

1 **Title**

2 Plasma-sensitive *E. coli* mutants reveal plasma resistance mechanisms

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10 **Key words (max. 6)**

11 disinfection, atmospheric pressure plasma, iron-sulphur cluster, antibacterial mechanism,
12 stress response, non-thermal plasma

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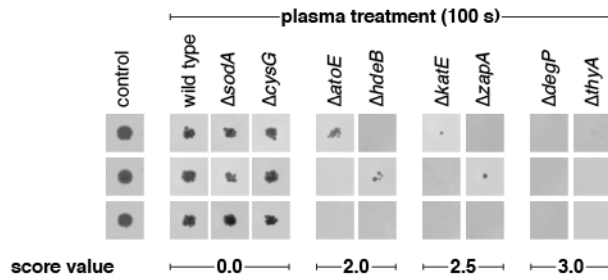
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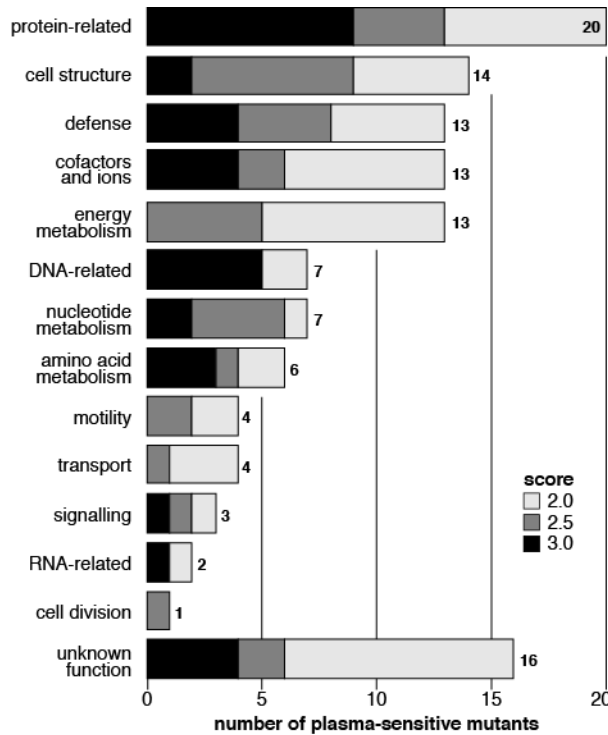
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21 **Supplementary Material**



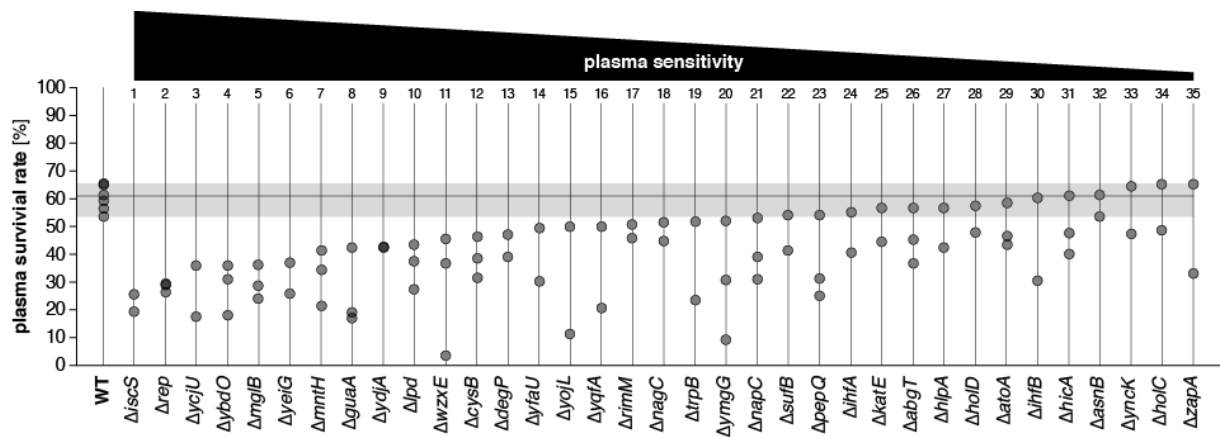
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 23 **Fig. S1** Examples of the grading of plasma sensitivity in the screening of the KEIO collection using the spot assay.
 24 Cells of an overnight culture were spotted onto LB agar (2 μ l) and exposed to the μ APPJ for 100 s. After incubation
 25 at 37 $^{\circ}$ C for 16 h, growth or the absence of growth was recorded. The wild type treated with gas only served as
 26 control. The results of three independent experiments were used (top, middle, bottom rows).



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 30 **Fig. S2** GO terminology of the 87 identified genes.

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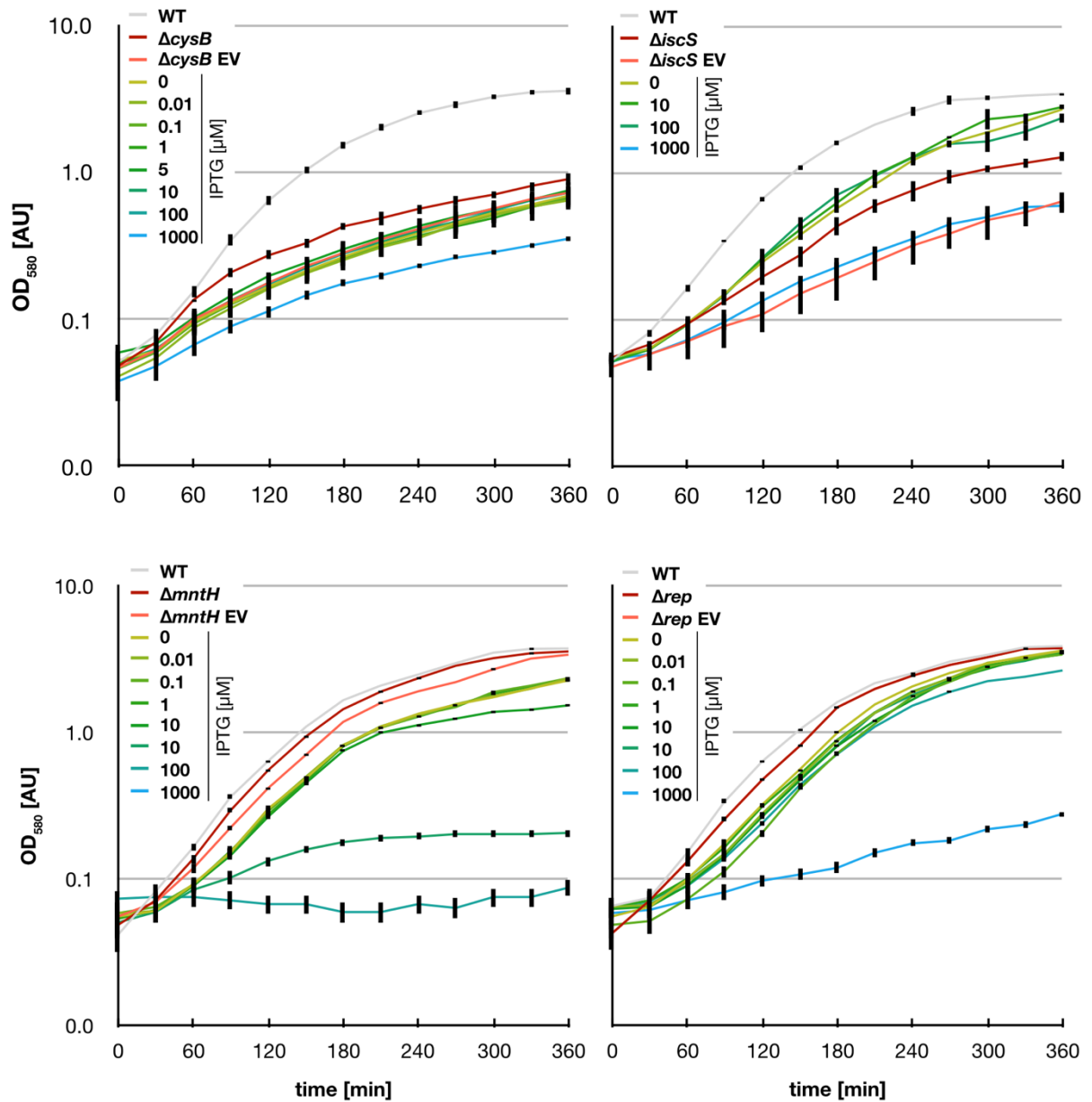


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34 **Fig. S3** Quantitation of plasma sensitivity. Low-density cell suspensions were placed onto a filter paper and
 35 exposed to the effluent of the μ APPJ with or without ignition of the plasma. After resuspending the cells in saline
 36 solution they were plated for CFU determination after overnight incubation. Data are shown for mutants with
 37 increased sensitivity compared to the wild type. Each data point represents one independent replicate.

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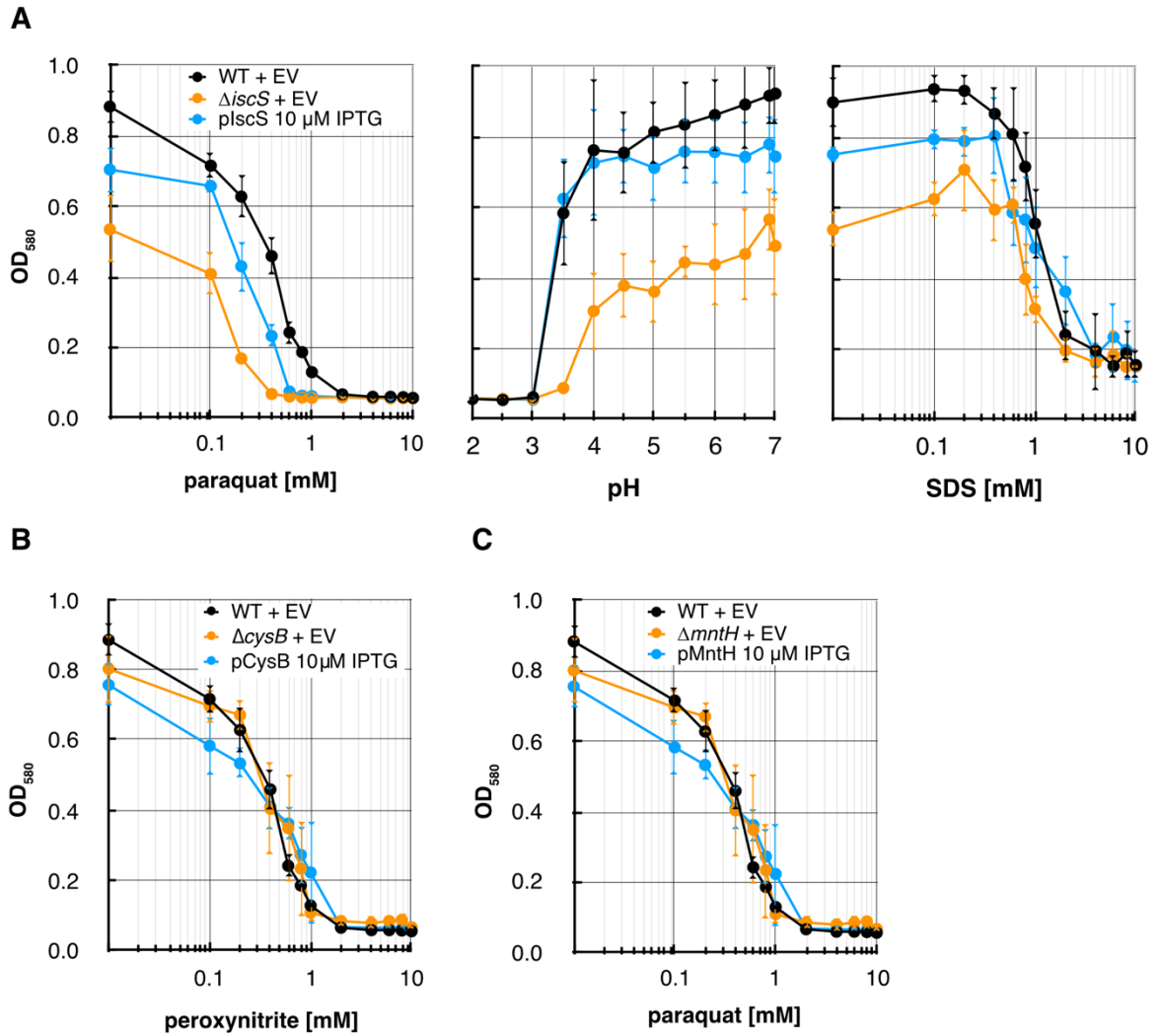
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41 **Fig. S4** Growth curves of the wild type, the four deletions strains analysed in detail, the deletion strains
 42 harbouring the empty vector pCA24N (EV), and the complemented strains with induction at different IPTG
 43 concentrations. The data reflect three independent biological experiments. Shown are the averages and standard
 44 deviations.

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47 **Fig. S5** Sensitivity of the complemented strains. Optical densities at 580 nm of the wild type, the deletion strains
 48 harbouring the empty vector pCA24N (EV), and the corresponding complemented strains are shown. Cells were
 49 incubated at different stressor concentrations for 16 h. Complementation was performed by induction with (A)
 50 10 μ M IPTG (*iscS*), (B) 1 μ M IPTG (*cysB*), or (C) 0.1 μ M IPTG (*mntH*). Data represent three independent
 51 experiments.

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