1 Supplementary Figure 1: E. coli BW25113 fluorescence signals for 143 fluorophores. 2 The signals for the non-dye control (Autofluorescence) and all fluorophores (log₁₀ values, Y-3 axis) are compiled for each of the 13 channels of the Intellicyt® flow cytometer (X-axis). The 4 median for autofluorescence (orange bars) is drawn for every channel. Colour coded 5 distribution of data by concentration of fluorophore (micromolar) are shown. The subtitles 6 contain our internal code (Fxxx). The number of biological replicates (*n*) for each fluorophore at a given concentration represent data from different experiments carried out over a period 7 8 of five months.

9 Supplementary Figure 2. Compilation of the fluorescence detected for 143 fluorescent molecules by the Intellicyt® in E. coli BW25113. A: Density distribution of the 10 11 fluorescence of the full set of fluorophores (blue dash line) using their individual optimal 12 excitation-emission channels at their optimal concentrations (as in Supplementary Table 1). 13 **B**: The signals for all fluorophores are compiled against autofluorescence for each channel. 14 Colour-coded distribution of data by concentration of fluorophore (micromolar) are shown. Orange bar: median value of autofluorescence (note that some dyes can quench 15 16 autofluorescence). C: Density distribution after autofluorescence data were subtracted. The 17 median fluorescence was reduced (blue dash line moved to the left) as a consequence of 18 the subtraction of the autofluorescence. D: New spread of the signals distribution for all fluorophores for the 13 channels after subtracting autofluorescence. 19 Colour-coded 20 distribution of data by concentration of fluorophore (micromolar).

Supplementary Figure 3: Dose-response fluorophore uptake. Regression analysis for each of the initial 47 fluorophores that had a two-fold or higher activity (in terms of median values of uptake) in *E. coli* BW25113. Locally weighted scatterplot smoothing was applied to fluorescence signal versus fluorophore concentration data. X-axis: the log₁₀ of the concentrations in micromolar. Y-axis: the log₁₀ of fluorescence signals. The subtitles contain the relevant channel for each fluorophore and our internal code (Fxxx).

Supplementary Figure 4: Effect of pH and CPZ on the uptake of 40 fluorophores. *E. coli* BW25113 (10^6 cells.mL⁻¹) were incubated with 3 µM fluorophores for 15 minutes at 37°C (red lines). Similar samples were incubated with both 3 µM fluorophores and 10µM CPZ under the same experimental conditions (blue lines). The incubations were carried out at different pH values: 6.0, 6.5, 7.0, 7.5, 8.0, and 8.5. X-axis: pH values. Y-axis: log_{10} of fluorescence signals after locally weighted scatterplot smoothing.

33 Supplementary Figure 5: Effect of yhjV deletion on fluorophore uptake in E. coli 34 **BW25113**. Fluorescence signals are presented as the log₁₀ of ratios of the fluorescence from the *yhjV* knock out cells ($\Delta yhjV$) over the fluorescence signals from the reference cells 35 (BW25113). X-axis: \log_{10} of the ratios ($\log_{10} = 0$). No difference between the two strain 36 37 would have a ratio of 1 ($\log_{10} = 0$). Boxplots are ordered by median (highest on top of the 38 plot). The legend lists the concentrations used for each fluorophore. The legend lists the 39 concentrations used for each fluorophore. The clearly related structures of the two most 40 differentiating dyes from the palette of 39 are shown.

41 Supplementary Figure 6: Differential uptake of 39 dyes between three different 42 knockout strain of *E. coli*. Experiments were performed as described in the legends to 43 Supplementary Figure 5 and 11 Figure 9, and in Table 2. Several dyes showing the largest 44 effect in $\Delta yihN$ are marked.



















Supplementary Figure 1

Supplementary Figure 3

Supplementary Figure 5 $\Delta yhjV$

Concentration (µM) 📫 2.5 📫 5.0 📫 7.5 📫 10.0

Signal ratio

Signal Ratio