

1 **APPLIED MICROBIOLOGY AND BIOTECHNOLOGY**

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3 Supplementary material for:

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5 **Characterisation of cytotoxicity and immunomodulatory effects of glycolipid biosurfactants on human**
6 **keratinocytes**

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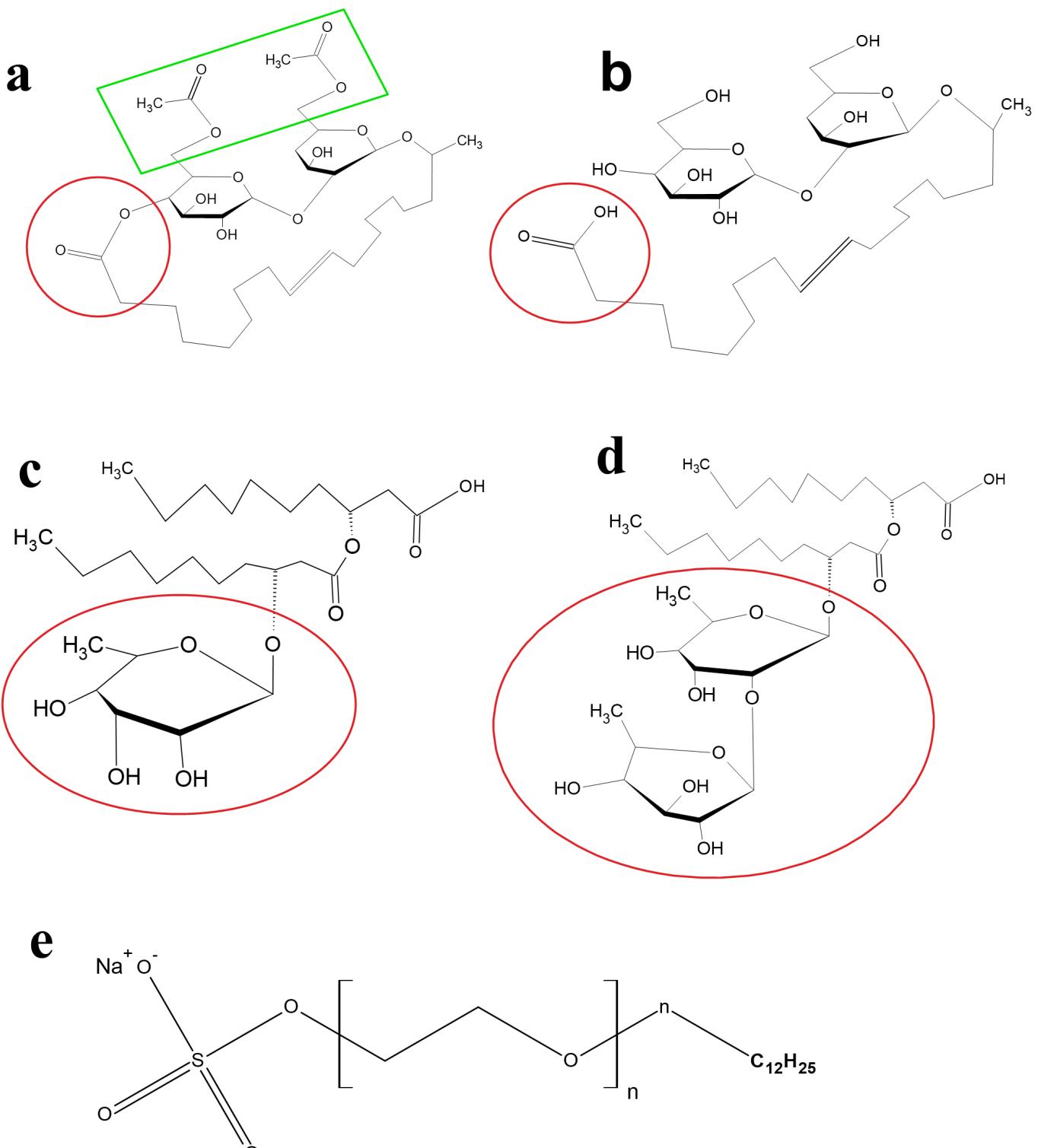
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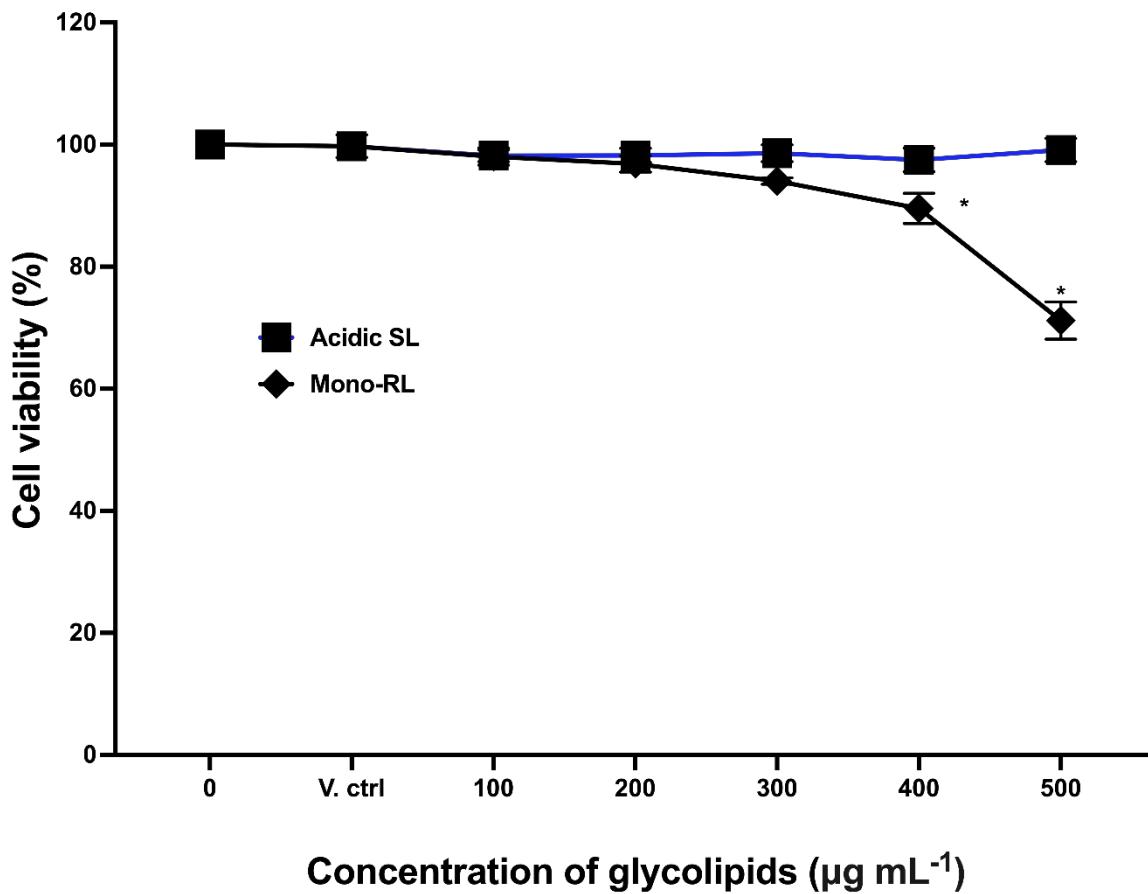
42 Supplementary Figures
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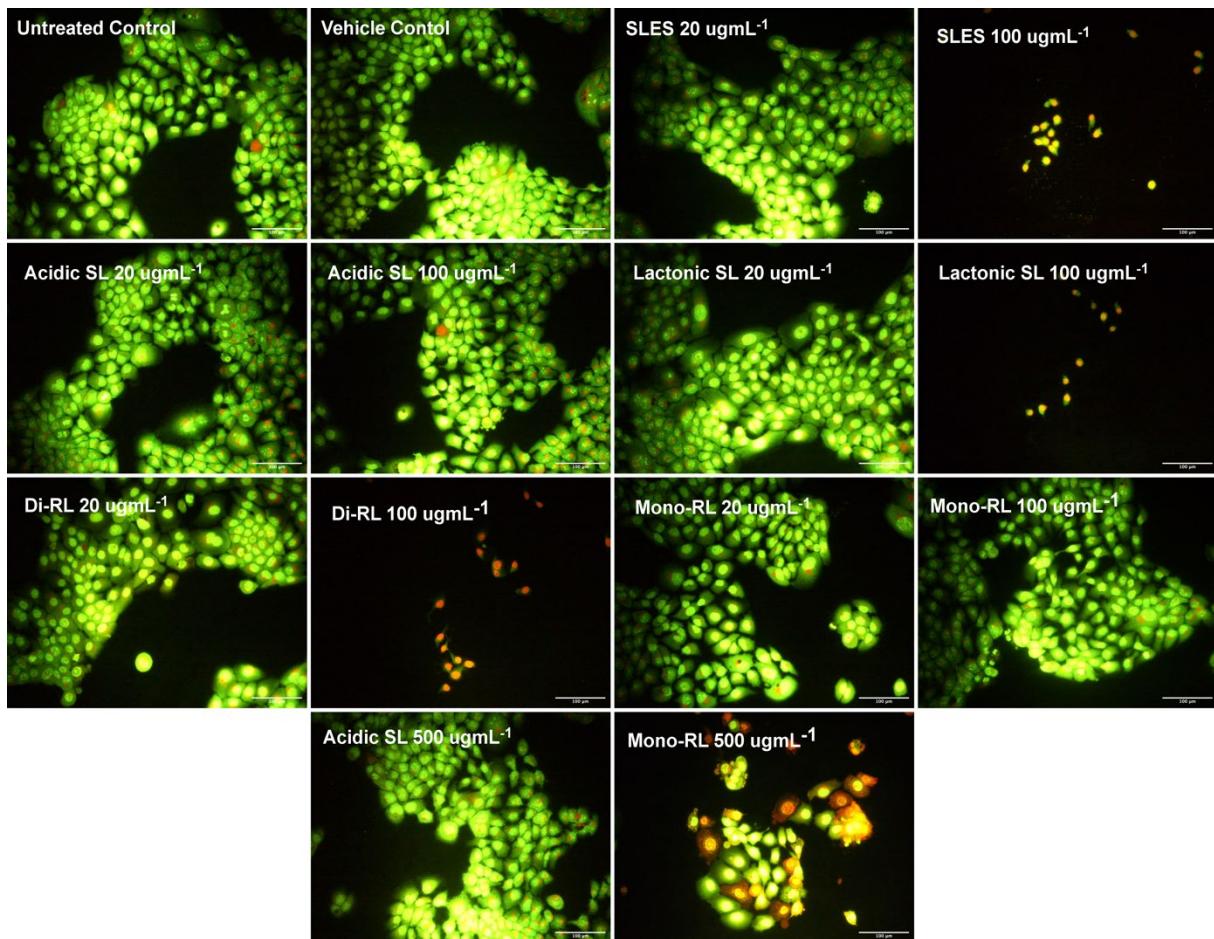
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Fig. S1

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84 Fig. S2
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102 **Fig. S3**

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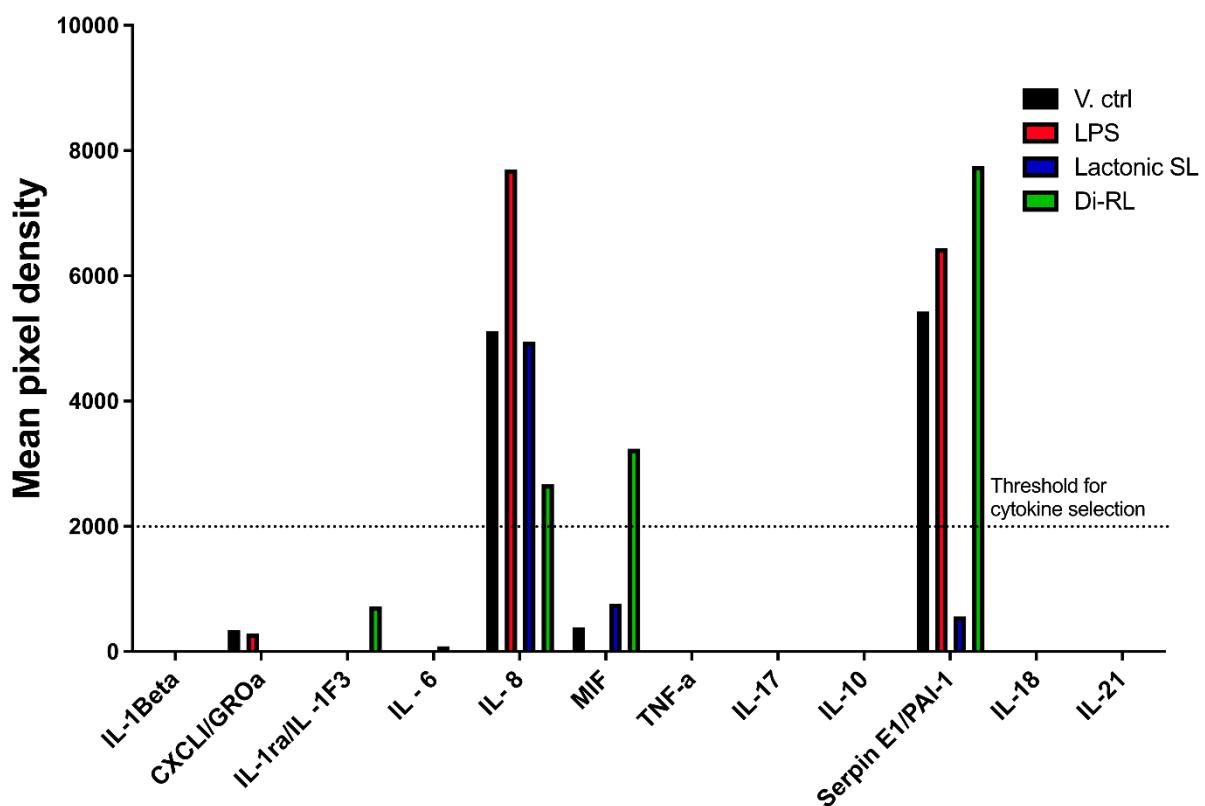
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137 Fig. S4
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146 **Supplementary figure legends**

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148 **Fig. S1** Chemical structure of (a) Lactonic SL (di-acetylated with hydroxy fatty acid tail esterified at the 6' or 6''
149 ends); (b) Acidic SL (non-acetylated with free fatty acid tail); (c) mono-RL (one rhamnose as hydrophilic moiety);
150 (d) di-RL (two rhamnose as hydrophilic moiety) and (e) SLES. Figure adapted from Banat et al., 2010 and created
151 with ChemSketch.com.

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153 **Fig. S2** The effects of Acidic SL and Mono-RL on the viability of human keratinocytes at treatment concentrations
154 between 100 and 500 $\mu\text{g mL}^{-1}$. Data are the mean results of three independent experiments with each treatment
155 group having six technical replicates, error bars represent standard error from the mean. Statistical significance
156 was determined using a one-way ANOVA followed by Dunnett's Multiple comparison Test (* = $p \leq 0.05$).

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158 **Fig. S3** The use of AO/PI dual staining technique to assess the morphological pattern of cell death induced by
159 surfactant treatment. Fluorescent microscopy images at 200 \times magnification of untreated HaCaT cells (Untreated
160 control); cells treated with 1% methanol (v/v) (Vehicle control); preparations of each glycolipid congener and
161 SLES at 20 and 100 $\mu\text{g mL}^{-1}$. For Acidic SL and Mono-RL, treatment concentration was increased up to 500 μg
162 mL^{-1} . Three images per well were randomly selected and processed with ImageJ Software for scale bar attachment
163 ; scale bar was set at 100 μm .

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165 **Fig. S4** Semi-quantitative pro-inflammatory cytokine profile of cell-free supernatant samples from HaCaT cells
166 treated with 1% methanol (v/v) (V. ctrl); 25 $\mu\text{g mL}^{-1}$ LPS (positive control for the assay); or LD₅₀ concentrations
167 of Lactonic SL and Di-RL.

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126 **Supplementary tables**

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128 **Table S1** Sequences of primer sets and product length of pro/ anti-inflammatory markers used for RT-qPCR analysis. Primers were designed using NCBI Primer-BLAST
129 software and sequences of primers were compared to the NCBI nucleotide data bases via BLASTn Search tool.

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Primer	Gene	Sequence (5' – 3')	Amplicon Length (bp)	Annealing Temperature (°C)
GAPDH SA-F	<i>GAPDH</i>	TCG ACA GTC AGC CGC ATC TT	80	49
GAPDH SA-R	<i>GAPDH</i>	CTC CGA CCT TCA CCT TCC CC	80	53
IL-8 SA-F	<i>CXCL8</i>	ACA CTG CGC CAA CAC AGA AA	117	47
IL-8 SA-R	<i>CXCL8</i>	TTC TCA GCC CTC TTC AAA AAC TTC	117	49
IL-RA SA-F	<i>IL1RN</i>	CCA AGA AGG CAA GAG CAA GCA	103	49
IL-RA SA-F	<i>IL1RN</i>	GCC CCA GCA GTT TAT GGG TT	103	49

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133 **Table S2** Critical micelle concentration (CMC) determination of microbial glycolipids and SLES. Surface activity
134 of glycolipids and SLES were analysed using KRUSS K10 ST digital tensiometer. CMC was estimated from 135
extrapolated intercepts of values plotted on both the X and Y -axes of XY graph. All surface tension measurements
136 were taken in triplicate.

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Surfactants	CMC (mg mL ⁻¹)	Surface tension (mN m ⁻¹)
Acidic SL	0.06	43.00
Lactonic SL	0.06	35.37
Mono - RL	0.03	41.60
Di-RL	0.06	28.70
SLES	0.66	32.20

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