

antibiotics_patE_Cp2_CPW_TAIL: elm correlation matrix

elm1: *_Bilophila	1.00	1.00	0.99	-0.22	1.00	-0.18	-0.18	-0.27	1.00	-0.18	1.00	1.00	-0.18	1.00	-0.12
elm2: *_	1.00	1.00	0.99	-0.21	1.00	-0.17	-0.17	-0.26	1.00	-0.17	1.00	1.00	-0.17	1.00	-0.17
elm3: *_Holdemania	0.99	0.99	1.00	-0.25	0.99	-0.20	-0.20	-0.31	0.99	-0.20	0.99	0.99	-0.20	0.99	0.00
elm4: *_Coprobacillus	-0.22	-0.21	-0.25	1.00	-0.21	0.97	0.97	0.59	-0.21	-0.21	-0.21	-0.21	0.97	-0.21	-0.21
elm5: *_	1.00	1.00	0.99	-0.21	1.00	-0.17	-0.17	-0.26	1.00	-0.17	1.00	1.00	-0.17	1.00	-0.17
elm6: *_Veillonella	-0.18	-0.17	-0.20	0.97	-0.17	1.00	1.00	0.65	-0.17	-0.17	-0.17	-0.17	1.00	-0.17	-0.17
elm7: *_Alistipes	-0.18	-0.17	-0.20	0.97	-0.17	1.00	1.00	0.65	-0.17	-0.17	-0.17	-0.17	1.00	-0.17	-0.17
elm8: *_Acinetobacter	-0.27	-0.26	-0.31	0.59	-0.26	0.65	0.65	1.00	-0.26	-0.26	-0.26	-0.26	0.65	-0.26	-0.26
elm9: *_	1.00	1.00	0.99	-0.21	1.00	-0.17	-0.17	-0.26	1.00	-0.17	1.00	1.00	-0.17	1.00	-0.17
elm10: *_	-0.18	-0.17	-0.20	-0.21	-0.17	-0.17	-0.17	-0.26	-0.17	1.00	-0.17	-0.17	-0.17	-0.17	-0.17
elm11: *_[Eubacterium]	1.00	1.00	0.99	-0.21	1.00	-0.17	-0.17	-0.26	1.00	-0.17	1.00	1.00	-0.17	1.00	-0.17
elm12: *_Porphyromonas	1.00	1.00	0.99	-0.21	1.00	-0.17	-0.17	-0.26	1.00	-0.17	1.00	1.00	-0.17	1.00	-0.17
elm13: *_Bulleidia	-0.18	-0.17	-0.20	0.97	-0.17	1.00	1.00	0.65	-0.17	-0.17	-0.17	-0.17	1.00	-0.17	-0.17
elm14: *_Rikenellaceae;Other	1.00	1.00	0.99	-0.21	1.00	-0.17	-0.17	-0.26	1.00	-0.17	1.00	1.00	-0.17	1.00	-0.17
elm15: *_	-0.12	-0.17	0.00	-0.21	-0.17	-0.17	-0.17	-0.26	-0.17	-0.17	-0.17	-0.17	-0.17	-0.17	1.00
	elm1	elm2	elm3	elm4	elm5	elm6	elm7	elm8	elm9	elm10	elm11	elm12	elm13	elm14	elm15