

Effects of tebuconazole application on soil microbiota and enzymes

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Table S1. Effect of tebuconazole on population numbers of microorganisms, 10ⁿ cfu kg⁻¹ soil d.m.

Object	Organotrophic bacteria 10 ¹⁰ cfu	Actinobacteria 10 ¹⁰ cfu	Fungi 10 ⁸ cfu
C	1.599±0.077 ^e	0.731±0.019 ^b	0.175±0.008 ^e
T1	1.670±0.006 ^d	0.987±0.011 ^a	0.327±0.006 ^d
T2	2.052±0.017 ^c	0.690±0.030 ^c	0.375±0.018 ^c
T3	2.242±0.018 ^b	0.630±0.027 ^{cd}	0.479±0.003 ^b
T4	2.575±0.024 ^a	0.630±0.028 ^{cd}	0.606±0.012 ^a
\bar{x}	2.028	0.734	0.392
r	0.936 [*]	-0.617	0.912

Tebuconazole doses in mg kg⁻¹: C – control soil, T1 – 0.01 mg, T2 – 0.1 mg, T3 – 0.5 mg, T4 – 1.0 mg; r – simple Pearson's correlation coefficient significant at $*p<0.05$, $n=20$; \bar{x} – arithmetic mean; \pm – standard deviation. Homogeneous groups designated with the same letters (a-e) were calculated separately for each group of microorganisms.

Table S2. Values of Shannon-Wiener index and Simpson index computed based on all OTU data.

Object	Taxa				
	Phylum	Class	Order	Family	Genus
Shannon-Wiener index					
C	1.718±0.006 b	2.419±0.007 b	2.928±0.005 b	3.025±0.006 c	1.835±0.01 2 ^a
T1	1.641±0.014 d	2.377±0.004 c	2.933±0.005 b	2.973±0.005 d	1.715±0.00 7 ^b
T2	1.802±0.004 a	2.552±0.008 a	3.012±0.007 a	3.063±0.006 b	1.825±0.00 6 ^a
T3	1.788±0.009 a	2.566±0.003 a	3.032±0.005 a	3.106±0.003 a	1.851±0.01 3 ^a
T4	1.690±0.007 c	2.419±0.006 b	2.902±0.006 c	3.034±0.008 c	1.828±0.01 1 ^a
\bar{x}	1.728	2.467	2.961	3.040	1.811
r	0.213	0.347	0.130	0.486	0.354
Simpson index					
C	0.737±0.006 ab	0.688±0.006 bc	0.748±0.010 c	0.655±0.006 a	0.355±0.01 6 ^a
T1	0.710±0.008 b	0.661±0.010 c	0.798±0.007 a	0.610±0.006 b	0.263±0.00 9 ^c
T2	0.764±0.006 a	0.718±0.013 a	0.763±0.006 bc	0.628±0.010 ab	0.330±0.00 7 ^a
T3	0.757±0.004 a	0.716±0.009 a	0.794±0.007 ab	0.623±0.008 b	0.298±0.01 2 ^b
T4	0.731±0.009 ab	0.700±0.008 ab	0.764±0.006 bc	0.634±0.005 ab	0.347±0.00 8 ^a
\bar{x}	0.740	0.697	0.773	0.630	0.319
r	0.257	0.534	0.205	-0.277	0.079

Tebuconazole doses in mg kg⁻¹: C – control soil, T1 – 0.01 mg, T2 – 0.1 mg, T3 – 0.5 mg, T4 – 1.0 mg; r – simple Pearson's correlation coefficient significant at * p <0.05, n=20; \bar{x} – arithmetic mean; \pm – standard deviation. Homogeneous groups designated with the same letters (a-d) were calculated separately for each group of microorganisms.

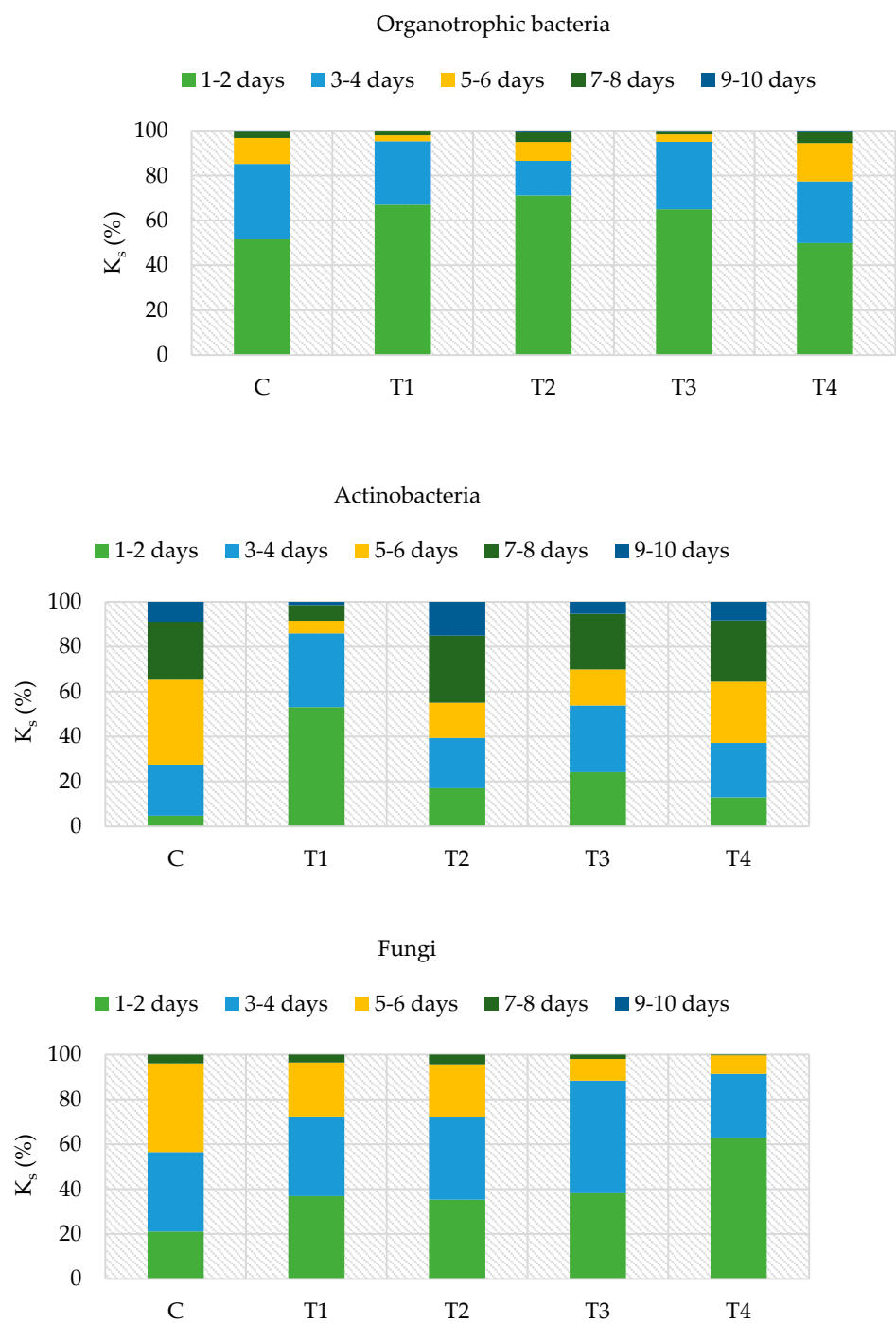


Figure S1. Increase in the abundance of microorganisms in various time intervals, in % (K_s). Tebuconazole doses in mg kg^{-1} : C – control soil, T1 – 0.01 mg, T2 – 0.1 mg, T3 – 0.5 mg, T4 – 1.0 mg.