

IBS_P1_metatranscriptomes_family_CORE: rank matrix & stability

	t.P1D1	t.P1D3	t.P1D5	t.P1D7	t.P1D14	t.P1D21	t.P1D28	t.P1D37	t.P1D42	Rank Stability Index (in %)
Lachnospiraceae	1	1	1	2	5	5	1	1	2	86.3
Bacteroidaceae	4	5	3	1	1	4	2	2	3	83.4
Ruminococcaceae	3	2	5	4	3	1	4	3	5	79.3
uc_Clostridiales	5	3	2	5	4	3	3	4	7	82.0
uc_Bacteria	7	6	4	3	2	6	5	5	1	79.3
Prevotellaceae	2	4	8	6	6	2	7	6	9	70.2
uc_Bacteroidales	9	7	7	9	8	7	10	11	4	75.3
Porphyromonadaceae	6	10	10	7	10	9	9	8	6	79.3
uc_Firmicutes	8	8	6	8	7	8	11	12	12	84.8
uc_Alphaproteobacteria	12	20	18	13	16	22	6	7	8	47.6
uc_Proteobacteria	16	15	13	15	11	14	8	9	10	71.4
uc_Bacteroidetes	14	9	12	12	13	13	12	14	11	77.9
Clostridiaceae	22	19	9	11	18	19	15	10	14	53.4
Puniceicoccaceae	13	12	23	17	9	18	18	16	17	51.4
Alcaligenaceae	15	16	19	10	15	10	20	23	19	49.5
uc_Clostridia	18	18	14	14	12	16	14	15	16	79.3
Rikenellaceae	11	13	15	16	23	12	16	17	15	59.7
Erysipelotrichaceae	10	11	11	18	26	23	17	13	20	53.4
Eubacteriaceae	23	26	24	19	14	11	21	20	22	58.6
uc_Burkholderiales	27	31	32	25	22	17	24	25	13	49.5
Peptostreptococcaceae	21	17	16	21	20	26	13	19	27	45.8
uc_Bacilli	17	14	17	22	24	30	19	18	31	45.8
uc_Betaproteobacteria	26	24	31	32	21	15	22	24	18	47.6
Veillonellaceae	20	28	21	24	17	20	32	27	25	43.1
uc_Opitutae	28	27	29	23	19	21	26	31	32	64.2
Staphylococcaceae	25	23	25	31	28	25	25	21	23	69.0
Peptococcaceae	19	21	28	26	27	24	30	22	30	52.4
Clostridiales_incertae_sedis_XIV	29	22	22	20	32	32	28	30	21	53.4
uc_Bacillales	24	30	20	28	30	28	27	26	28	57.5
Desulfovibrionaceae	30	25	27	27	25	29	29	32	26	69.0
uc_Sphingobacteriales	32	32	30	29	29	31	23	28	29	72.7
uc_Actinomycetales	31	29	26	30	31	27	31	29	24	65.4