## **Supporting Information**

## Effect of micellar and reverse micellar interface on solute location: 2,6-pyridinedicarboxylate in CTAB micelles and CTAB and AOT reverse micelles

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## **Supporting Information Available**

**Figure S1.** The 400 MHz <sup>1</sup>H NMR spectra of H<sub>3</sub>dipic<sup>+</sup> cation in CF<sub>3</sub>COOD and D<sub>2</sub>O mixture. Weight percent of CF<sub>3</sub>COOD is shown next to each spectrum. Spectra were referenced against internal DSS. Hammett acidity functions ( $H_0$ ) for non-deutarated CF<sub>3</sub>COOH and H<sub>2</sub>O mixtures with identical weight percentages shown in the figure are equal to 0.0 (12%), -0.5 (46%), -1.0 (74%), -1.5 (81%), -2 (86%), and -2.8 (96%).<sup>55</sup> The actual  $H_0$  values may be different due to isotopic effects.

Figure S2. pH-variable UV spectra of 0.1 mM of  $H_2$ dipic in 30 mM NaCl (a), and 4 mM of CTAB (b).

**Figure S3.** Molar absorbtivity of pyridine dicarboxylic acids in monocationic, neutral, monoanionic and dicationic form. Data taken from Ref.29.

Figure S4. The 300 MHz <sup>1</sup>H NMR spectrum of 20 mM CTAB in H<sub>2</sub>O with proton assignment.

**Figure S5.** <sup>1</sup>H chemical shift of dipic protons is shown as a function of aqueous stock solution pH for CTAB micellar solutions ( $H_a$  and  $H_b$ ) and aqueous solution ( $H^{aq}_a$  and  $H^{aq}_b$ ). Data points for micellar solutions were taken from spectra shown in Figure 1 and aqueous dipic chemical shifts adjusted for the common reference standard taken from Ref. 8.

**Figure S6.** <sup>1</sup>H chemical shift of dipic protons is shown as a function of aqueous stock solution pH for CTAB reverse micellar solutions ( $H_a$  and  $H_b$ ) and aqueous solution ( $H^{aq}_a$  and  $H^{aq}_b$ ). Data points for reverse micellar solutions were taken from spectra shown in Figure 11 and aqueous dipic chemical shifts adjusted for the common reference standard taken from Ref. 8.

**Figure S7.** <sup>1</sup>H NMR chemical shifts of dipic<sup>2-</sup> as a function of CTAB concentration. The data were obtained from spectra shown in Figure 5.

**Table S1.** <sup>1</sup>H NMR chemical shifts of dipic<sup>2-</sup> and H<sub>2</sub>dipic in various media.

Table S2. Sizes of the micellar and reverse micellar systems.

## References

- (61) Spitzer, U. A.; Toone, T. W.; Stewart, R., Can. J. Chem. 1976, 54, 440-447.
- (62) Dorshow, R.; Briggs, J.; Bunton, C.A.; Nicoll, D.F., J. Phys. Chem. 1982, 86, 2388-2395.
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**Figure S1.** The 400 MHz <sup>1</sup>H NMR spectra of H<sub>3</sub>dipic<sup>+</sup> cation in CF<sub>3</sub>COOD and D<sub>2</sub>O mixture. Weight percent of CF<sub>3</sub>COOD is shown next to each spectrum. Spectra were referenced against internal DSS. Hammett acidity functions ( $H_0$ ) for non-deutarated CF<sub>3</sub>COOH and H<sub>2</sub>O mixtures with identical weight percentages shown in the figure are equal to 0.0 (12%), -0.5 (46%), -1.0 (74%), -1.5 (81%), -2 (86%), and -2.8 (96%).<sup>61</sup> The actual  $H_0$  values may be different due to isotopic effects.



**Figure S2.** pH-variable UV spectra of 0.1 mM of  $H_2$ dipic in 30 mM NaCl (a), and 4 mM of CTAB (b).



**Figure S3.** Molar absorbtivity of pyridine dicarboxylic acids in monocationic, neutral, monoanionic and dicationic form. Data were obtained from a table in the Ref. 29 and used to construct this figure.



Figure S4. The 300 MHz <sup>1</sup>H NMR spectrum of 20 mM CTAB in H<sub>2</sub>O with proton assignment.



**Figure S5.** <sup>1</sup>H chemical shift of dipic protons is shown as a function of aqueous stock solution pH for CTAB micellar solutions ( $H_a$  and  $H_b$ ) and aqueous solution ( $H^{aq}_a$  and  $H^{aq}_b$ ). Data points for micellar solutions were taken from spectra shown in Figure 3 and aqueous dipic chemical shifts were taken from Ref 8. The 0.10 ppm were subtracted from chemical shifts in aqueous solution to adjust to the common reference.



**Figure S6.** <sup>1</sup>H chemical shift of dipic protons is shown as a function of aqueous stock solution pH for CTAB reverse micellar solutions ( $H_a$  and  $H_b$ ) and aqueous solution ( $H^{aq}_a$  and  $H^{aq}_b$ ). Data points for reverse micellar solutions were taken from spectra shown in Figure 7 and aqueous dipic chemical were taken from Ref. 8. The 0.10 ppm were subtracted from chemical shifts in aqueous solution to adjust to the common reference.



**Figure S7.** <sup>1</sup>H NMR chemical shifts of dipic<sup>2-</sup> as a function of CTAB concentration. The data were obtained from spectra shown in Figure 4a.



**Figure S8.** <sup>1</sup>H NMR chemical shifts of Hdipic<sup>-</sup> as a function of CTAB concentration. The data were obtained from spectra shown in Figure 4b.

System	δ(H <sub>a</sub> ),	$\delta(H_b),$
	ppm	ppm
Micellar and reverse micellar		
Na <sub>2</sub> dipic in H <sub>2</sub> O/0.1 mM CTAB (pH 6.6)	7.92 <sup><i>a</i></sup>	$7.92^{a}$
Na <sub>2</sub> dipic in H <sub>2</sub> O/20 mM CTAB (pH 6.8)	$7.98^{a}$	7.66 <sup><i>a</i></sup>
Na <sub>2</sub> dipic in H <sub>2</sub> O (pH 6.7)/0.1 M CTAB/0.5 M 1-pentanol, $w_0$ =6	8.10 <sup>b</sup>	$7.87^{b}$
Na <sub>2</sub> dipic in H <sub>2</sub> O (pH 6.7)/0.1 M CTAB/0.5 M 1-pentanol, $w_0$ =20	$8.07^{b}$	$7.88^{b}$
Aqueous		
Na <sub>2</sub> dipic in H <sub>2</sub> O (pH 6.4)	8.005 <sup>c</sup>	8.027 <sup>c</sup>
[N(CH <sub>3</sub> ) <sub>4</sub> ] <sub>2</sub> dipic in H <sub>2</sub> O (pH 7)	8.00 <sup>c</sup>	8.00 <sup>c</sup>
H <sub>2</sub> dipic in H <sub>2</sub> O (pH 1.3)	8.27 <sup>c</sup>	8.38 <sup>c</sup>
Organic solvent		
Na <sub>2</sub> dipic in CD <sub>3</sub> OD	8.10 <sup>b</sup>	7.93 <sup><i>b</i></sup>
[N(CH <sub>3</sub> ) <sub>4</sub> ] <sub>2</sub> dipic in CD <sub>3</sub> OD	7.91 <sup><i>b</i></sup>	$7.82^{b}$
H <sub>2</sub> dipic in CD <sub>3</sub> OD	8.23 <sup><i>b</i></sup>	8.38 <sup>b</sup>

**Table S1.** <sup>1</sup>H NMR chemical shifts of dipic<sup>2-</sup> and H<sub>2</sub>dipic in various media

<sup>*a*</sup>referenced against coaxial capillary with C<sub>6</sub>D<sub>6</sub> and TMS, 35°C; <sup>*b*</sup>referenced against internal TMS, 24°C; <sup>*c*</sup>referenced against internal DSS, 24°C.

**Table S2.** Sizes of the micellar and reverse micellar systems.

System Diameter		Aggregation	
	(nm)	number	
CTAB micelles	5.8 <sup><i>a</i></sup>	~100 <sup><i>a,b</i></sup>	
CTAB reverse micelles w <sub>0</sub> =6	$5.0^{c}$	-	
AOT reverse micelles w <sub>0</sub> =6	$5.6^{d}$	$50^d$	

<sup>*a*</sup>Ref. 62, <sup>*b*</sup>ref. 63 at 35°C, <sup>*c*</sup>ref. 38, <sup>*d*</sup>ref. 64.