

**Supplemental Information**

**Quantitative Studies of an RNA Duplex Electrostatics by Ion Counting**

**Magdalena Gebala and Daniel Herschlag**

## Supplemental Information

# Quantitative studies of an RNA duplex electrostatics by ion counting

Magdalena Gebala<sup>1</sup>, Daniel Herschlag<sup>1,2,3,\*</sup>

<sup>1</sup> Department of Biochemistry, Stanford University, Stanford, CA, 94305 USA

<sup>2</sup> Department of Chemistry, Stanford University, Stanford, CA, 94305 USA

<sup>3</sup> ChEM-H Institute, Stanford University, Stanford, CA, 94305 USA

## Overview of the Supplementary Information

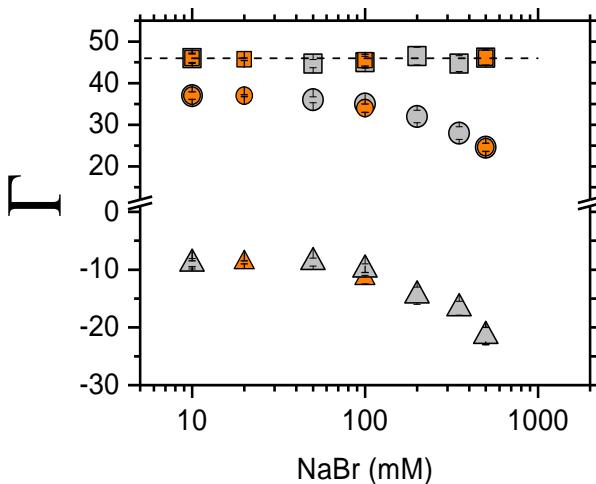
In this supplementary information, we provide a table with preferential ion interaction coefficients  $\Gamma_i$  (e.g. the number of associated ions,  $i = \text{Na}^+$  or  $\text{Br}^-$ ), around 24-bp RNA, 24-bp and 23-bp DNA (Table S1) for NaBr; a table summarizing the fraction of charge neutralization from attraction of  $\text{Na}^+$  measured by ASAX and BE-ICP-MS around dsRNA and dsDNA (Table S2); a table with preferential ion interaction coefficients from competition experiments between  $\text{Na}^+:\text{Mg}^{2+}$  and  $\text{Cs}^+:\text{Mg}^{2+}$  around 24-bp RNA. The results are consistent with observations in the main text and support the conclusions described therein.

**Table S1:** Experimentally determined preferential interaction coefficients ( $\Gamma_i$ ) for NaBr around 24-bp RNA, 24-bp DNA, and 23-bp DNA

C [M]	24 bp RNA			24 bp DNA			23 bp DNA		
	$\Gamma_{\text{Na}^+}$	$\Gamma_{\text{Br}^-}$	total	$\Gamma_{\text{Na}^+}$	$\Gamma_{\text{Br}^-}$	total	$\Gamma_{\text{Na}^+}$	$\Gamma_{\text{Br}^-}$	total
0.01	39 ± 1.0	-6.0 ± 0.7	45.0 ± 1.2	37.0 ± 1.0	-9.0 ± 0.5	46.0 ± 1.0	36.0 ± 0.2	-8.0 ± 0.3	44.0 ± 0.3
0.02	39.5 ± 0.5	-6.5 ± 1.0	46.0 ± 1.0	37.0 ± 0.2	-8.75 ± 0.2	46.0 ± 0.3	35.0 ± 0.3	-8.6 ± 0.2	44.0 ± 0.3
0.10	37.0 ± 1.0	-8.0 ± 1.2	45.0 ± 1.6	34.0 ± 1.0	-11.5 ± 1.0	45.5 ± 1.4	32.0 ± 1.0	-12.0 ± 1.0	44.0 ± 1.0
0.12	36.0 ± 1.0	-10.0 ± 1.5	46.0 ± 1.8	-	-	-	-	-	-
0.26	-	-	-	-	-	-	27.0 ± 1.5	-17.5 ± 1.0	44.5 ± 1.8
0.50	31.0 ± 1.5	-13 ± 2.0	44.0 ± 2.5	24.6 ± 1.0	-21.5 ± 1.5	46.0 ± 1.8	18.0 ± 0.5	-26 ± 0.5	44.0 ± 0.5
0.65	-	-	-	-	-	-	-	-	-

**Table S2.** Interaction coefficients for NaBr around 24-bp DNA obtained previously in reference (3)

C [M]	NaBr		
	$\Gamma_{Na^+}$	$\Gamma_{Br^-}$	total
0.010	$37.0 \pm 0.9$	$-9.0 \pm 0.9$	$46 \pm 1.3$
0.050	$36.0 \pm 0.7$	$-8.7 \pm 0.7$	$44.7 \pm 1.0$
0.100	$35.0 \pm 1.0$	$-10 \pm 1.0$	$45 \pm 1.4$
0.200	$32.0 \pm 1.5$	$-14.5 \pm 1.5$	$46.5 \pm 2.0$
0.350	$28.0 \pm 1.5$	$-16.7 \pm 1.2$	$44.7 \pm 2.0$
0.500	$24.6 \pm 1.0$	$-21.5 \pm 1.5$	$46.1 \pm 1.8$



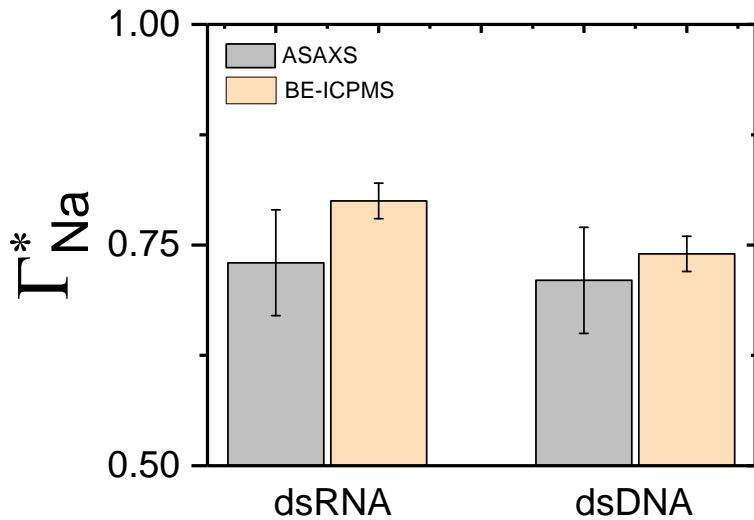
**Figure S1.** Comparison of current (orange symbols) and previous (grey symbols) ion counting results for association of NaBr around 24-bp DNA from BE-ICP-MS measurements. Data point in grey are from reference (3) and values are given in Table S2.

**Table S3:** Experimentally determined fraction of charge neutralization ( $\Gamma_{Na}^*$ ) for  $Na^+$  around dsRNA and dsDNA at 100 mM monovalent salt concentration.

C [M]	dsRNA		ds DNA	
	$\Gamma_{Na}^{*ASAXS}$	$\Gamma_{Na}^{*BE-ICPMS}$	$\Gamma_{Na}^{*ASAXS}$	$\Gamma_{Na}^{*BE-ICPMS}$
0.1	$0.73 \pm 0.06^{(a)}$	$0.80 \pm 0.02$	$0.71 \pm 0.06^{(b)}$	$0.74 \pm 0.02$

a) Data taken from reference 1

b) Data taken from reference 2



**Figure S2.** Comparison of experimentally determined fraction of charge neutralization ( $\Gamma_{Na}^*$ ) for  $Na^+$  around dsRNA and dsDNA from ASAXS and BE-ICP-MS. Data from Table S2.

**Table S4:** Experimentally determined preferential interaction coefficients and  $\alpha$  value for **NaBr** and **CsBr** around 24-bp RNA in the presence of 6 mM  $MgBr_2$ .

NaBr	24-bp RNA				CsBr	24-bp RNA				
	C [M]	$\Gamma_{Na^+}$	$\Gamma_{Mg^{2+}}$	$\Gamma_{Br^-}$		C [M]	$\Gamma_{Cs^+}$	$\Gamma_{Mg^{2+}}$	$\Gamma_{Br^-}$	total
0.00	0	$21.0 \pm 0.5$	$-3.0 \pm 1.0$	$46.0 \pm 1.0$	0	0	$21.0 \pm 0.5$	-4.0	$46.0 \pm 0.7$	
0.0015	$0.6 \pm 0.5$	$22.0 \pm 0.5$	$-2.0 \pm 0.5$	$46.8 \pm 0.8$	0.02	$3.6 \pm 1.0$	$19.0 \pm 0.5$	$-5.4 \pm 1.0$	$47.0 \pm 1.5$	
0.01	$2.9 \pm 0.8$	$20.0 \pm 0.4$	$-3.0 \pm 1.6$	$45.6 \pm 1.4$	0.03	$8.0 \pm 1.0$	$16.0 \pm 0.3$	$-5.0 \pm 0.6$	$45.6 \pm 1.2$	
0.02	$3.9 \pm 0.6$	$18.5 \pm 0.4$	$-5.4 \pm 1.4$	$46.5 \pm 1.6$	0.06	$10.0 \pm 0.5$	$16.0 \pm 0.6$	$-4.0 \pm 1.0$	$46.0 \pm 1.2$	
0.03	$6.5 \pm 1.0$	$17.4 \pm 0.4$	$-4.7 \pm 1.8$	$46.0 \pm 2.0$	0.10	$11.5 \pm 1.0$	$13.5 \pm 0.8$	$-7.0 \pm 1.0$	$45.6 \pm 1.6$	
0.05	$9.3 \pm 2.0$	$15.5 \pm 1.4$	$-6.0 \pm 1.2$	$46.3 \pm 2.7$	0.2	$19.0 \pm 0.9$	$8.1 \pm 1.4$	$-10.0 \pm 1.0$	$45.0 \pm 1.7$	
0.08	$10.0 \pm 1.0$	$13.5 \pm 1.0$	$-9.0 \pm 1.0$	$46.0 \pm 1.7$		-	-	-	-	
0.11	$12.5 \pm 2.0$	$11.0 \pm 1.3$	$-11.0 \pm 1.0$	$46.0 \pm 2.7$		-	-	-	-	
0.20	$19.8 \pm 1.5$	$7.2 \pm 0.4$	$-13.3 \pm 1.3$	$47.5 \pm 2.0$		-	-	-	-	
					$\alpha_{Na}^* = 17.0 \pm 1.7$					
					$\alpha_{Cs}^* = 18.3 \pm 2.5$					

\*Defined in the main text

**Table S5:** Poisson Boltzmann calculations of preferential interaction coefficients and  $\alpha$  value for monovalent salt (MX) around 24-bp DNA in the presence of 6 mM divalent salt ( $\text{MX}_2$ ).

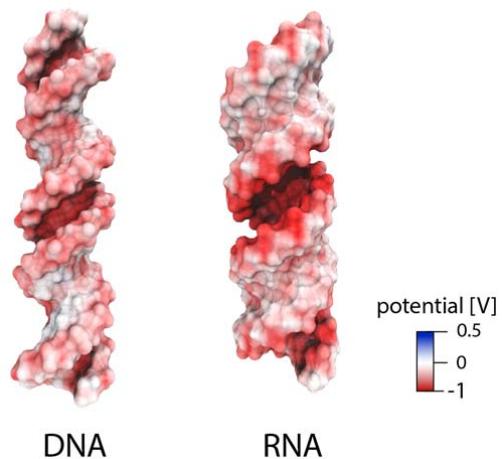
MX C [M]	24-bp DNA			
	$\Gamma_{M^+}$	$\Gamma_{M^{2+}}$	$\Gamma_{X^-}$	total
0.00	0	21.14	-3.7	46.0
0.005	2.5	19.5	-4.5	46.0
0.01	4.5	18.2	-5.1	46.0
0.02	7.53	16.16	-6.15	46.0
0.03	9.85	14.6	-6.95	46.0
0.04	11.73	13.32	-7.63	46.0
0.045	12.5	12.8	-7.95	46.0
0.05	13.23	12.26	-8.25	46.0
0.06	14.45	11.36	-8.83	46.0
0.08	16.45	9.85	-9.84	46.0
0.10	17.9	8.73	-10.7	46.0
0.15	20.0	6.68	-12.6	46.0
0.20	21.0	5.37	-14.26	46.0
0.30	21.27	3.79	-17.15	46.0
$\alpha_{M^+}^* = 7.5$				

\*Defined in the main text

**Table S6:** Poisson Boltzmann calculations of preferential interaction coefficients and  $\alpha$  value for monovalent salt (MX) around 24-bp RNA in the presence of 6 mM divalent salt ( $\text{MX}_2$ ).

MX C [M]	24-bp RNA			
	$\Gamma_{M^+}$	$\Gamma_{M^{2+}}$	$\Gamma_{X^-}$	total
0.00	0	21.45	-3.1	46.0
0.001	0.44	21.13	-3.3	46.0
0.01	3.5	19.1	-4.3	46.0
0.02	5.86	17.5	-5.14	46.0
0.03	7.62	16.32	-5.74	46.0
0.04	9.0	15.4	-6.2	46.0
0.05	10.24	14.5	-6.76	46.0
0.08	12.84	12.6	-7.96	46.0
0.10	14.13	11.62	-8.64	46.0
0.20	17.9	8.3	-11.5	46.0
0.3	19.5	6.3	-13.9	46.0
0.5	19.72	3.93	-18.43	46.0
$\alpha_{M^+}^* = 13.3$				

\*Defined in the main text



**Figure S3.** Poisson-Boltzmann calculations of electrostatic surface potential of the DNA and RNA duplexes. Calculations were carried out as described in the main text and Figure 3 in the main text.

#### REFERENCES:

1. Pabit, S. A., S. P. Meisburger, L. Li, J. M. Blose, C. D. Jones, and L. Pollack. 2010. Counting Ions around DNA with Anomalous Small-Angle X-ray Scattering. *J. Am. Chem. Soc.* 132(46):16334-16336.
2. Kirmizialtin, S., S. A. Pabit, S. P. Meisburger, L. Pollack, and R. Elber. 2012. RNA and Its Ionic Cloud: Solution Scattering Experiments and Atomically Detailed Simulations. *Biophys. J.* 102(4):819-828.
3. Gebala, M., G. M. Giambasu, J. Lipfert, N. Bisaria, S. Bonilla, G. Li, D. M. York, and D. Herschlag. 2015. Cation-anion interactions within the nucleic acid ion atmosphere revealed by ion counting. *J. Am. Chem. Soc.* 137(46):14705-14715