

SUPPLEMENTARY INFORMATION

Time dependence of cisplatin-induced duplex dissociation of 15-mer RNAs and mature miR-146a

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EXPERIMENTAL DETAILS

Chemicals and solutions

All oligonucleotides (5'-CUUCUUGGUUCUCUU-3' and 5'-AAGAGAACCAAGAAG-3' (RNA-1), 5'-CUUCUUGUGUCUCUU-3' and 5'-AAGAGACACAAGAAG-3' (RNA-2), 5'-CCUCUGAAAUUCAGUUCUUCAG-3' and 5'-UGAGAACUGAAUCCAUGGGUU-3' (miR-146a)) were purchased from IBA GmbH (IBA Nucleic Acids Synthesis, Göttingen, Germany) of HPLC grade quality. Cisplatin (*cis*-[Pt(NH₃)₂(Cl)₂], (**1**)) and dimethylformamide (DMF) was obtained from Sigma-Aldrich. Sodium dihydrogen phosphate monohydrate (NaH₂PO₄ • H₂O), disodium hydrogen phosphate dehydrate (Na₂HPO₄ • 2H₂O), sodium perchlorate monohydrate (NaClO₄ • H₂O) and silver nitrate (AgNO₃) was obtained from Merck. Ammonium acetate (CH₃CO₂NH₄) was purchased from Scharlau and acetonitrile (CH₃CN, HPLC grade) from LAB-Scan. All water used was of Millipore quality and autoclaved before use.

Kinetic studies

Monoaquated cisplatin (*cis*-[Pt(NH₃)₂Cl(OH₂)⁺], (**1a**)) was prepared by dissolving 3.9 mg of **1** in 1.30 ml DMF, by vortexing (1400 rpm) over night at room temperature. AgNO₃ solution was prepared by dissolving 2.3 mg AgNO₃ in 1.38 ml DMF, by vortexing (2200 rpm) during 5 min at room temperature. 0.98 mole equivalents of AgNO₃ were added to the **1** solution. The mixture was incubated at 37 °C during shaking (170 rpm) for 24 h in the dark. Precipitated silver chloride was removed by two consecutive centrifugations (during 3h, and 2 h and 30 min) at room temperature. The supernatant was transferred to new eppendorf tubes after each centrifugation step. Stock solutions of **1a** were stored in a desiccator together with silica gel (the particles (1 – 3 mm) contained a moisture indicator) and were kept in the dark at room temperature.

Kinetic studies were performed on a Varian Cary 4000 spectrophotometer, equipped with a thermal control unit. The oligonucleotide strands for RNA-1, RNA-2 and miR-146a were mixed separately and the measurements were conducted with a total strand concentration (*C*_T) of 3.0 μM at two different P_i-buffer conditions (the 5-fold concentrated stock solution was prepared according to *Current protocols in molecular biology*¹); *C*_{Pi} = 20 mM (pH 5.8 and *C*_{Na+} = 29 mM) and *C*_{Pi} = 20 mM, supplemented with NaClO₄ (pH 5.7 and *C*_{Na+} = 129 mM). The complementary oligonucleotides were heated to 90 °C and allowed to hybridize by slow cooling (0.5 °C/min) to 20 °C. The hybridized samples were heated to 38 °C and **1a** was added (the final concentrations were 7.5, 15.0, 22.5, 30.0 and 45.0 μM). The change in absorbance was measured at λ = 260 nm during 19 h after addition of **1a**.

Thermal melting studies

Thermal melting studies were performed on a Varian Cary 4000 spectrophotometer, equipped with a thermal control unit. The oligonucleotide strands for RNA-1, RNA-2 and miR-146a were mixed separately and the measurements were conducted with *C*_T = 3.0 μM at two different P_i-buffer conditions (the 5-fold concentrated stock solution was prepared according to *Current protocols in molecular biology*¹); *C*_{Pi} = 20 mM (pH 5.8 and *C*_{Na+} = 29 mM) and *C*_{Pi} = 20 mM, supplemented with NaClO₄ (pH 5.7 and *C*_{Na+} = 129 mM). The complementary oligonucleotides were heated to 90 °C and allowed to hybridize by slow cooling (0.5 °C/min) to 20 °C. The thermal melting points were evaluated by the first derivative method, using the Cary WinUV software. Data points were collected every 0.20 °C and the equidistant data was used for the Savitzky-Golay calculations². These calculations give rise to a minimized signal noise, by obtaining a new derivative point out of a determined number of surrounding data points. The standard deviation of independent measurements of *T*_m, was estimated to 0.3 °C based on 8 consecutive measurements.

¹ *Current Protocols in Molecular Biology*, 1996, A.2.1-A.2.8.
² A. Savitzky, M. J. E. Golay, *Anal. Chem.*, 1964, **36**, 1627-1639.

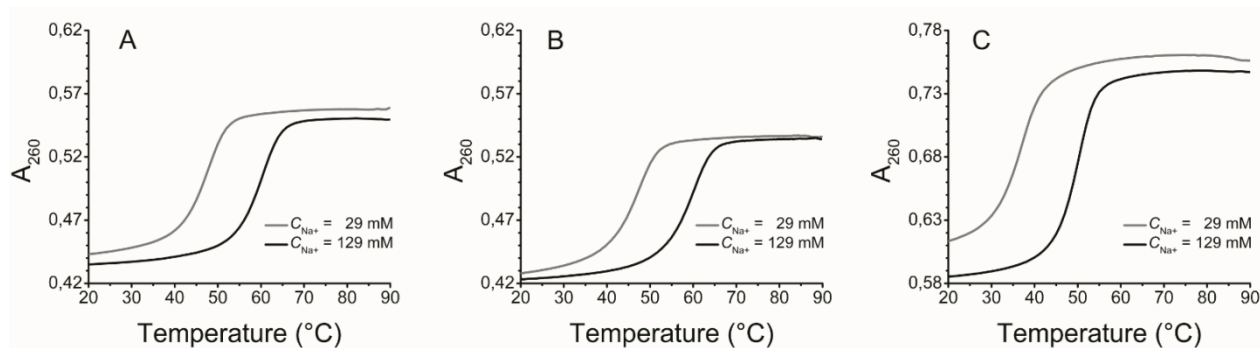


Figure S1. Thermal melting curves obtained for the native RNA duplexes (A) RNA-1, (B) RNA-2 and (C) miR-146a. All measurements were conducted with $C_T = 3.0 \mu\text{M}$ at two different buffer conditions; $C_{\text{Na}^+} = 29 \text{ mM}$ (—; 20 mM $\text{Na}_2\text{HPO}_4/\text{NaH}_2\text{PO}_4$ pH 5.8) and $C_{\text{Na}^+} = 129 \text{ mM}$ (—; 20 mM $\text{Na}_2\text{HPO}_4/\text{NaH}_2\text{PO}_4$ pH 5.7, supplemented with 100 mM NaClO_4).

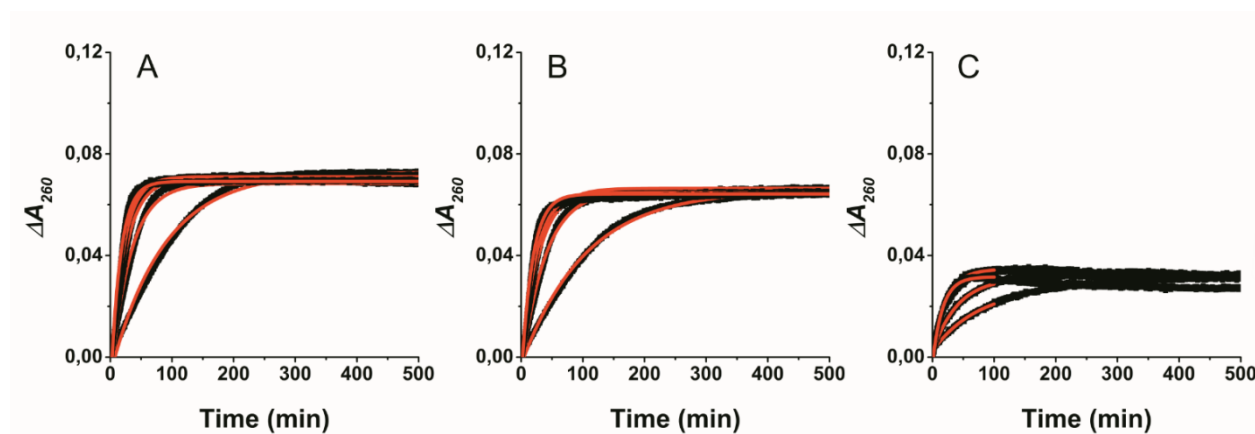


Figure S2. Absorbance change as a function of time after addition of **1a** ($C_{1a} = 7.5 - 45.0 \mu\text{M}$) to (A) RNA-1, (B) RNA-2 and (C) miR-146a. All measurements were conducted with $C_T = 3.0 \mu\text{M}$ at 38°C in buffered solution; $C_{\text{Na}^+} = 29 \text{ mM}$ ($20 \text{ mM Na}_2\text{HPO}_4/\text{NaH}_2\text{PO}_4$ pH 5.8).

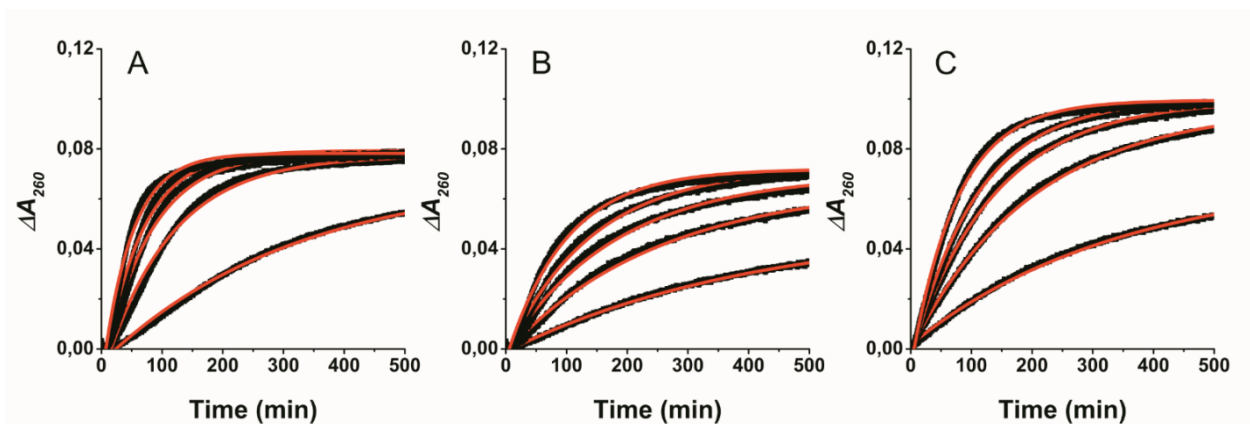


Figure S3. Absorbance change as a function of time after addition of **1a** ($C_{1a} = 7.5 - 45.0 \mu\text{M}$) to (A) RNA-1, (B) RNA-2 and (C) miR-146a. All measurements were conducted with $C_T = 3.0 \mu\text{M}$ at 38°C in buffered solution; $C_{\text{Na}^+} = 129 \text{ mM}$ (20 mM $\text{Na}_2\text{HPO}_4/\text{NaH}_2\text{PO}_4$ pH 5.7, supplemented with NaClO_4).

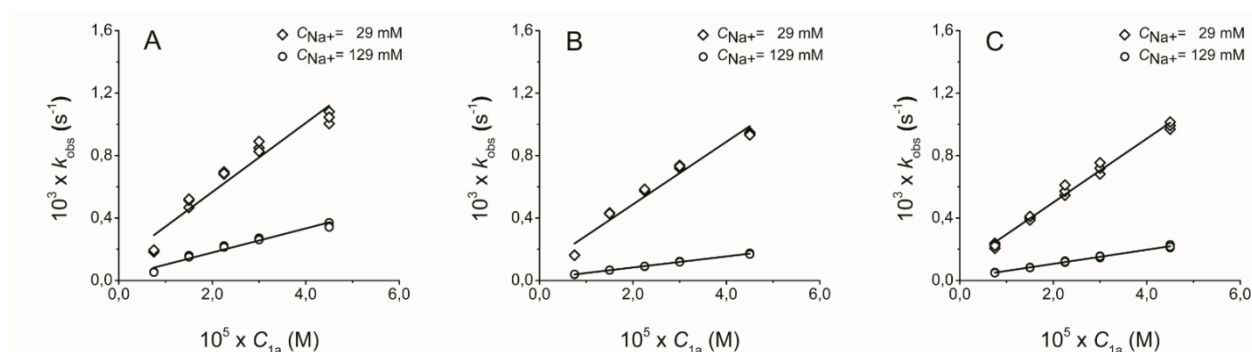


Figure S4. Observed pseudo-first-order rate constants (k_{obs}) plotted as a function of **[1a]** ($C_{1a} = 7.5 - 45.0 \text{ }\mu\text{M}$) for (A) RNA-1, (B) RNA-2 and (C) miR-146a. All measurements were conducted with $C_T = 3.0 \text{ }\mu\text{M}$ and regression lines were fitted to data obtained at two different buffer conditions; $C_{\text{Na}^+} = 29 \text{ mM}$ ($\blacklozenge$; 20 mM $\text{Na}_2\text{HPO}_4/\text{NaH}_2\text{PO}_4$ pH 5.8) and $C_{\text{Na}^+} = 129 \text{ mM}$ ($\circ$; 20 mM $\text{Na}_2\text{HPO}_4/\text{NaH}_2\text{PO}_4$ pH 5.7, supplemented with 100 mM NaClO_4).