

antibiotics_h_patD_pre_TAIL: elm correlation matrix

elm1: _*	1.00	0.97	-0.24	-0.24	-0.24	-0.26	-0.26	-0.26	-0.17	1.00	-0.17	-0.16	-0.17	1.00	-0.16
elm2: *_Erysipelotrichaceae;Other	0.97	1.00	-0.30	-0.30	-0.30	-0.33	-0.33	-0.33	-0.21	0.97	-0.21	-0.21	-0.21	0.97	-0.21
elm3: _*	-0.24	-0.30	1.00	-0.08	0.73	-0.37	-0.37	0.93	-0.24	-0.24	-0.24	0.88	0.32	-0.24	0.88
elm4: *_Betaproteobacteria;Other;Other;Other	-0.24	-0.30	-0.08	1.00	0.19	-0.37	0.50	0.06	-0.24	-0.24	-0.24	-0.24	0.32	-0.24	-0.24
elm5: *_Coprobacillus	-0.24	-0.30	0.73	0.19	1.00	-0.37	0.06	0.50	-0.24	-0.24	-0.24	0.88	-0.24	-0.24	0.88
elm6: _*	-0.26	-0.33	-0.37	-0.37	-0.37	1.00	0.30	-0.40	0.65	-0.26	0.65	-0.26	-0.26	-0.26	-0.26
elm7: *_Actinomyces	-0.26	-0.33	-0.37	0.50	0.06	0.30	1.00	-0.40	0.65	-0.26	0.65	-0.26	-0.26	-0.26	-0.26
elm8: *_Alistipes	-0.26	-0.33	0.93	0.06	0.50	-0.40	-0.40	1.00	-0.26	-0.26	-0.26	0.65	0.65	-0.26	0.65
elm9: *_Lactococcus	-0.17	-0.21	-0.24	-0.24	-0.24	0.65	0.65	-0.26	1.00	-0.17	1.00	-0.17	-0.17	-0.17	-0.17
elm10: *_Weissella	1.00	0.97	-0.24	-0.24	-0.24	-0.26	-0.26	-0.26	-0.17	1.00	-0.17	-0.17	-0.17	1.00	-0.17
elm11: *_Rothia	-0.17	-0.21	-0.24	-0.24	-0.24	0.65	0.65	-0.26	1.00	-0.17	1.00	-0.17	-0.17	-0.17	-0.17
elm12: *_Anaerotruncus	-0.16	-0.21	0.88	-0.24	0.88	-0.26	-0.26	0.65	-0.17	-0.17	-0.17	1.00	-0.17	-0.17	1.00
elm13: _*	-0.17	-0.21	0.32	0.32	-0.24	-0.26	-0.26	0.65	-0.17	-0.17	-0.17	-0.17	1.00	-0.17	-0.17
elm14: *_Firmicutes;Other;Other;Other;Other	1.00	0.97	-0.24	-0.24	-0.24	-0.26	-0.26	-0.26	-0.17	1.00	-0.17	-0.17	-0.17	1.00	-0.17
elm15: *_Holdemania	-0.16	-0.21	0.88	-0.24	0.88	-0.26	-0.26	0.65	-0.17	-0.17	-0.17	1.00	-0.17	-0.17	1.00
	elm1	elm2	elm3	elm4	elm5	elm6	elm7	elm8	elm9	elm10	elm11	elm12	elm13	elm14	elm15