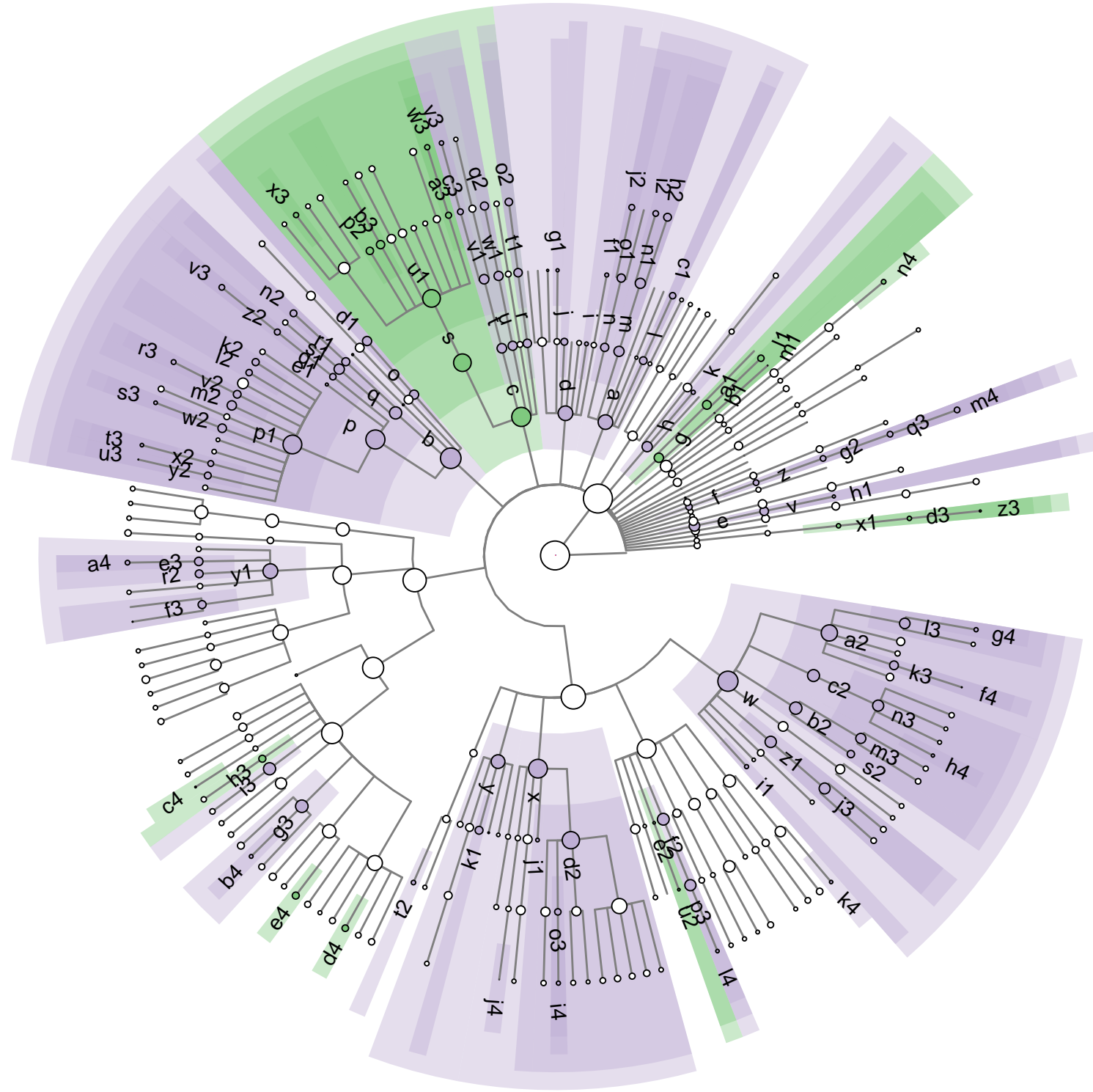


Groups

- AF
- FB



Taxa

a: p__Acidobacteria	z: c__Spirochaetes	y1: o__Bacillales	x2: f__Streptomycetaceae	w3: g__Butyricimonas
b: p__Actinobacteria	a1: c__Mollicutes	z1: o__Caulobacterales	y2: f__Streptosporangiaceae	x3: g__Paraprevotella
c: p__Bacteroidetes	b1: c__RF3	a2: o__Rhizobiales	z2: f__Bifidobacteriaceae	y3: g__Flavobacterium
d: p__Chloroflexi	c1: o__iii1_15	b2: o__Rhodospirillales	a3: f__[Odoribacteraceae]	z3: g__Candidatus_Proteochlamydia
e: p__Gemmatimonadetes	d1: o__Acidimicrobiales	c2: o__Sphingomonadales	b3: f__Rikenellaceae	a4: f__Bacillaceae_g__Bacillus
f: p__Spirochaetes	e1: o__Micrococcales	d2: o__Burkholderiales	c3: f__Flavobacteriaceae	b4: g__Proteiniclasticum
g: p__Tenericutes	f1: o__[Roseiflexales]	e2: o__Pasteurellales	d3: f__Parachlamydiaceae	c4: g__Anaerofustis
h: p__TM7	g1: o__B07_WMSP1	f2: o__Xanthomonadales	e3: f__Bacillaceae	d4: g__Coproccoccus
i: c__Ellin6529	h1: o__Ellin5290	g2: o__Spirochaetales	f3: f__Planococcaceae	e4: g__Oscillospira
j: c__Thermomicrobia	i1: o__Ellin329	h2: f__Acidobacteriaceae	g3: f__Clostridiaceae	f4: g__Afipia
k: c__TM7_1	j1: o__SC_I_84	i2: f__Koribacteraceae	h3: f__Peptococcaceae	g4: g__Rhodoplanes
l: c__Acidobacteria_6	k1: o__Myxococcales	j2: f__Solibacteraceae	i3: f__Veillonellaceae	h4: g__Kaistobacter
m: c__Acidobacteriia	l1: o__RF39	k2: f__Geodermatophilaceae	j3: f__Caulobacteraceae	i4: g__Burkholderia
n: c__Solibacteres	m1: o__ML615J_28	l2: f__Intrasporangiaceae	k3: f__Bradyrhizobiaceae	j4: g__Nitrosomonas
o: c__Acidimicrobiia	n1: o__Acidobacteriales	m2: f__Micromonosporaceae	l3: f__Hyphomicrobiaceae	k4: g__Psychrobacter
p: c__Actinobacteria	o1: o__Solibacterales	n2: f__Gaiellaceae	m3: f__Rhodospirillaceae	l4: g__Rhodanobacter
q: c__Thermoleophilia	p1: o__Actinomycetales	o2: f__Chitinophagaceae	n3: f__Sphingomonadaceae	m4: g__Treponema
r: c__[Saprospirae]	q1: o__Bifidobacteriales	p2: f__S24_7	o3: f__Burkholderiaceae	n4: g__Akkermansia
s: c__Bacteroidia	r1: o__Gaiellales	q2: f__Sphingobacteriaceae	p3: f__Xanthomonadaceae	
t: c__Flavobacteriia	s1: o__Solirubrobacterales	r2: f__Paenibacillaceae	q3: f__Spirochaetaceae	
u: c__Sphingobacteriia	t1: o__[Saprospirales]	s2: f__Acetobacteraceae	r3: g__Arthrobacter	
v: c__Gemmatimonadetes	u1: o__Bacteroidales	t2: f__Campylobacteraceae	s3: g__Nocardioideae	
w: c__Alphaproteobacteria	v1: o__Flavobacteriales	u2: f__Pasteurellaceae	t3: g__Streptomyces	
x: c__Betaproteobacteria	w1: o__Sphingobacteriales	v2: f__Micrococcaceae	u3: g__Planotetrastroma	
y: c__Deltaproteobacteria	x1: o__Chlamydiales	w2: f__Nocardioideae	v3: g__Bifidobacterium	