

# Supporting Information

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## SI Materials and Methods

Flat-bottom, nonsterile, clear polystyrene 96-well plates were from Thermo Fisher Scientific. Decanoic acid was from Fluka/Sigma, glucose from Thermo Fisher Scientific, xylose from Calbiochem/EMD Millipore, phosphate-buffered saline (PBS) from Mediatech, Inc., and hypoxanthine from Acros Organics. All other chemicals were from Sigma. All solutions were prepared in 18 M $\Omega$ -cm water.

**Decanoic Acid Solutions.** Decanoic acid was dissolved, with heating, in 190 mM NaOH to yield a 180-mM solution. This stock then was diluted to obtain 80 mM decanoic acid with or without 100 mM bicine (diluted from a 1-M stock solution) or 100–300 mM NaCl (diluted from a 4-M stock solution). The pH was adjusted by adding HCl, typically from 0.5- or 1-M solutions.

**Imaging Vesicles.** All samples contained 10  $\mu$ M rhodamine 6G and were placed between two coverslips sealed with vacuum grease. In experiments involving temperature changes, the bottom coverslip was coupled with thermal paste (Omega Engineering) to the microscope stage. Temperature control of the stage was achieved with a Wavelength controller connected to a Peltier device and a thermistor temperature probe with a manufacturer-quoted accuracy of 0.02  $^{\circ}$ C (Wavelength Electronics). Epifluorescence microscopy was performed with a 60 $\times$  or 10 $\times$  air objective on a Nikon Y-FL microscope with a CoolSNAP HQ CCD camera (Photometrics).

**Dialysis.** Adenine, uracil, or thiouracil was dissolved to 15 mM in solutions of either 180 mM decanoic acid/190 mM NaOH or 180 mM acetic acid/190 mM NaOH. Solutions were brought to the same temperature and then titrated with HCl to the same pH, typically about 7.8. Next, 0.4 mL of each titrated solution was placed in a Slide-A-Lyzer with a cutoff of 3.5 kDa (Pierce/Thermo Fisher Scientific). Two solutions were dialyzed side by side in beakers containing 200 mL of a solution of 180 mM acetic acid/190 mM NaOH adjusted to pH 7.8. Samples of 200  $\mu$ L were withdrawn from the dialysis buffer at intervals of 1 or 2.5 min, dried down with a centrifugal vacuum evaporator (SPD121P SpeedVac; Thermo Fisher Scientific), and resuspended in 30  $\mu$ L water. The absorbance of each sample at 260 nm was measured with a NanoDrop 1000 spectrophotometer (Thermo Fisher Scientific).

Absorbance data were fit to a linear function, with best-fit slopes and associated fit uncertainties determined using MATLAB software (MathWorks). Uncertainties stated for individual experiments (i.e., the two discrete experiments shown in Fig. 1B) reflect only the uncertainty in the fit. We set the criterion that differences in slopes are significant only if they are greater than zero with a probability of  $P < 0.05$  (equivalent to a difference in slope at least twice as large as the experimental uncertainty). For example, it is significant that the rate of release of adenine was  $24 \pm 5\%$  lower from decanoic acid than from acetic acid, because  $24/5 \sim 5$ , i.e., a 5-sigma change. Similarly, it is insignificant that the rate of release of uracil was  $8 \pm 7\%$  greater from decanoic acid than from acetic acid, because  $8/7 \sim 1$ . Uncertainties quoted in the main text for the mean difference between the rates of release from decanoic vs. acetic acids, over several experiments, account for both the uncertainty in fit and variation among replicates.

**Langmuir Trough Studies.** In Langmuir monolayer studies, the surface pressure is determined using a balance to measure forces on

a plate of filter paper or platinum partially immersed in the subphase. The force on the plate results from three components: the buoyant force of water, force of gravity, and surface tension of water. The only one of these terms that changes during our experiments is the surface tension. Our studies were performed as in ref. 1 using a NIMA trough with a subphase temperature of 22  $^{\circ}$ C. The subphase contained either 10 mM or no (control) adenine in PBS, which was prepared from a 10 $\times$  stock solution; the inclusion of adenine did not change the pH detectably. Stearic acid in chloroform was deposited at the air–water interface using a Hamilton syringe. Ample time (10 min) was allowed for chloroform to evaporate before data were taken.

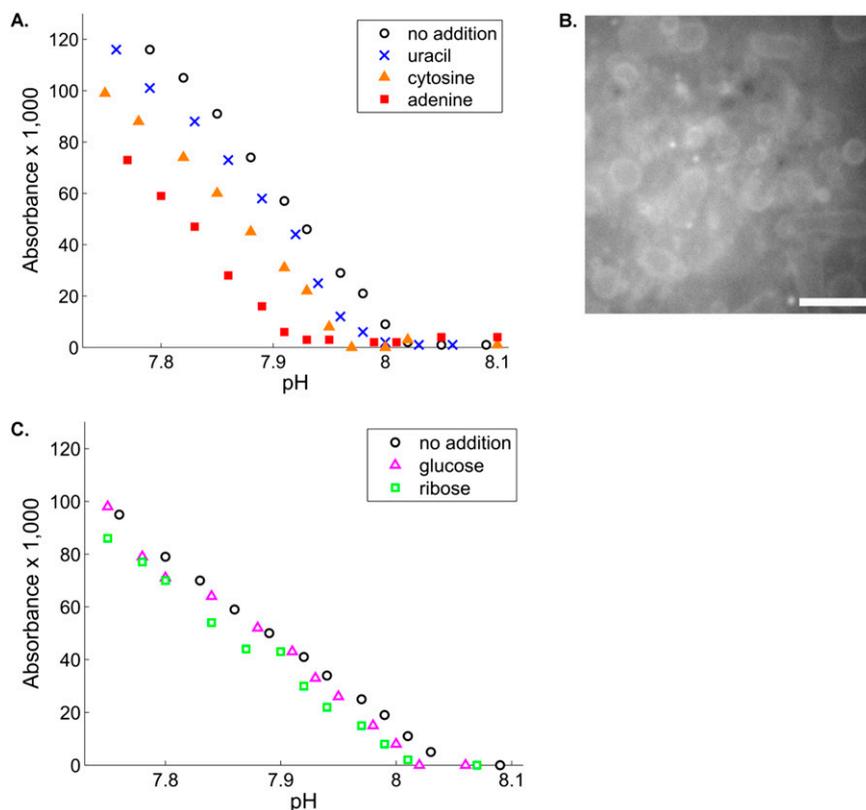
**Filtration Assay.** Bases were dissolved in 180 mM decanoic acid/pH 8.25; at the concentrations used,  $\leq 0.3$  mM, the bases did not alter pH detectably. Solutions of guanine and xanthine, which have low solubility, were centrifuged at 3,000  $\times g$  for 10 min in conical-bottomed tubes, and the supernates were used for the assay. All base solutions then were treated equally. Typically, 2 mL was placed in an Amicon Ultra-4 3K filter (Millipore) and centrifuged at 3,000  $\times g$  for 10 min in a Sorvall Legend RT swinging-bucket centrifuge (Thermo Fisher Scientific). Aliquots of the starting solution (taken before centrifugation), the retentate (after gentle agitation to dislodge aggregates on surfaces), and the filtrate then were measured for absorbance at 280 nm for 2,6-diaminopurine, 300 nm for 2-aminopurine, 250 nm for hypoxanthine, 242 nm for pyrimidine, or 260 nm in all other cases. To confirm that decanoic acid micelles are retained in the retentate, we used a pinacyanol chloride assay for aggregated lipids (2); we found that the retentate contained over 10-fold more aggregated decanoic acid than the filtrate.

The decrease in concentration of base in the filtrate relative to the starting material was used as the measure of base retained with the micelles. This decrease generally was of the same magnitude as the increase in base concentration in the retentate, and was more reproducible. This agreement provides evidence against nonspecific loss of base on the surfaces of the centrifuge tube and filter. Further evidence against nonspecific loss came from control experiments in which bases were dissolved in a 20-mM decanoic acid solution, which is below the critical micelle concentration. The amount of base retained in these experiments generally was only 0–2% of the starting concentration, as expected if retention of base in the experiments with 180 mM decanoic acid is primarily the result of binding to micelles.

**NaCl-Induced Flocculation in Test Tubes.** NaCl was added to 80-mM decanoic acid/pH 7.60–7.65 solutions to a final concentration of 300 mM by diluting from a 4-M stock solution, and the solutions were vortexed briefly immediately after the addition. Including 30 mM adenine in the solutions altered the pH only slightly, lowering it about 0.03 units, and adenine inhibited flocculation equally well when bicine was included to eliminate pH changes. A Canon PowerShot SD600 camera was used to take photographs of samples within test tubes in a rack on a black mat. In the case of samples above room temperature, photographs were taken promptly upon removal from the indicated temperature.

**NaCl-Induced Flocculation in 96-Well Plates.** Bases and sugars were dissolved in a solution of 80 mM decanoic acid/100 mM bicine/pH 7.9. The inclusion of bicine ensured that the effects of bases or sugars on flocculation were not the result of changes in pH. Typically, 19  $\mu$ L of 4 M NaCl was added to 231  $\mu$ L of the test





**Fig. 52.** Nucleobases and ribose lower the pH at which vesicles form in decanoic acid solutions when the pH is decreased by titrating with HCl. (A) Nucleobases at 30 mM lower the pH below which 80 mM decanoic acid/100 mM NaCl forms vesicles. The density of vesicles is related to the turbidity of the solution (the absorbance at 490 nm). Results shown are representative of 10 experiments with adenine and 4 with cytosine and uracil. (B) Epifluorescence microscopy shows that the turbidity corresponds with the presence of vesicles. Here, a solution of 80 mM decanoic acid/100 mM NaCl was titrated with HCl to pH 7.8, and 10  $\mu$ M rhodamine 6G was added for imaging. Scale bar, 10  $\mu$ m. (C) Ribose at 120 mM lowers the pH required for 80 mM decanoic acid/100 mM NaCl to form vesicles, whereas glucose has minimal effect. The density of vesicles is related to the turbidity of the solution (the absorbance at 490 nm). Results shown are representative of four experiments with ribose and three with glucose.





