IBS_h_C1_metatranscriptomes_family: rank matrix & stability

uc_Bacteria	8	1	3	1	1	1	1	4	1
Lachnospiraceae	2	2	1	2	3	3	2	1	2
Ruminococcaceae	1	4	2	4	2	2	2 5	2	3
Bacteroidaceae	4	6	2 5	3	4	4	3	2	1 2 3 4
uc Clostridiales	5	3	4	5	5	5	4	5	5
Prevotellaceae	3	5	6	5 6	6	6	6	5 8	5
Prevolenaceae			0	0				9	9
uc_Firmicutes	9	7	8	8	8	8	7	7	7
Porphyromonadaceae	6	9	9	9	7	9	9	9	9
uc_Bacteroidales	7	13	7	10	10	10	11	6	10
uc Clostridia	14	8	11	12	11	11	15	11	11
Puniceicoccaceae	10	14	27	15	9	14	10	10	8
uc Alphaproteobacteria	19	15	45	7	23	7	8	18	23
uc Proteobacteria	17	12	14	11	15	12	12	14	17
Rikenellaceae	īí	22	16	18	12	21	21	13	18
Alcaligenaceae	12	11	12	17	17	16	18	15	14
	15		15		13			12	
uc_Bacteroidetes		21		14		15	16		13
- Victivallaceae	41	19	13	22	19	13	17	24	16
Clostridiaceae	18	10	22	16	31	17	13	21	21
Peptostreptococcaceae	30	20	24	13	27	19	14	22	12
Erysipelotrichaceae	16	25	17	21	18	20	22	16	15
Veillonellaceae	21	18	20	19	14	22	20	26	19
Verrucomicrobiaceae	99	24	10	29	16	28	19	30	45
Peptococcaceae	26	17	26	20	26	18	30	19	20
uc Burkholderiales	20	26	19	42	24	35	45	25	25
Aerococcaceae	13	58	48	37	69	67	71	17	62
uc Bacilli	38	28	28	24	30	24	25	20	27
	50	16	30		34				
Enterobacteriaceae	56			74		78	80	67	37
uc_Opitutae	92	23	38	58	20	36	35	54	24
Desulfovibrionaceae	28	27	29	47	25	23	54	32	30
Clostridiales_incertae_sedis_XIV	25	34	57	30	29	29	26	28	28
uc_Verrucomicrobia	97	33	25	62	22	38	24	55	31
Methanobacteriaceae	64	31	18	80	83	55	84	75	78
Coriobacteriaceae	27	43	43	31	21	39	53	47	22
Bifidobacteriaceae	23	38	35	38	33	42	32	27	26
uc Betaproteobacteria	39	41	23	36	51	41	38	29	29
Comamonadaceae	34	32	21	72	41	75	52	46	55
Thermoanaerobacteraceae	83	30	87	23	43	25	65	36	35
uc Deltaproteobacteria	32	29	32	33	45	48	46	94	33
								94 44	
uc_Gammaproteobacteria	40	37	94	28	65	26	28		44
uc_Bacillales	88	40	34	35	36	30	44	37	39
uc_Lactobacillales	91	42	37	44	40	49	67	23	53
Synergistaceae	31	55	85	41	28	47	36	50	41
uc_Actinomycetales	86	57	90	32	44	27	27	43	40
Clostridiales incertae sédis XIII	24	47	42	53	38	74	33	34	36
Staphylococcaceae	80	45	82	26	39	33	29	91	34
uc Archaea	87	95	91	25	63	34	31	52	38
uc Desulfovibrionales	33	36	46	43	46	44	94	38	47
Bacteroidales incertae sedis	22	70	51	68	72	52	75	31	66
	57	44	36	39	42	54	56	40	32
Eubacteriaceae	37			27	54	32	64	40 49	
Streptomycetaceae		93	84						91
	~	, C. C. D. 24	~		^{t,} C _Z O ₃			[¢] CZ _O 5	_^
	t (2/2)	~;		SV.	Q	Ŝ	Q.	Q.	Q
	C'	77	77	7	C [*]	77	77	C'	C'
	₩:		ب		₩.	Ų	ب	<	₩

Rank Stability Index (in %)