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## Supplementary Material for Random Design of Microfluidics

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## Role of diffusion constant in chip behavior

To determine whether solutes with different diffusion coefficients still behave as predicted in our randomly-designed microfluidic chips, we simulated each of the 10,513 random chip designs using three different model solutes: Na<sup>+</sup> (representative of ions), fluorescein (representative of small molecules), and bovine serum albumin (BSA; representative of proteins). The simulation results from all three solutes are summarized in Supplementary Figure 1. The similarity of the results from each solute confirm that our library of randomly-designed chips can be used to predict the behavior of a wide range of different solutes from ions to proteins.



Supplementary Figure 1: Fluid velocity vs. concentration for each outlet in each of the 10,513 chip designs in our simulation library, using the diffusion coefficients of three different solutes: Na<sup>+</sup> ( $1.33 \times 10^{-9} \text{ m}^2/\text{s}$ ), fluorescein ( $4.25 \times 10^{-10} \text{ m}^2/\text{s}$ ), and bovine serum albumin (BSA;  $6.38 \times 10^{-11} \text{ m}^2/\text{s}$ ). Each solute behaves similarly in each chip design, meaning that our randomly-designed chips will still function as predicted regardless of the specific solute used.