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Article:

Sales de Barros Santos, H., Dame-Teixeira, N., Hitomi Nagano, M. et al. (3 more authors) (2023) Acid tolerance of *Lactobacillus* spp. on root carious lesions: a complex and multifaceted response. *Archives of Oral Biology*, 156. 105820. ISSN 0003-9969

<https://doi.org/10.1016/j.archoralbio.2023.105820>

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Table S1 mRNA raw read count attributed to *Lactobacillus* spp. genomes in sound root surfaces

Genomes	Samples									
	SRS_8	SRS_9	SRS_10	SRS_12	SRS_15	SRS_16	SRS_19	SRS_20	SRS_21	SRS_22
<i>L. acidophilus</i>	0	0	0	0	0	0	0	0	0	0
<i>L. brevis</i>	0	0	0	0	0	0	0	0	0	0
<i>L. buchneri</i>	0	0	0	0	0	0	0	0	0	0
<i>L. casei</i>	0	0	0	0	0	0	0	0	0	0
<i>L. crispatus</i>	0	0	0	0	0	0	0	0	0	0
<i>L. curvatus</i>	0	0	0	0	0	0	0	0	0	0
<i>L. delbrueckii</i>	0	0	0	0	0	0	0	0	0	0
<i>L. fermentum</i>	0	0	0	7.48	0	0	0	0	0	0
<i>L. gasseri</i>	0	0	0	8.26	0	0	0	0	0	0
<i>L. jensenii</i>	0	0	0	0	0	0	0	0	0	0
<i>L. johnsonii</i>	0	0	0	0	0	0	0	0	0	0
<i>L. paracasei</i>	0	0	0	1.67	0	0	0	0	0	0
<i>L. plantarum</i>	0	0	0	0	0	0	0	0	0	0
<i>L. rhamnosus</i>	0	0	0	0	0	0	0	0	0	0
<i>L. salivarius</i>	0	0	0	0	0	0	0	0	0	0

Data normalized considering total number of transcripts divided by the total number of genes in each sample.