

Additional Table S3 - Average nucleotide identities (ANI) between *Williamsia* sp. ARP1 and nearest actinomycete genomes.

Species	<i>Williamsia</i> sp. ARP1	<i>Williamsia</i> sp. D3	<i>Gordonia bronchialis</i> DSM 43234	<i>Gordonia polyisoprenivorans</i> VH2
<i>Williamsia</i> sp. ARP1	-	72.37	72.95	72.71
<i>Williamsia</i> sp. D3	83.97	-	71.60	71.50
<i>Gordonia bronchialis</i> DSM 43234T	84.34	84.21	-	84.84
<i>Gordonia polyisoprenivorans</i> VH2T	84.17	84.19	84.84	-

Upper right, ANIb score calculated with BLAST; lower left, ANIm calculated with mummer.