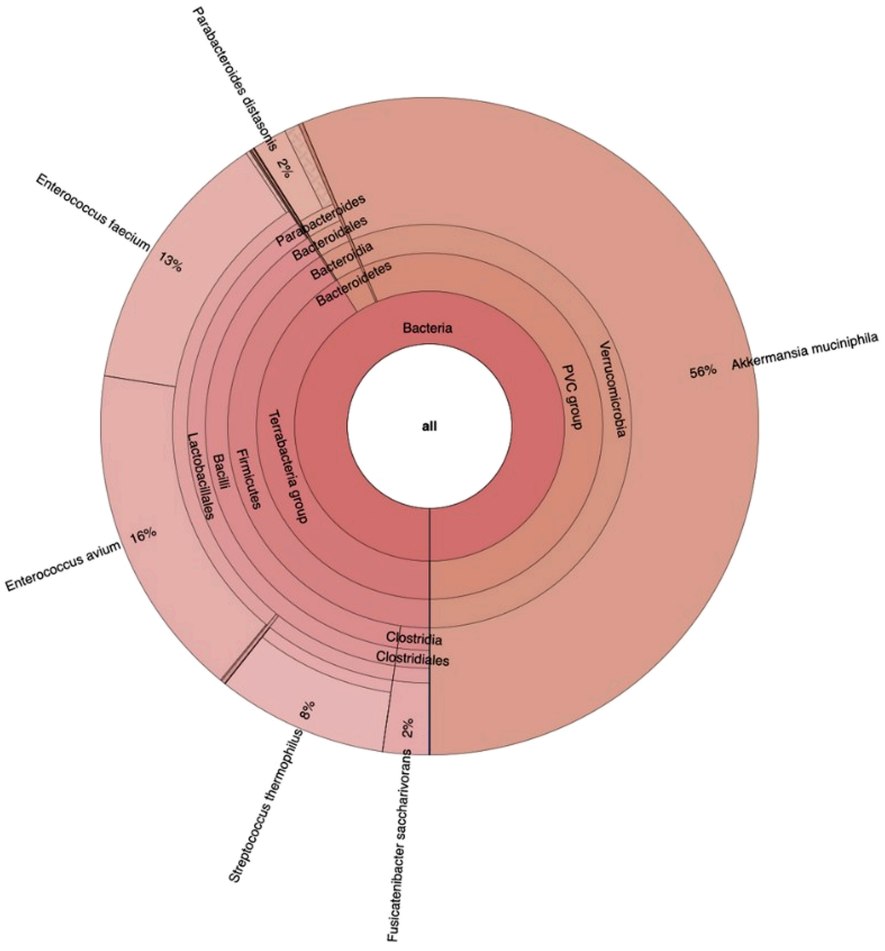
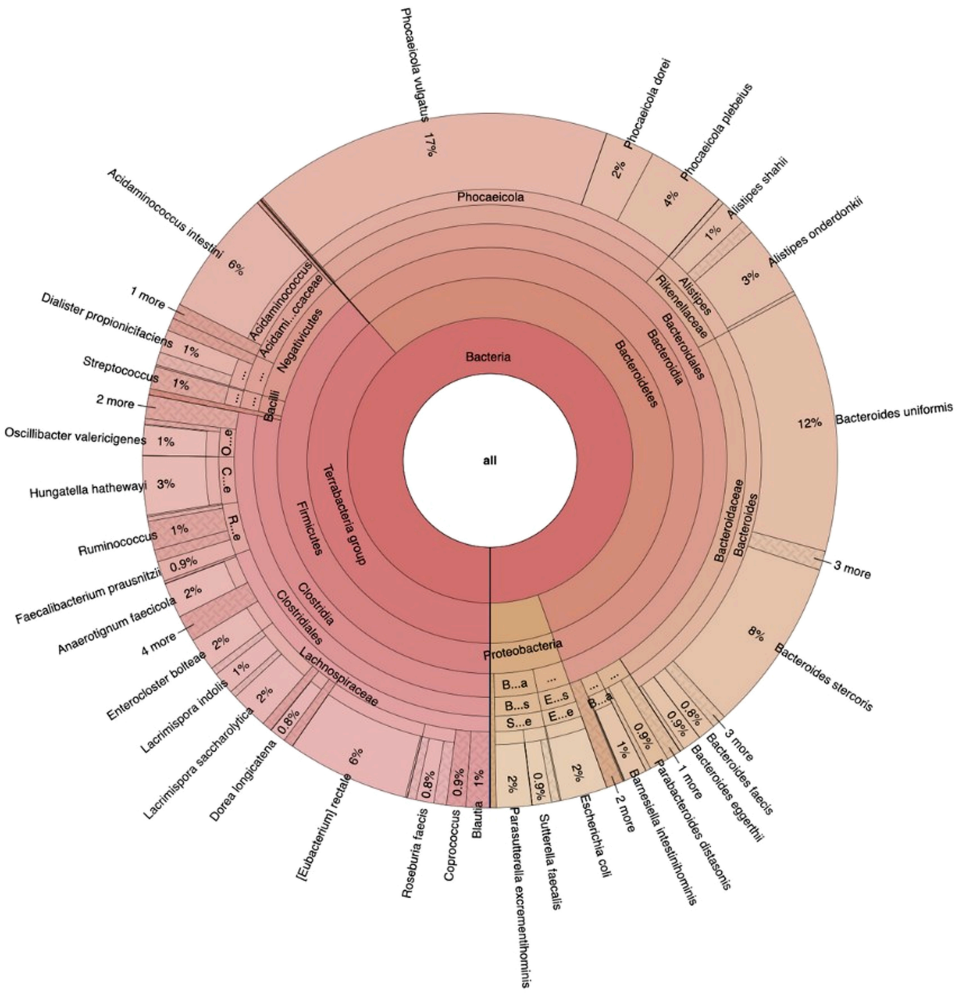
Microbial composition in patients with PDAC



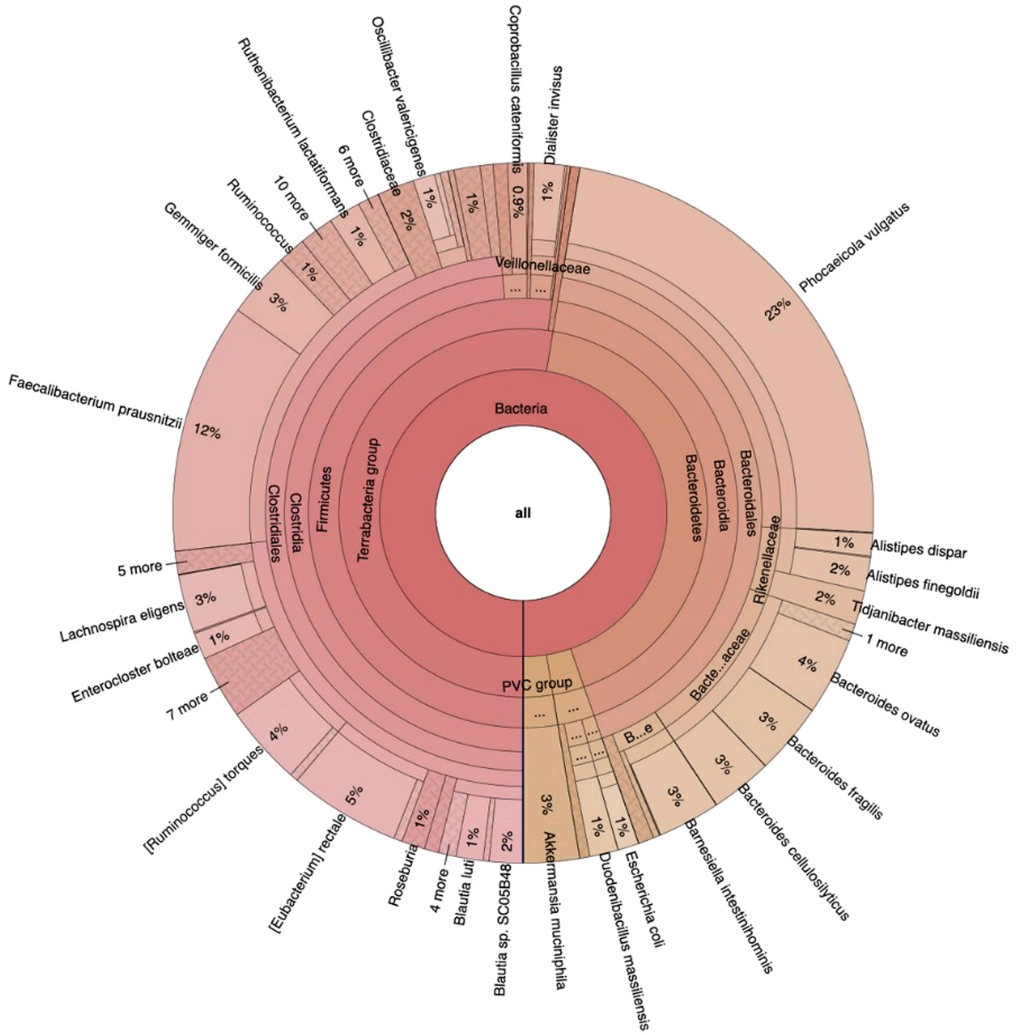
IKPA1



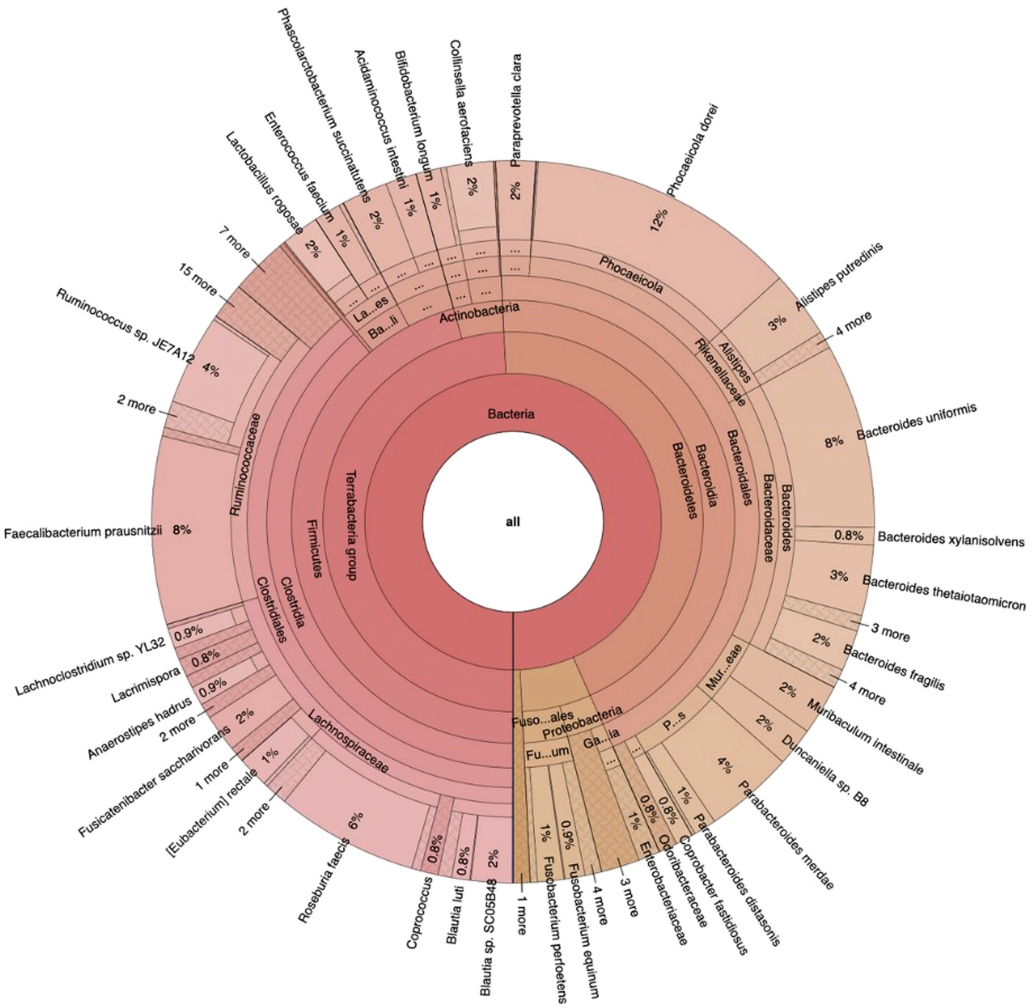
IKPA4



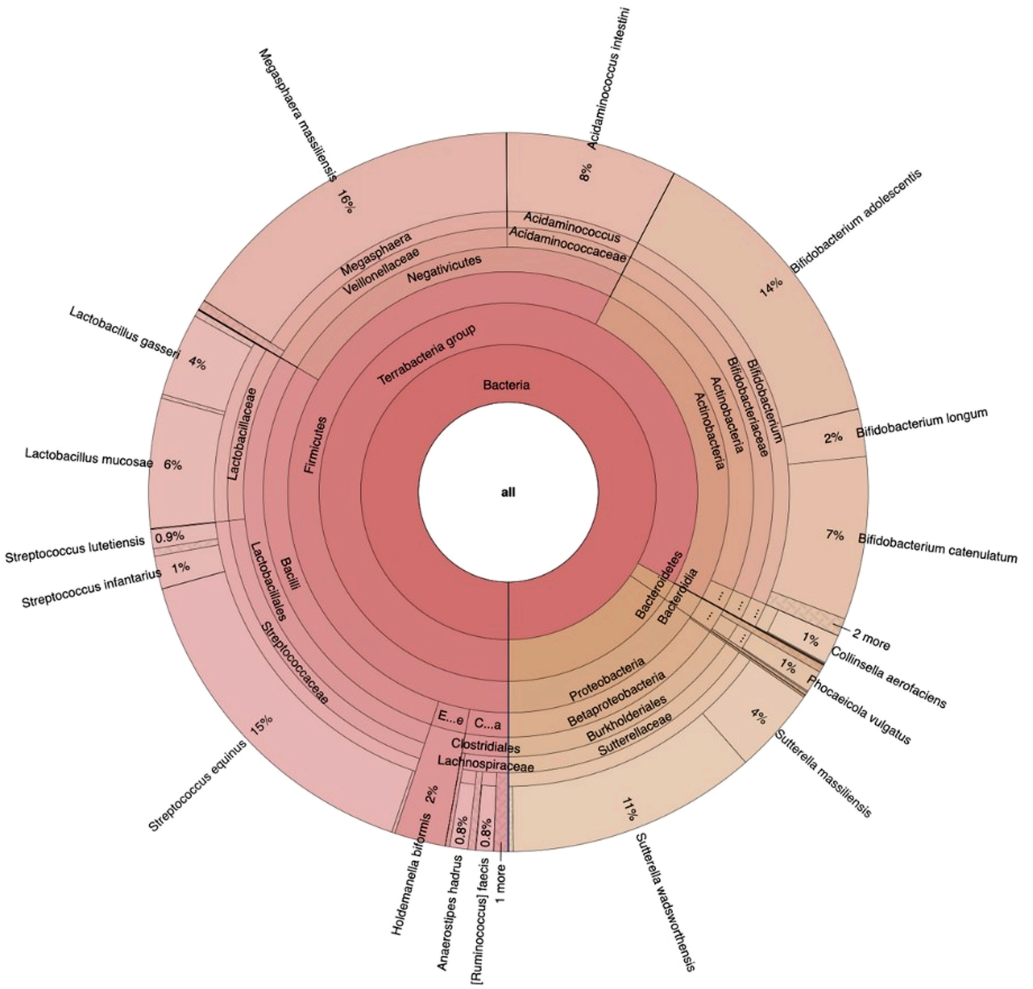
IKPA9



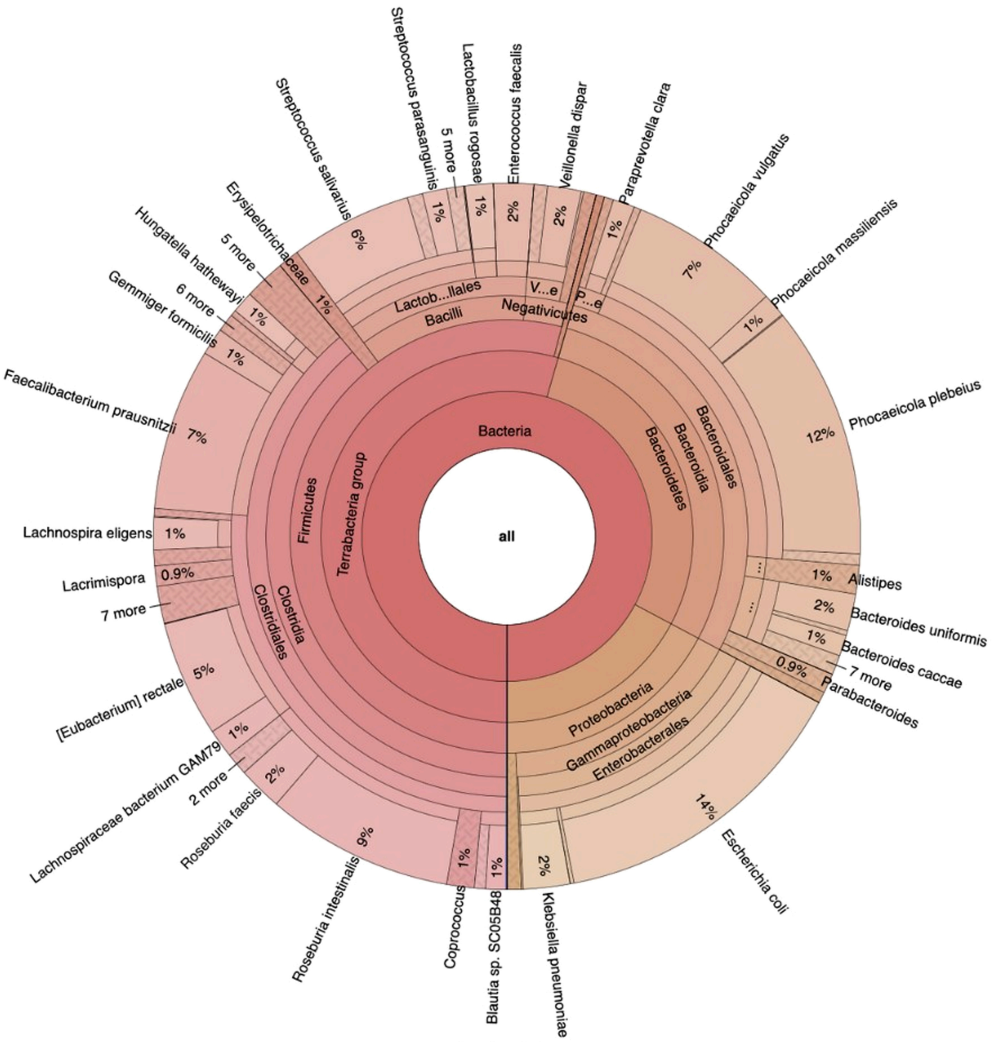
IKPA11



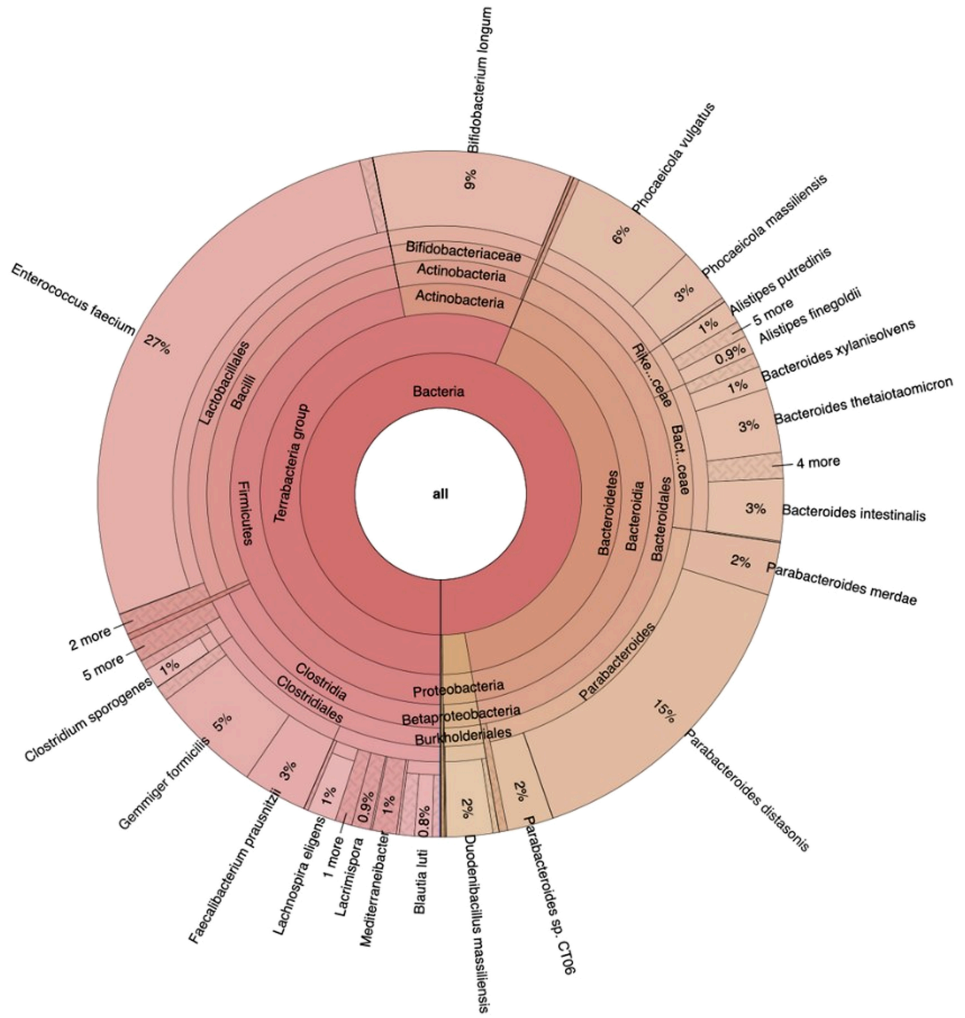
IKPA14



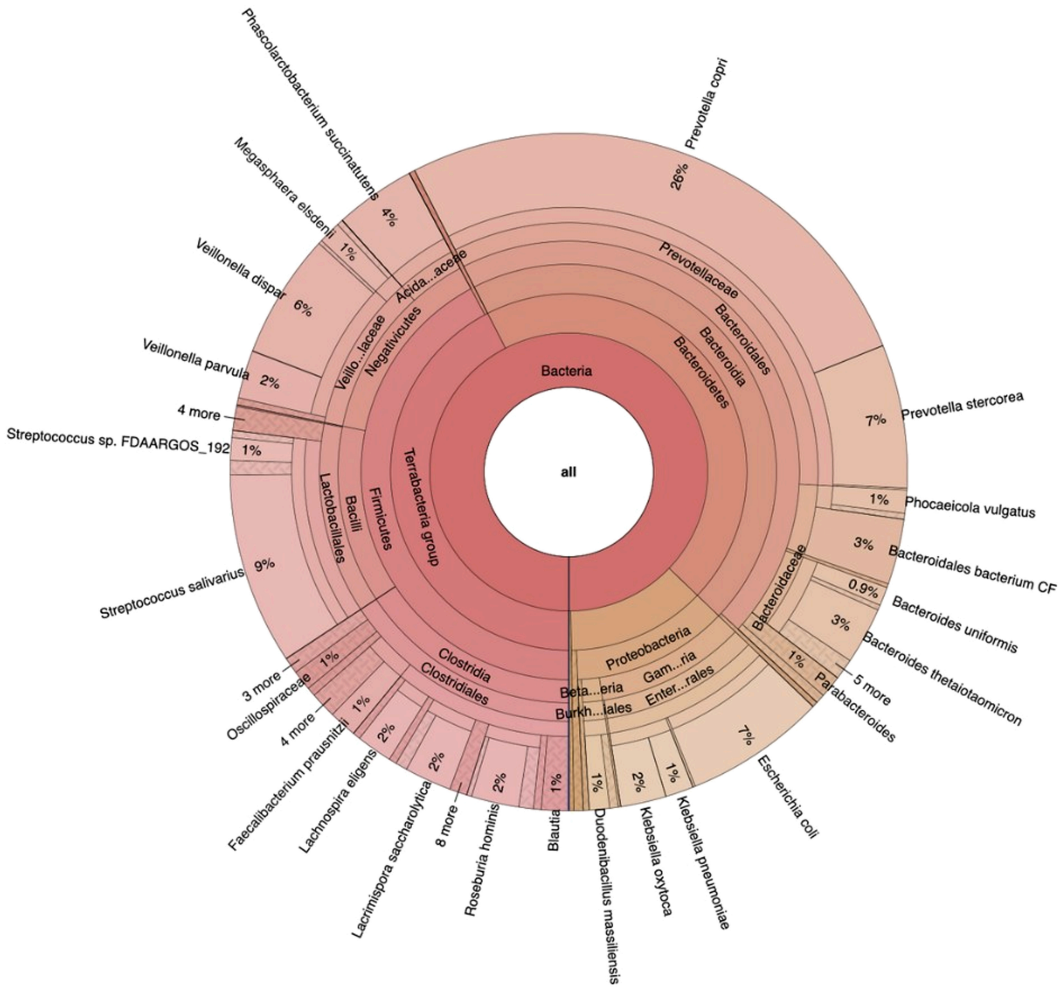
IKPA15



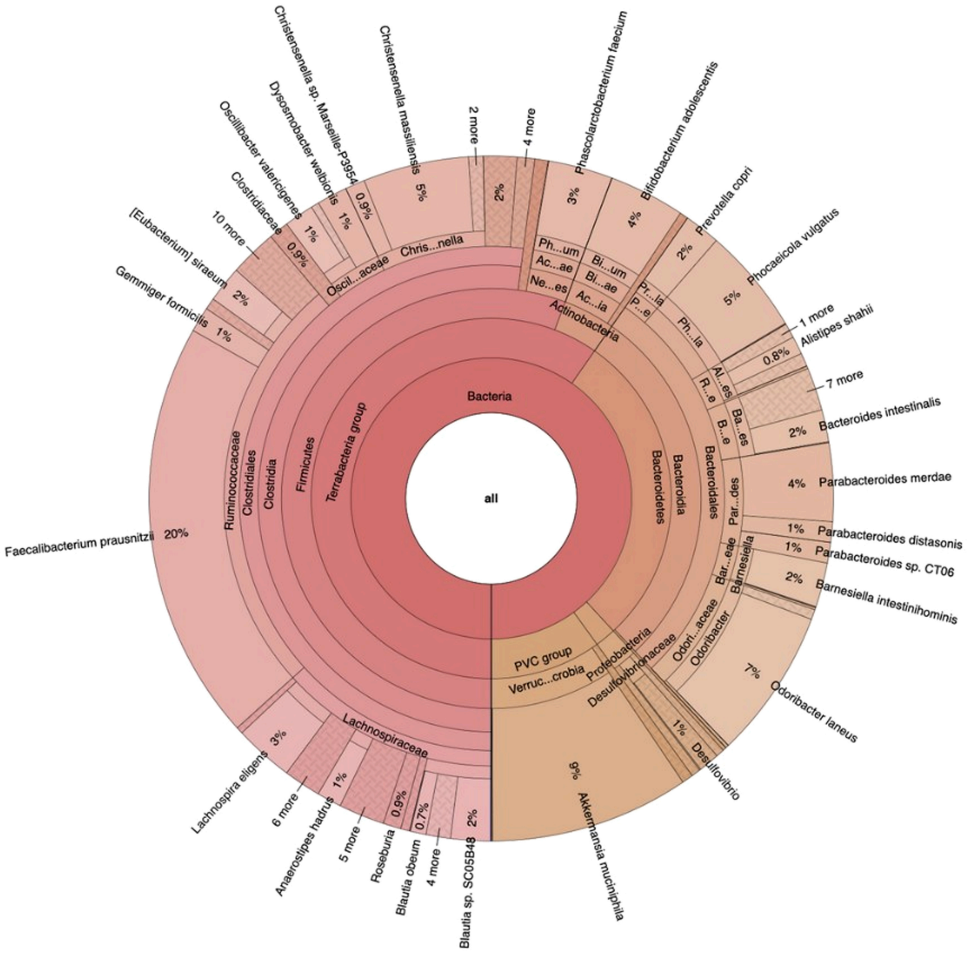
IKPA22



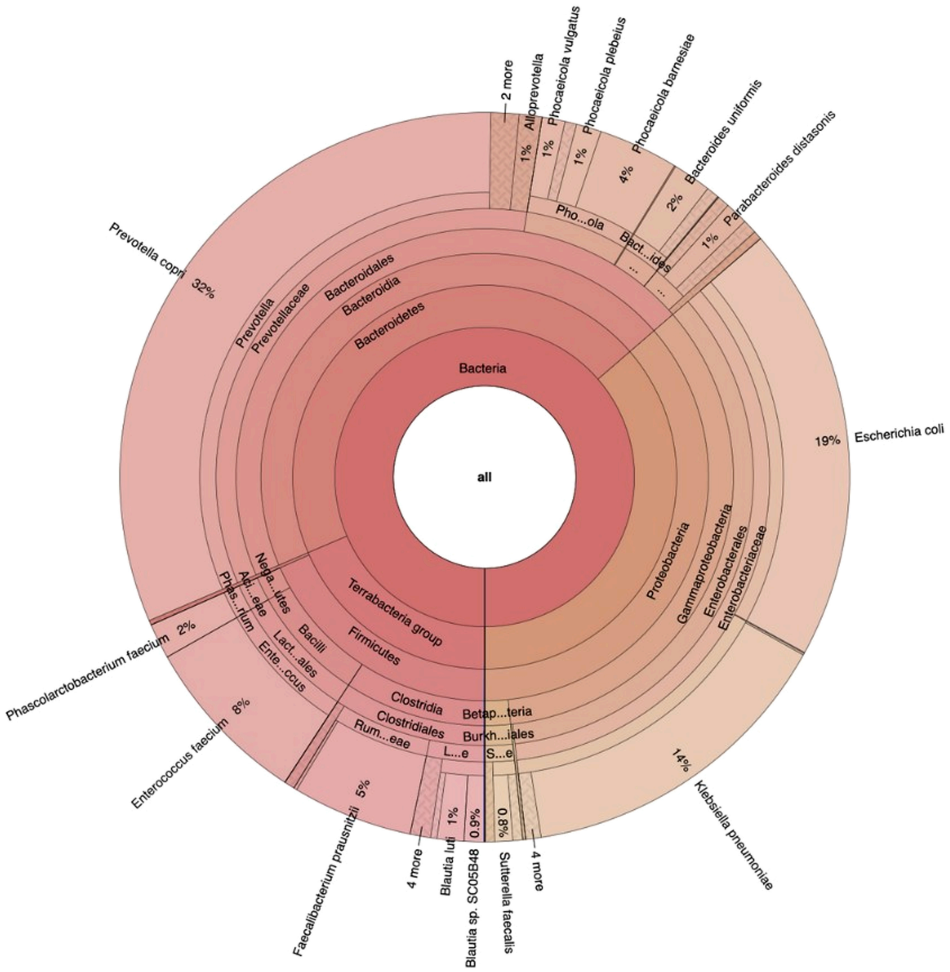
IKPA30



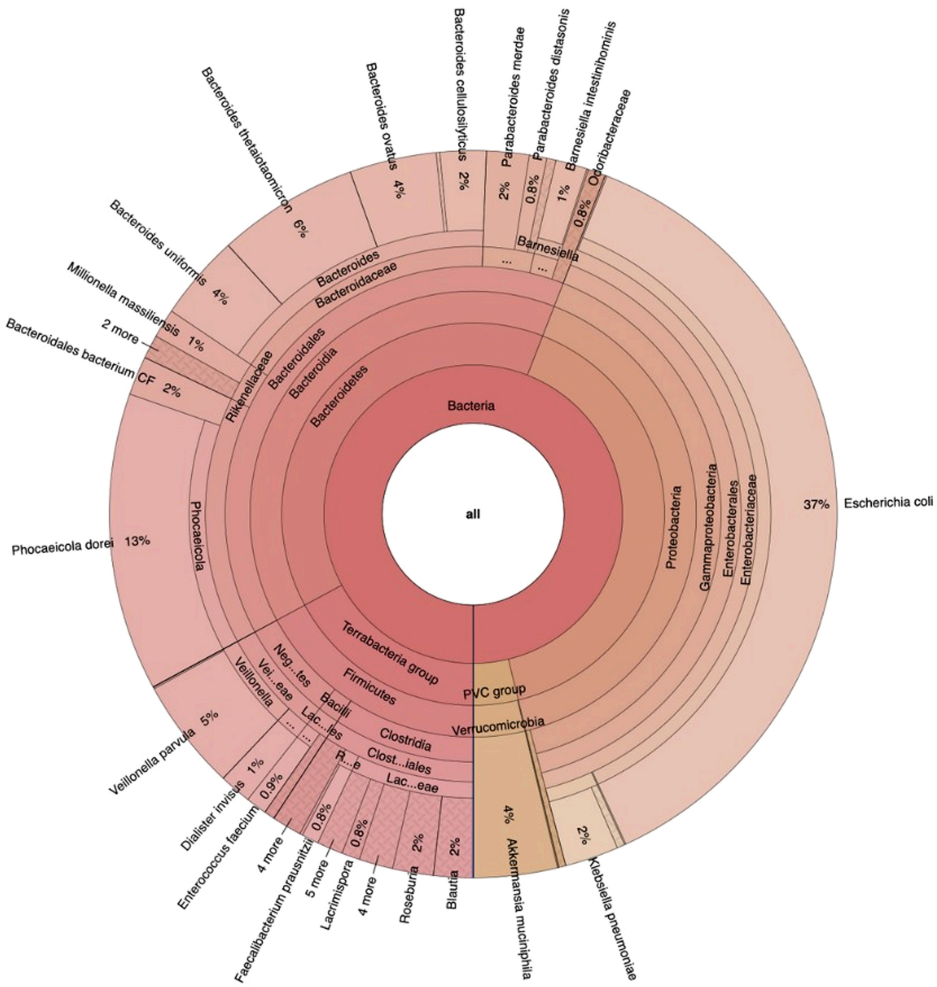
PLPA4



PLPA10

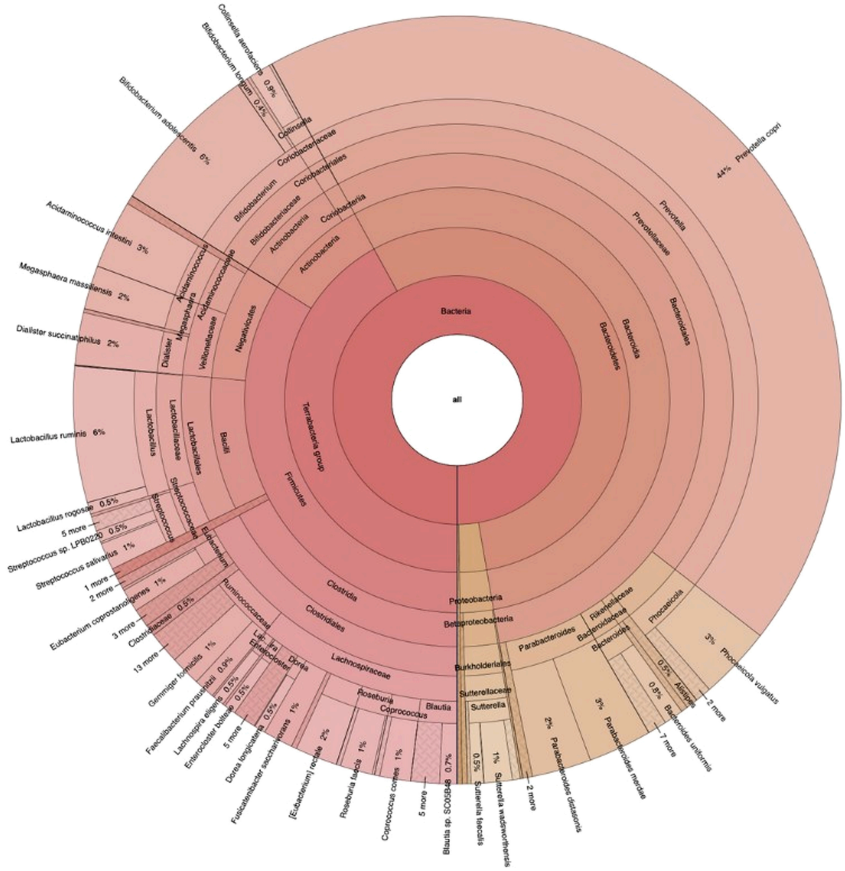


PLPA11

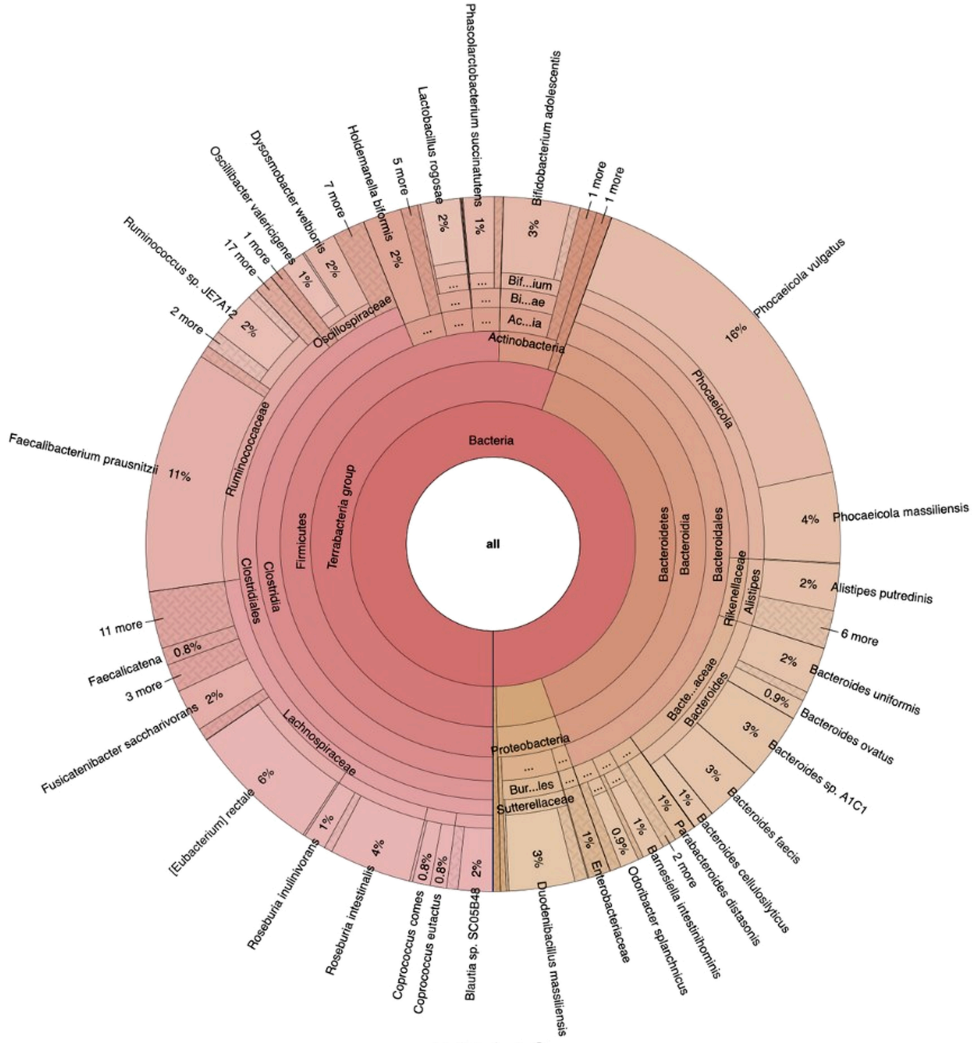


PLPA24

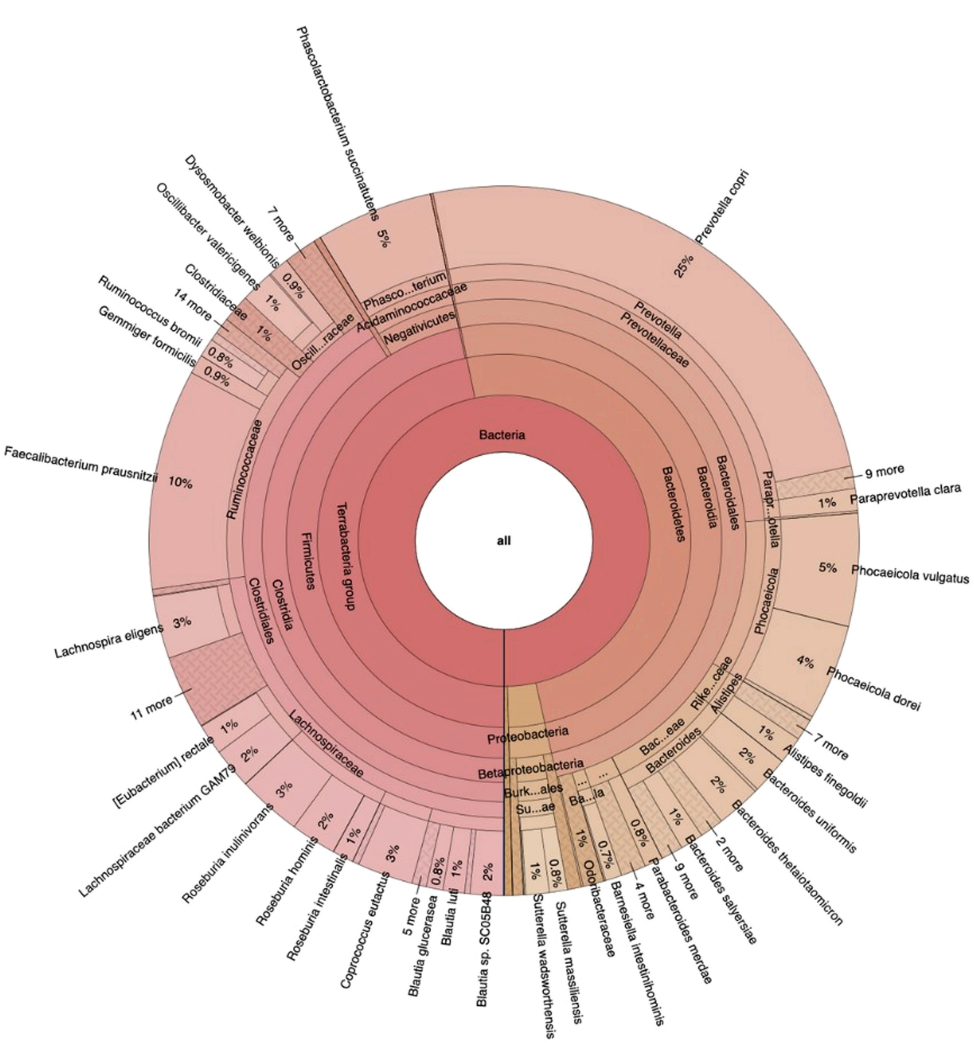
Microbial composition in patients with CP



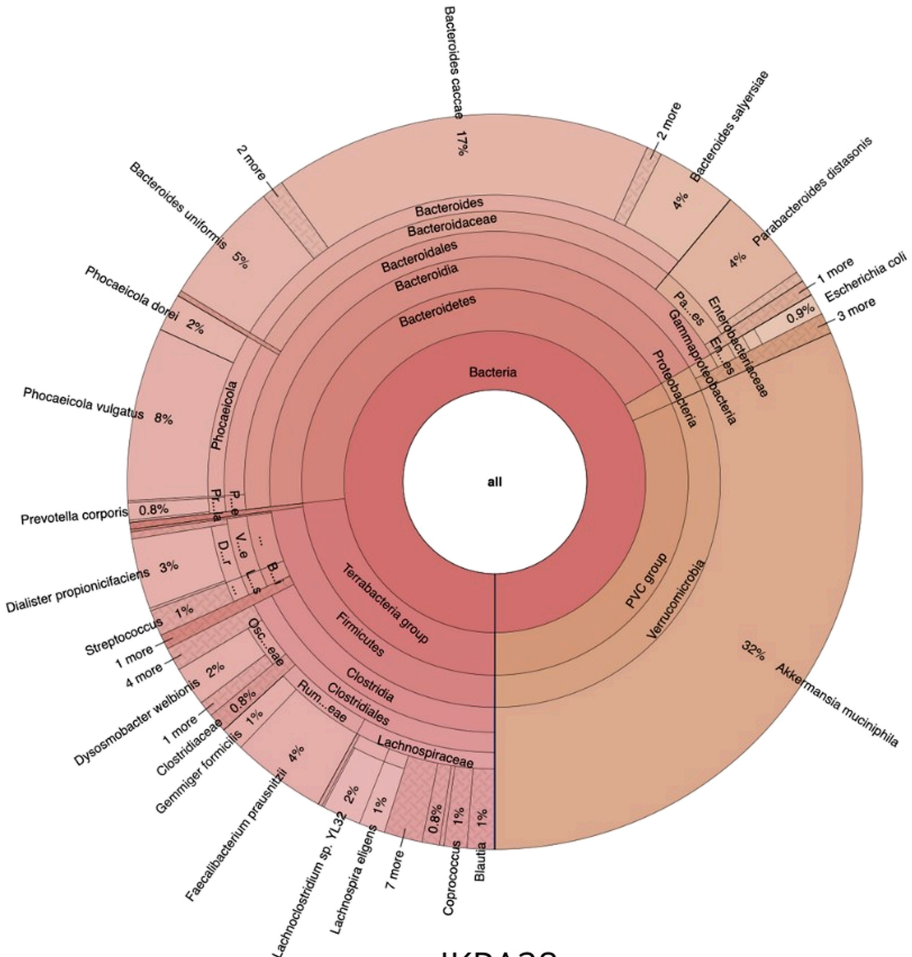
IKPA16



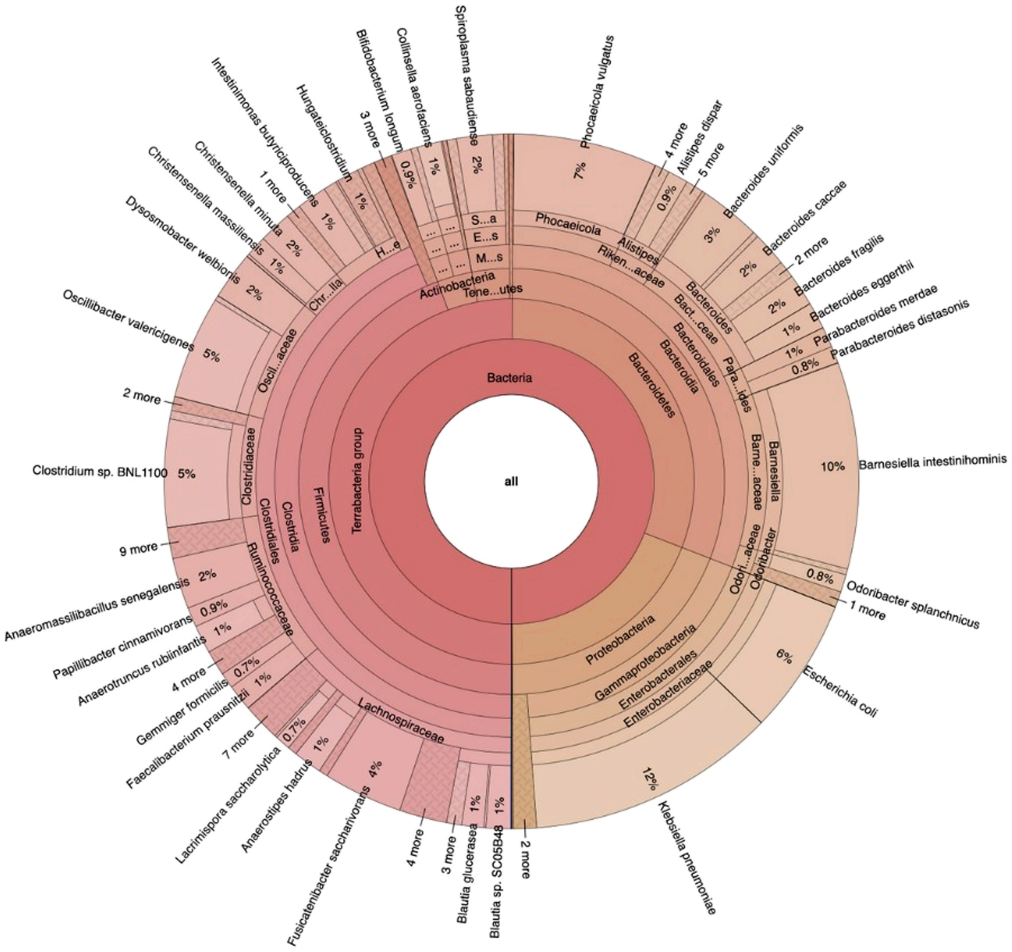
IKPA18



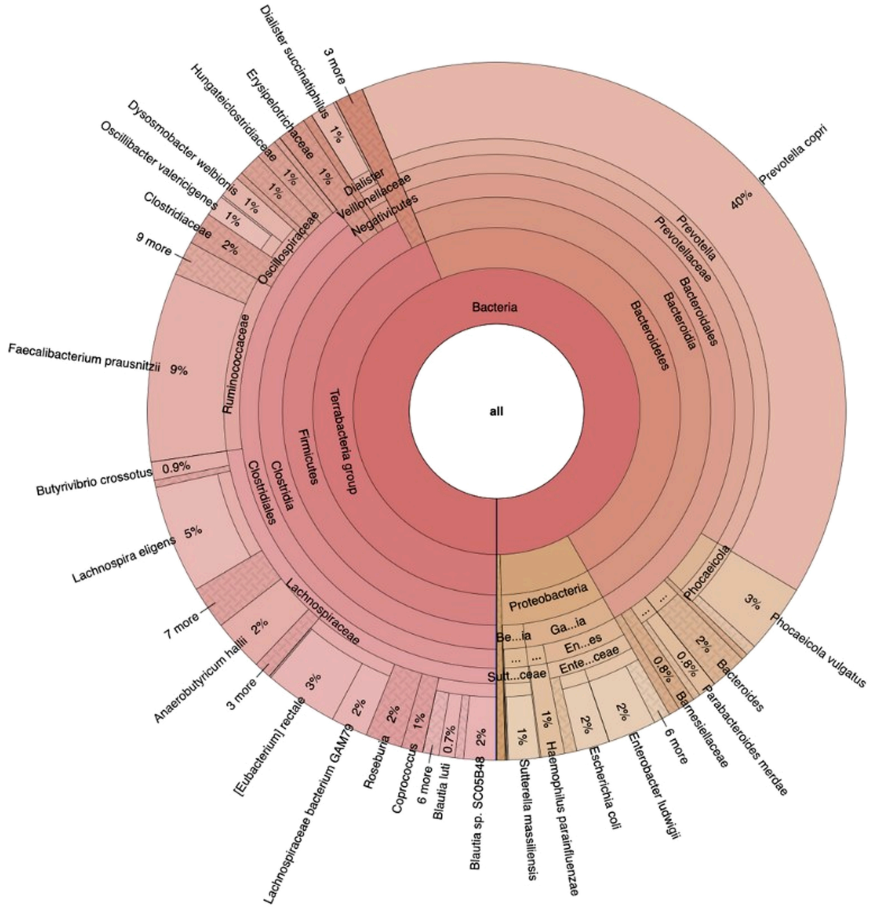
IKPA19



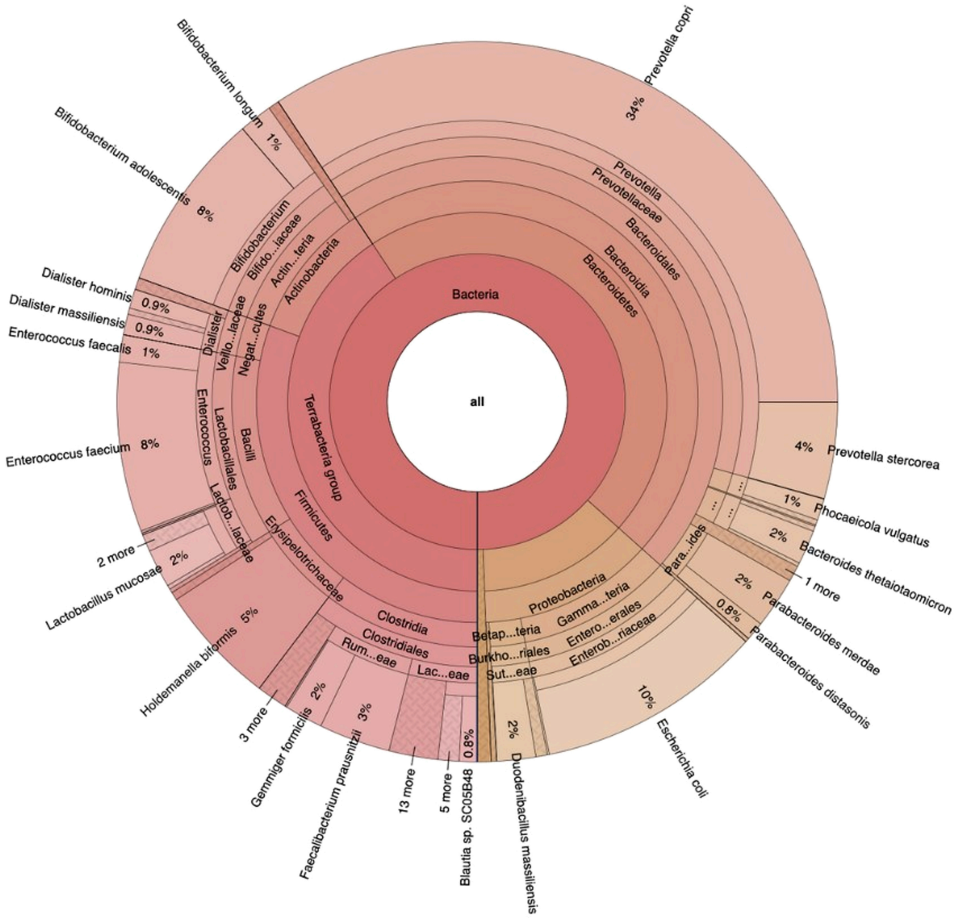
IKPA28



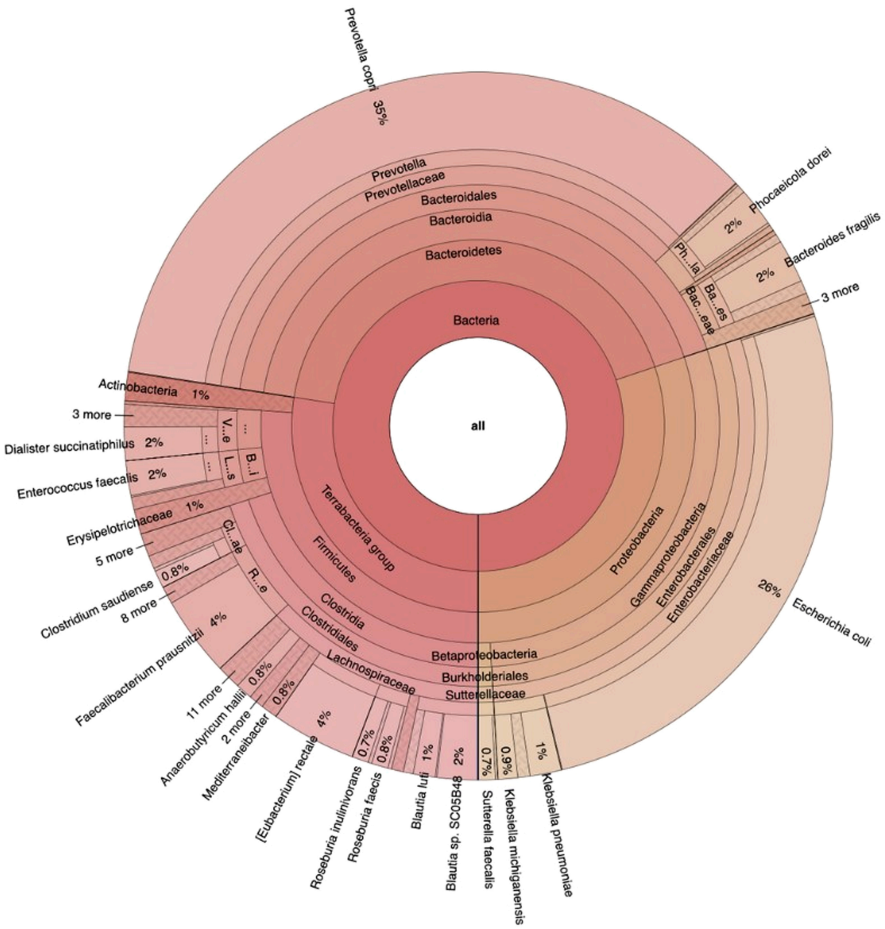
IKPA37



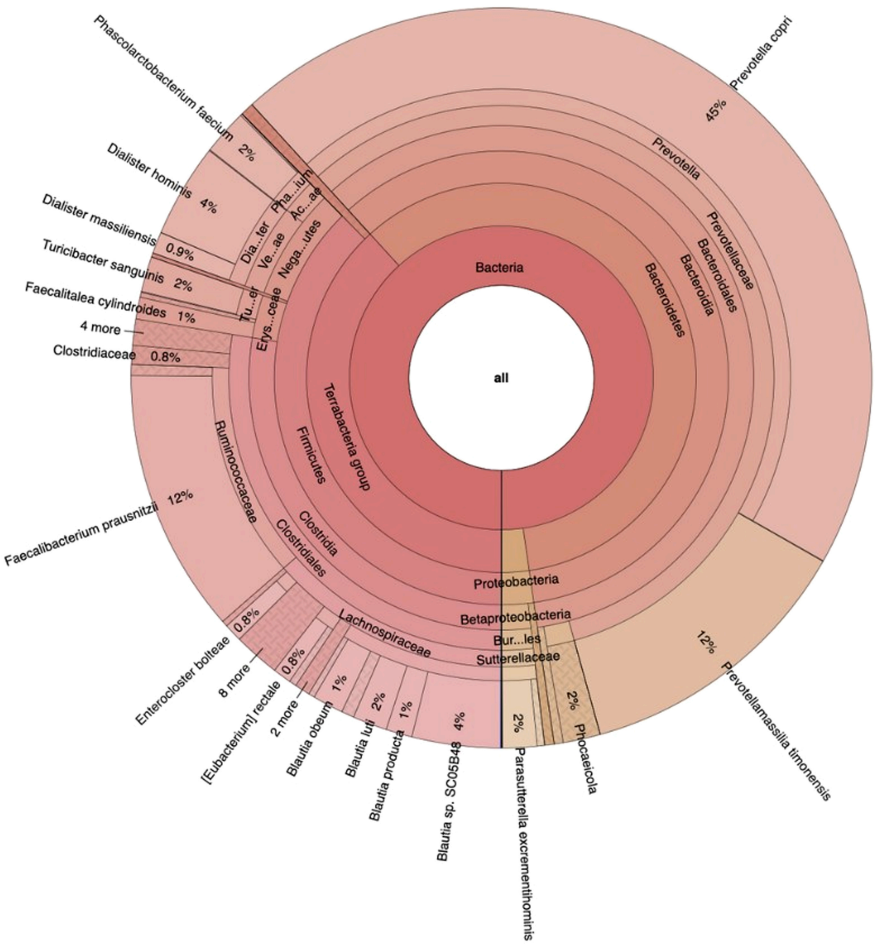
IKPA48



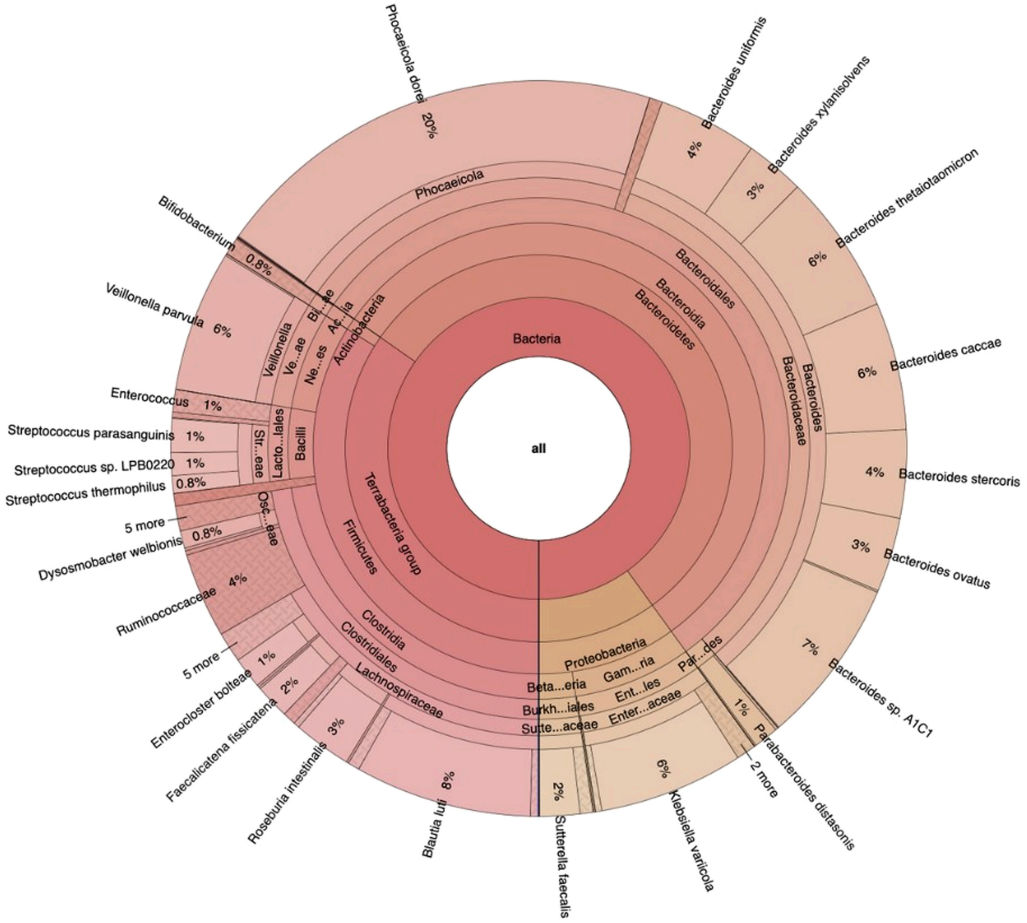
IKPA50



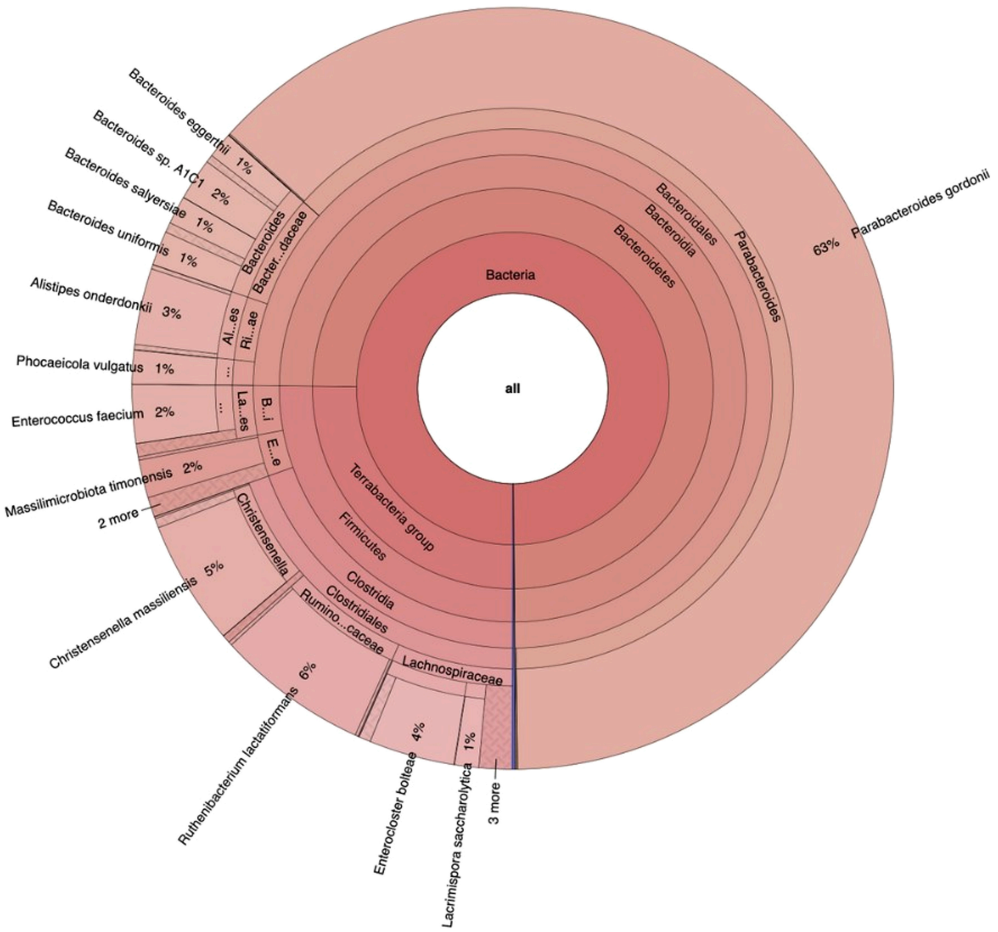
IKPA51



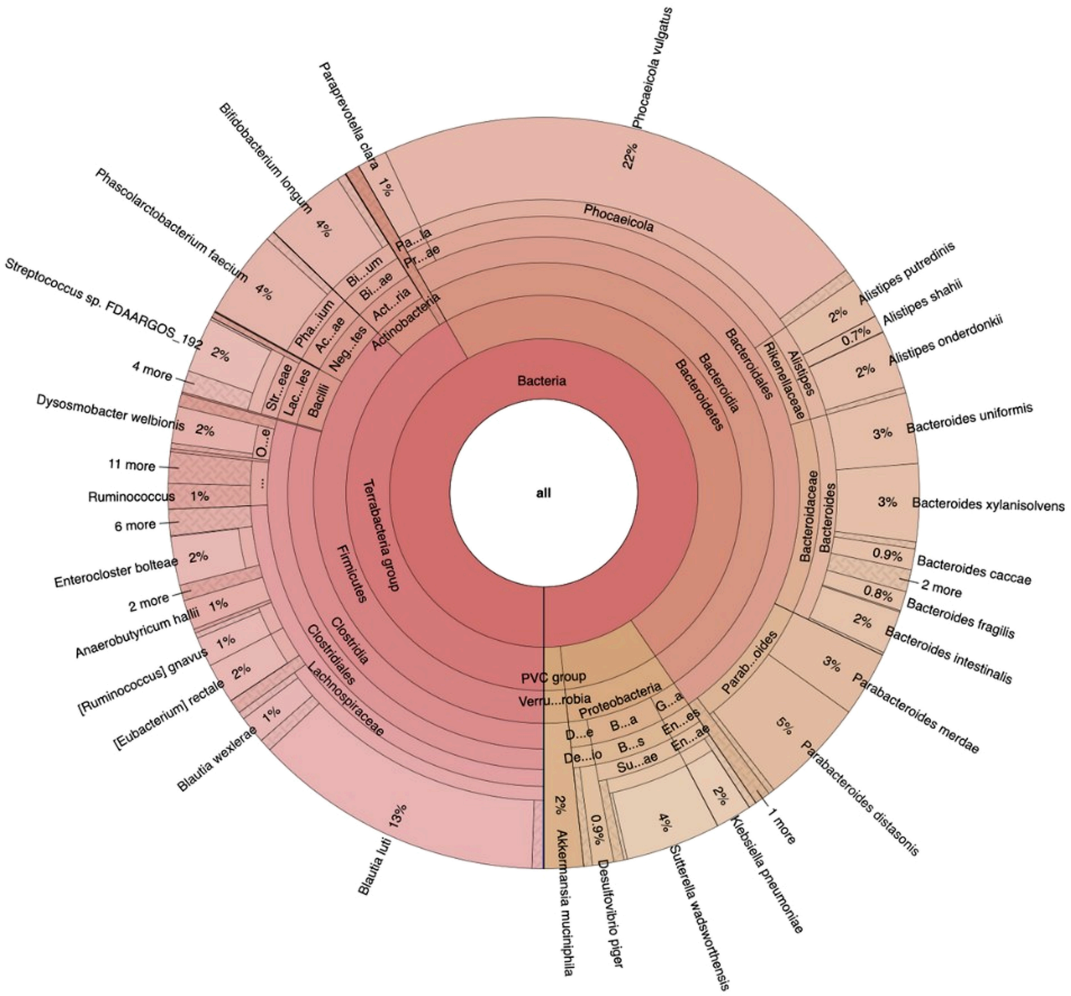
IKPA59



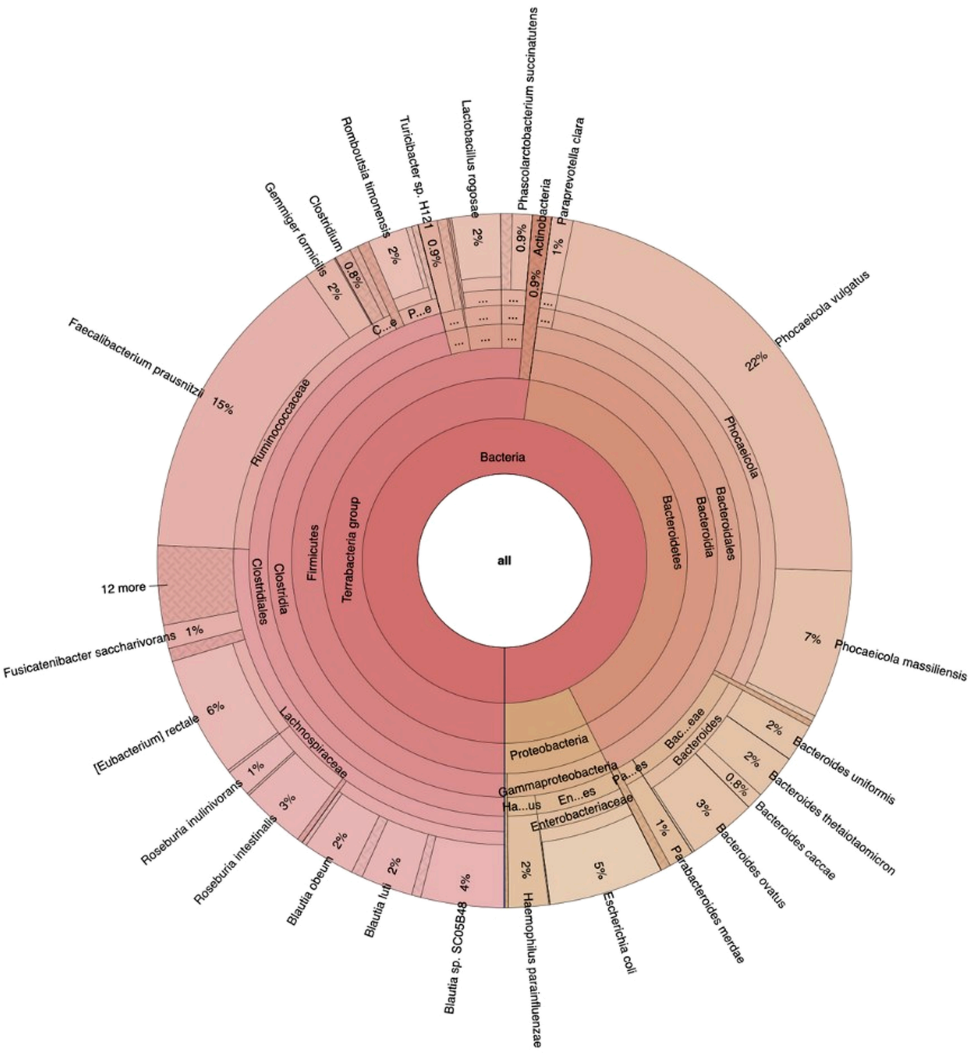
IKPA65



IKPA94

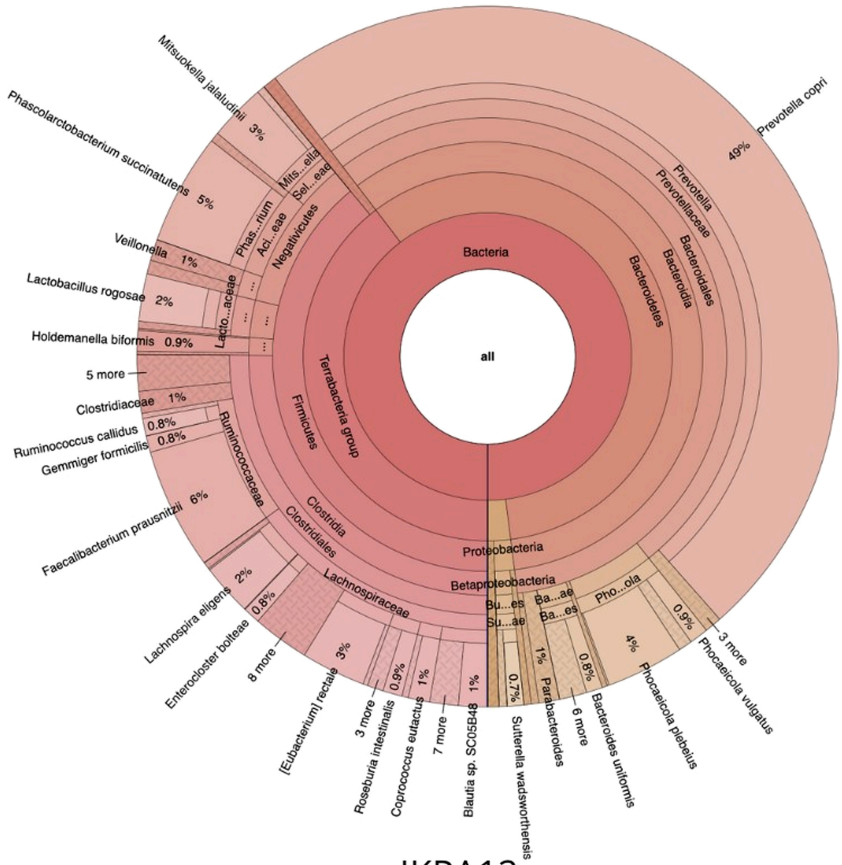


IKPA109

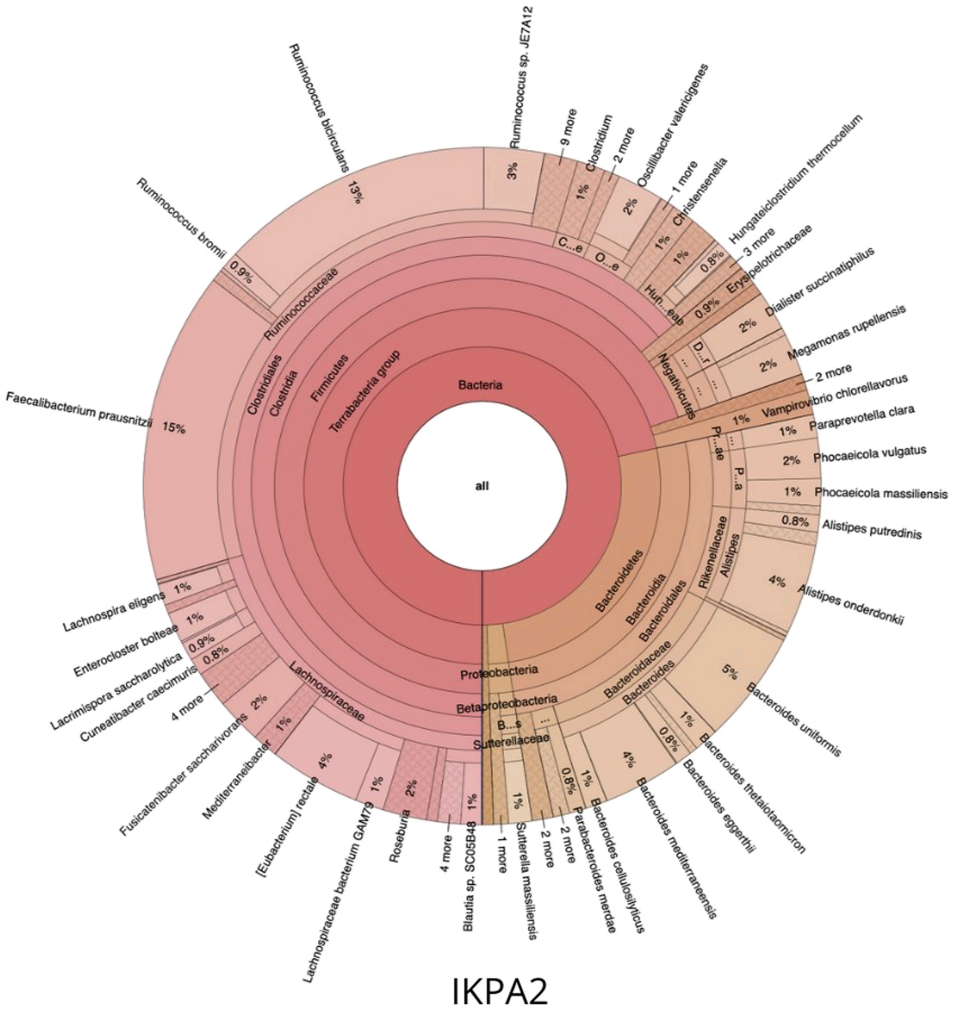


IKPA111

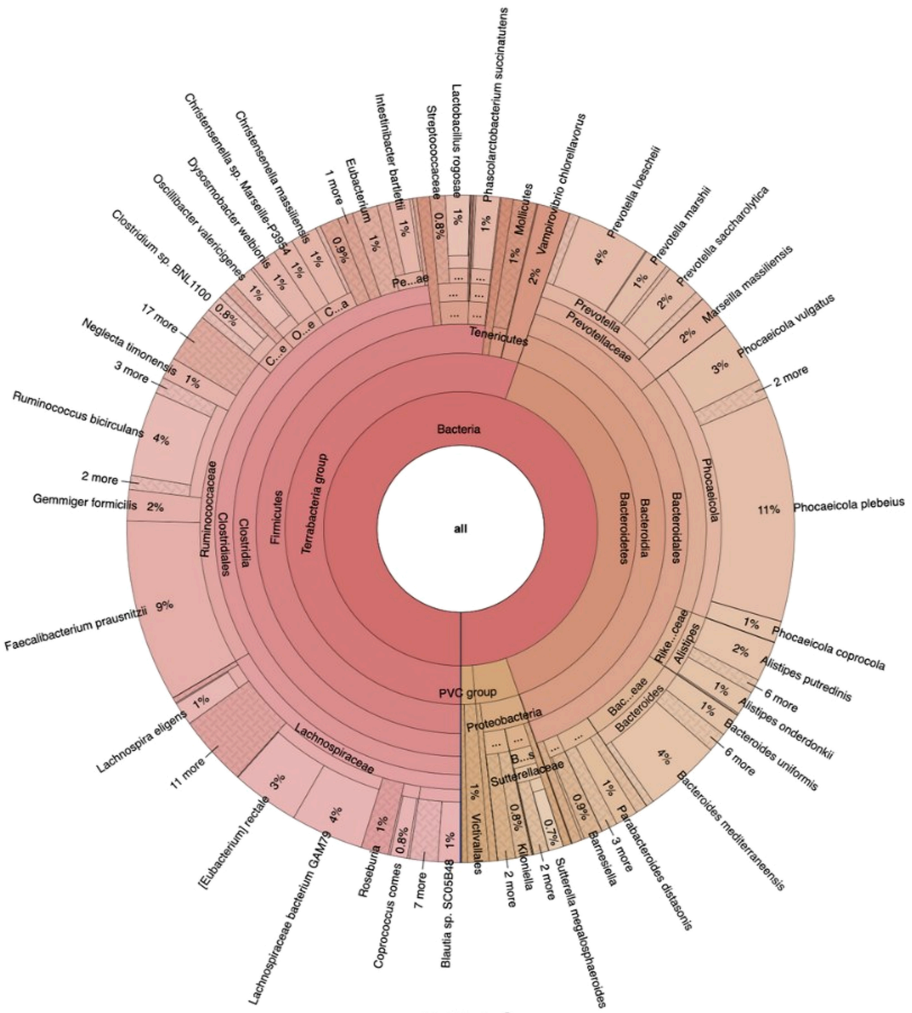
Microbial composition in healthy individuals



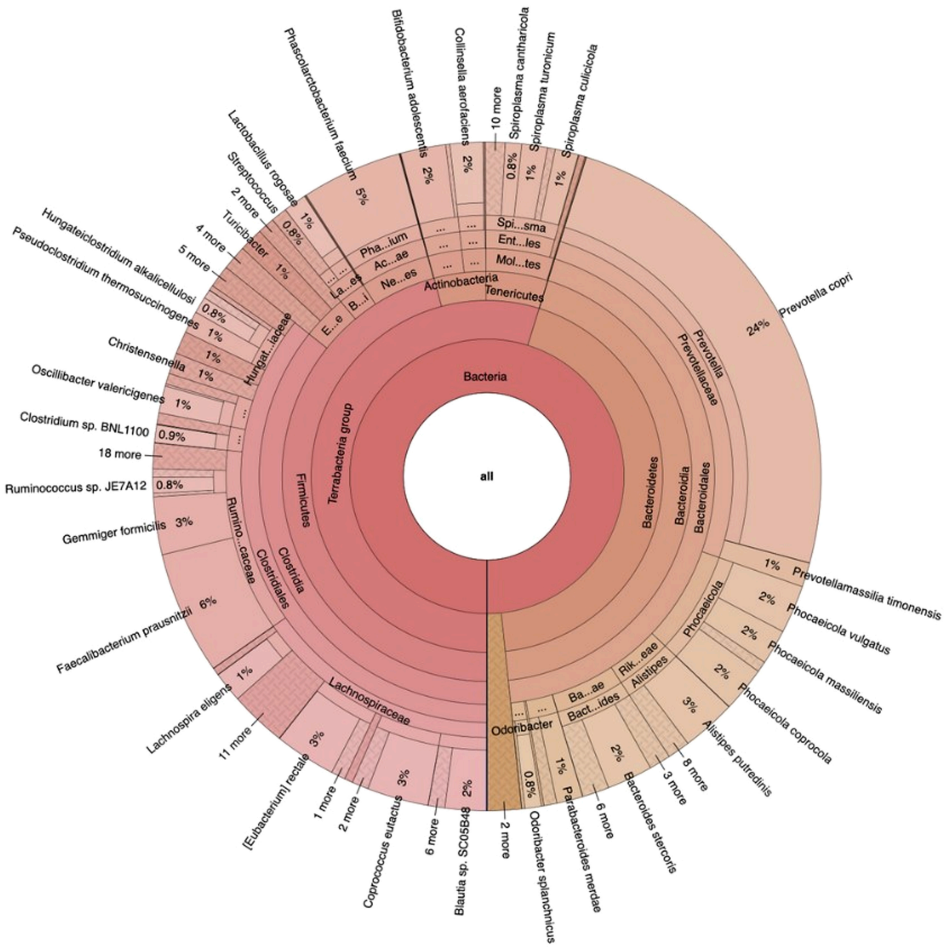
IKPA12


all
Bacteria
Firmicutes
Terrabacteria group
Clostridia
Clostridiales
Ruminococcaceae
Bacteroidetes
Bacteroidia
Bacteroidales
Rikenellaceae
Alistipes
Bacteroidaceae
Bacteroides
Proteobacteria
Betaproteobacteria
Sutterellaceae
Lachnospiraceae
Negativicutes
Ruminococcus sp. JE7A12
Ruminococcus bicirculans
Ruminococcus bromii
Faecalibacterium prausnitzii 15%
Lachnospira eligens 1%
Enterocloster bolteae 1%
Lacrimispora saccharolytica 0.9%
Cuneatibacter caecimuris 0.8%
4 more
Fusicatenibacter saccharivorans 2%
Mediterraneibacter 1%
[Eubacterium] rectale 4%
Lachnospiraceae bacterium GAM73 1%
Roseburia 2%
4 more
Blautia sp. SC05B48 1%
1 more
Sutterella massiliensis 1%
2 more
Parabacteroides merdae 0.8%
Bacteroides cellulosilyticus 1%
Bacteroides mediterraneensis 4%
Bacteroides eggerthii 0.8%
Bacteroides thetaiotaomicron 1%
Bacteroides uniformis 5%
Alistipes onderdonkii 4%
Alistipes putredinis 0.8%
Phocaeicola massiliensis 1%
Phocaeicola vulgatus 2%
Paraprevotella clara 1%
Vampirovibrio chlorellavorus 1%
2 more
Megamonas rupellensis 2%
Dialister succinatiphilus 0.9%
Erysipelotrichaceae 0.8%
3 more
Hungateiclostridium thermocellum 1%
Christensenella 1%
1 more
Oscillibacter valericigenes 2%
Clostridium 2%
2 more
9 more 1%
3%

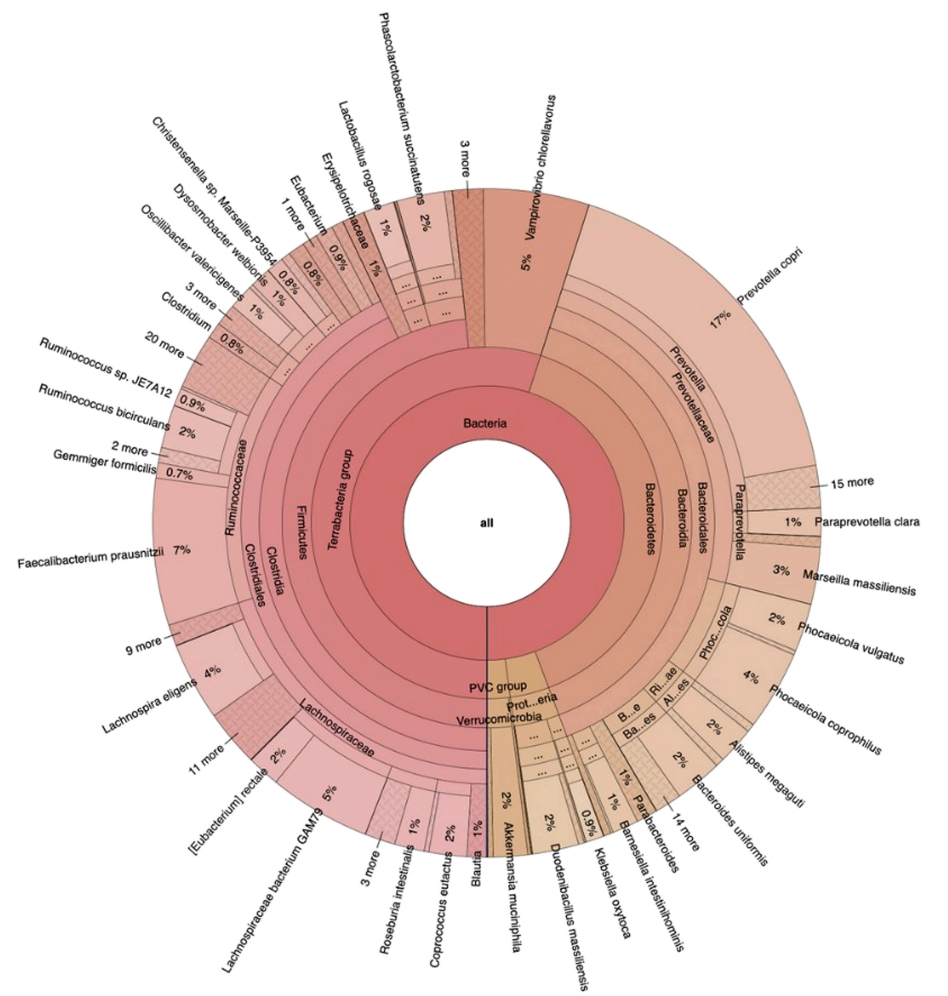
IKPA2


all
Bacteria
Firmicutes
Terrabacteria group
Clostridia
Clostridiales
Ruminococcaceae
Bacteroidetes
Bacteroidia
Bacteroidales
Phocaeicola
Prevotellaceae
Prevotella
Tenericutes
Mollicutes
PVC group
Proteobacteria
Sutterellaceae
Lachnospiraceae
Ruminococcus bicirculans 4%
Faecalibacterium prausnitzii 9%
Lachnospira eligens 1%
11 more
[Eubacterium] rectale 3%
Lachnospiraceae bacterium GAM73 4%
Roseburia 1%
Coprococcus comes 0.8%
7 more
Blautia sp. SC05B48 1%
Victivallales 1%
2 more
Klebsiella 0.8%
Sutterella megalosphaeroides 0.7%
2 more
Barnesiella 0.9%
3 more
Parabacteroides distasonis 1%
Bacteroides mediterraneensis 4%
Bacteroides uniformis 1%
6 more
Alistipes onderdonkii 1%
Alistipes putredinis 2%
6 more
Phocaeicola coprocola 1%
Phocaeicola plebeius 11%
2 more
Phocaeicola vulgatus 3%
Marseilla massiliensis 2%
Prevotella saccharolytica 2%
Prevotella marshii 1%
Prevotella loescheii 4%
Vampirovibrio chlorellavorus 2%
Phascolarctobacterium succinatutens 1%
Lactobacillus rogosae 1%
Streptococcaceae 0.8%
Intestinibacter bartlettii 1%
Eubacterium 0.9%
1 more
Christensenella massiliensis 1%
Christensenella sp. Marseille-P3054 1%
Dysosmobacter welbionis 1%
Oscillibacter valericigenes 1%
Clostridium sp. BNL1100 0.8%
17 more
Neglecta timonensis 1%
3 more
Gemmiger formicilis 2%
2 more

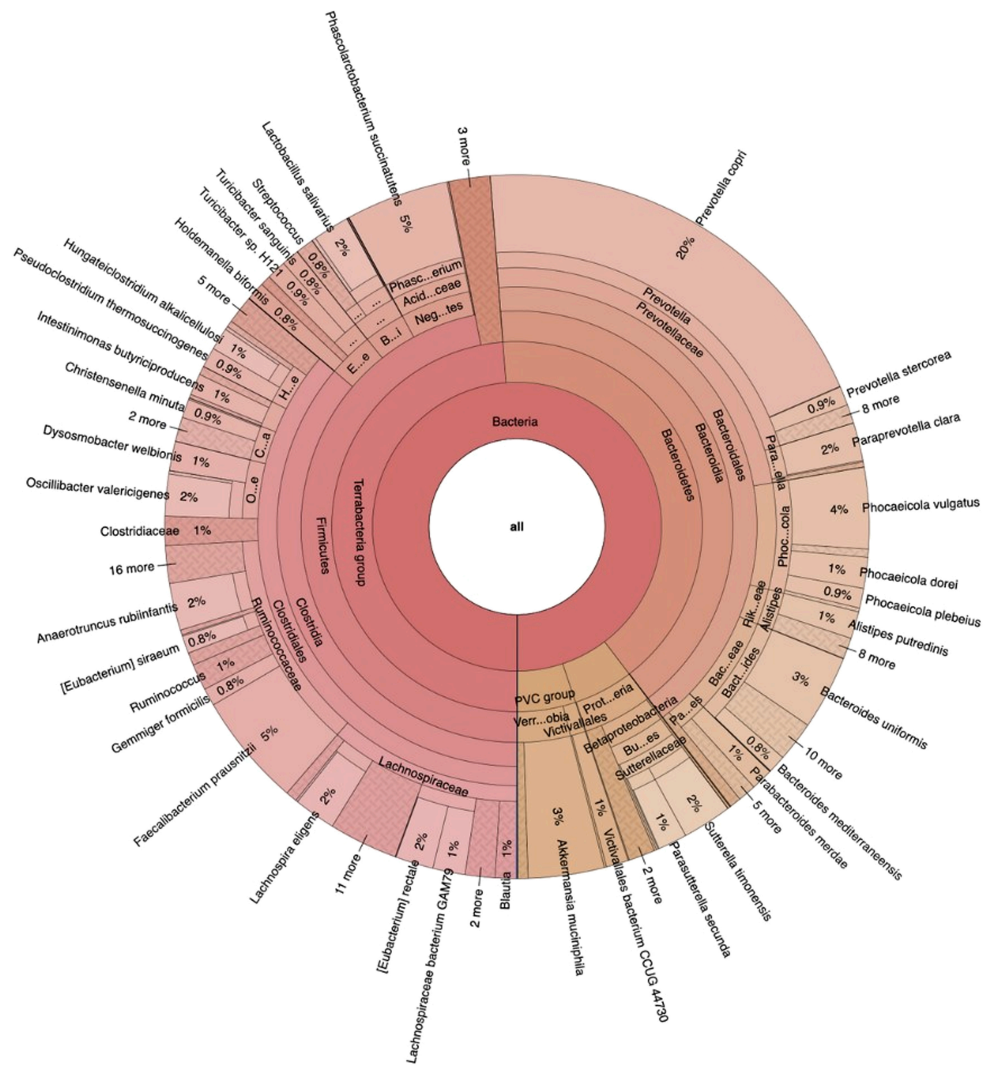
IKPA3


all
Bacteria
Terrabacteria group
Firmicutes
Clostridia
Clostridiales
Rumino...ceae
Lachnospiraceae
Bacteroidetes
Bacteroidia
Bacteroidales
Prevotellaceae
Prevotella
Actinobacteria
Tenericutes
Prevotella copri 24%
Faecalibacterium prausnitzii 6%
Gemmiger formicilis 3%
Ruminococcus sp. JE7A12 0.8%
18 more
Clostridium sp. BNL1100 0.9%
Oscillibacter valericigenes 1%
Christensenella 1%
Hungat. laceae
Turicibacter 1%
4 more
Hungateiclostridium alkalicellulosi
Pseudoclostridium thermosuccinogenes 0.8%
5 more
Lactobacillus rogosae 1%
Streptococcus 0.8%
2 more
Phascolarctobacterium faecium 5%
Bifidobacterium adolescentis 2%
Collinsella aerofaciens 2%
10 more
Spiroplasma citri 1%
Spiroplasma taiwanense 1%
Spiroplasma culicicola 1%
Spi...sma
Ent...les
Mol...tes
Pha...ium
Ac...ae
Ne...es
La...ea
B...i
E...e
Lachnospira eligens 1%
11 more
[Eubacterium] rectale 3%
1 more
2 more
Coprococcus eutactus 3%
6 more
Blautia sp. SC05B48 2%
2 more
Odoribacter 2%
Odoribacter splanchnicus 0.8%
Parabacteroides merdae 1%
6 more
Bacteroides stercoris 2%
3 more
8 more
Alistipes putredinis 3%
Phocaeicola coprocola 2%
Phocaeicola massiliensis 2%
Phocaeicola vulgatus 2%
Prevotellamassilia timonensis 1%
Alistipes
Rik...eae
Ba...ae
Bact...ides

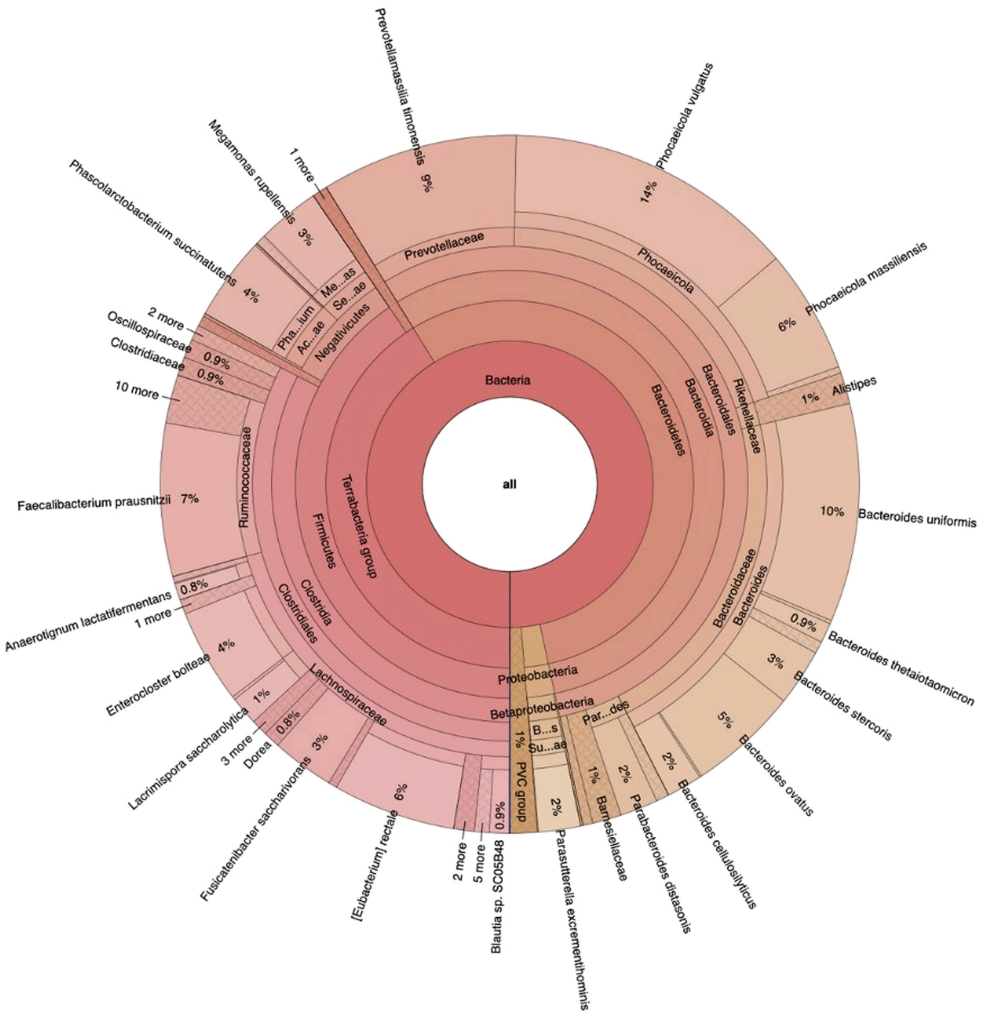
IKPA5


all
Bacteria
Terrabacteria group
Firmicutes
Clostridia
Clostridiales
Ruminococcaceae
Lachnospiraceae
Bacteroidetes
Bacteroidia
Bacteroidales
Prevotellaceae
Prevotella
Paraprevotella
PVC group
Verrucomicrobia
Prevotella copri 17%
Faecalibacterium prausnitzii 7%
9 more
Lachnospira eligens 4%
11 more
[Eubacterium] rectale 2%
Lachnospiraceae bacterium GAM79 5%
3 more
Roseburia intestinalis 1%
Coprococcus eutactus 2%
Blautia 1%
Akkermansia muciniphila 2%
Duodenibacillus massiliensis 2%
Klebsiella oxytoca 0.9%
Barnesiella intestinihominis 1%
Parabacteroides 1%
14 more
Bacteroides uniformis 2%
Alistipes megaguti 2%
Phocaeicola coprophilus 4%
Phocaeicola vulgatus 2%
Marseilia massiliensis 3%
Paraprevotella clara 1%
15 more
Prot...eria
Al...es
Ri...ae
Ba...es
B...e
Phoc...cola
Ruminococcus sp. JE7A12 0.9%
Ruminococcus bicirculans 2%
2 more
Gemmiger formicilis 0.7%
20 more
Clostridium 0.8%
3 more
Oscillibacter valericigenes 1%
Dysosmobacter welbionis 0.8%
Christensenella sp. Marseille-P3954 0.8%
1 more
Eubacterium 0.9%
Erysipelotrichaceae 1%
Lactobacillus rogosae 1%
Phascolarctobacterium succinatutens 2%
3 more
Vampirovibrio chlorellavorus 5%

IKPA6



IKPA8



IKPA10