Table S1. Accuracy of DNA extraction in accordance with Best Practice Recommendations for Internal Validation of DNA Extraction Methods [52].

Accuracy	VN141R	MD141	SD151	CVN291	MANUAL
Specificity (%) *	1	1	1	1	1
Sensibility (%) **	0.92	0.89	0.88	0.92	1
Repeatability (CV%) ***	19.9	16.0	19.8	13.8	24.6

* Specificity=(True Negatives (D))/(True Negatives (D)+False Positives (B)); ** Sensibility=(True Positives (A))/(True Positives (A)+False Negatives (C)); *** CV%=(SD Yield DNA/mean Yield DNA)×100. The acceptability criteria were set at CV≤25%.

Table S2. Primers, probes and references [2,44,48,49,64].

Target	Forward Primer	Reverse Primer	Probe	Reference
Propionibacterium spp.	GCGTGAGTGACGGTAA TGGGTA	TTCCGACGCGATCAAC CA	AGCGTTGTCCGGATTTATTGGG CG	Elshi et al, 2000
Corynebacterium spp.	ACCATCGACGAAGGGT TTCT	TCACCGCTACACCAGG AATT	GGCGATACGGGCATAACTTG	Piana et al., 2021
Bacteroides vulgatus	AAAGAAGTTCCTAGCG CTGC	GCTGGCTTTCTTTTACG CCT	TCTGAAAAACGTGCTCAGGC	Valeriani et al., 2018
Enterococcus spp.	AGAAATTCCAAACGAA CTTG	CAGTGCTCTACCTCCATTGGTTCTCTCCGAAATAGCTTT AGGGCTA		Valeriani et a., 2018
Staphylococcus epidermidis	CAACTCGATGCAAATC AGCAA	GAACCGCATAGCTCCC TGC	TACTACGCTGGTGGAACTTCAA ATCGTTATCG	Chung et al., 2016
Staphylococcus aureus	ACACCTGAAACAAAGC ATCC	CGCTAAGCCACGTCCA TATT	TGGTCCTGAAGCAAGTGCATTT ACGA	Giampaoli et al., 2012
Streptococcus salivarius	GATGCCAAGGGTGAAG TTGT	GAGCCATCAGGATTCG TAGC	TGGCGAACAGACGATCAACGG	Giampaoli et al., 2012
Streptococcus mutans	CGGTTCTCAGCAAGAC ATGA	ATGGTACCCAATCCGC AATA	TGCAGTTAAAGCTCTGCATAA AAGCGG	Giampaoli et al., 2012

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