

IBS_P2 Metatranscriptores_CORE: rank matrix & stability

Bacteroidaceae	2	6	1	1	1	3	7	5	2	4	7	2	4	4	63.6
uc_Bacteria	1	2	6	5	4	2	2	2	8	1	1	1	1	5	69.5
uc_Alphaproteobacteria	3	1	12	15	7	5	1	1	7	3	6	5	3	15	42.2
Lachnospiraceae	4	4	4	2	2	1	4	4	1	5	5	6	2	2	78.0
uc_Proteobacteria	5	3	13	14	9	4	3	3	10	2	3	8	5	9	46.6
uc_Clostridiales	7	5	8	3	5	6	6	6	4	7	2	3	7	3	63.6
Ruminococcaceae	6	7	5	12	6	7	5	9	5	11	8	7	6	11	53.7
Clostridiaceae	12	15	20	13	14	14	9	17	15	6	4	4	14	1	37.5
uc_Bacteroidales	9	13	3	6	3	8	14	8	9	9	15	9	11	10	45.8
Porphyromonadaceae	10	9	2	10	15	13	10	7	6	14	14	15	12	17	50.5
Rikenellaceae	8	8	9	17	18	10	8	10	3	12	9	10	10	16	49.7
uc_Bacteroidetes	15	10	7	8	8	12	15	12	12	13	12	11	13	12	70.5
uc_Firmicutes	14	12	11	11	10	11	11	13	14	10	10	12	8	8	78.0
Erysipelotrichaceae	13	14	16	4	12	9	16	11	16	8	13	14	9	14	36.2
Peptostreptococcaceae	11	11	21	7	13	16	12	15	17	19	11	13	16	6	36.2
Prevotellaceae	16	18	10	9	11	15	19	16	11	15	20	17	18	18	54.6
uc_Gammaproteobacteria	20	16	23	20	21	18	13	22	23	16	17	18	15	20	48.1
Alcaligenaceae	19	22	15	22	16	22	22	18	13	24	22	16	22	13	33.2
uc_Clostridia	17	17	14	21	20	21	18	19	18	17	16	23	19	19	65.5
Pasteurellaceae	23	23	24	16	22	24	23	20	20	22	21	20	23	7	52.9
Bifidobacteriaceae	18	20	17	18	17	17	17	14	19	18	23	19	17	22	63.6
uc_Bacilli	22	21	22	23	19	19	20	21	21	20	19	22	20	21	79.1
uc_Sphingobacteriales	21	19	19	24	24	20	21	23	24	21	18	24	24	24	68.5
Staphylococcaceae	24	24	18	19	23	23	24	24	22	23	24	21	21	23	74.7
	t_P2D1	t_P2D3.1	t_P2D3.2	t_P2D4	t_P2D5	t_P2D7	t_P2D14	t_P2D21	t_P2D27	t_P2D28	t_P2D35	t_P2D42	t_P2D49	t_P2D56	

Rank Stability Index (in %)