diets_h_pl_9_pre: rank matrix & stability

	diets_ii_pi_9_pre. rank matrix & stability		
p Bacteroidetes;*;g Bacteroides	1	1	1
p Firmicutes;*;g	2 6 5 4 3 7 9	2	2
pProteobacteria;*;g	6	4	3 4 6 13 7 9
p_Firmicutes;*;g	5	3 5	4
p Firmicutes;*;g Faecalibacterium	4	5	6
p Bacteroidetes;*;g	3	14	13
p Firmicutes;*;g	7	6	7
p Firmicutes;*;g Blautia	9	9	9
p Proteobacteria;*;g Sutterella	12	6 9 8 7	12
p_Firmicutes;*;g_Lachnospira	11	7	18
p Actinobacteria;*;g Bifidobacterium	14	16	5
p Firmicutes;*;Other	15	11	10
pFirmicutes;*;gRuminococcus	16	10	15
p_Firmicutes;*;g_Coprococcus	21	12	11
pfirmicutes;*;Other	19	13	14
p Firmicutes:*:a Dialister	19 24	22	8
pVerrucomicrobia;*;gAkkermansia	8	44	8 24
p Proteobacteria;*;Other	8 10	20	27
p_Bacteroidetes;*;g_Parabacteroides	18	19	17
pFirmicutes;*;gOscillospira	13	18	30
p Bacteroidetes:*:g Butyricimonas	17	17	30 33
pBacteroidetes;*;gButyricimonas pProteobacteria;*;gDesulfovibrio	26	21	16
pFirmicutes;*;Other	20	28	28
p_Bacteroidetes;*;g_	22	23	21
pFirmicutes;*;gRoseburia	41	15	22
pFirmicutes;*;g	41 27	27	21 22 20
Other;*;Other	25 29	24	31
p_Firmicutes;*;g_	29	46	19
pFirmicutes;*;gMegasphaera	23	42	38
pFirmicutes;*;gPhascolarctobacterium	37	25	26
pFirmicutes;*;gDorea	30	29	32
p Firmicutes:*:a Clostridium	64	26	23
pProteobacteria;*;gTrabulsiella	34	36	23 29 41
pFirmicutes;*;gStreptococcus	28	45	41
p Firmicutes;*;Other	28 40	48	25
p_Proteobacteria;*;g_Bilophila	32	35	40
pFirmicutes;*;g[Ruminococcus]	32 36	30	34
p Bacteroidetes:*:a Odoribacter I	35	31	42
p_Actinobacteria;*;g_Collinsella	42	32	35
p_Proteobacteria;*;g_Serratia	31	95	95
pFirmicutes;*;gLeuconostoc	39	33	54
p_Firmicutes;*;g_	33	47	46
pFirmicutes; * ;gAnaerostipes	44	37	39
pProteobacteria;*;g	54	34	39 48 47 45
pFirmicutes;*;g	45	38	47
pProteobacteria;*;gErwinia	68	39	45
p_Firmicutes;*;g_Turicibacter	57	74	36
p Firmicutes;*;g	76	79	36 37
p Firmicutes;*;g Enterococcus	38	70	83
p Proteobacteria;*;Other	46	43	68

Rank Stability Index (in %