

Supplementary Material. Table S11.- Bacterial OTUs with significant differential abundances between healthy bulk (HB) and diseased bulk (DB) soils. OTUs with a positive LFC are highlighted on a yellow background while OTUs with a negative LFC are highlighted on a blue background. LFC: log2-fold changes; SE: standard error; *p*-values were adjusted with the False Discovery Rate (FDR) method; unclas.: unclassified

	LFC	LFC (SE)	Wald statistic	<i>p</i> -value	FDR	Phylum	Kingdom	Class	Order	Family	Genus
Otu01429	5,306	1,179	4,501	6,77E-06	0,027	Bacteria	Bacteria_unclas.	Bacteria_unclas.	Bacteria_unclas.	Bacteria_unclas.	Bacteria_unclas.
Otu00893	3,841	0,786	4,885	1,04E-06	0,008	Bacteria	Patescibacteria	Saccharimonadia	Saccharimonadales	Saccharimonadales_unclas.	Saccharimonadales_unclas.
Otu00097	1,305	0,242	5,400	6,67E-08	0,001	Bacteria	Chloroflexi	Ktedonobacteria	C0119	C0119_fa	C0119_ge
Otu01273	-5,891	1,310	-4,496	6,93E-06	0,027	Bacteria	Chloroflexi	Anaerolineae	Anaerolineales	Anaerolineaceae	uncultured