Taylor parameters results

`cmplxcruncher v1.1rc12`

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1 counts.genus.individuals.hA

Metadata	V	β	R^2	V_{st}	β_{st}
h_A01	0.41 ± 0.06	0.840 ± 0.020	0.917	2.5 ± 0.8	1.0 ± 0.9
h_A02	0.39 ± 0.06	0.845 ± 0.021	0.931	2.4 ± 0.8	1.2 ± 1.0
h_A03	0.164 ± 0.022	0.784 ± 0.018	0.925	-0.65 ± 0.28	-1.6 ± 0.9
h_A04	0.235 ± 0.029	0.821 ± 0.016	0.945	0.3 ± 0.4	0.1 ± 0.7
h_A05	0.40 ± 0.06	0.845 ± 0.018	0.933	2.4 ± 0.7	1.2 ± 0.9
h_A06	0.156 ± 0.021	0.803 ± 0.020	0.924	-0.75 ± 0.28	-0.7 ± 0.9
h_A07	0.213 ± 0.027	0.806 ± 0.018	0.939	-0.01 ± 0.35	-0.6 ± 0.8
h_A08	0.28 ± 0.04	0.832 ± 0.018	0.933	0.9 ± 0.5	0.6 ± 0.9
h_A09	0.29 ± 0.04	0.816 ± 0.019	0.930	1.0 ± 0.5	-0.1 ± 0.9
h_A10	0.179 ± 0.029	0.791 ± 0.024	0.906	-0.4 ± 0.4	-1.3 ± 1.1
E01	0.43 ± 0.06	0.849 ± 0.019	0.930	2.8 ± 0.8	1.4 ± 0.9
E02	0.246 ± 0.029	0.814 ± 0.017	0.933	0.4 ± 0.4	-0.2 ± 0.8
E03	0.185 ± 0.023	0.789 ± 0.017	0.933	-0.37 ± 0.30	-1.4 ± 0.8
E04	0.33 ± 0.04	0.841 ± 0.018	0.939	1.6 ± 0.6	1.1 ± 0.9
E05	0.224 ± 0.026	0.805 ± 0.016	0.930	0.14 ± 0.34	-0.6 ± 0.8
E06	0.29 ± 0.04	0.823 ± 0.019	0.929	1.1 ± 0.5	0.2 ± 0.9
E07	0.28 ± 0.04	0.824 ± 0.019	0.936	0.9 ± 0.5	0.3 ± 0.9
E08	0.139 ± 0.020	0.777 ± 0.021	0.894	-0.98 ± 0.27	-2.0 ± 1.0
E09	0.186 ± 0.022	0.796 ± 0.016	0.944	-0.35 ± 0.28	-1.0 ± 0.7
E10	0.28 ± 0.04	0.819 ± 0.017	0.923	0.9 ± 0.5	0.0 ± 0.8
I01	0.46 ± 0.07	0.860 ± 0.019	0.937	3.3 ± 1.0	2.0 ± 0.9
I02	0.38 ± 0.06	0.841 ± 0.019	0.925	2.2 ± 0.7	1.0 ± 0.9
I03	0.27 ± 0.04	0.816 ± 0.018	0.936	0.7 ± 0.5	-0.1 ± 0.8
I04	0.21 ± 0.05	0.832 ± 0.032	0.785	0.0 ± 0.6	0.6 ± 1.5
I05	0.34 ± 0.05	0.837 ± 0.019	0.928	1.7 ± 0.6	0.9 ± 0.9
I06	0.244 ± 0.033	0.810 ± 0.018	0.937	0.4 ± 0.4	-0.4 ± 0.8
I07	0.54 ± 0.08	0.871 ± 0.019	0.930	4.3 ± 1.0	2.5 ± 0.9
I08	0.240 ± 0.032	0.812 ± 0.018	0.938	0.3 ± 0.4	-0.3 ± 0.9
I09	0.38 ± 0.06	0.839 ± 0.020	0.932	2.2 ± 0.7	0.9 ± 0.9
I10	0.215 ± 0.029	0.800 ± 0.019	0.927	0.0 ± 0.4	-0.9 ± 0.9

Table 1: Taylor parameters for the dataset counts.genus.individuals.hA. The healthy population is described by $\bar{V} = 0.21 \pm 0.08, \bar{\beta} = 0.819 \pm 0.021$.