



**Effect of buffer composition on PNA-RNA hybridization
studied in the microfluidic microarray chip**

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23 **Abstract:** Herin we report that Peptide Nucleic Acid sequences (PNAs) can be used as the probe
24 species for detection of RNA and that a microfluidic microarray (MMA) chip can be used as the
25 platform for detection of hybridizations between immobilized PNA probes and RNA targets. The
26 RNA targets used are derived from influenza A sequences. This paper discusses the optimization of
27 the two probe technologies used for RNA detection and investigates how the composition of the probe
28 buffer or the content of the hybridization solution can influence the overall results. Our data shows
29 that the PNA probe is a better choice over the DNA probe when there is low salt in the probe buffer
30 composition. Furthermore, we have shown that the absence of salt (NaCl) in the hybridization buffer
31 does not hinder the detection of RNA sequences. The results conclude that PNA probes are superior
32 to DNA probes in term of sensitivity and adaptability, as PNA immobilization and PNA-RNA
33 hybridization are less affected by salt content in the reaction buffers unlike DNA probes.

34 **Key words:** peptide nucleic acid (PNA), influenza viral RNA, probe buffer, hybridization buffer, salt,
35 formamide, microfluidic microarray (MMA).

37 Introduction

38 Peptide nucleic acids (PNAs) are DNA analogs in which the negatively charged deoxyribose
39 phosphate backbone is replaced by an electrically neutral peptide-like backbone.¹⁻² Since the PNA
40 probes are not negatively charged, unlike traditional DNA probes, they should be less affected by the
41 salt content of the buffer in which these probes are used.³ When PNAs are used in the assembly of
42 microarrays, it would be useful to better understand the effects of variation in salt concentration, in
43 both the probe buffer used for printing the PNA probes on solid substrates and the hybridization
44 buffer used for reaction between the probes and the target nucleic acids.

45 In this paper, PNA probes designed to detect a RNA sequence related to influenza A are
46 employed in the construction of a microfluidic microarray (MMA) chip. The influenza A virus is the
47 causative agent of yearly epidemics and occasional pandemics⁴ and numerous microfluidic
48 technologies have been developed to detect the presence of influenza A viral RNAs in clinical
49 samples. Devices employed in these technologies include microfluidic chips,⁵⁻¹⁵ integrated
50 Microsystems,¹⁶⁻²⁰ and lateral flow strips.^{8, 21-22} The detection methodologies include RT-qPCR,^{15, 17,}
51 ^{20, 23-26} electrophoretic immunoassays,²⁷ PCR-capillary electrophoresis,^{11-14, 18, 29}
52 immunochromatography,^{8, 19, 21-22} microarray,^{5, 9, 16, 30-32} and electrochemical approaches.^{4, 7, 28, 33}
53 Among the microarray detection methods, oligonucleotide probes are most commonly used.^{5, 9, 16, 30-32}
54 Whilst PNA probes have been employed to detect various influenza virus strains,³⁴⁻³⁶ and one group
55 has employed PNA probes in the microfluidic chip to detect the viral DNA⁷ to date, there are no
56 details provided about the buffer compositions in the use of the PNA probes. Therefore, we conducted
57 a study to investigate the effect of buffer composition on PNA probe immobilization and on PNA-
58 RNA hybridization. We believe it is informative to report this study even though, generally, influenza
59 A viral RNAs are detected indirectly after their conversion to cDNA.

60 In our group, we have previously employed the MMA chip for detection of DNA³⁷⁻³⁹ and
61 RNA targets.⁴⁰ The microfluidic microarray chip offers several advantages to other designs (e.g. plain
62 microarrays) in that it can be constructed without an expensive robotic spotting system, and used for
63 multiple samples. The use of the chip can also offer fast hybridization rates, and achieve cost-
64 effectiveness.⁴¹⁻⁴⁵ The schematic diagrams for the use of the MMA chip are shown in Fig. 1 A-C. The
65 construction of the device follows several steps. Firstly, a PDMS channel plate is sealed with a glass
66 slide that is arrayed with lines consisting the probe molecules (Fig. 1A). Secondly, after removing the
67 first PDMS plate, the glass slide is sealed against a second PDMS plate to admit target samples for

68 hybridization (Fig. 1B). Thirdly, the PDMS plate is removed and the glass slide is fluorescently
69 scanned for detection (Fig. 1C). An image of the actual experiment used in this study and consisting
70 of 16 probes hybridized with 16 targets is shown in Fig. 1D.

71 In this paper, we describe the principle of detection of PNA-RNA hybridizations in the MMA
72 chip. This chip has the capability of studying various buffer conditions simultaneously using multiple
73 samples. We report the effect of compositions of reaction buffers (*i.e.* probe buffer and hybridization
74 buffer) on PNA-RNA hybridization results.

75

76 **Experimental**

77 **Nucleic acid sequences**

78 RNA target and DNA probe sequences were synthesized by Integrated DNA Technologies
79 (Coralville IA). PNA probes were purchased from PNA Bio, Inc. (Thousand Oaks, CA). The
80 sequences of the nucleic acid probes as well as RNA targets utilized in this study are shown in Table
81 1.

82 Three probes were used: two were PNA probes (P-PNA, P-PNA-inf) and one was DNA probe
83 (P-DNA). Both PNA probes are 13-mer PNA probe sequences utilizing an 'O monomer' spacer
84 molecule (2-(2-aminoethylethoxy) ethoxy acetic acid) located at the N-terminal position for
85 attachment to the chip surface. The P-PNA probe has the same base sequence as the central region of
86 a 30-mer DNA probe (P-DNA), utilising an amino group with a C6 linker located at the 5' end for
87 surface attachment. This DNA probe was previously used for analyzing RNA samples⁴⁰. The P-PNA-
88 inf probe is a 13-mer PNA probe that was previously reported to probe influenza A viral RNA.^{34, 50}

89 Two target RNA sequences were used. One is complementary to the PNA probe (T-RNA),
90 whilst the other is designed to mimic an influenza A viral RNA target sequence (T-RNA-inf). Both

91 RNA target sequences were labeled by biotin at the 5' termini. The sequence of T-RNA was so
92 designed that the binding region (section underlined in the T-RNA sequence in Table 1) is
93 complementary to both the PNA probe (P-PNA) and the central region of the oligonucleotide probe
94 (underlined in P-DNA).

95 The influenza A virus consists of single-stranded RNAs⁴⁶⁻⁴⁷. The RNA molecule comprises
96 eight segments that encode for different proteins necessary for transcription, replication and viral
97 assembly⁴⁸⁻⁴⁹. The sequence of T-RNA-inf was obtained from the analysis of the 5' untranslated
98 region (UTR) which was conserved among each of the eight RNA segments.^{34, 50}

99 PNA-RNA or DNA-RNA hybridizations were determined by imaging of the Cy5 tag (emits at
100 670 nm when excited at 633 nm using fluorescence spectroscopy) incorporated by conjugation of
101 streptavidin-Cy5 to the 5' biotin moiety on the RNA target sequences.

103 **Materials**

104 Glutaraldehyde (25%), 3-aminopropyltriethoxysilane (APTES), sodium dodecyl sulfate (SDS),
105 sodium borohydride, and bovine serum albumin were obtained from Sigma-Aldrich and were used
106 without further purification. Formamide was purchased from Merck. Ultrapure water (18 MΩ cm)
107 was obtained from a water purification system (Easy pure RF, Dubuque, IA). Streptavidin-Cy5 (1
108 mg/μL) was purchased from Invitrogen, Life Technologies. Phosphate buffered saline (PBS), saline
109 sodium citrate (SSC), sodium citrate, sodium bicarbonate, concentrated sulfuric acid, hydrogen
110 peroxide (30 %) and ethanol (95 %) were all reagent-grade materials.

111 The 2% APTES solution was prepared by mixing 2.4 mL of APTES in 117.6 mL of 95% ethanol.
112 The 5% glutaraldehyde solution was prepared by mixing 24 mL of 25% glutaraldehyde, 50 mL of
113 20X PBS and 46 mL of water.

114 The plain glass microscope slides (3" x 2") were purchased from Fisher Scientific (Ottawa, ON).

115

116 **Functionalization of Glass Slides**

117 Four glass slides were washed with detergent (Sparkleen) and rinsed with pure water, as
118 previously described⁵¹⁻⁵². In summary, the glass slides were submerged in piranha solution (3:1
119 mixture of concentrated H₂SO₄, 30% H₂O₂) for 15 min. at 80 °C. After being rinsed with distilled
120 water, ethanol 95%, then distilled water, the glass slides were dried under N₂. They were then
121 submerged in a 2% APTES solution (under a N₂ gas environment) for 20 min. and then rinsed with
122 100% ethanol. Dispensing of APTES liquid was performed by a 10"-long needle together with a
123 compensating nitrogen gas balloon. One observation is that the liquid should be yellow or light brown
124 at this point and if the color turns dark and/or particles are found in the liquid it should be discarded.

125 The glass slides were subsequently baked in the oven for 1 h at 120 °C in air. Thereafter, the glass
126 slides were submerged in a 5% glutaraldehyde solution in the refrigerator for 1 h and then rinsed with
127 pure water. Finally, the glass slides were N₂-dried and stored in the dark at 4 °C before probe printing.

128

129 **Probe Immobilization**

130 A 2" x 2" PDMS channel plate consisting of microchannels (200-µm wide and 35-µm deep) was
131 fabricated as described elsewhere.^{37, 52} The 1.5-mm thick channel plate was sealed against the
132 aldehyde-functionalized glass slide to form the MMA chip. Then, 0.6 µL of nucleic acid probes
133 prepared in the probe buffer (0.15 M sodium bicarbonate with 1 M NaCl, unless noted otherwise)
134 were added into the inlet reservoirs of the MMA chip using a micropipette. The probe solutions were
135 filled through the channels by applying suction at the outlets followed by incubation at room
136 temperature for 1 h. After the microchannels had been rinsed with 1 µL of a wash solution (0.15 M

137 sodium bicarbonate with 1 M NaCl), the PDMS channel plate was peeled away from the surface.
138 Afterward, the glass slide was treated with the sodium borohydride solution (50mg NaBH₄, 15mL
139 PBS 1X, 5mL ethanol 95%, 150 μL SDS 1.5%) for 20 min. The glass chip was then rinsed with pure
140 water for 2 min. and then subsequently dried under N₂. The prepared glass slide was submerged in the
141 blocking solution (10 mL 10X SSC, 150 μL SDS 1.5%, 20mg BSA and made up to 20mL with pure
142 water) to inhibit non-specific binding. The glass chip incorporating the printed probe was then rinsed
143 with deionized water for 2 min. and finally dried as above.

144

145 **Hybridization and Detection**

146 The glass slide with probe line arrays was covered with a second 2'' x 2'' PDMS channel plate
147 to form the MMA chip assembly. The straight channels were aligned orthogonal to the printed probe
148 lines on the slide. Hybridization reactions occur at the intersections between the channels and the
149 probe lines. The target RNA samples were diluted in the hybridization buffer (1X SSC, unless noted
150 otherwise, with 0.15% SDS). After the target solutions were denatured at 95 °C for 5 min, they were
151 quickly cooled in ice just prior to their hybridization to immobilized probes. The target solutions (0.6
152 μL) were added to the inlet reservoirs of the MMA chip using a micropipette. Target solutions in
153 different reservoirs were then drawn into the channels simultaneously by suction. Probe-target
154 hybridization was carried out in an oven at 40 °C for incubation for 30 min. The microchannels were
155 rinsed with 1μL of hybridization buffer immediately after incubation. SA-Cy5 solutions (0.6 μL, 1
156 mg/mL) were added to the inlet reservoirs using a micropipette, and the solutions were then drawn
157 into the channels by suction simultaneously. After a further incubation step (30 min. at room
158 temperature), the microchannels were rinsed immediately with 1 μL of wash buffer (1X PBS and
159 0.2% Tween 20)

160 For analysis, the glass slide was imaged on a confocal laser fluorescent scanner (Typhoon
161 Trio+ variable mode imager) purchased from GE Healthcare (Baie d'Urfé, QC). The scan conditions
162 were spatial resolution: 25 μm , excitation wavelength: 633 nm, emission wavelength: 670 nm, and
163 photomultiplier tube voltage: 600V. The scanned image was analyzed by the ImageQuant 5.2
164 software. In the data quantification procedure, lines were drawn manually to cover the rectangular
165 hybridization patches on the glass slide, and the average fluorescent intensity of the patch was
166 recorded in the relative fluorescent unit (RFU).

167

168 **Results and Discussion**

169 **Effect of probe buffer composition**

170 RNA targets were hybridized to DNA and PNA probes in the presence of different amounts of salt
171 in the probe buffers. The hybridization results are shown in Figure 2A. Here, the RNA targets were
172 applied along the vertical direction, intersecting with the probe lines created in the horizontal
173 orientation. Experiments were conducted using two concentrations of RNA targets (T-RNA). Stronger
174 fluorescent signals were observed for the DNA-based probes (P-DNA) when prepared in the probe
175 buffer containing the high salt concentration (1 M NaCl) rather than the low salt one (0.1 M NaCl). It
176 is assumed this observation arises as the DNA probe possesses a negatively charged backbone and
177 accordingly it requires the high ionic strength conditions provided by the 1M NaCl solution to reduce
178 the Columbic repulsion between the negatively charged backbone and the glass slide during the probe
179 immobilization step as previously reported.⁵⁵⁻⁵⁶ Another observation is that much stronger signals are
180 observed for the PNA-based probes (P-PNA) rather than the DNA-based probes (P-DNA). As shown
181 in Fig. 2B, enhancement in signal intensity of 521% was found for the 10-nM RNA target, and 317%

182 for the 20-nM target, when P-PNA rather than P-DNA was the probe using 0.1 M NaCl as the probe
183 buffer.

184 Furthermore, there was considerably less influence of salt concentration on RNA hybridization
185 efficiencies when using the PNA-based probes (P-PNA). For instance, the hybridization signals
186 (depicted in Fig. 2B) were similar for the 10-nM RNA target when the PNA probes were immobilized
187 using buffers of different salt contents (i.e. the signal was decreased by just 28.8% when 0.1 M
188 instead of 1 M NaCl was used). On the other hand, the DNA-based probes were less robust in which
189 the signal decreases were higher when the probe buffer contained 1 M instead of 0.1 M NaCl (i.e. the
190 signals were decreased by 83.9% and 66.4% for 20-nM and 10-nM RNA, respectively). These data
191 supports the notion that PNA probes are more robust than DNA probes in terms of sensitivity and
192 adaptability.³⁴⁻³⁶

194 **Effect of hybridization buffer composition**

195 Hybridizations involving RNA targets generally require a high temperature that functions to
196 remove secondary structures of the single-stranded nucleic acid molecules. Nevertheless, formamide
197 can also be used to achieve the same function allowing hybridization to be conducted at more
198 reasonable temperatures.⁵³⁻⁵⁴

199 The results of RNA hybridizations conducted in formamide-containing hybridization buffer
200 are shown in Fig.3A. Stronger signal intensities are observed for T-RNA targets when prepared in the
201 hybridization buffer containing 30% formamide (Fig. 3B) indicating the utility of formamide to
202 minimize any secondary structural motifs inherent in the target RNA sequences. Such a signal
203 enhancement was observed even when the PNA probes (P-PNA) were printed at a low-salt probe

204 buffer containing 0.1 M NaCl. Similar observations were obtained in Fig. 3A for the RNA target (T-
205 RNA-inf) which is related to the influenza A viral RNA sequence.

206

207 Although 1X SSC (saline-sodium citrate) is the buffer commonly used for hybridization
208 experiments, we tested the capacity of our system to deliver reasonable results when using low-salt
209 hybridization buffers such as C (15 mM sodium citrate) in the presence of 30% formamide. A
210 comparison of hybridizations obtained in SSC and C buffers is illustrated in Fig. 4A. It is found that
211 intensities with signal-to-noise ratio greater than 2 are still observed using hybridization buffers
212 prepared in the low-salt conditions such as C buffer without NaCl added, suggesting that the need of
213 high salt for reducing Columbic repulsion is not as critical when PNA probes, rather than DNA
214 probes, are used. In fact, using PNA probes signals are still observable against the T-RNA-inf target
215 sequence even at 5nM using low-salt hybridization buffers (i.e. C buffer) are used (Fig. 4B).

216

217 **Conclusion**

218 The microfluidic microarray (MMA) chip has been employed to study the effects of buffer
219 compositions on PNA-RNA hybridization experiments. The presence of salt in the probe buffer and
220 hybridization buffer does result in better signals for the detection of RNA targets, although
221 hybridization still occurred to a reasonable extent in the absence of NaCl. Moreover, the presence of
222 formamide in the hybridization buffer offers better RNA hybridization when conducted at a relatively
223 low temperature of 40 °C. The MMA chip results support the notion that the PNA probe is more
224 useful than the DNA probe in low ionic strength conditions with respect to assay sensitivity and
225 experimental adaptability for RNA detection.

226

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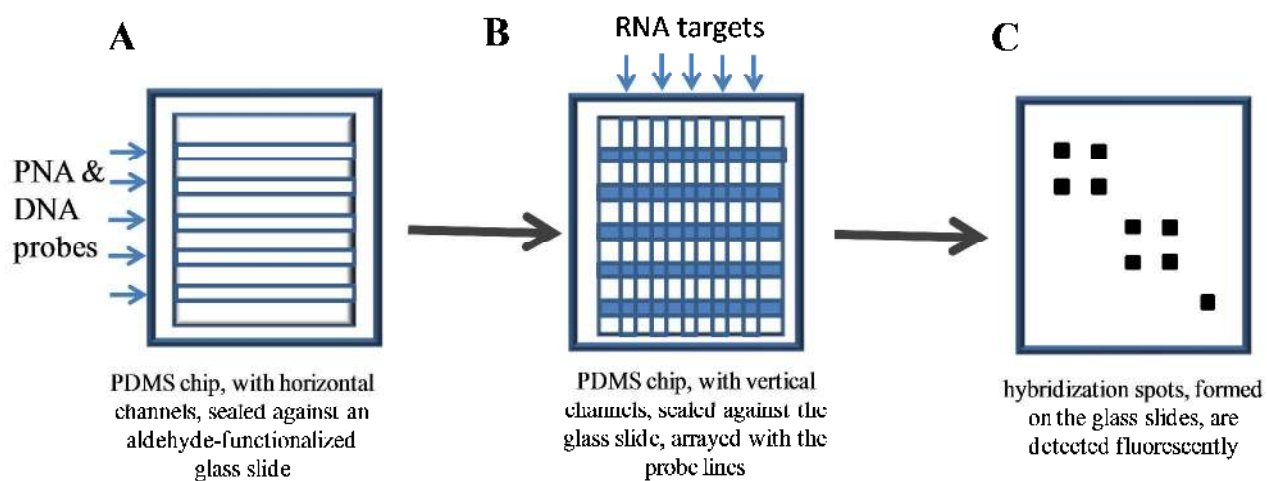
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339 **Table 1.** Sequences of probes (P) and targets (T) used to study PNA-RNA hybridizations

Identifier	Sequence
Probes	
P-DNA	H ₂ N-C6-CTGTATTGAGTTGTATCGTGTGGTGTATTT
P-PNA	H ₂ N-O-O-AGTTGTATCGTGT
P-PNA-inf	H ₂ N-O-O-CCTTGTTTCTACT
Targets	
T-RNA	biotin-ACGGAGCGCAAAAUACACC <u>ACACGAUACAACU</u> CAAUACAGUCGACGCCUA
T-RNA-inf	biotin- <u>AGUAGAAACAAGG</u> CCUGCUUUUG

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341 **Fig. 1.** Detection procedure using the microfluidic microarray (MMA) chip. The schematic diagrams
 342 of printing the probe array (A), hybridization with the RNA targets (B), detection of fluorescent
 343 results on the glass slide surface (C); an image of the experimental results (D).



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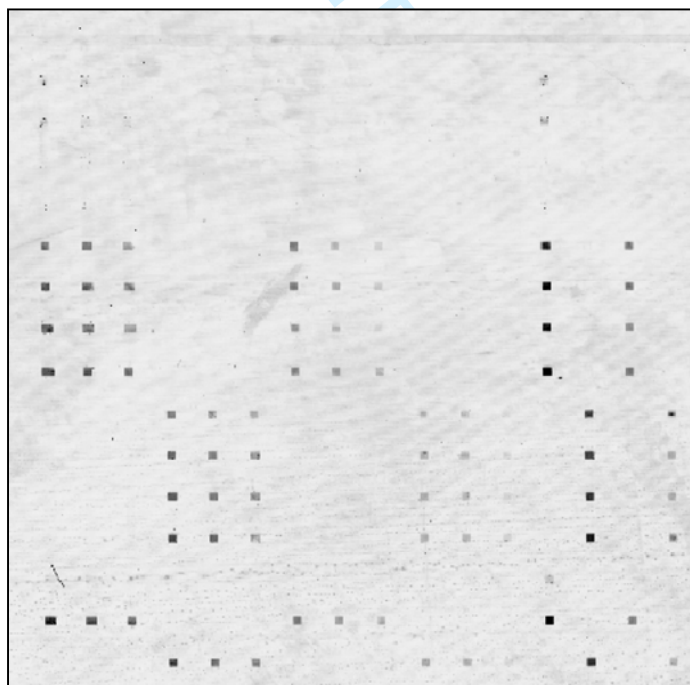
346

D

16 targets



16 probes →

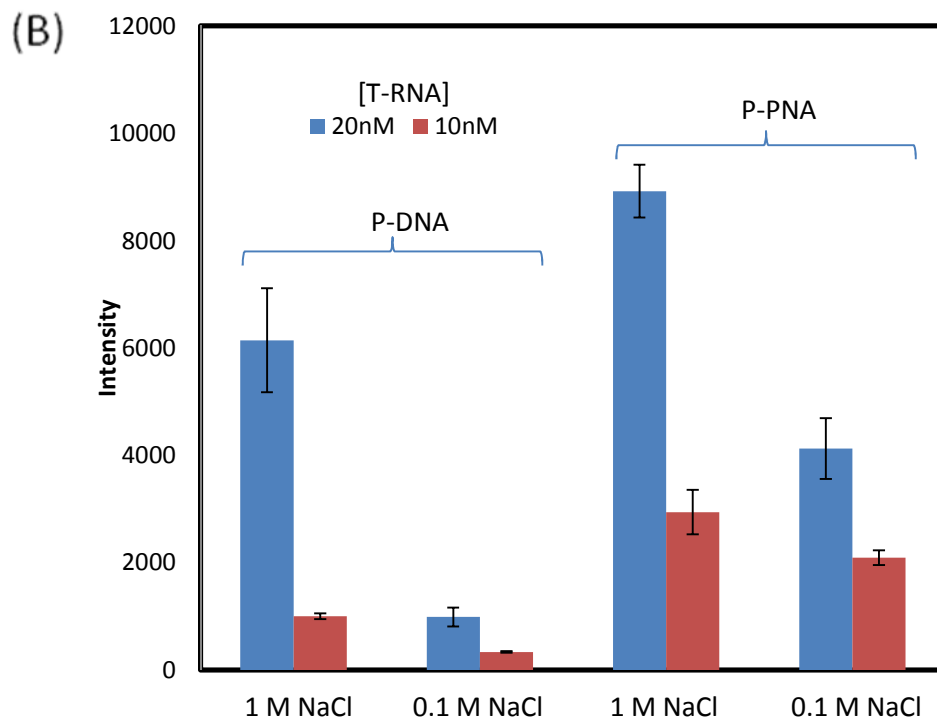
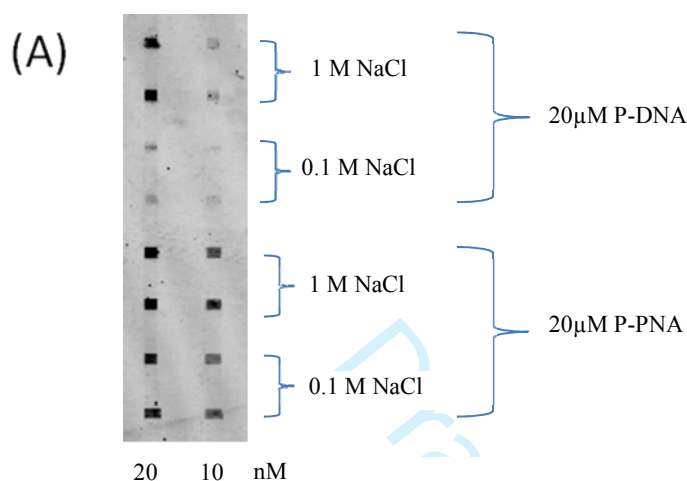


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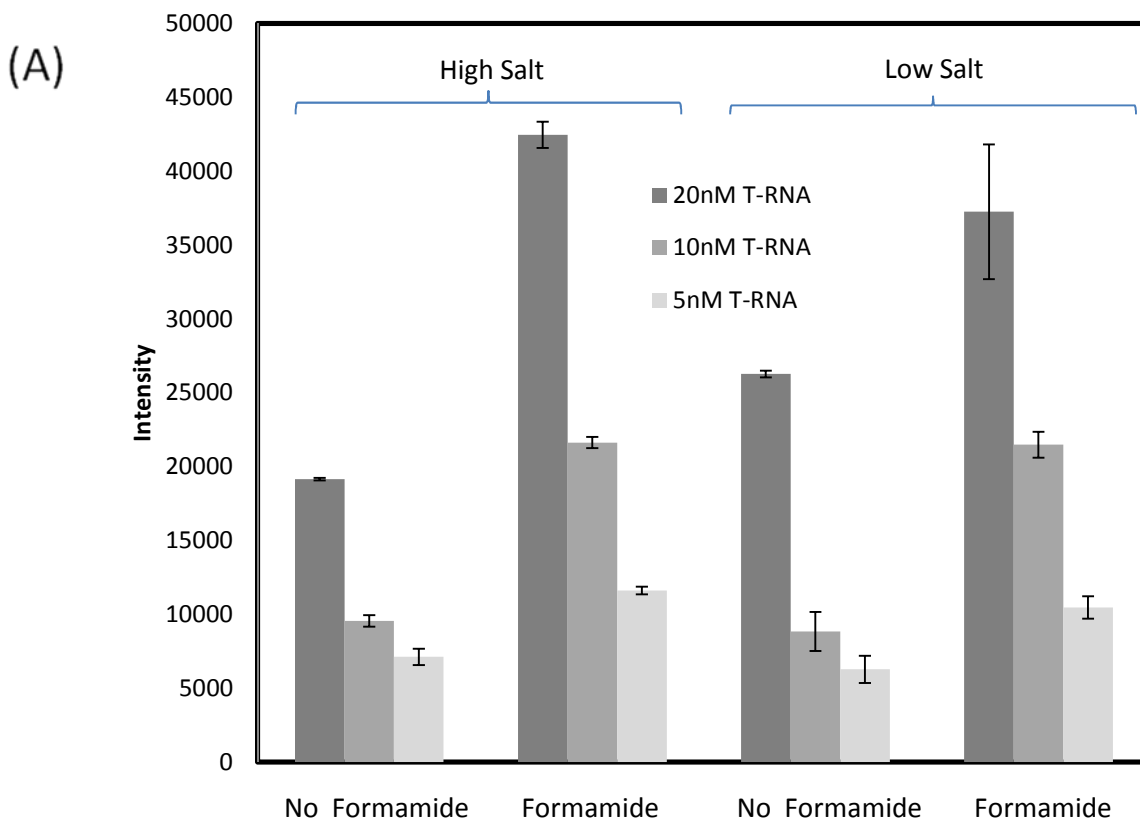
350 **Fig. 2.** (A) Image and (B) background-subtracted intensity of hybridization of T-RNA with the
351 complementary DNA probe (P-DNA) and PNA probe (P-PNA) in different low-salt buffer (0.1 M
352 NaCl) and high-salt spotting buffer (1 M NaCl). The probe solutions contain P-DNA (20 μ M), 0.15 M
353 sodium bicarbonate, and low/high salt content. The target solutions contain T-RNA (20 nM or 10
354 nM), 1X SSC (saline-sodium citrate) buffer, and 0.15% SDS.



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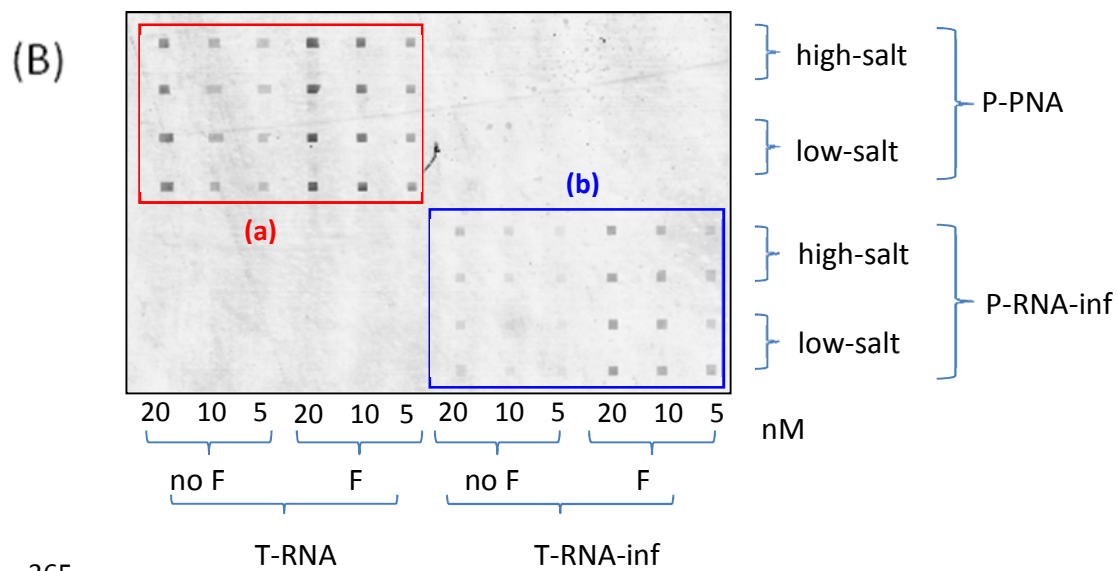
356

357 **Fig. 3.** (A) Background-subtracted intensity and (B) images of hybridization of (a) T-RNA with the
358 complementary P-PNA probe and (b) T-RNA-inf with the complementary P-PNA-inf probe; both
359 probes were prepared in different low-salt (0.1M NaCl) and high-salt (1 M NaCl) probe buffers
360 containing 0.15 M NaHCO₃. The targets were prepared in the 1X SSC buffer with formamide (F) or
361 without formamide (no F), and 0.15% SDS. The concentration of formamide is 30%. The
362 concentrations of T-RNA and T-RNA-inf are 5, 10, and 20 nM.



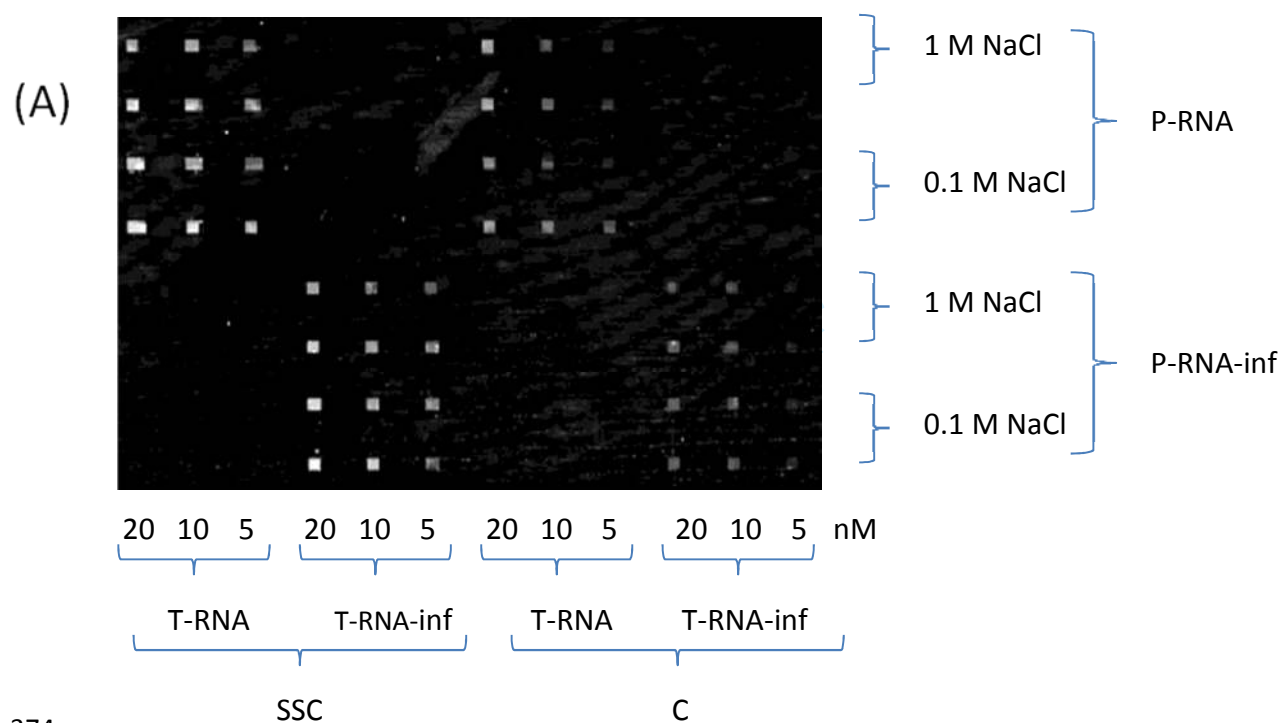
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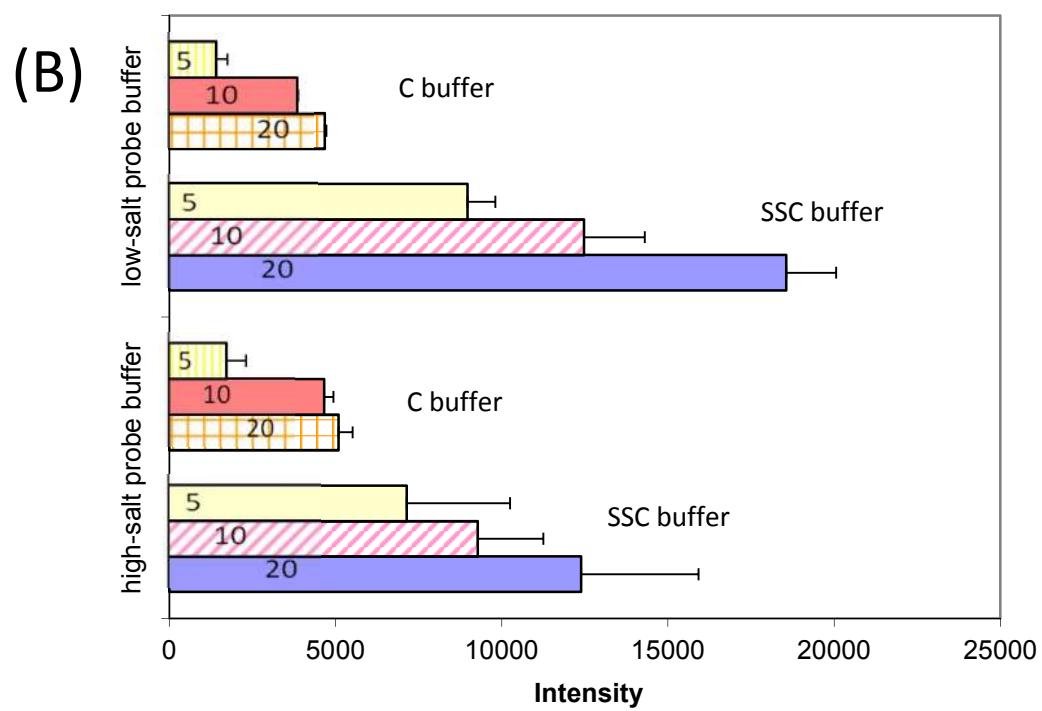
Draft

366 **Fig. 4.** (A) Images of hybridizations of T-RNA with the complementary PNA probe (P-RNA), and of
 367 T-RNA-inf with the complementary PNA probe (P-RNA-inf). The images are shown with white
 368 patches and black background for clarity. The probes were prepared in low-salt (0.1 M NaCl) and
 369 high-salt (1 M NaCl) probe buffers containing 0.15 M NaHCO₃. The target solutions were prepared in
 370 1X SSC solution or 15 mM C solutions, which all contained 30% formamide. (B) Background-
 371 subtracted intensity of hybridization of T-RNA-inf with the complementary P-PNA-inf probe when
 372 two probe buffers and two hybridization buffers are used. The numbers represent the T-RNA-inf
 373 concentration in nM.



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