

Supplementary material

Manuscript title: “Fractional Inhibitory Concentration of bio-actives from agricultural waste disassembles biofilms and quenches virulence of nosocomial pathogens” by **Srividhya et al.**,

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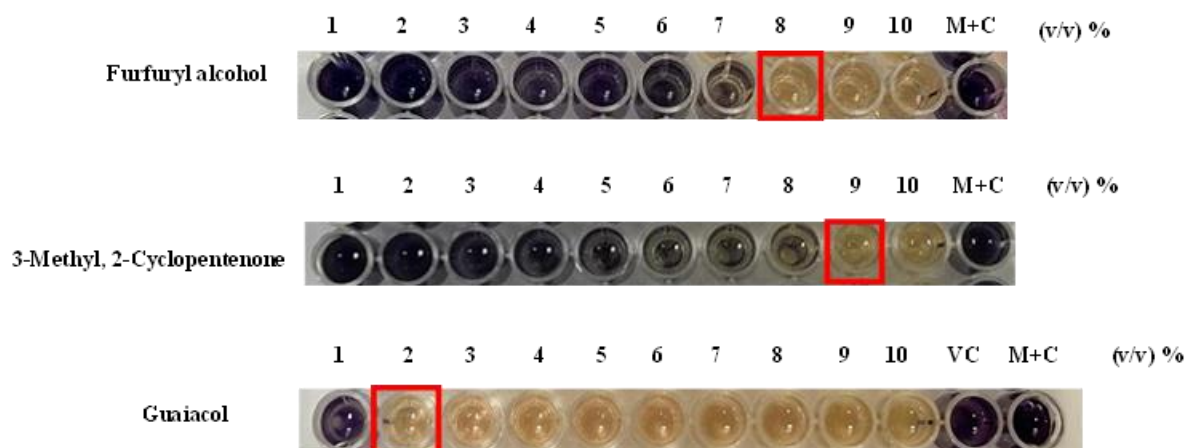


Figure A1. Minimum inhibitory concentration (MIC) of compounds Furfuryl alcohol, 3-Methyl, 2-Cyclopentenone and Guaiacol at 8%, 9% and 2% respectively highlighted in red box.

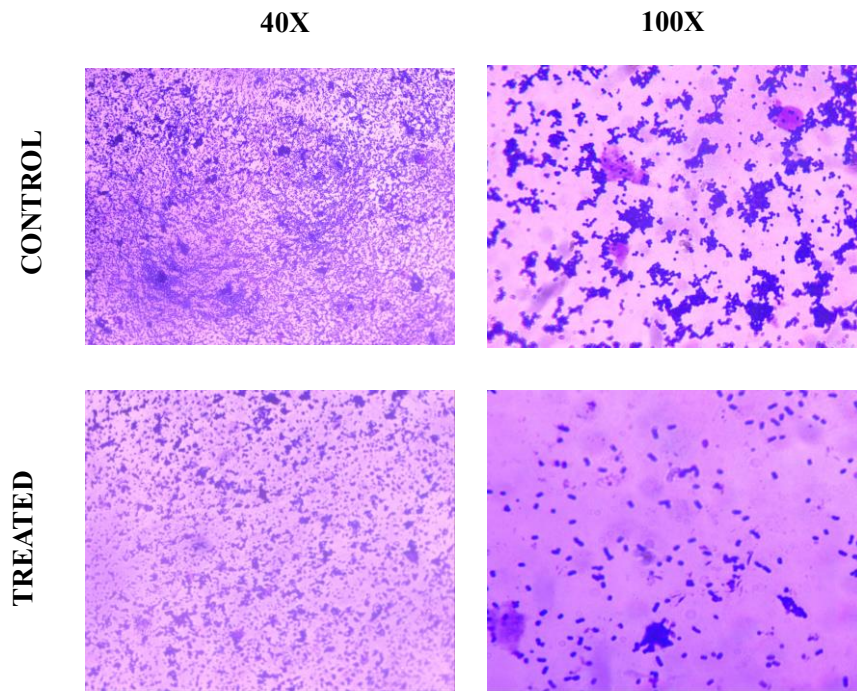


Figure A2. Light microscopic visualisation of biofilm eradication observed in the control and BIC treated sample at 40X and 100X magnifications

Table A3. The Minimum inhibitory concentration (MIC) of compounds Furfuryl alcohol, 3-Methyl, 2-Cyclopentenone and Guaiacol at 8%, 9% and 2% against the mixed species.

COMPOUNDS	MIC (%)
Furfuryl alcohol	8
3-Methyl, 2-Cyclopentenone	9
Guaiacol	2

Table A4. Primer's list of MRSA, *A. baumannii* and *C. auris* along with along with their role

S.No	Gene	Sequence (5'-3')	Role
1	<i>agrAC</i> (F)	CTGATAATCCCTTATGAGGTGC	QS virulence factors
	<i>agrAC</i> (R)	CGATGCATAGCAGTGTTTC	
2	<i>icaA</i> (F)	ACACTTGCTGGCGCAGTCAA	EPS adhesion
	<i>icaA</i> (R)	TCTGGAACCAACATCCAACA	
3	<i>icaD</i> (F)	ATGGTCAAGCCCAGACAGAG	EPS adhesion
	<i>icaD</i> (R)	AGTATTTTCAATGTTTAAAGCA	
4	<i>crtM</i> (F)	ATCCAGAACCACCCGTTTTT	Staphyloxanthin inhibition
	<i>crtM</i> (R)	GCGATGAAGGTATTGGCATT	
5	<i>crtN</i> (F)	GATGAAGCTTTGACGCAACA	Staphyloxanthin inhibition
	<i>crtN</i> (R)	TTCGCATGATACGTTTGCTC	
6	<i>fnbA</i> (F)	ATCAGCAGATGTAGCGGAAG	Fibronectin binding gene
	<i>fnbA</i> (R)	TTTAGTACCGCTCGTTGTCC	
7	<i>fnbB</i> (F)	AAGAAGCACCGAAAACGTGTG	Fibronectin binding gene
	<i>fnbB</i> (R)	TCTCTGCAACTGCTGTAACG	
8	<i>sspB</i> (F)	CCAGCAAATTGTTGTTGTGCTAG	Autoinducer signalling molecule
	<i>sspB</i> (R)	AAGCCAAAGCCGATTACACTC	
9	<i>csu A/B</i> (F)	CAGCAGCAACAGGTGGCAATA	Initial attachment
	<i>csu A/B</i> (R)	AAGGTTTGTACGTGCAGCATCA	
10	<i>csuE</i> (F)	GCTTGGCTTTAGCAAACATGACC	Initial attachment
	<i>csuE</i> (R)	ATTGCCATCAGGCCCCGCTA	
11	<i>bfmS</i> (F)	ACCGCCCGTAATCCGAAC	Abiotic surface
	<i>bfmS</i> (R)	TGAACTTATTCCACCGCCTTTA	
12	<i>bfmR</i> (F)	GTTTAACCGTTTGTCTGTG	Abiotic surface
	<i>bfmR</i> (R)	GTGGTTGAACTGGTTTCG	
13	<i>pgaB</i> (F)	AAGAAAATGCCTGTGCCGACCA	Structural integrity
	<i>pgaB</i> (R)	GCGAGACCTGCAAAGGGCTGAT	
14	<i>ompA</i> (F)	CGCTTCTGCTGGTGCTGAAT	Fibronectin binding gene
	<i>ompA</i> (R)	CGTGCAGTAGCGTTAGGGTA	
15	<i>abaR</i> (F)	ATGGAAAGTTGGCAAGAG	Surface motility
	<i>abaR</i> (R)	CTACAAAAGCCCTAGCATTAC	
16	<i>abaI</i> (F)	ATGAATATTATTGCTGGA	Surface motility
	<i>abaI</i> (R)	CTACACATCAATCAAGCA	
17	16S rRNA (F)	ACTCCTACGGGAGGCAGCAG	Housekeeping gene
	16S rRNA (R)	ATTACCGCGGCTGCTGG	
18	<i>erg11</i> (F)	GAAAGAGAACCATTACCAGG	Azole resistance
	<i>erg11</i> (R)	AGGAATCGACGGATCAC	
19	<i>cdr</i> (F)	TGGTGCCATGACTCCTGCTA	Efflux gene
	<i>cdr</i> (R)	CCATCGAGACCAACCCAACA	
20	<i>efg</i> (F)	CCAGGGTGCTGCTAATG	Biofilm formation
	<i>efg</i> (R)	GGGTGAAGGGTGAACCTGAACC	
21	<i>hgc</i> (F)	GCTTCCTGCACCTCATCAAT	Morphogenesis and biofilm formation
	<i>hgc</i> (R)	AGCACGAGAACCAGCGATAC	
22	ITS (F)	TCCGTAGGTGAACCTGCGG	Housekeeping gene

	ITS (R)	TCCTCCGCTTATTGATATGC	
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