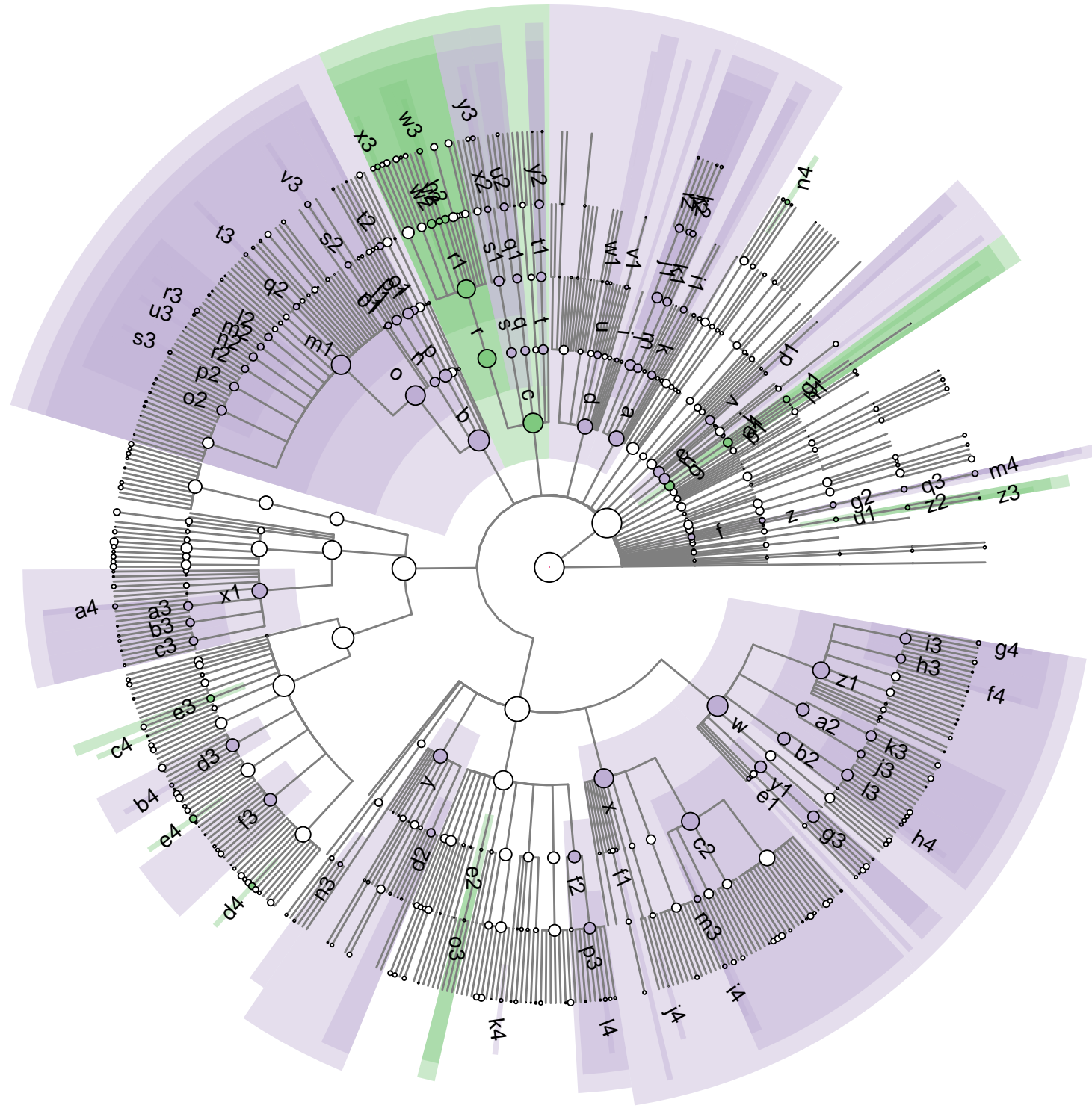


## Groups

■ AF  
■ FB



## Taxa

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