

Species	Similarity	Genes	antiSMASH	ClusterFinder
Mycobacterium tuberculosis str. Haarlem/NITR202	-		<input type="checkbox"/>	<input type="checkbox"/>
Mycobacterium tuberculosis RGTB423	80%		<input type="checkbox"/>	<input type="checkbox"/>
Mycobacterium intracellulare MOTT-64	68%		<input checked="" type="checkbox"/>	<input type="checkbox"/>
Mycobacterium avium subsp. hominissuis TH135	67%		<input checked="" type="checkbox"/>	<input type="checkbox"/>
Mycobacterium avium 104	64%		<input checked="" type="checkbox"/>	<input type="checkbox"/>
Mycobacterium tuberculosis CDC1551	63%		<input type="checkbox"/>	<input type="checkbox"/>
Mycobacterium indicus pranii MTCC 9506	61%		<input checked="" type="checkbox"/>	<input type="checkbox"/>
Mycobacterium sp. MOTT36Y	58%		<input checked="" type="checkbox"/>	<input type="checkbox"/>
Mycobacterium tuberculosis EAI5/NITR206	55%		<input type="checkbox"/>	<input type="checkbox"/>
Mycobacterium intracellulare MOTT-02	54%		<input checked="" type="checkbox"/>	<input type="checkbox"/>
Mycobacterium_kansasii_ATCC_12478	51%		<input type="checkbox"/>	<input checked="" type="checkbox"/>
		Class-specific          Redox          Regulatory          Group Transfer          Transport          Unknown		