kwashiorkor_gen_h_h10B: rank matrix & stability

	kwasilioikoi_ge	II_II_IITOD. I alik II	iati ix & Stability
Prevotella	1	1	4
Faecalibacterium	5	3 2	1
Bifidobacterium	2	2	8 2
Ruminococcus	14	17	2
Bacteroides	7	4	$\bar{3}$
Lactobacillus	3	10	9
Eubacterium	22	27	3 9 5
Megasphaera	4	18	33
Streptococcus	8	6	11
Roseburia	24	20	6
Mitsuokella	11	8	10
Collinsella	9	5	14
Blautia	6	9	13
Alistipes	142	76	7
Dorea	10	76	16
Akkermansia	10	138	16
			21
Veillonella	12	14	21
Haemophilus	13	11	30
Clostridium	15	16	17
Coprococcus	30	30	15
Dialister	16	19	23
Parabacteroides	26	33	18
Sutterella	21	13	26
Catenibacterium	19	22	27
Butyricimonas	128	126	19
Éscherichia	28	12	63
Leuconostoc	58	35	20
Senegalemassilia	17	23	31
Lactococcus	102	15	71
Butyrivibrio	31	46	22
Neisseria	20	26	29
Acidaminococcus	27	21	29 37
Campylobacter	23	28	32
Rothia	33	66	24
Oscillibacter	36	40	25
Weissella	265	24	28
Mastadenovirus	18	211	237
Anaerostipes	29	25	58
Enterococcus	32	29	44
Helicobacter	25	110	40
Subdoligranulum	55	37	34
Peptoclostridium	38	57	35
Gordonibacter	37	32	50
Holdemania	59	63	36
Pseudoflavonifractor	51	43	38
Odoribacter	35	38	43
Olsenella	49	36	47
Klebsiella	197	31	147
Barnesiella	43	34	73
Aggregatibacter	43 39	39	191
Aggregatibacter			
	\circ	8	8

Rank Stability Index (in %)