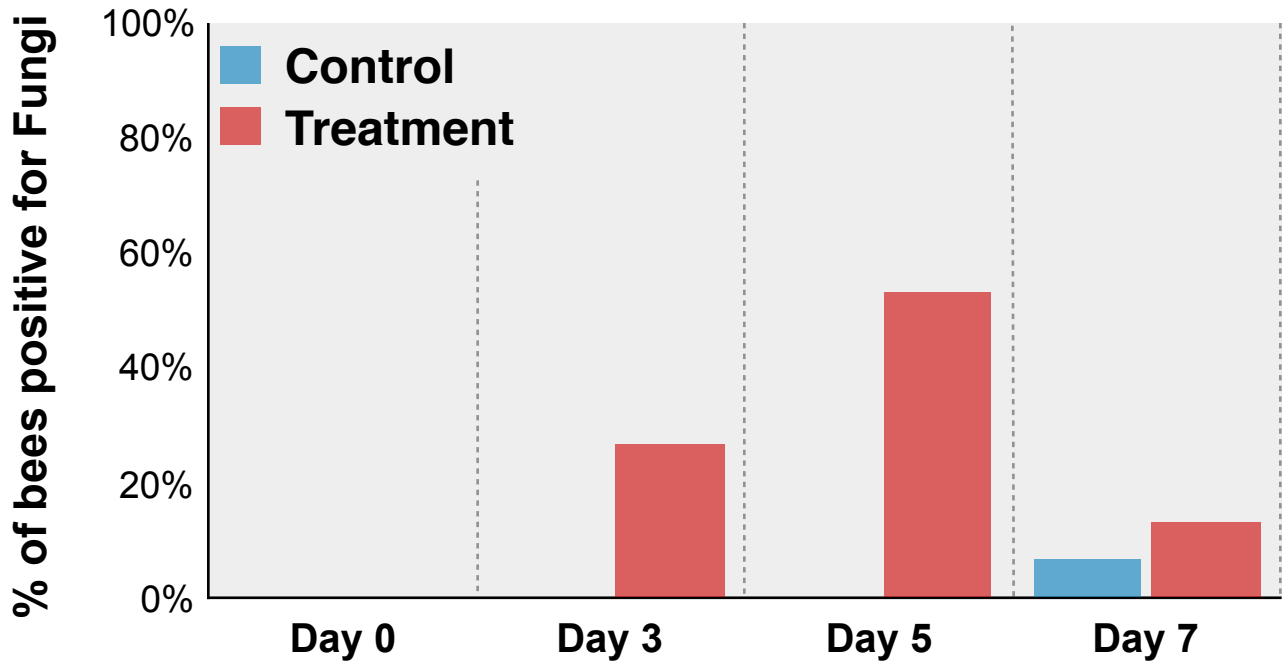


A

PCR with universal fungal primers

**B**

Clones	Sample	Species*	Kingdom
T2 D3 1	Treatment bee 2 day 3	<i>Metschnikowia koreensis</i>	Fungi
T2 D3 2	Treatment bee 2 day 3	<i>Metschnikowia koreensis</i>	Fungi
T2 D3 3	Treatment bee 2 day 3	<i>Metschnikowia koreensis</i>	Fungi
T2 D3 1	Treatment bee 2 day 3	<i>Metschnikowia koreensis</i>	Fungi
T6 D3 1	Treatment bee 6 day 3	<i>Lachancea thermotolerans</i>	Fungi
T3 D5 1	Treatment bee 3 day 5	<i>Starmerella meliponinorum</i>	Fungi
T10 D5 1	Treatment bee 10 day 5	<i>Nothoscordum bivalve</i>	Plantae
T10-D5 2	Treatment bee 10 day 5	<i>Nothoscordum texanum</i>	Plantae
T13 D5 1	Treatment bee 13 day 5	<i>Metschnikowia koreensis</i>	Fungi
T1 D7 1	Treatment bee 1 day 7	<i>Elaeagnus macrophylla</i>	Plantae
C1 D7 1	Control bee 1 day 7	<i>Nothoscordum bivalve</i>	Plantae
C1 D7 2	Control bee 1 day 7	<i>Nothoscordum bivalve</i>	Plantae
C1 D7 3	Control bee 1 day 7	<i>Nothoscordum bivalve</i>	Plantae
C1 D7 4	Control bee 1 day 7	<i>Nothoscordum bivalve</i>	Plantae
C1 D7 5	Control bee 1 day 7	<i>Nothoscordum bivalve</i>	Plantae
C1 D7 6	Control bee 1 day 7	<i>Nothoscordum bivalve</i>	Plantae
C1 D7 7	Control bee 1 day 7	<i>Nothoscordum bivalve</i>	Plantae

* signifies >97% sequence identity to the given species