

Determination of the minimum inhibitory concentrations of the antibacterial oregano and thyme essential oils in the presence of polysorbate 80

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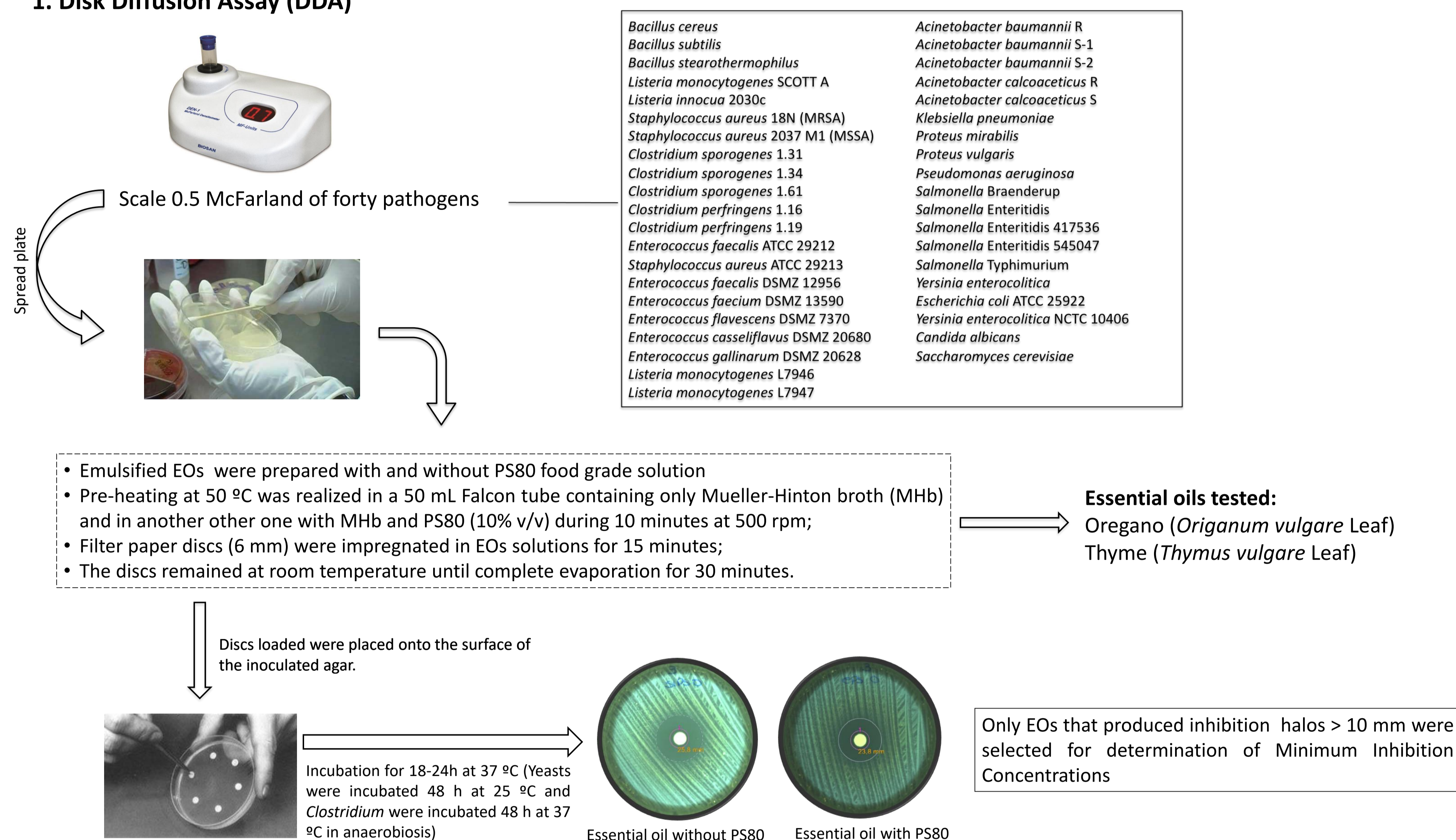
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Introduction

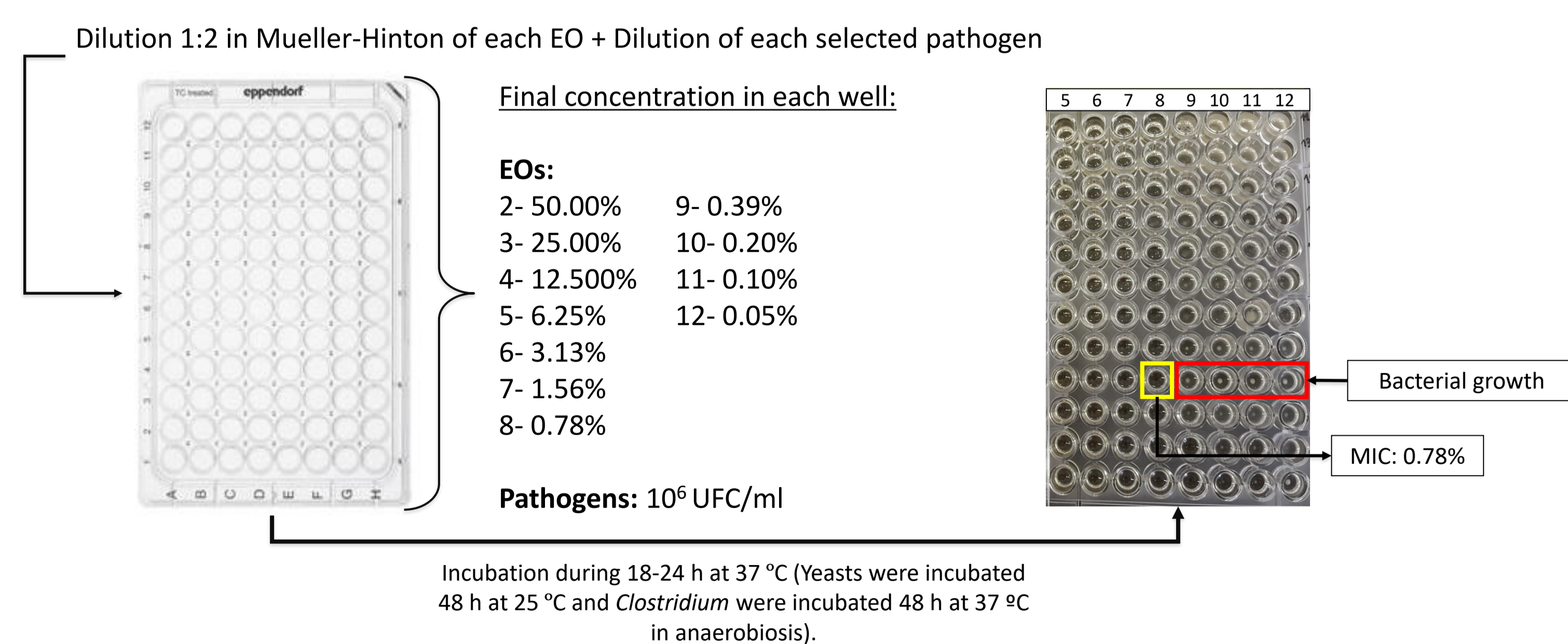
Plant-derived essential oils (EOs) have attracted the interest of the scientific community in recent years regarding their safety, as natural ingredients, as well as their application in food products (Alizadeh-Behbahani and Tabatabaei-Yazdi 2015; Fournomiti et al., 2015). Essential oils have shown remarkable antimicrobial potential against spoilage and pathogenic microorganisms. However, different antimicrobial activities of similar EOs may be observed when different methods are used (Bajalan et al., 2016). This study aims to evaluate and demonstrate the influence of polysorbate 80 (PS80) on the antimicrobial activity and minimum inhibitory concentrations (MICs) of oregano and thyme EOs against forty microorganisms.

Methods

1. Disk Diffusion Assay (DDA)



2. Minimum Inhibition Concentration (MIC)



3. Statistical analysis

The comparison of the antimicrobial activity of EOs against each microorganism was carried out by one-way analysis of variance (ANOVA). The Tukey-Kramer test was used to determine the significant differences ($p < 0.05$) among group means.

Results

Table 1. Zones of growth Inhibition (mm; mean \pm standard deviation) showing antimicrobial activity (including the disk diameter of 6.0 mm) and Minimum Inhibitory Concentration (MIC) (results are expressed in % of EO) of oregano and thyme EOs with and without polysorbate against different microorganisms

	DDA with PS80		DDA without PS80		MIC with PS80		MIC without PS80	
	Oregano	Thyme	Oregano	Thyme	Oregano	Thyme	Oregano	Thyme
<i>Bacillus cereus</i>	37.7 \pm 0.5	28.4 \pm 2.8	39.7 \pm 3.7	36.1 \pm 2.2	12.50	3.13	0.05	0.20
<i>B. stearothersophilus</i>	34.2 \pm 0.5	26.6 \pm 1.5	35.1 \pm 2.9	41.0 \pm 4.9	12.50	6.25	0.39	0.39
<i>B. subtilis</i>	22.9 \pm 0.4	18.6 \pm 0.5	24.4 \pm 2.3	23.2 \pm 1.1	12.50	6.25	0.05	0.10
<i>Clostridium sporogenes</i> 1.31	20.9 \pm 0.1	25.7 \pm 0.7	39.7 \pm 1.1	13.5 \pm 0.9	6.25	6.25	0.05	0.20
<i>Clostridium sporogenes</i> 1.34	17.3 \pm 2.6	19.0 \pm 0.8	37.4 \pm 0.0	18.8 \pm 1.7	6.25	12.50	0.05	0.20
<i>Clostridium sporogenes</i> 1.61	25.8 \pm 0.1	28.1 \pm 0.4	11.1 \pm 0.4	14.1 \pm 1.8	3.13	6.25	0.05	0.20
<i>Clostridium perfringens</i> 1.16	20.7 \pm 3.9	20.9 \pm 2.1	12.1 \pm 1.0	22.0 \pm 1.0	3.13	3.13	0.05	0.10
<i>Clostridium perfringens</i> 1.19	18.9 \pm 1.6	20.4 \pm 1.1	37.4 \pm 0.0	16.1 \pm 2.1	3.13	6.25	0.05	0.10
<i>E. faecalis</i> ATCC 29212	21.7 \pm 5.2	10.8 \pm 0.2	29.6 \pm 3.5	28.9 \pm 1.1	12.50	12.50	0.05	0.20
<i>E. faecalis</i> DSMZ 12956	23.1 \pm 2.1	10.1 \pm 0.5	36.6 \pm 2.3	16.8 \pm 2.0	12.50	6.25	0.05	0.20
<i>E. faecium</i> DSMZ 13590	36.3 \pm 0.5	16.7 \pm 3.2	34.5 \pm 2.3	28.3 \pm 4.4	6.25	6.25	0.05	0.20
<i>E. flavescens</i> DSMZ 7370	40.2 \pm 3.6	37.4 \pm 1.1	37.2 \pm 0.8	30.1 \pm 0.4	6.25	6.25	0.10	0.05
<i>E. gallinarum</i> DSMZ 20628	30.4 \pm 0.0	26.3 \pm 0.4	29.6 \pm 2.4	28.7 \pm 3.2	6.25	6.25	0.05	0.20
<i>E. casseliflavus</i> DSMZ 20680	27.9 \pm 1.1	17.3 \pm 2.1	28.0 \pm 2.5	26.8 \pm 1.1	6.25	6.25	0.10	0.10
<i>L. monocytogenes</i> 7946	31.1 \pm 0.2	31.1 \pm 0.8	38.4 \pm 2.5	36.0 \pm 1.4	0.39	3.13	0.05	0.05
<i>L. monocytogenes</i> 7947	31.5 \pm 3.3	32.1 \pm 2.4	35.9 \pm 2.4	33.7 \pm 5.9	0.39	0.78	0.05	0.05
<i>L. monocytogenes</i> SCOTT A	14.0 \pm 2.4	11.4 \pm 1.7	14.8 \pm 0.8	13.0 \pm 2.8	3.13	6.25	0.05	0.10
<i>Listeria innocua</i> 2030c	34.4 \pm 0.8	29.4 \pm 2.5	39.5 \pm 3.3	32.1 \pm 1.7	3.13	6.25	0.05	0.10
<i>St. aureus</i> ATCC	30.8 \pm 1.1	28.4 \pm 3.6	28.8 \pm 1.3	29.9 \pm 2.5	3.13	6.25	0.05	0.05
<i>St. aureus</i> 18N (MRSA)	35.1 \pm 3.2	30.5 \pm 0.4	30.0 \pm 1.6	34.7 \pm 3.0	6.25	6.25	0.05	0.10
<i>St. aureus</i> 2037 M1 (MSSA)	30.2 \pm 0.5	29.3 \pm 2.6	31.0 \pm 0.1	30.0 \pm 1.6	0.78	3.13	0.05	0.10
<i>A. baumannii</i>	36.5 \pm 0.1	30.1 \pm 3.5	37.4 \pm 4.0	32.6 \pm 3.7	1.56	1.56	0.05	0.05
<i>A. baumannii</i>	36.8 \pm 0.0	28.1 \pm 2.7	37.7 \pm 3.5	38.3 \pm 3.3	1.56	1.56	0.05	0.05
<i>A. baumannii</i>	35.8 \pm 0.8	30.1 \pm 0.6	33.5 \pm 0.0	32.4 \pm 0.0	0.78	1.56	0.05	0.05
<i>A. calcoaceticus</i>	29.8 \pm 1.8	26.0 \pm 1.3	29.4 \pm 1.5	30.4 \pm 1.2	0.78	1.56	0.05	0.05
<i>A. calcoaceticus</i>	32.5 \pm 1.8	30.1 \pm 0.8	33.0 \pm 1.8	31.4 \pm 1.4	1.56	1.56	0.05	0.05
<i>Escherichia coli</i> ATCC	33.0 \pm 1.9	24.7 \pm 0.5	33.9 \pm 1.8	31.9 \pm 2.7	3.13	3.13	0.10	0.20
<i>Klebsiella pneumoniae</i>	16.6 \pm 2.8	16.8 \pm 3.3	24.7 \pm 2.1	20.4 \pm 3.5	6.25	3.13	0.10	0.20
<i>Proteus mirabilis</i>	32.4 \pm 2.7	28.6 \pm 0.4	30.1 \pm 2.8	36.7 \pm 1.8	1.56	3.13	0.05	0.05
<i>Proteus vulgaris</i>	23.6 \pm 1.0	16.0 \pm 1.8	24.9 \pm 2.1	22.6 \pm 2.8	3.13	12.50	0.10	0.20
<i>Pseudomonas aeruginosa</i>	22.2 \pm 0.7	17.8 \pm 1.6	21.1 \pm 2.1	18.5 \pm 0.2	6.25	12.50	0.05	0.05
<i>S. Braenderup</i>	23.3 \pm 1.8	16.8 \pm 0.4	25.1 \pm 1.6	22.1 \pm 2.5	3.13	12.50	0.10	0.10
<i>S. Enteritidis</i>	23.4 \pm 0.0	18.8 \pm 2.1	23.3 \pm 0.5	20.9 \pm 1.5	3.13	12.50	0.10	0.10
<i>S. Enteritidis</i> 417536	17.0 \pm 0.7	16.1 \pm 1.3	19.6 \pm 1.7	20.5 \pm 2.5	1.56	3.13	0.05	0.05
<i>S. Enteritidis</i> 545047	16.3 \pm 3.0	18.4 \pm 3.4	19.8 \pm 0.3	20.0 \pm 0.2	3.13	6.25	0.05	0.05
<i>S. Tiphymurium</i>	30.7 \pm 1.3	24.6 \pm 0.4	33.3 \pm 3.0	29.9 \pm 1.8	1.56	3.13	0.10	0.10
<i>Yersinia enterocolitica</i>	36.1 \pm 3.2	31.2 \pm 1.3	40.4 \pm 4.0	47.8 \pm 3.3	1.56	3.13	0.05	0.10
<i>Yersinia enterocolitica</i> NCTC10406	22.1 \pm 1.1	18.6 \pm 1.5	23.3 \pm 2.5	21.5 \pm 1.0	1.56	3.13	0.05	0.10
<i>Candida albicans</i>	44.5 \pm 0.4	35.7 \pm 0.4	55.3 \pm 2.6	50.7 \pm 2.8	3.13	0.78	0.05	0.10
<i>Saccharomyces cerevisiae</i>	44.5 \pm 4.5	26.7 \pm 4.4	50.0 \pm 1.8	37.5 \pm 0.1	0.78	1.56	0.05	0.05

- For the Disk Diffusion Assay, comparing both techniques, it was observed that in most microorganisms no differences were seen in the size of the halo;
- However, some differences were observed for MIC determination: MICs of each EO were different in the presence and absence of PS80, for all microorganisms.

Conclusions

The PS80 was used to facilitate the homogenization of EOs, but this may turn the antimicrobial compounds in EOs less available, with no direct contact with the target organisms limiting the microbial reduction capacity. This study reinforces the need for using defined standard methods for the *in vitro* determination of MICs.

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