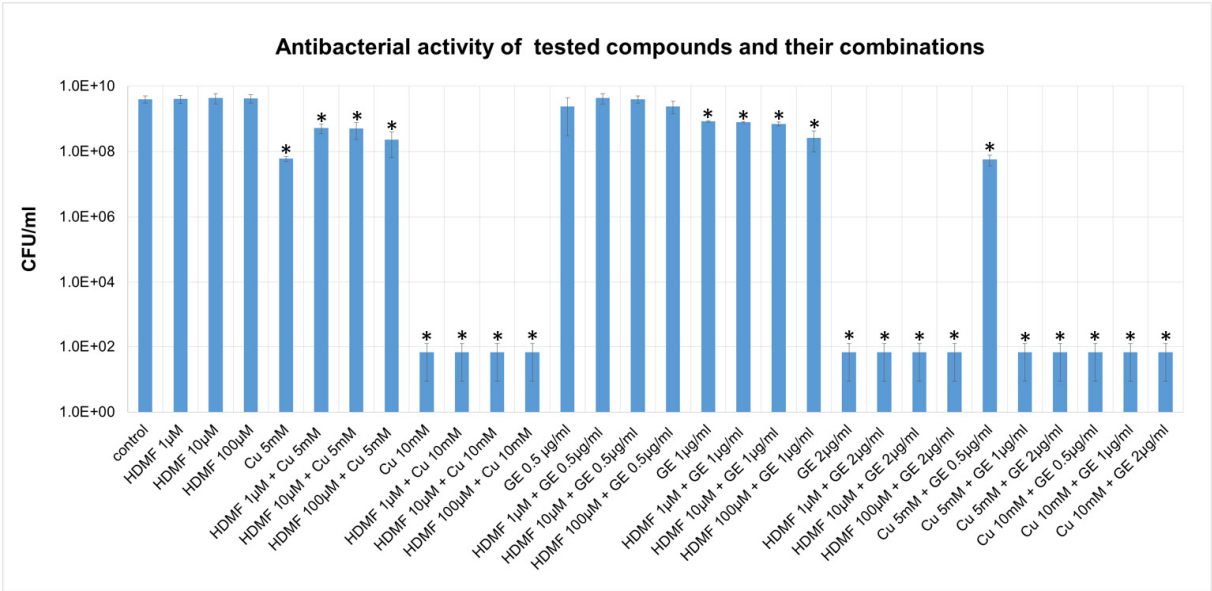


# The *in vitro* anti-pseudomonal activity of Cu<sup>2+</sup>, strawberry furanone, gentamicin, and lytic phages alone and in combination: pros and cons

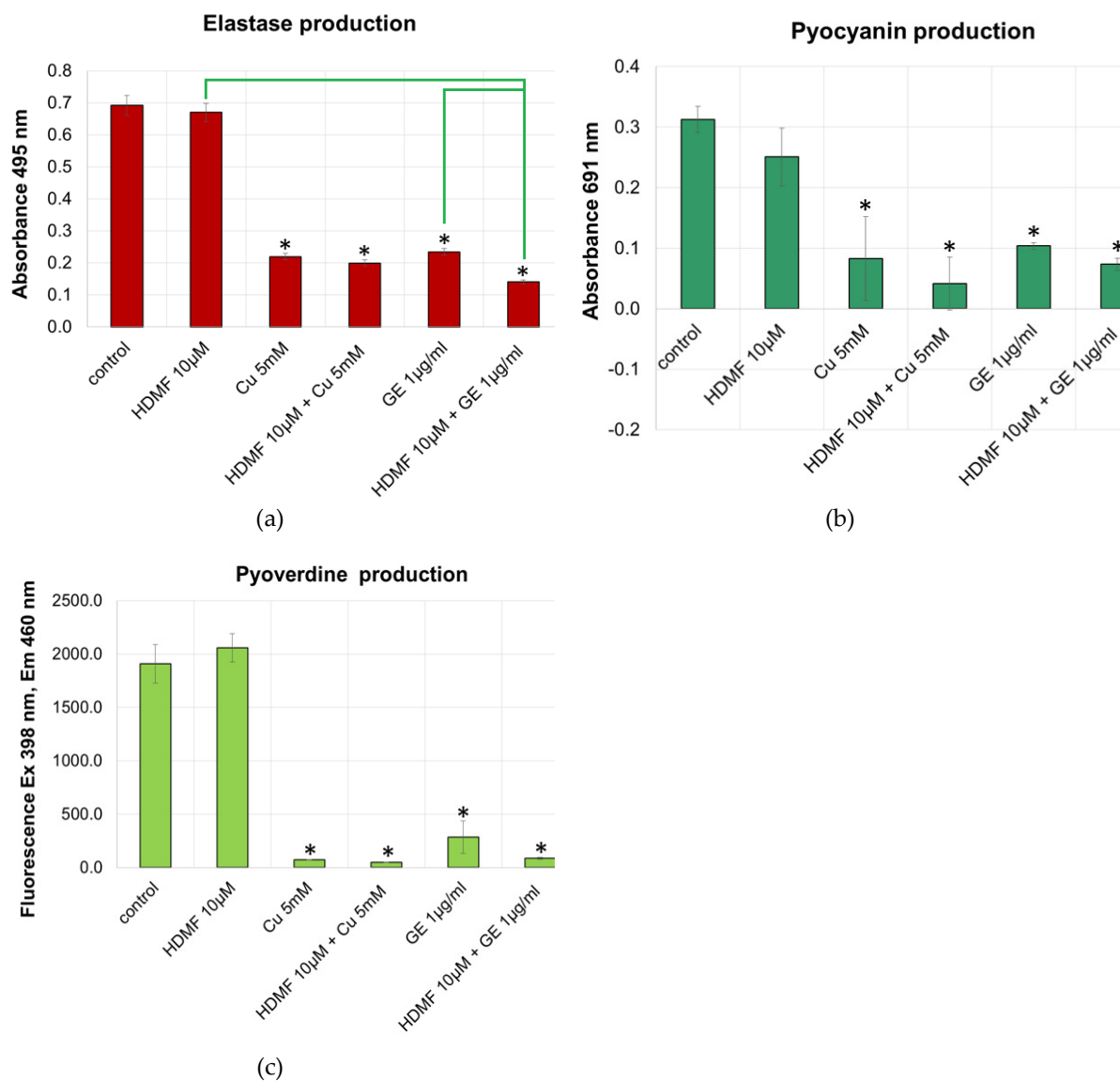
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**Figure S1.** The impact of Cu<sup>2+</sup>, HDMF, and GE on 20 h growth of *P. aeruginosa* PAO1; \*statistically significant differences according to PAO1 untreated control (*P*<0.05)



**Figure S2.** The impact of  $\text{Cu}^{2+}$ , HDMF, and GE on *P. aeruginosa* PAO1 virulence determinants production: (a) elastase production; (b) pyocyanin production; (c) pyoverdine production; \*statistically significant differences according to PAO1 control ( $P < 0.05$ ), brackets indicate statistically significant differences between agent combinations ( $P < 0.05$ ).

**Table S1.** Statistical analysis of the significance of differences between tested combinations of antibacterial and anti-virulent components (P<0.05).

	Figure 1b		Figure 1c		Figure 1d	
	HDMF 10μM GE 1μg/ml	HDMF 10μM Cu 5mM	HDMF 10μM GE 1μg/ml	HDMF 10μM Cu 5mM	HDMF 10μM GE 1μg/ml	HDMF 10μM Cu 5mM
HDMF 10μM	Y	Y	Y	Y	Y	Y
Cu 5mM		N		N		N
GE 1μg/ml	Y		N		N	

Y-statistically significant with P<0.05; N-statistically not significant;

**Table S2.** The impact of Cu<sup>2+</sup>, HDMF and GE alone and in combination on antibiotic sensitivity patterns of *P. aeruginosa* PAO1.

Tested antibiotic	PAO1 (control)	HDMF 10 μM	Cu 5 mM	HDMF 10μM Cu 5mM	GE 0,5 μg/ml	HDMF 10μM GE 0,5μg/ml	Cu 5mM GE 0,5μg/ml
CTX	24-25 <sup>S</sup>	-	-	-	-	-	-
CAZ	29-31 <sup>S</sup>	-	-	-	-	-	-
PIP	31-32 <sup>S</sup>	-	-	-	-	-	-
TZP	33-35 <sup>S</sup>	-	-	-	-	-	-
IPM	27-29 <sup>S</sup>	-	-	-	-	-	-
CN	18-19 <sup>S</sup>	-	-	-	-	-	-
AK	24-25 <sup>S</sup>	-	-	-	-	-	-
CIP	36-38 <sup>S</sup>	-	-	-	-	-	-

Numbers indicate diameters of zones of inhibition in mm, S-sensitive according to EUCAST recommendations ([https://www.eucast.org/ast\\_of\\_bacteria/](https://www.eucast.org/ast_of_bacteria/)), – no change observed,

**Table S3.** The starters used for genetic confirmation of phage culture purity.

Phage	Genome size	GenBank accession number	Starter sequence	Melting temperature [°C]
KT28	66,381 bp	KP340287	(F) CTTCGACGCCGACCGGGAATC	60.2
			(R) GATATGCTACGCCGCCCATTTGTG	58.8
KTN4	279,593 bp	KU521356	(F) CTAACCTAAATATATTGACGCTGTATCG	53.7
			(R) CCGCTATATACTTGTAGATAACGC	54
LUZ19	43,548 bp	NC_010326	(F) CTAACCTAAATATATTGACGCTGTATCG	62
			(R) CAACCTCCAGCCAATCTCAATAAAAATT	57.3