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Author Correction: Reprogramming *Escherichia coli* for the production of prenylated indole diketopiperazine alkaloids

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In Figure 2, panel 2C should not be shown. The correct Figure 2 appears below as Fig. 1.

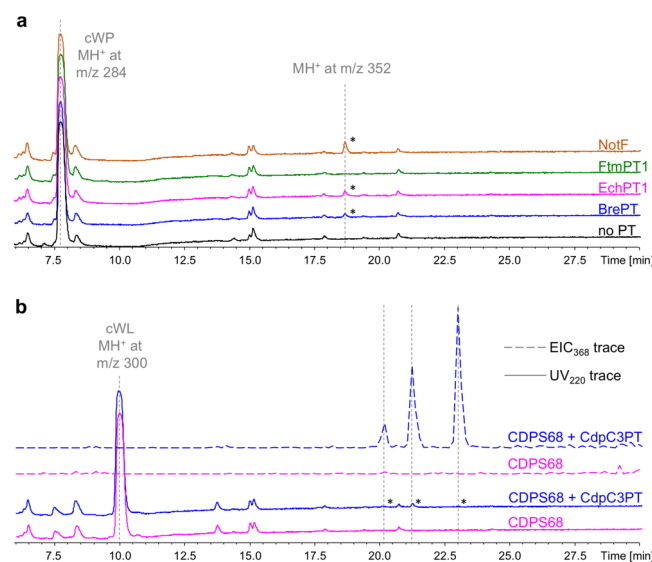


Figure 1. LC-MS/MS analysis of metabolite production by recombinant *E. coli*. Samples corresponding to 50 μ l of culture supernatants were analysed. (a) SPE-treated bacterial supernatants of cultures of BL21AI expressing CDPS74 and BrePT (blue), EchPT1 (pink), FtmPT1 (green), NotF (orange) or CDPS74 alone (black) were analysed. UV traces recorded at 220 nm are shown between 6 and 30 min with the absorbance scale set from 0 to 700 mU. Asterisks highlight specific peaks for which the MS data are indicated. (b) SPE-treated bacterial supernatants of cultures of BL21AI expressing CDPS68 and CdpC3PT (blue) or CDPS68 alone (pink) were analysed. UV chromatograms recorded at 220 nm (UV₂₂₀, plain lines) and extracted ion current at m/z 368 (EIC₃₆₈, dotted lines) are shown between 6 and 30 min. The Y-axis of the UV₂₂₀ traces was set from 0 to 700 mU and that of the EIC₃₆₈ traces from 0 to 3,380,000.



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