Supplementary Materials



Figure S1. TIEs of mixed standards (including cytosine; uracil; thymine; adenine; hypoxanthine; guanine; xanthine; thymidine; cytidine; uridine; cordycepin; adenosine; 2'-deoxyguanosine; inosine and guanosine.) with different ammonium acetate concentrations. (a) 20 mM, (b) 25 mM, (c) 30 mM, (d) 35 mM, (e) 40 mM. The concentration of each diluted standard was 40 μ g/mL.



Figure S2. Cont.



Figure S2. TIEs of mixed standards at different pH values. (a) pH9.5, (b) pH9.6, (c) pH 9.7,
(d) pH 9.8, (e) pH 9.9, (f) pH 10.0, (g) pH 10.1, (h) pH 10.2, (i) pH10.3, (j) pH10.4,
(k) pH10.5. The concentration of each diluted standard was 40 μg/mL.



Figure S3. TIEs of mixed standards with different ratio of organic modifier (methanol) in buffer. (a) 0%, (b) 0.5%, (c) 1%, (d) 2%, (e) 5%, (f) 10%. The concentration of each diluted standard was 40 μ g/mL.

Peak #	Compound ID	Formula	$[M+H]^+$	[M+Na] ⁺	[2M+H] ⁺	[M + K] ⁺	Product Ions
1	cytosine	$C_4H_5N_3O$	112.0507	-	223.0939	-	-
2	uracil	$C_4H_4N_2O_2$	113.0348	135.0171	-	151.0606	-
3	thymine	$C_5H_6N_2O_2$	127.0504	-	-	-	-
4	adenine	$C_5H_5N_5$	136.0618	-	-	-	-
5	hypoxanthine	$C_5H_4N_4O$	137.0459	159.0274	-	175.0018	-
6	guanine	$C_5H_5N_5O$	152.0569	174.0384	-	190.0123	-
7	xanthine	$C_5H_4N_4O_2$	153.0405	175.0230	-	-	-
8	thymidine	$C_{10}H_{14}N_2O_5$	243.0976	265.0799	-	281.0536	$[M-C_5H_8O_3+H]^+$ 127.0504
9	cytidine	$C_{9}H_{13}N_{3}O_{5}$	244.0933	266.0756	-	-	-
10	uridine	$C_{9}H_{12}N_{2}O_{6}$	245.0776	267.0589	-	283.0327	-
11	cordycepin	$C_{10}H_{13}N_5O_3$	252.1092	274.0914	-	-	-
12	adenosine	$C_{10}H_{13}N_5O_4$	268.1037	290.0860	-	-	-
13	2'-deoxyguanosine	$C_{10}H_{13}N_5O_4$	268.1037	290.0860	-	306.0598	$[M-C_5H_8O_3+H]^+$ 152.0567
14	inosine	$C_{10}H_{12}N_4O_5$	269.0881	291.0702	-	307.0436	-
15	guanosine	$C_{10}H_{13}N_5O_5$	284.0987	306.0753	-	322.0539	[M+H-C ₅ H ₈ O ₄] ⁺ 152.0569

Table S1. CE-ESI-TOF/MS accurate mass measurements of 15 nucleobase and nucleoside compounds.

Note: Extracted ion electropherograms (EIEs) of 15 nucleobase and nucleoside standards are shown in Figure 1. Peak eluting earlier to peak 3 (in Figure 1) is a fragmentation ion of peak 8 (in Figure 1). The exact masses of peak 3 ($[M+H]^+$) and the fragmentation ion of peak 8 ($[M-C_5H_8O_3+H]^+$) were consistent with the same elemental composition of $C_5H_7N_2O_2$. Peak eluting after peak 9 (in Figure 1) is the C_{13} isotope peak of peak 8 ($[M+H]^+$). The exact masses of peak 9 ($[M+H]^+$) and the C_{13} isotope peak of peak 8 ($[M+H]^+$) were similar.