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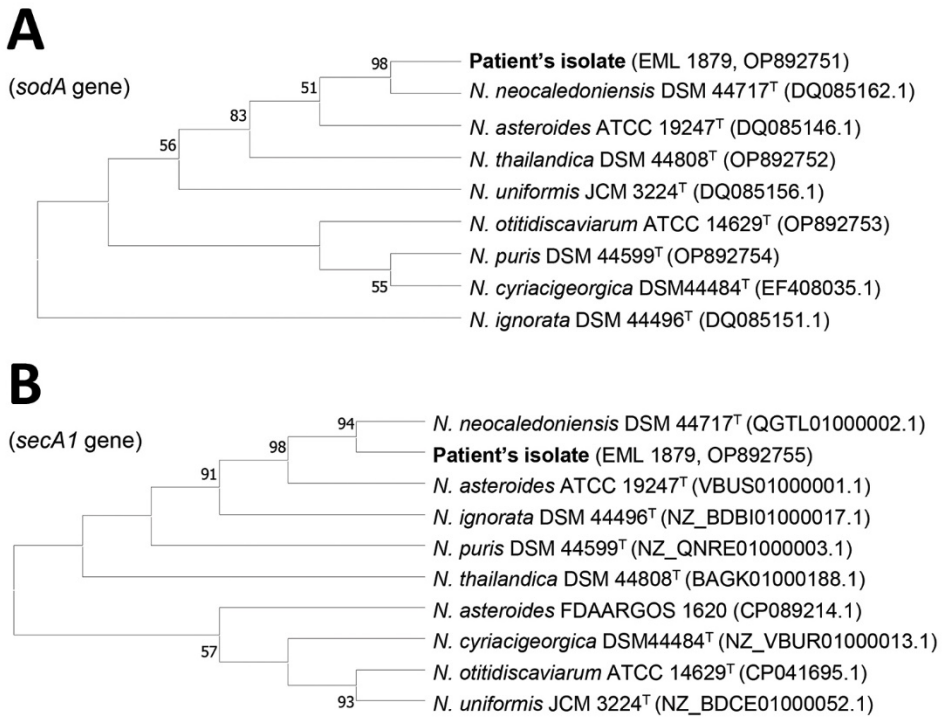
Nocardia neocaledoniensis as Rare Cause of Spondylodiscitis

Appendix

Appendix Table. Primers used in this study for the DNA sequencing

Gene and primers	Sequences (5′ →3′)	Reference
16S rRNA		(*1*,*2*)
VIV8F	AGAGTTTGATCMTGGCTCAG	
V1V8R	GGGCGGWGTGTACAAGGC	
V3V4F	CGGCCCAGACTCCTACGGGAGGCAGCA	
V3-V4R	GCGTGGACTACCAGGGTATCTAATCC	
Noc1F	GCTTAACACATGCAAGTCG	
Noc2R	GAATTCCAGTCTCCCCTG	
sodA		(*3*)
SODF	CACCAYWSCAAGCACCA	
SODR	CCTTGACGTTCTGGTACTG	
secA1		(*4*)
M13F	GTAAAACGACGACCAGGACAGYGAGTGGATGGGYCGSGTGCACCG	
M13R	CAGGAAACAGCTATGCGGCGGACGATGTAGTCCTTGTC	



Appendix Figure. Alignment of 386 bp of the *sodA* (A) and 468 bp of the *secA1* genes (B) of the patient's isolate (strain EML 1879) with the *sodA* and *secA1* genes sequences of a set of reference strains of *Nocardia* species according to the Sánchez-Herrera's instructions (*5*). The evolutionary history

was inferred using the Neighbor-Joining method, Kimura's two-parameter model and bootstrap of 1000. Branches corresponding to partitions reproduced in less than 50% bootstrap replicates were collapsed.

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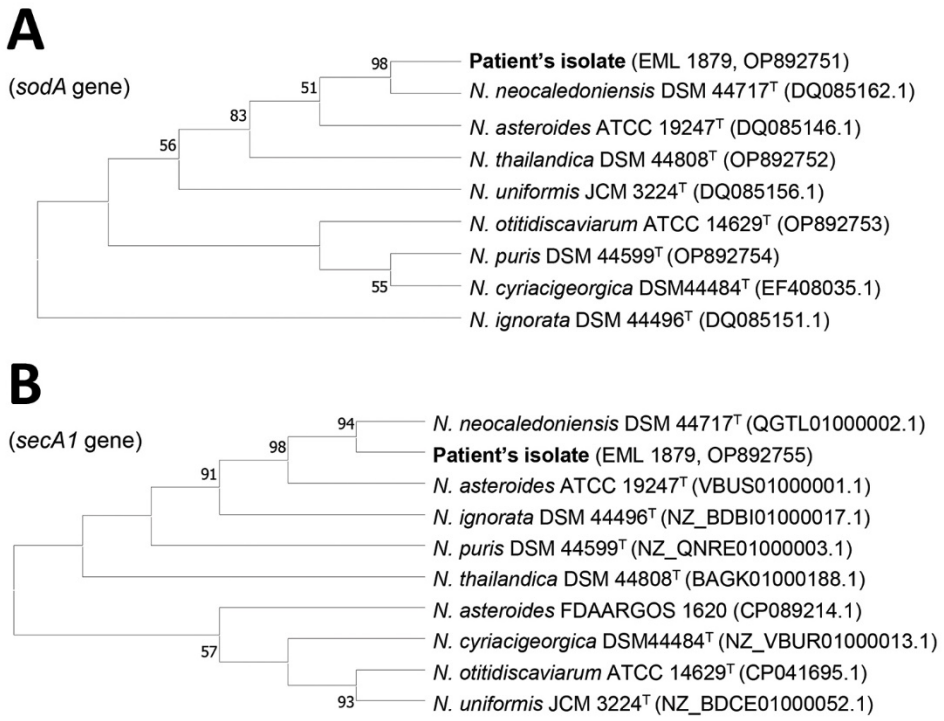
Noc2R	GAATTCCAGTCTCCCCTG	
sodA		(*3*)
SODF	CACCAYWSCAAGCACCA	
SODR	CCTTGACGTTCTGGTACTG	
secA1		(*4*)
M13F	GTAAAACGACGACCAGGACAGYGAGTGGATGGGYCGSGTGCACCG	
M13R	CAGGAAACAGCTATGCGGCGGACGATGTAGTCCTTGTC	



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References

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