

Shinella sumterensis

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Table 1: Complete list of names proposed in the current register list.

Proposed Taxon	Etymology	Description	Parent Taxon	Type	Registry URL
Species <i>Shinella sumterensis</i>	[sum.ter.en'sis] N.L. fem. adj. <i>sumterensis</i> , in recognition of Sumter National Forest, South Carolina, United States, the location from which the type strain was isolated	Cells are Gram-negative staining cocci arranged in long chains, facultatively anaerobic, motile, catalase and oxidase positive. Colonies are small, pinkish-white, glistening, smooth, with entire margins after growth for 24 h on levoglucosan isolation medium at 37°C. Growth was observed on arabinose, cellobiosan, fructose, galactose, glucose, <i>myo</i> -inositol, lactose, levoglucosan, mannitol, mannose, sorbitol, sucrose, and xylose. The DNA G+C content is 63.4 mol%. The type strain (MEC087T = DSM 27294T = ATCC TSD-106) was isolated from silty topsoil heavy in organic matter from a campfire pit (Russell Homestead, SC; GPS location 34.91107, −83.17371). The accession number for the genome assembly is MYFN00000000 , and the 16S rRNA sequence is locus tag B5M41_026195.	<i>Shinella</i>	NCBI Assembly: GCF_004514425.2 Ts	seqco.de/i:49625