antibiotics_patD_Cp2_WPC_TAIL: elm correlation matrix

	elm1	elm2	elm3	elm4	elm5	elm6	elm7	elm8	elm9	elm10	elm11	elm12	elm13	elm14	elm15
elm1:_*_[Eubacterium]	1.00	1.00	1.00	-0.26	1.00	-0.19	1.00	-0.24	1.00	-0.17	1.00	-0.26	-0.25	1.00	-0.25
elm2:_*_Odoribacter	1.00	1.00	1.00	-0.26	1.00	-0.19	1.00	-0.24	1.00	-0.17	1.00	-0.26	-0.26	1.00	-0.26
elm3:_*_Bacteroidales;Other;Other	1.00	1.00	1.00	-0.27	1.00	-0.19	1.00	-0.25	1.00	-0.17	1.00	-0.27	-0.27	1.00	-0.27
elm4:_*_Clostridium	-0.26	-0.26	-0.27	1.00	-0.26	-0.30	-0.26	-0.37	-0.26	0.65	-0.26	-0.40	-0.40	-0.26	-0.40
elm5:_*_	1.00	1.00	1.00	-0.26	1.00	-0.19	1.00	-0.24	1.00	-0.17	1.00	-0.26	-0.26	1.00	-0.26
elm6:_*_	-0.19	-0.19	-0.19	-0.30	-0.19	1.00	-0.19	0.29	-0.19	-0.19	-0.19	0.62	-0.30	-0.19	0.62
elm7:_*_	1.00	1.00	1.00	-0.26	1.00	-0.19	1.00	-0.24	1.00	-0.17	1.00	-0.26	-0.26	1.00	-0.26
elm8:_*_Actinomyces	-0.24	-0.24	-0.25	-0.37	-0.24	0.29	-0.24	1.00	-0.24	-0.24	-0.24	0.93	0.50	-0.24	0.06
elm9:_*_Erysipelotrichaceae;Other	1.00	1.00	1.00	-0.26	1.00	-0.19	1.00	-0.24	1.00	-0.17	1.00	-0.26	-0.26	1.00	-0.26
elm10:_*_Mogibacterium	-0.17	-0.17	-0.17	0.65	-0.17	-0.19	-0.17	-0.24	-0.17	1.00	-0.17	-0.26	-0.26	-0.17	-0.26
:_*_Bacteria;Other;Other;Other;Other;Other	1.00	1.00	1.00	-0.26	1.00	-0.19	1.00	-0.24	1.00	-0.17	1.00	-0.26	-0.26	1.00	-0.26
elm12:_*_Weissella	-0.26	-0.26	-0.27	-0.40	-0.26	0.62	-0.26	0.93	-0.26	-0.26	-0.26	1.00	0.30	-0.26	0.30
elm13:_*_	-0.25	-0.26	-0.27	-0.40	-0.26	-0.30	-0.26	0.50	-0.26	-0.26	-0.26	0.30	1.00	-0.26	0.30
elm14:_*_Prevotella	1.00	1.00	1.00	-0.26	1.00	-0.19	1.00	-0.24	1.00	-0.17	1.00	-0.26	-0.26	1.00	-0.26
elm15:_*_Lactobacillus	-0.25	-0.26	-0.27	-0.40	-0.26	0.62	-0.26	0.06	-0.26	-0.26	-0.26	0.30	0.30	-0.26	1.00