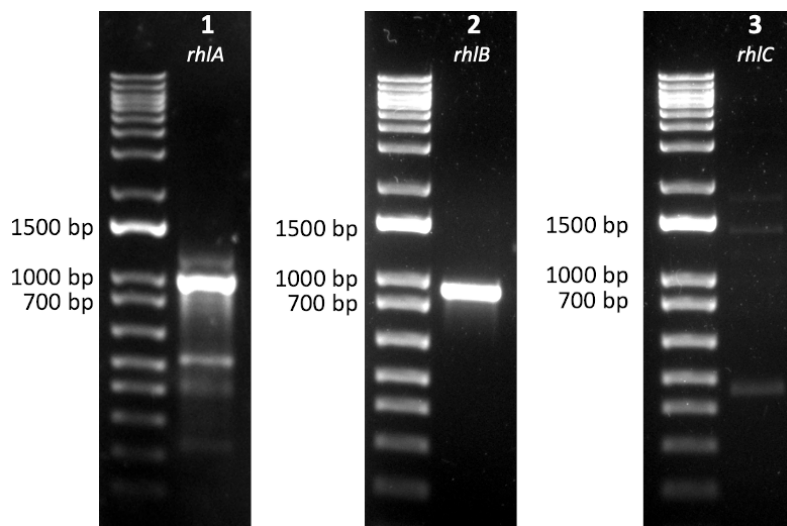
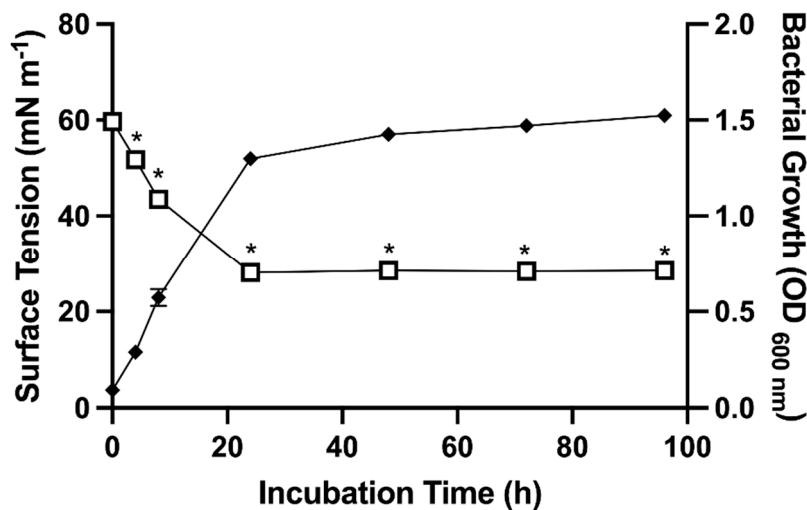


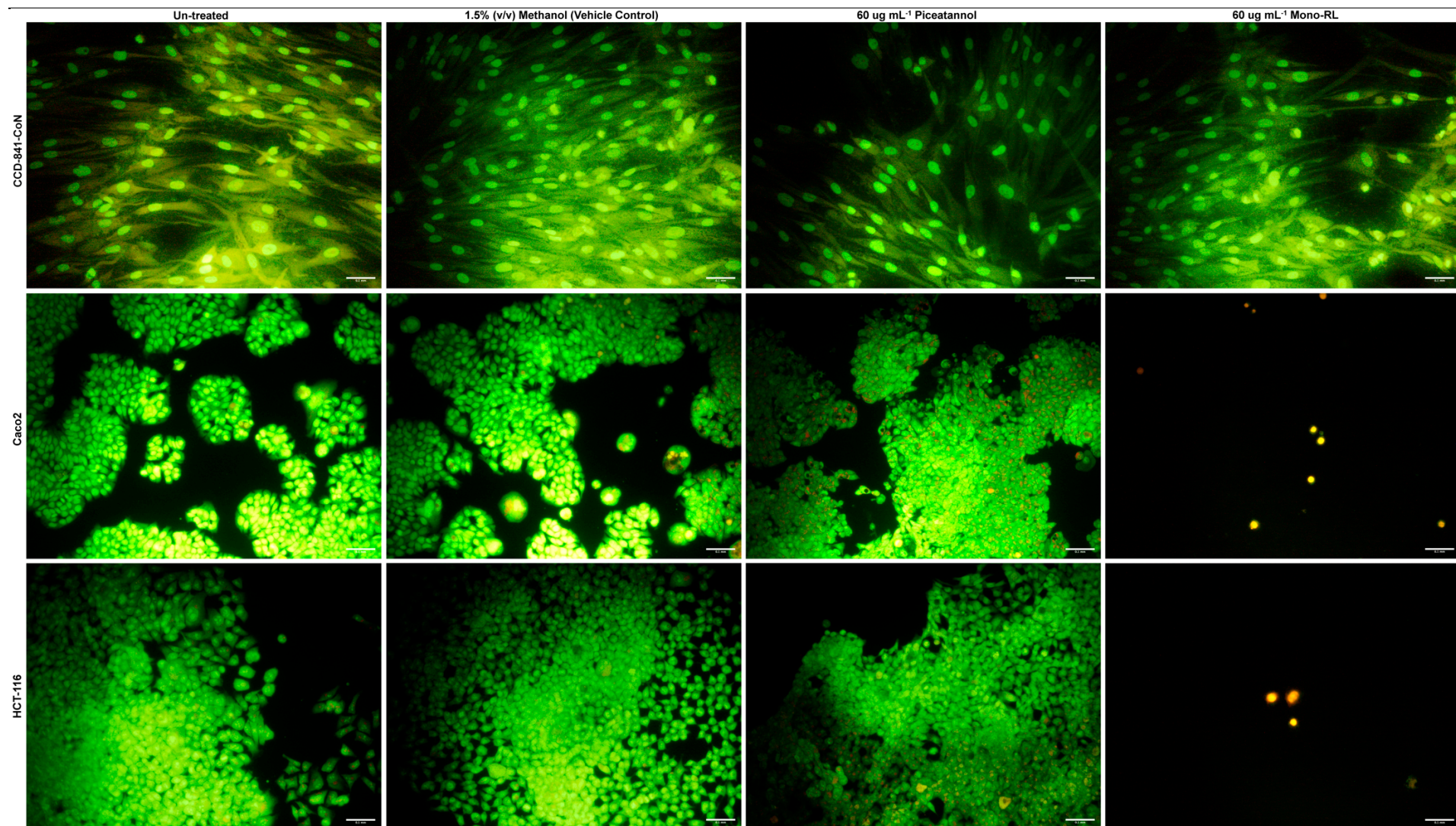
## Supplementary Materials: Mono-Rhamnolipid Biosurfactants Synthesized by *Pseudomonas aeruginosa* Detrimentally Effect Colorectal Cancer Cells



**Figure S1.** Amplified DNA fragments resulting from the PCR screening for RL biosynthesis genes *rhlA*, *rhlB* and *rhlC* in *P. aeruginosa* PAO1  $\Delta rhlC$ . The lack of an amplification product at  $\approx 800$  bp for *rhlC* is confirmation of mutation of this gene within the strain.



**Figure S2.** Growth and surface tension measurements of *P. aeruginosa* PAO1  $\Delta rhlC$  over 96 h. Bacterial growth of in NB supplemented with 1% (*v/v*) glycerol ( $\blacklozenge$ ). Surface tension measurement of cell-free supernatant samples obtained from cultures over time ( $\square$ ). Error bars represent standard deviation from the mean ( $n = 3$ ). Analysis of surface tension measurements was by two-way ANOVA with post hoc. comparison to surface tension at  $t = 0$  h ( $* = p \leq 0.05$ ).



**Figure S3.** Images of AO/ PI-stained cell lines. CCD-841-CoN, Caco2 and HCT-116 either un-treated or following 24 h of treatment with 1.5% (*v/v*) methanol (vehicle control); 60  $\mu\text{g mL}^{-1}$  piceatannol; or 60  $\mu\text{g mL}^{-1}$  mono-RL. Cells were imaged at 200 $\times$  magnification, scale bar = 100  $\mu\text{m}$ .

**Table S1.** PCR primer pairs used in this study and their melting temperatures. Oligonucleotide primers were obtained in lyophilized form from *Eurofins Genomics*, Germany, reconstituted in nuclease free distilled H<sub>2</sub>O at a concentration of 100 µM and utilized in each PCR at a working concentration of 0.5 µM.

Gene Amplified	Primer Pair	Sequence (5'-3')	Melting Temp. (°C)
16S rRNA	27-F	AGAGTTTGATCMTGGCTCAG	56.3
	1492-R	CGGTTACCTTGTTACGACTT	55.3
	907-R *	CCGTCAATTCMTTTRAGTTT	51.1
<i>rhlA</i>	<i>rhlA</i> -F	AGACGTACTCGTAGACCGGC	61.4
	<i>rhlA</i> -R	AAGGACGACGAGGTGGAAAT	57.3
<i>rhlB</i>	<i>rhlB</i> -F	CCCGTAGTTCTGCATCTGGT	59.3
	<i>rhlB</i> -R	AACTGCAACGCTTTCTCGAT	55.3
<i>rhlC</i>	<i>rhlC</i> -F	GTCCACGTGGTCGATGAAC	58.8
	<i>rhlC</i> -R	CCTGGTCGACTCGTCATTCT	59.3

\* Primer 907-R used as internal primer for 16S rRNA sequencing