

IBS_P1_metatranscriptomes_family_TAIL: rank matrix & stability

	1	1	4	6	14	8	11	9	5	
Bacteroidales_incertae_sedis	1	1	4	6	14	8	11	9	5	59.8
Opitutaceae	18	17	19	17	1	1	22	21	17	40.4
uc_Rhodospirillales	30	30	30	30	30	30	9	7	2	59.8
Planctomycetaceae	20	19	1	19	23	19	6	22	19	19.9
Comamonadaceae	13	10	13	2	5	12	16	14	1	41.3
Streptococcaceae	22	22	2	22	9	22	7	24	22	10.5
uc_Entomoplasmatales	28	2	27	28	27	27	28	29	28	33.1
uc_Desulfovibrionales	27	27	26	27	3	2	27	28	27	34.7
Rhodobacteraceae	21	20	7	20	2	20	23	4	20	10.2
Alphaproteobacteria_incertae_sedis	9	4	9	5	13	7	2	1	4	49.9
Aeromonadaceae	2	3	8	4	12	6	10	8	3	52.0
Carnobacteriaceae	11	8	3	10	18	10	3	2	9	41.3
Parvularculaceae	19	18	20	18	8	18	5	3	18	33.9
uc_Caulobacteriales	26	26	25	26	11	26	1	27	26	16.6
uc_Flavobacteriales	29	28	28	3	28	28	29	6	29	11.1
Leuconostocaceae	17	16	18	16	4	5	21	20	16	47.9
Burkholderiaceae	6	7	11	9	17	4	14	12	8	43.1
Lactobacillaceae	3	15	17	15	22	17	20	19	15	50.9
Shewanellaceae	4	21	21	21	24	21	24	23	21	58.6
Enterococcaceae	15	12	6	12	20	14	4	16	12	33.9
Bacteroidetes_incertae_sedis	5	5	5	7	15	9	12	10	6	63.4
Synergistaceae	23	23	22	23	25	23	25	5	23	41.3
Bartonellaceae	10	6	10	8	16	3	13	11	7	40.4
uc_Archaea	25	25	24	25	26	25	8	26	25	46.9
Clostridiales_incertae_sedis_XI	12	9	12	1	19	11	15	13	10	36.2
Entomoplasmataceae	7	13	15	13	21	15	18	17	13	55.2
uc_Mollicutes	8	29	29	29	29	29	30	30	30	67.1
Deferribacteraceae	14	11	14	11	6	13	17	15	11	56.3
Fusobacteriaceae	16	14	16	14	7	16	19	18	14	57.5
Syntrophomonadaceae	24	24	23	24	10	24	26	25	24	53.1
	t.P1D1	t.P1D3	t.P1D5	t.P1D7	t.P1D14	t.P1D21	t.P1D28	t.P1D37	t.P1D42	Rank Stability Index (in %)