Supplementary Material

Selective Inhibition of MBNL1·CCUG Interaction by Small Molecules Toward Potential Therapeutic Agents for Myotonic Dystrophy Type 2 (DM2)

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Additional biophysical studies for ligand 1:



Figure S1. Job analysis of the stoichiometry of the DNA complex with ligand **1** with total concentrations of 60 μ M (\blacktriangle) and 120 μ M (\blacklozenge).



Figure S2. (a) CD spectra showing the change in CD signal upon ligand **1** binding. (b) A plot of CD signal at 288 nm versus the mole ratio of ligand to duplex indicating the saturation of binding sites at about 3 equivalents of ligand **1**.

Melting curves for RNA used:



Figure S4. Melting curves for RNA duplex containing two mismatches and their corresponding single strands.



Figure S5. Melting curves for RNA containing a single mismatch and their corresponding single strands.

EMSA:



Figure S6. EMSA of ligand **1** in the presence of 100 nM tRNA. Control lane 1 (C1): RNA only. Control lane 2 (C2): RNA + MBNL1.

RNA stability experiment:

Method: Labeled (CCUG)₆ RNA was annealed by incubation at 95 °C for 1 min and then placed on ice for 20 min in a buffer containing 66 mM NaCl, 6.7 mM MgCl₂ and 27 mM Tris-Cl (pH = 7.5). Ligands dissolved in DMSO were then added to the RNA at a volume ratio of 1:9. The final reaction volume was 20µl with 1 nM RNA and 200 μ M ligands. The reactions were incubated at room temperature for 1–4 h before loading on a 15% polyacrylamide-urea denaturing gel. Electrophoresis was performed at 200 V and the gel was visualized by PhosphorImaging.



Figure S7. These studies showed that ligands **1–3** do not cleave the target RNA.

ITC curves for ligand 3:





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¹H NMR of Ligand 2:



¹³C NMR of Ligand 2:



¹H NMR of Ligand 3:



¹³C NMR of Ligand 3:

