

Supplementary Materials

Supplementary Table 1: Bacterial Strains and Growth Media

Strain	Growth Media
<i>B. cepecia</i>	Cation-adjusted Mueller-Hinton Broth II
<i>E. coli</i> WT BW25113	Luria Broth
<i>E. coli</i> CFT073	Luria Broth
<i>C. difficile</i>	Reinforced Clostridial Medium
<i>P. acnes</i>	Reinforced Clostridial Medium
<i>H. influenza</i>	Cation-adjusted Mueller-Hinton Broth II
<i>V. cholerae</i>	Luria Broth
<i>A. baumannii</i> AB17978	Luria Broth
<i>A. baumannii</i> AB5075	Luria Broth
<i>E. coli</i> UPEC CFT073	Gutnick Minimal Media
<i>E. coli</i> UPEC J96	Gutnick Minimal Media
<i>E. coli</i> BW25113	Gutnick Minimal Media
<i>E. coli</i> NCM3722	Gutnick Minimal Media
<i>S. aureus</i> MRSA COL	Luria Broth
<i>S. aureus</i> MRSA USA300	Luria Broth
<i>M. fortuitum</i>	Cation-adjusted Mueller-Hinton Broth II
<i>N. gonorrhoeae</i>	Cation-adjusted Mueller-Hinton Broth II
<i>S. aureus</i> VanA	Cation-adjusted Mueller-Hinton Broth II
<i>S. aureus</i> VISA	Cation-adjusted Mueller-Hinton Broth II
<i>S. aureus</i> MRSE	Cation-adjusted Mueller-Hinton Broth II
<i>S. epidermidis</i>	Cation-adjusted Mueller-Hinton Broth II
<i>B. subtilis</i> W168	Luria Broth
<i>E. coli</i> IptD4213	Luria Broth
<i>E. faecium</i>	Cation-adjusted Mueller-Hinton Broth II

<i>S. pneumoniae</i>	Cation-adjusted Muller-Hinton Broth II with 5% lysed horse blood
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Supplementary Table 2: The 14 features evaluated in BCP analysis

Features of BCP Analysis
Cell Area
Cell Length
Cell Width
Cell Eccentricity
Cell Perimeter
Nucleoid Area
Ratio of Nucleoid to Cell
Nucleoid Eccentricity
DNA Length
DNA Width
DNA Perimeter
Mean Sytox intensity
Mean FM4-64 intensity
Mean Dapi intensity

Supplementary Figure Legends

Figure S1. SCH-79797 is bactericidal against *S. aureus*. Colony forming units (CFU ml⁻¹) after 3-hour treatment of *S. aureus* MRSA USA300 with 1% DMSO, 1X MIC SCH-79797 and 5X MIC novobiocin. Each data point represents 3 independent samples and 3 technical replicates. Mean ± s.d. are shown.

Figure S2. SCH-79797 is an effective antibiotic in an infection model of *G. mellonella* by *A. baumannii*. A-B. The percent survival of non-infected *G. mellonella* wax worms after treatment with 2µl/larva of 100% DMSO, 67µg/larva SCH-79797, 6µg/larva gentamicin, and

67µg/larva rifampicin. Data in (A) represents a typical cohort (n = 12) from a biological triplicate and the pooled results are presented in (B). Mantel-cox statistics for the cohort were calculated with PRISM. C. The percent survival of *A. baumannii* infected *G. mellonella* wax worms after treatment with 67µg/larva SCH-79797, 6µg/larva gentamicin, and 67µg/larva rifampicin. Data represents the pooled results from a biological triplicate. D. The average survival of drug-treated *A. baumannii* infected *G. mellonella* wax worms relative to larvae treated with DMSO. Data represents the pooled results from a biological triplicate.

Figure S3. SCH-79797 is not prone to resistance in both *S. aureus* and *A. baumannii*. A. Fold increase in resistance of *S. aureus* MRSA USA300 to SCH-79797, novobiocin, trimethoprim, and nisin after 25 days of serial passaging in 0.5X MIC of each drug and plotted on a log₂ scale. Resistance was confirmed by remeasuring MIC's from aliquots of each passage that were collected and stored at -80°C. B. Fold increase in resistance of *A. baumannii* AB17978 to SCH-79797 and gentamicin after 5 days of serial passaging in 0.5X MIC of each drug and plotted on a log₂ scale. Resistance was confirmed as described above. The experiment was performed in duplicate and identical results were found in both cases.

Figure S4. Thermal stability of DHFR increases after SCH-79797 and Trimethoprim treatment. A-B. The relative thermal stability of DHFR after treatment of whole cell and cell lysate samples with (A) SCH-79797 and (B) trimethoprim. Changes in thermal stability were determined by measuring changes in the abundance of DHFR across 10 different temperatures ranging from 42-72°C and 4 drug concentrations and a vehicle control.

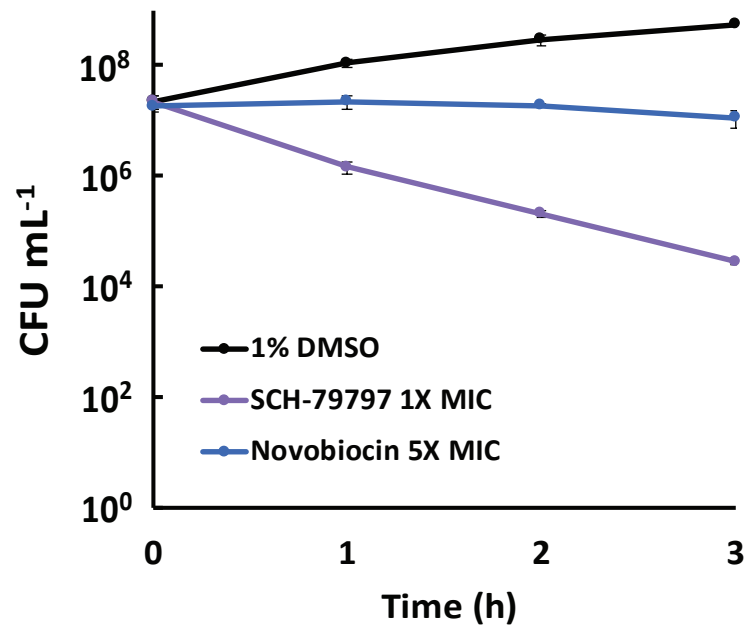
Figure S5. CRISPRi mutants not involved in folate metabolism are not sensitized to SCH-79797. A. The growth of CRISPRi *B. subtilis* knockdown mutants relative to a DMSO-treated control after SCH-79797 treatment. Bacterial growth was measured for 14h and the final optical density (OD₆₀₀) of each condition was plotted against drug concentration. Each data point represents 2 independent replicates. Mean ± s.d. are shown.

Figure S6. Treatment with ampicillin, rifampicin, and novobiocin does not disrupt membrane integrity. A. Flow cytometry analysis of the membrane potential and permeability of *E. coli* IptD4213 cells after 15 min. incubation with 2X MIC ampicillin, rifampicin, novobiocin.

Figure S7. SCH-79797 disrupts *B. subtilis* W168 membrane integrity. A. Flow cytometry analysis of the membrane potential and permeability of *B. subtilis* W168 cells after 15 min. incubation with 1% DMSO, 1X MIC SCH-79797, 2X MIC SCH-79797.

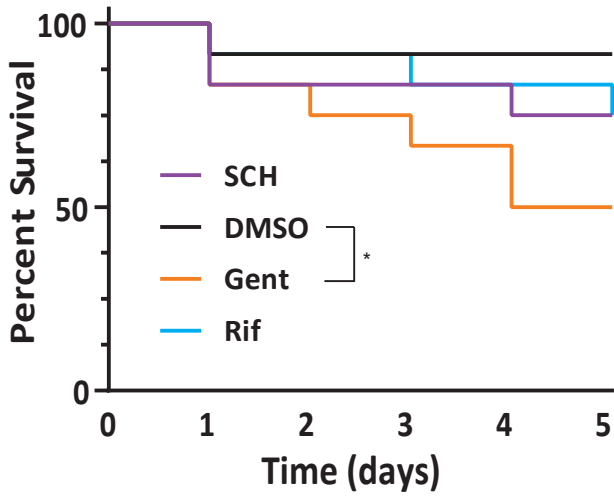
Supplementary Figure 1

A

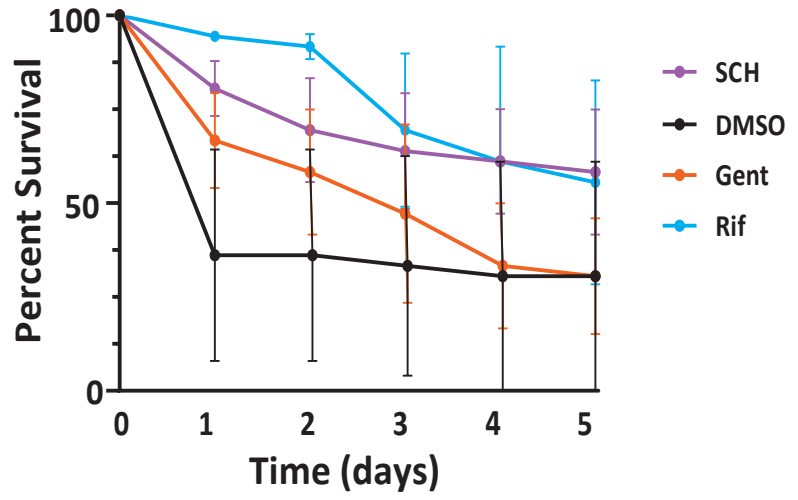


Supplementary Figure 2

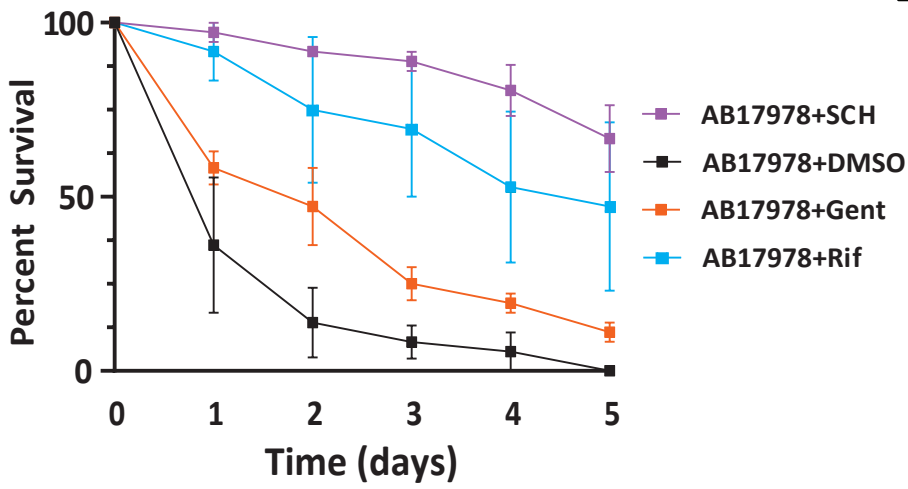
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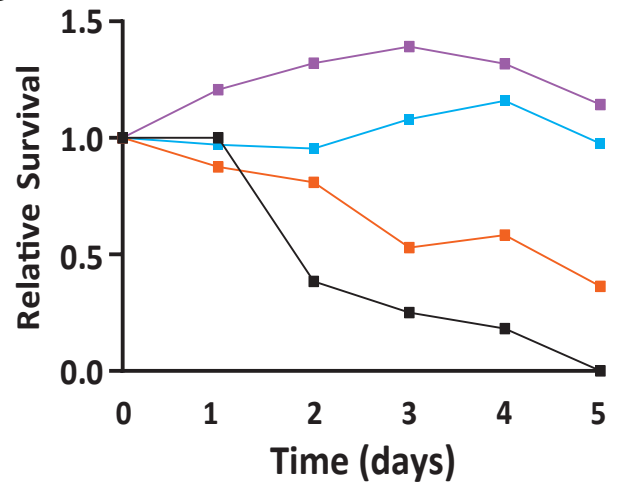
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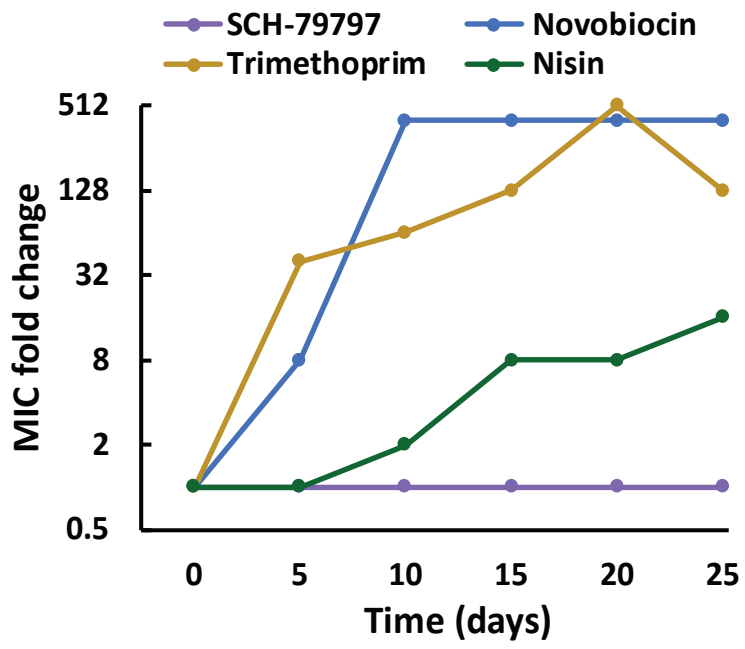
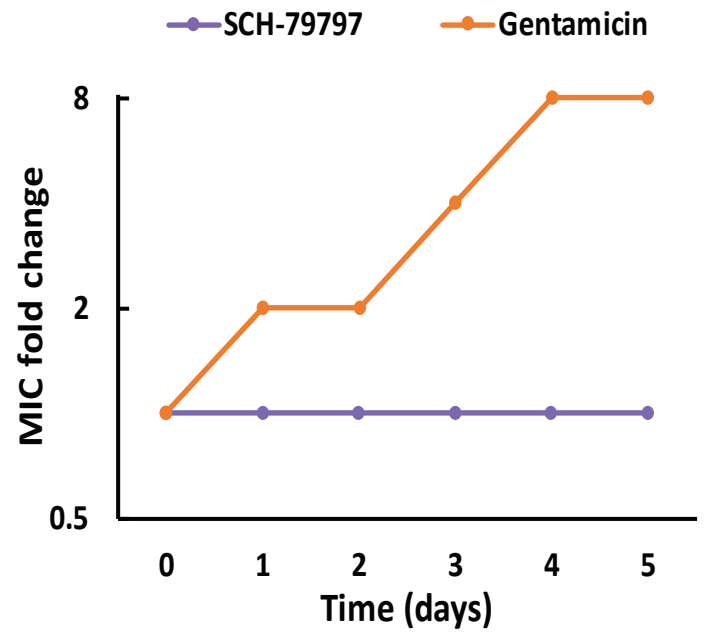
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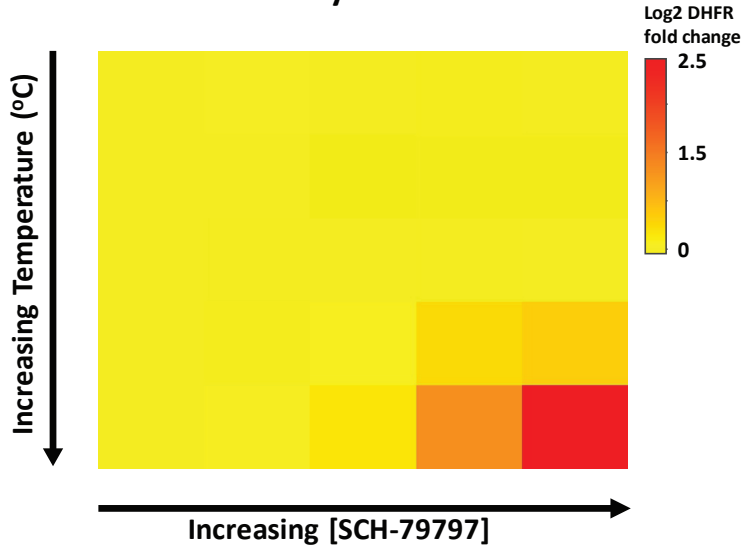
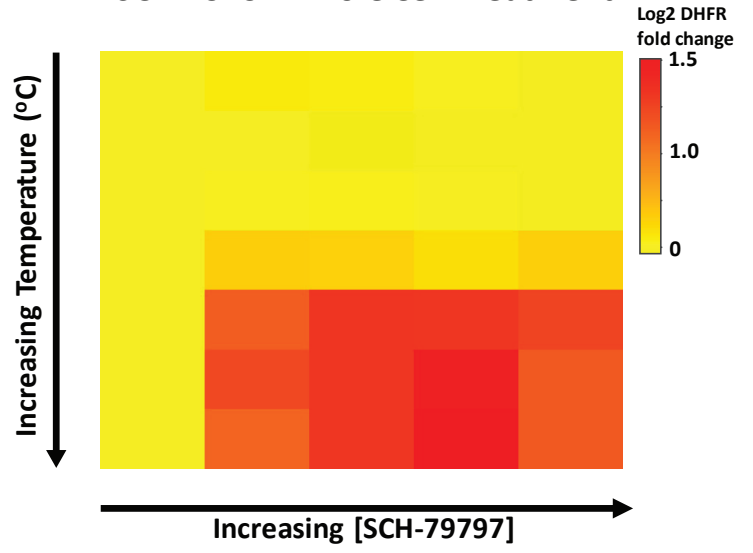
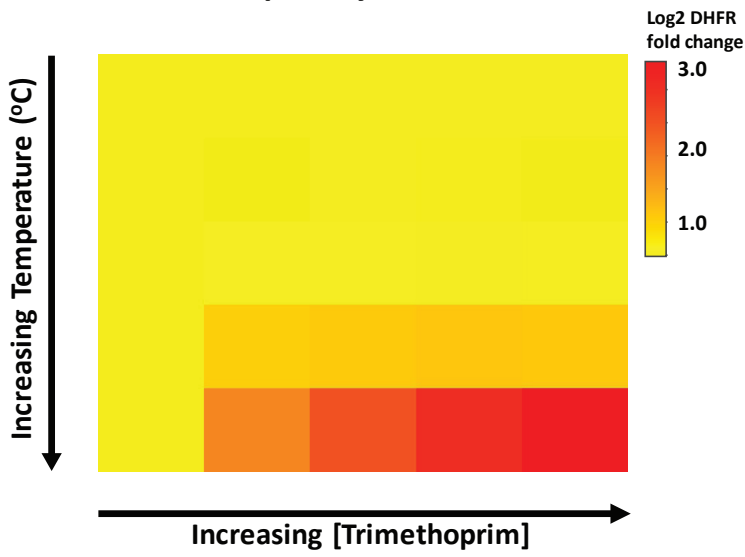
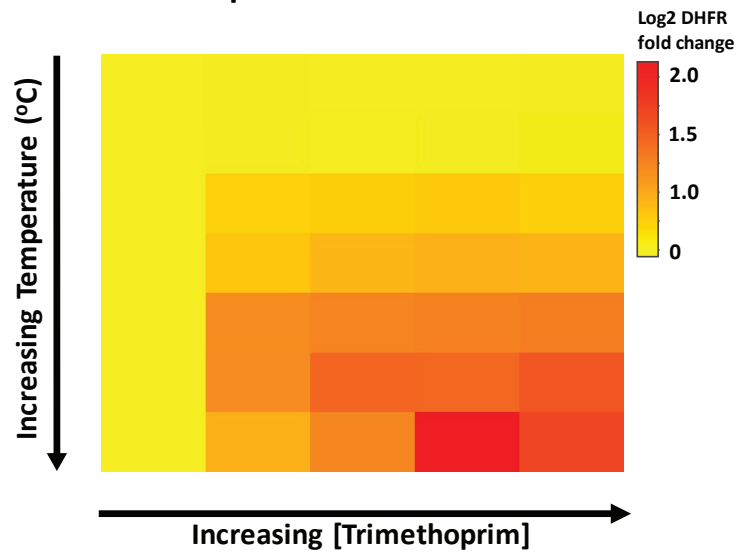
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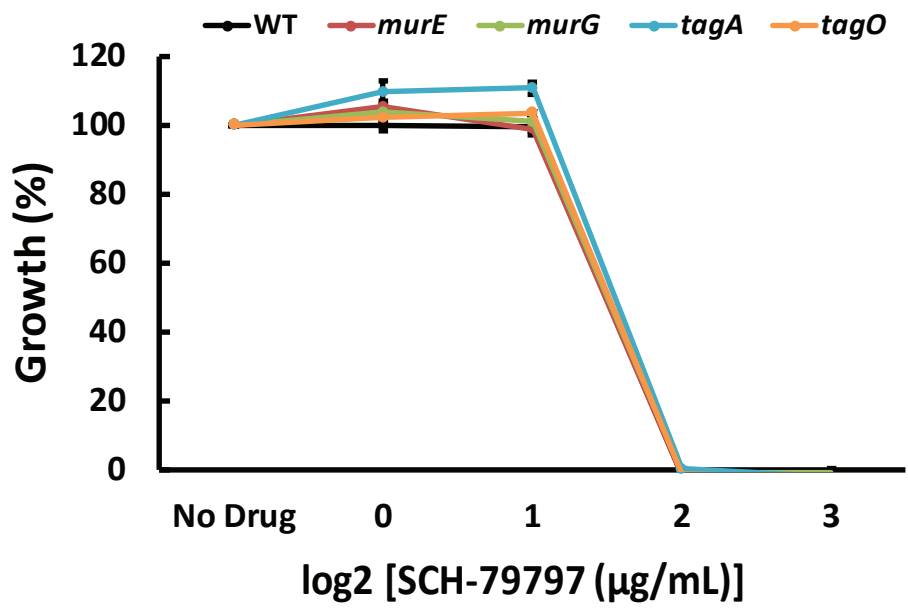
Supplementary Figure 3

A**B**

Supplementary Figure 4

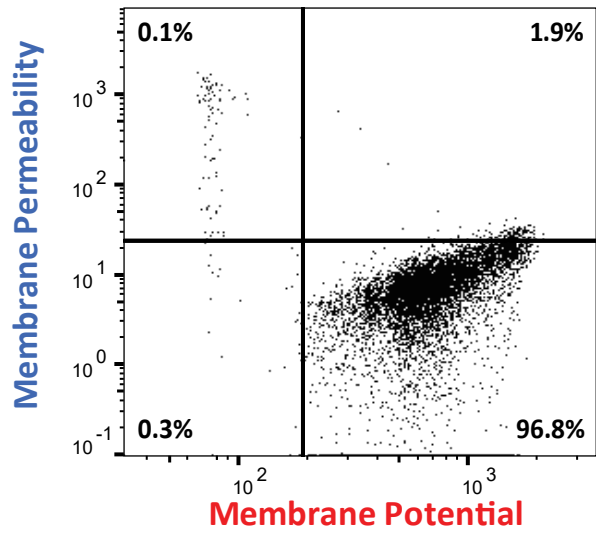
A**SCH-79797 Lysate Treatment****SCH-79797 Whole Cell Treatment****B****Trimethoprim Lysate Treatment****Trimethoprim Whole Cell Treatment**

Supplementary Figure 5

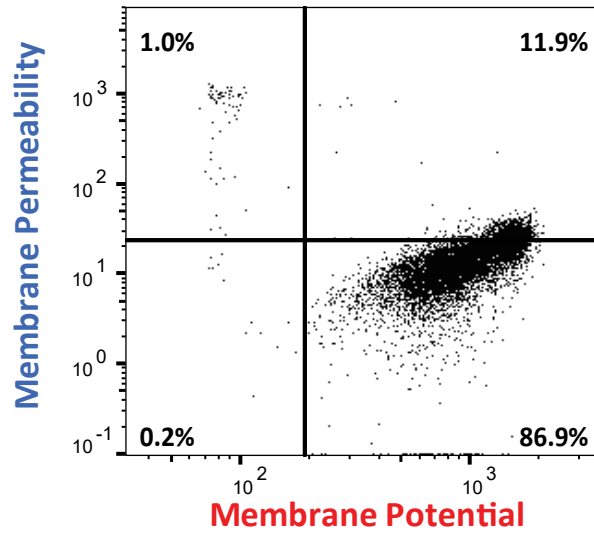


Supplementary Figure 6

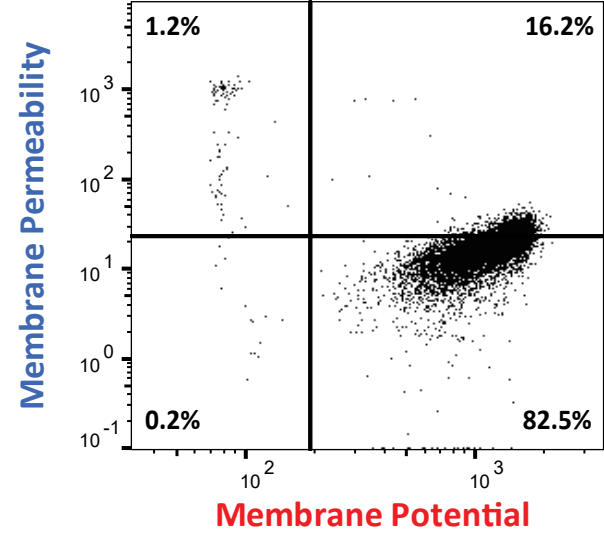
Ampicillin 2X MIC



Rifampicin 2X MIC



Novobiocin 2X MIC



Supplementary Figure 7

