

Anandamide alters the membrane properties, halts the cell division and prevents drug efflux in multidrug resistant *Staphylococcus aureus*

Shreya Banerjee, Ronit Vogt Sionov, Mark Feldman, Reem Smoum,
Raphael Mechoulam, and Doron Steinberg

Suppl. Table 1. Primers used for real-time PCR of *S. aureus*

Gene	Forward Primer	Reverse Primer	Ref
16S rRNA	CCAGCAGCCGCGGTAAT	CGCGCTTTACGCCAATA	1
<i>agrA</i>	AAAGTTGCAGCGATGGATTT	ATGGGCAATGAGTCTGTGAG	1
<i>agrC</i>	CATTCGCGTTGCATTTATTG	CCTAAACCACGACCTTCACC	2
<i>atl</i>	GACCCTGCTATTGTCCAACC	CGCTGATTGATTAGCACGAG	3
<i>aur</i>	CAAAAGAGTGATGCGGTCAA	AGGTGCATGAACACCATCAA	2
<i>capA</i>	CAGTTAAAGTCGCACCAA	GAACCCAATACAGGCAAT	1
<i>clfA</i>	ACCCAGGTTTCAGATTCTGGCAGCG	TCGCTGAGTCGGAATCGCTTGCT	1
<i>cna</i>	AATAGAGGCGCCACGACCGT	GTGCCTTCCCAAACCTTTTGAGCA	1
<i>crtM</i>	ATCCAGAACCACCCGTTTTT	GCGATGAAGGTATTGGCATT	2
<i>crtN</i>	GATGAAGCTTTGACGCAACA	TTCGCATGATACGTTTGCTC	2
<i>divIB</i>	CGACTCGATTGATGAGGAAAC	TTGTGCTTACGTCTTAACTTCC	3
<i>divIC</i>	ATGCGTGTTGTTTCGTAGGC	CGCTCCTGTGCATCAATATC	3
<i>divIVA</i>	GATCGCGTTTCCGTATGTTAG	TCAAGCGTCACTTGTTGAGC	3
<i>ebps</i>	GGTGCAGCTGGTGCAATGGGTGT	GCTGCGCCTCCAGCCAAACCT	1
<i>eno</i>	TGCCGTAGGTGACGAAGGTGGTT	GCACCGTGTTTCGCTTCGAACT	1
<i>ezrA</i>	ATTGGTGAGGCAGCAAGTC	GCTATATGGTTGTGCGCTTG	3
<i>fib</i>	CGTCAACAGCAGATGCGAGCG	TGCATCAGTTTTTCGCTGCTGGTTT	1
<i>fmhB</i>	AAGCGAGGTACGACAGTAGAACG	CATCTCCATCTTCATGCAACGCA	4
<i>fnbA</i>	AAATTGGGAGCAGCATCAGT	GCAGCTGAATTCCCATTTTC	1
<i>fnbB</i>	AAGAAGCACCGAAAACGTGTG	TCTCTGCAACTGCTGTAACG	2
<i>ftsA</i>	GGCGAGAAATTTCACAATGG	TGTGTCTTTGATTGCTTGTCG	3
<i>ftsH</i>	GATTGATGCTGTTGGTCGTC	CCGAAACCATCCATTTCAA	3
<i>ftsL</i>	GATACGCGAGGAAAGATTGC	ATCGTTCTCAAGGCTCATCC	3
<i>ftsZ</i>	AAAGCTGCAGAGGAATCTCG	TTTAGCAACGACTGGTGCTG	3
<i>ftsW</i>	CTTGTGTGTGTTGGGATTGC	TTGATGATCCACCAAAGCTG	3
<i>glyA</i>	CTACAAACTCACAGCCAC	GTATCGGAAGCGGTTATG	5
<i>gmk</i>	CCATCTGGAGTAGGTAAGG	CTACGCCATCAACTTCAC	5
<i>gtf</i>	TGGTGACGCCGAAGGACTC	GCAGCACGAGCAGGAACAC	1
<i>gyrA</i>	TGGCCCAAGACTTTAGTTATCGTTATCC	TGGGGAGGAATATTTGTAGCCATACCTAC	6
<i>gyrB</i>	GGTGCTGGGCAAATAACAAGT	TCCCACACTAAATGGTGCAA	2
<i>hla</i>	AATGAATCCTGTCGCTAATGCCGC	CTGAAGGCCAGGCTAAACCACCTT	1
<i>hld</i>	TAATTAAGGAAGGAGTGATTTCAATG	TTTTTAGTGAATTTGTTCACTGTGTC	1
<i>icaA</i>	ACACTTGCTGGCGCAGTCAA	TCTGGAACCAACATCCAACA	7
<i>icaB</i>	ATGGTCAAGCCCAGACAGAG	AGTATTTTCAATGTTTAAAGCA	7
<i>icaC</i>	CTTGGGTATTTGCACGCATT	GCAATATCATGCCGACACCT	1
<i>kdpA</i>	ATTGTTCCGTTTATTGTCC	CATCATACTGCCATTTCT	1
<i>lytA</i>	TGGTGGTGCAAAGTTCATTC	TATCTGCCAGCGAATGTC	3
<i>lytM</i>	TCAGCAAGTAAAGCGACAGC	TTTCAGGCATTGCATAGTCG	3
<i>lytN</i>	TGCCAATGACACCATTAGTAGAAC	ACCGTCGAAATCCCATCC	3
<i>mecA</i>	AACCGAAGATAAAAAAGAACC	GTCCGTAACCTGAATCAGC	1
<i>mecR1</i>	GTGCTCGTCTCCACGTTAATTCCA	GACTAACC GAAGAAGTCGTGTCAG	4
<i>mdeA</i>	CTTTCAGGTTACCTTGTGAATATTTAAAC	ATCAATAGGTACTTTAATTGTAGTTCCAAC	1
<i>mepA</i>	ATGGTATAGGTTTCTTGTACTGGTATG	AATGATAATTGCACCTTGTAATAATGGC	1
<i>mgrA</i>	AGTACAATCTAACATACC	TTGCGATAAAGAAGAAGC	1
<i>norA</i>	GACATTTACCAAGCCATCAA	TGCCATAAATCCACCAATCC	1

<i>norB</i>	GCTACACCATCAACAGATACAGCAA	ACTCAATGCGACGCCAAA	1
<i>norC</i>	TGGGTTGGAGATGGATTTTC	ACAATTAGCCCTGCAACGTC	1
<i>nuc</i>	GCGATTGATGGTGATACGGTT	AGCCAAGCCTTGACGAACTAAAGC	2
<i>opp1C</i>	CGTGCCTGTGATGTTATGTTGGCA	ITCGGCACCCATTCCAAACAATGC	4
<i>oppF</i>	GGTGGGCAACGTCAGCGTGTA	GGACACTGCCTCGTCGCAAACA	4
<i>pbp2</i>	CGTTTCTACGAACATGGCGCAC	GGCACCTCAGAACCAAATCCACCA	4
<i>proC</i>	GGCAGGTATTCCGATTGA	CCAGTAACAGAGTGTCCAAC	5
<i>psmα</i>	TATCAAAAGCTTAATCGAACAATTC	CCCCTTCAAATAAGATGTTTCATATC	2
<i>qacA/B</i>	GGTGCTTTAATAATGCC	CCAGTCCAATCATGCCTGC	1
<i>recF</i>	AGTTATAGACACGGCACG	GCGTCGTCTTATTTGAGG	5
<i>rho</i>	GGAAGATACGACGTTTCAGAC	GAAGCGGGTGGAAAGTTTA	5
<i>RNAII</i>	TATGAATAAATGCGCTGATGATATACCACG	TTTTAAAGTTGATAGACCTAAACCACGACC	6
<i>RNAIII</i>	GCACTGAGTCCAAGGAACTAACTCT	AGCCATCCCAACTTAATAACCATGT	2
<i>rpoB</i>	TCCTGTTGAACGCGCATGTAA	GCTGGTATGGCTCGTGATGGTA	8
<i>saeR</i>	AAGTGGCGACCATTACAT	CATTATTGCCTCAAATACGT	1
<i>saeS</i>	TGCCAATACCTTCATCGCTAA	CAATATCGAACGCCACTTGA	7
<i>sarA</i>	GACATACATCAGCGAAAA	TACGTTGTTGTGCATTAA	1
<i>sdrM</i>	CAACATGGCATTGGTTATTCTAC	ACAGCTGTTGGTTTAATAAAGC	1
<i>sepA</i>	GAAGTATGTACGATAACCTATTATATTATG GC	AAAGTCGCGCCTCTAAAATATGC	1
<i>sepF</i>	GCGGTACTGTTTATGCAATCG	TGGTCTGTAATGCTTCCAGCTAC	3
<i>sle1</i>	TCAGGATCTGCAACAACGAC	CCTTTACCAATTTTCAGCACGAC	3
<i>spsB</i>	CCAGCAAATTGTTGTTGTGCTAG	AAGCCAAAGCCGATTCACTC	7

References

1. Salaheen, S., Peng, M., Joo, J., Teramoto, H. & Biswas, D. Eradication and sensitization of methicillin resistant *Staphylococcus aureus* to methicillin with bioactive extracts of berry pomace. *Front Microbiol* **8**, 253, doi:10.3389/fmicb.2017.00253 (2017).
2. Valliammai, A. *et al.* 5-Dodecanolide interferes with biofilm formation and reduces the virulence of Methicillin-resistant *Staphylococcus aureus* (MRSA) through up regulation of agr system. *Sci Rep* **9**, 13744, doi:10.1038/s41598-019-50207-y (2019).
3. Wassmann, C. S., Højrup, P. & Klitgaard, J. K. Cannabidiol is an effective helper compound in combination with bacitracin to kill Gram-positive bacteria. *Sci Rep* **10**, 4112, doi:10.1038/s41598-020-60952-0 (2020).
4. Chung, P. Y., Chung, L. Y. & Navaratnam, P. Transcriptional profiles of the response of methicillin-resistant *Staphylococcus aureus* to pentacyclic triterpenoids. *PLoS One* **8**, e56687, doi:10.1371/journal.pone.0056687 (2013).
5. Theis, T., Skurray, R. A. & Brown, M. H. Identification of suitable internal controls to study expression of a *Staphylococcus aureus* multidrug resistance system by quantitative real-time PCR. *J Microbiol Methods* **70**, 355-362, doi:10.1016/j.mimet.2007.05.011 (2007).
6. Eleaume, H. & Jabbouri, S. Comparison of two standardisation methods in real-time quantitative RT-PCR to follow *Staphylococcus aureus* genes expression during *in vitro* growth. *J Microbiol Methods* **59**, 363-370, doi:10.1016/j.mimet.2004.07.015 (2004).
7. Sethupathy, S., Vigneshwari, L., Valliammai, A., Balamurugan, K. & Pandian, S. K. L-Ascorbyl 2, 6-Dipalmitate inhibits biofilm and virulence in methicillin-resistant *Staphylococcus aureus* and prevents triacylglyceride accumulation in *Caenorhabditis elegans*. *RSC Adv* **7**, 23392-23406 doi:10.1039/c7ra02934a (2017).
8. Baroja, M. L. *et al.* The SaeRS Two-Component System Is a Direct and Dominant Transcriptional Activator of Toxic Shock Syndrome Toxin 1 in *Staphylococcus aureus*. *J Bacteriol* **198**, 2732-2742, doi:10.1128/jb.00425-16 (2016).