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**Table 1.** Total number of *Lactobacillus* spp. genes, total number of all differently expressed genes (DEGs; % in relation to total number of genes), total number of genes potentially associated with acid tolerance (% in relation to total number of genes), number of acid tolerance-associated DEGs between sound root and carious root surfaces (% in relation to the number of acid tolerance genes) and the percentage of acid tolerance-associated DEGs in relation to all DEGs:

Genome	Total number of genes	Total number of all DEGs	Total number acid tolerance-associated genes	Number of acid tolerance-associated DEGs*	Percentage of acid tolerance-associated DEGs in relation to all DEGs
<i>L. acidophilus</i>	1,832	108 (5.89)	116 (6.33)	33 (28.44)	30.55
<i>L. brevis</i>	2,185	26 (1.18)	103 (4.71)	11 (10.69)	42.30
<i>L. buchneri</i>	2,383	62 (2.60)	117 (4.90)	16 (13.67)	25.80
<i>L. casei</i>	2,765	21 (0.75)	142 (5.13)	1 (0.70)	4.76
<i>L. crispatus</i>	1,934	1,294 (66.9)	96 (4.96)	91 (94.79)	7.03
<i>L. curvatus</i>	1,960	34 (1.73)	107 (5.45)	3 (2.80)	8.82
<i>L. delbrueckii</i>	1,808	23 (1.27)	110 (6.08)	12 (10.90)	52.17
<i>L. fermentum</i>	1,946	1,535 (78.87)	125 (6.42)	98 (78.4)	6.38
<i>L. gasseri</i>	1,772	1,426 (80.47)	108 (6.09)	92 (85.18)	6.45
<i>L. jensenii</i>	1,405	7 (0.49)	95 (6.76)	4 (4.21)	57.14
<i>L. johnsonii</i>	1,804	214 (11.86)	96 (5.32)	41 (42.70)	19.15
<i>L. paracasei</i>	2,764	71 (2.56)	180 (6.51)	17 (9.44)	23.94
<i>L. plantarum</i>	2,883	20 (0.69)	87 (3.01)	4 (9.4.69)	20.0
<i>L. rhamnosus</i>	2,745	1,663 (60.58)	131 (4.77)	114 (87.02)	6.85
<i>L. salivarius</i>	1,876	1,364 (72.70)	124 (6.60)	116 (93.54)	8.50

\*All acid tolerance associated DEGs showed Log2fold change >1 indicating an overexpression in carious root surfaces

