antibiotics\_h\_patE\_pre\_TAIL: elm correlation matrix

·	<i>3</i> 11C1	DIO	رادع	_' '	pac	P	'	1 / \11			COII	CIG	CIOI		
elm1:_*_	1.00	-0.20	-0.25	-0.17	0.65	-0.26	-0.26	1.00	-0.17	-0.26	-0.17	-0.26	-0.17	-0.17	-0.17
elm2:_*_Bifidobacterium	-0.20	1.00	-0.17	-0.20	-0.31	-0.31	-0.31	-0.20	-0.20	0.61	-0.20	0.61	0.99	-0.20	-0.20
elm3:_*_Porphyromonas	-0.25	-0.17	1.00	0.76	-0.39	0.39	-0.39	-0.25	0.76	0.39	-0.25	0.39	-0.25	-0.25	-0.25
elm4:_*_1855D	-0.17	-0.20	0.76	1.00	-0.26	0.65	-0.26	-0.17	1.00	0.65	- <b>0.1</b> 7	0.65	- <b>0.1</b> 7	-0.17	- <b>0.1</b> 7
elm5:_*_Actinomyces	0.65	-0.31	-0.39	-0.26	1.00	-0.40	0.30	0.65	-0.26	-0.40	-0.26	-0.40	-0.26	-0.26	-0.26
elm6:_*_Rothia	-0.26	-0.31	0.39	0.65	-0.40	1.00	-0.40	-0.26	0.65	0.30	-0.26	0.30	-0.26	-0.26	-0.26
elm7:_*_	-0.26	-0.31	-0.39	-0.26	0.30	-0.40	1.00	-0.26	-0.26	-0.40	0.65	-0.40	-0.26	0.65	0.65
elm8:_*_Rhodoferax	1.00	-0.20	-0.25	- <b>0.1</b> 7	0.65	-0.26	-0.26	1.00	- <b>0.1</b> 7	-0.26	-0.17	-0.26	-0.17	- <mark>0.1</mark> 7	-0.17
elm9:_*_Enterobacteriaceae;Other	-0.17	-0.20	0.76	1.00	-0.26	0.65	-0.26	-0.17	1.00	0.65	-0.17	0.65	- <b>0.1</b> 7	- <mark>0.1</mark> 7	-0.17
elm10:_*_SMB53	-0.26	0.61	0.39	0.65	-0.40	0.30	-0.40	-0.26	0.65	1.00	-0.26	1.00	0.65	-0.26	-0.26
elm11:_*_[Prevotella]	-0.17	-0.20	-0.25	-0.17	-0.26	-0.26	0.65	-0.17	- <b>0.1</b> 7	-0.26	1.00	-0.26	- <b>0.1</b> 7	1.00	1.00
elm12:_*_Paraprevotella	-0.26	0.61	0.39	0.65	-0.40	0.30	-0.40	-0.26	0.65	1.00	-0.26	1.00	0.65	-0.26	-0.26
elm13:_*_Lachnobacterium	-0.17	0.99	-0.25	-0.17	-0.26	-0.26	-0.26	-0.17	-0.17	0.65	-0.17	0.65	1.00	-0.17	-0.17
elm14:_*_Clostridium	-0.17	-0.20	-0.25	-0.17	-0.26	-0.26	0.65	-0.17	-0.17	-0.26	1.00	-0.26	-0.17	1.00	1.00
elm15:_*_	-0.17	-0.20	-0.25	-0.17	-0.26	-0.26	0.65	-0.17	-0.17	-0.26	1.00	-0.26	-0.17	1.00	1.00
	Pulys	Sulso	8/11/3	sup <sub>4</sub>	54/18	94/10	<4/1/9	84/12	64/10	oru <sub>10</sub>	Trups	?tu <sub>/o</sub>	elm13	eln14	elm <sub>15</sub>