

Additional Table S2 - Orthologous gene comparison of *Williamsia* sp. ARP1 and three other 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	4,438	2,768	2,726	2,818
<i>Williamsia</i> sp. D3	75.53	4,930	2,778	2,836
<i>Gordonia bronchialis</i> DSM 43234T	75.17	74.04	4,616	3,282
<i>Gordonia polyisoprenivorans</i> VH2T	74.84	73.89	77.07	4,945

Upper right, numbers of found orthologous genes; lower left, average nucleotide identities of orthologous genes. Bold numbers indicate the numbers of genes for each genome.