kwashiorkor\_gen\_h26B: rank matrix & stability

	KWWGS	,, ,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,	_9~''_''	LOD. Tui	ik illaci	in a sta	Dilicy
Prevotella	1	1	1	2	1	1	1
Faecalibacterium	2	4	2	6	2	2	9
Eubacterium	14	5 7	4	3	2 5 4	3 5	9 5 6
Bacteroides	4	7	6	ĭ	ă	5	6
Streptococcus	Ž	3	3	8	3	8	3
Bifidobacterium	3	8	5	16	7	4	3 4
Lactobacillus	6	29	28	41	6	6	2
							11
Veillonella	10	2	11	10	16	10	Ĭī
Blautia	5	9	9	4	12	9	8
Dorea	8	10	10	5	9	7	7
Roseburia	15	6	7	9	8	13	16
Ruminococcus	16	12	12	7	10	11	15
Collinsella	17	14	14	17	15	16	10
Phascolarctobacterium	253	45	8	12	42	67	25
Clostridium	13	13	17	15	13	15	14
Haemophilus	12	11	25	14	30	19	20
Helicobacter	188	204	27	59	11	12	94
Sutterella	9	19	13	11	14	25	43
Coprococcus	24	21	18	19	22	14	13
Escherichia	11	22	15	13	35	24	219
Anaerostipes	18	18	21	18	18	20	21
Senegalemassilia	23	16	16	22	20	28	17
Seriegalemassilla	126	53	154	120	46	140	12
Catenibacterium		20	37		46 21		18
Dialister	20			26		18	
Campylobacter	122	31	48	38	19	17	19
Parabacteroides	21	24	26	25	28	27	22
Acidaminococcus	22	25	45	33	26	23	24
Brachyspira	115	144	147	109	17	22	41
Akkermansia	93	15	22	62	27	34	170
Alistipes	95	35	19	21	71	54	47
Neisseria	30	50	53	20	222	26	27
Subdoligranulum	25	30	24	23	24	31	42
Butyrivibrio	61	26	31	24	25	29	29
Weissella	34	306	23	306	307	64	23
Mastadenovirus	217	228	225	206	208	21	250
Enterococcus	32	27	20	159	51	43	38
Megasphaera	40	17	118	208	81	68	39
Actinomyces	42	23	42	35	90	41	32
Peptoclostridium	31	34	34	42	29	39	33
Megamonas	219	111	106	70	23	30	251
Oscillibacter	28	33	41	31	31	33	37
Rothia	26 26	33 38	40	54	268		31
						292	67
Aggregatibacter	91	28	130	30	93	36	
Odoribacter	37	48	244	228	39	268	28
Hallella	186	203	205	180	67	71	26
Alloprevotella	99	32	112	94	100	153	30
Salmonella	19	279	33	270	271	69	285
Barnesiella	111	40	32	55	32	42	45
Anaerotruncus	48	37	137	99	40	32	49
Klebsiella	29	44	30	188	190	37	160
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Rank Stability Index (in %