

IBS_h_C1_metatranscriptomes_family_TAIL: elm correlation matrix

elm1:Methanobacteriaceae	1.00	-0.21	0.92	-0.25	0.99	-0.20	0.04	-0.16	-0.18	-0.05	0.72	0.04	-0.14	-0.15	-0.17
elm2:Bacteroidales_incertae_sedis	-0.21	1.00	-0.26	0.63	-0.18	-0.21	-0.18	0.26	-0.22	-0.26	-0.27	-0.18	-0.34	0.92	-0.28
elm3:Lactobacillaceae	0.92	-0.26	1.00	-0.25	0.94	-0.15	-0.19	-0.21	-0.15	-0.08	0.66	-0.19	-0.36	-0.19	-0.25
elm4:Microbacteriaceae	-0.25	0.63	-0.25	1.00	-0.22	0.53	-0.22	-0.24	-0.21	-0.18	-0.33	-0.22	-0.42	0.72	0.32
elm5:Actinomycetaceae	0.99	-0.18	0.94	-0.22	1.00	-0.18	-0.13	-0.14	-0.15	-0.18	0.75	-0.12	-0.24	-0.12	-0.23
elm6:uc_Rhizobiales	-0.20	-0.21	-0.15	0.53	-0.18	1.00	-0.18	-0.20	-0.16	-0.11	-0.27	-0.18	-0.36	-0.18	0.82
elm7:Pseudomonadaceae	0.04	-0.18	-0.19	-0.22	-0.13	-0.18	1.00	-0.14	-0.15	0.83	-0.19	1.00	0.64	-0.13	0.28
elm8:Burkholderiales_incertae_sedis	-0.16	0.26	-0.21	-0.24	-0.14	-0.20	-0.14	1.00	-0.17	-0.20	-0.13	-0.14	-0.22	-0.14	-0.25
elm9:Leuconostocaceae	-0.18	-0.22	-0.15	-0.21	-0.15	-0.16	-0.15	-0.17	1.00	-0.08	-0.23	-0.15	0.48	-0.15	-0.28
elm10:Micrococcaceae	-0.05	-0.26	-0.08	-0.18	-0.18	-0.11	0.83	-0.20	-0.08	1.00	-0.27	0.83	0.44	-0.18	0.12
elm11:Bacillaceae	0.72	-0.27	0.66	-0.33	0.75	-0.27	-0.19	-0.13	-0.23	-0.27	1.00	-0.19	-0.01	-0.19	-0.34
elm12:Vibrionaceae	0.04	-0.18	-0.19	-0.22	-0.12	-0.18	1.00	-0.14	-0.15	0.83	-0.19	1.00	0.64	-0.12	0.28
elm13:Flavobacteriaceae	-0.14	-0.34	-0.36	-0.42	-0.24	-0.36	0.64	-0.22	0.48	0.44	-0.01	0.64	1.00	-0.24	-0.05
elm14:Rhizobiaceae	-0.15	0.92	-0.19	0.72	-0.12	-0.18	-0.13	-0.14	-0.15	-0.18	-0.19	-0.12	-0.24	1.00	-0.23
elm15:uc_Rhodospirillales	-0.17	-0.28	-0.25	0.32	-0.23	0.82	0.28	-0.25	-0.28	0.12	-0.34	0.28	-0.05	-0.23	1.00
	elm1	elm2	elm3	elm4	elm5	elm6	elm7	elm8	elm9	elm10	elm11	elm12	elm13	elm14	elm15