

***Supporting Information for:***

Long term storage of miRNA at room and elevated temperatures in silica sol-gel matrix.

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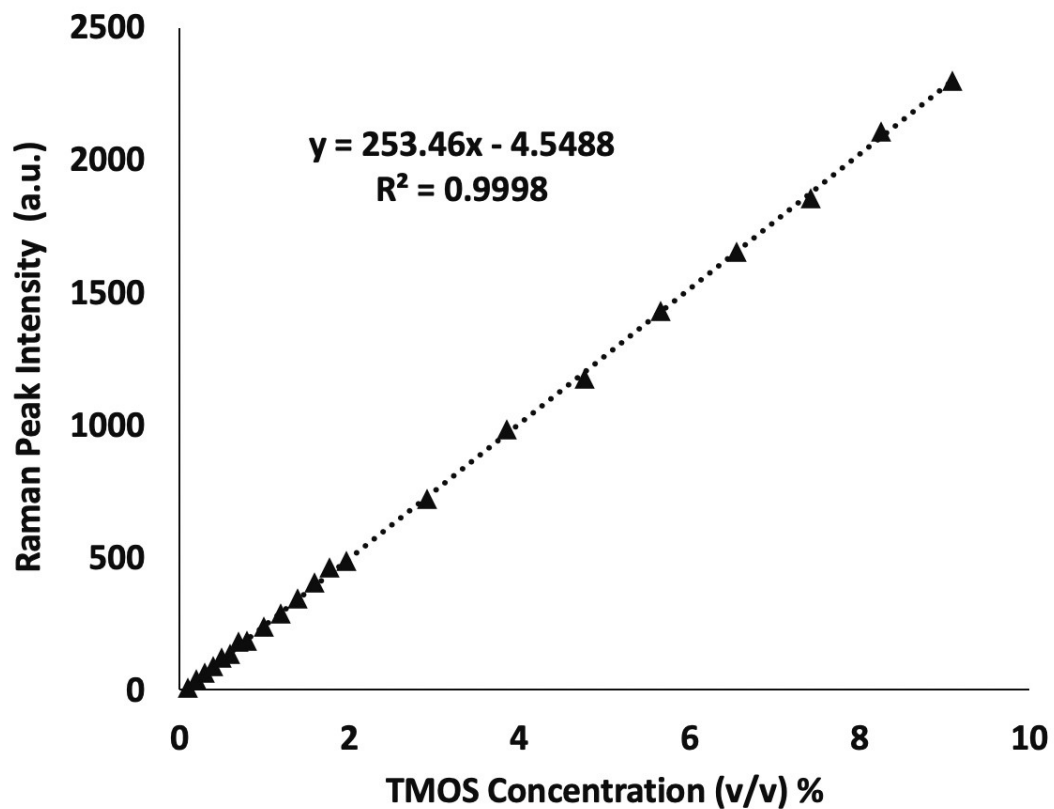


Figure. S1. Calibration Curve of Raman Peak Intensity Vs. TMOS Concentration (v/v) %: A significant increase in the raman peak intensities of methanol is demonstrated with the increase in TMOS concentrations in (0.5 - 10) v/v % range.

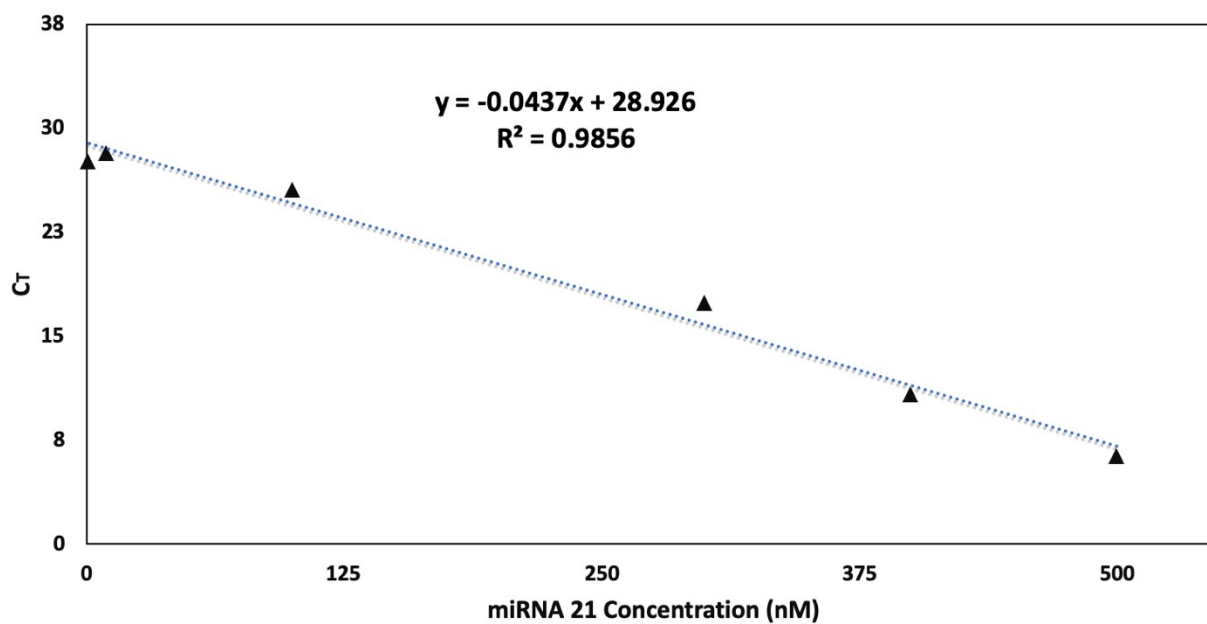


Figure. S2. miRNA 21 concentration (nM) calibration curve in CaRGOS using qRT-PCR analysis: Formulation parameters used were 0.5 v/v % CaRGOS, Low-salt Tris EDTA buffer and nuclease free water.

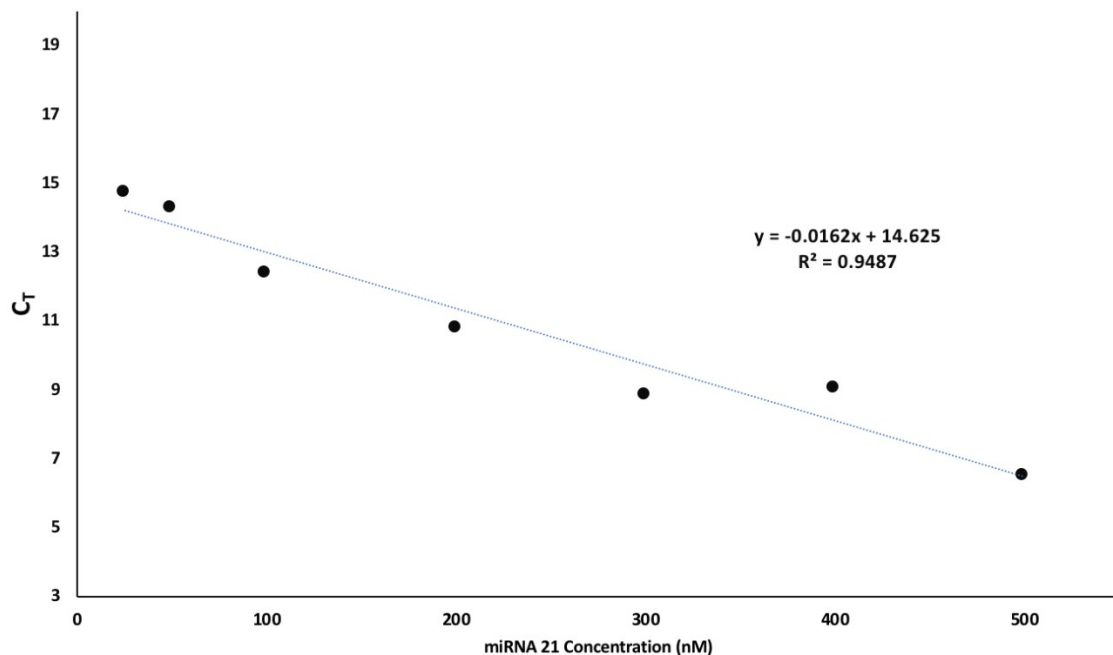


Figure.S3. miRNA 21 concentration (nM) calibration curve in nuclease free water using qRT-PCR analysis: Formulation parameters used were low-salt buffer and nuclease free water.

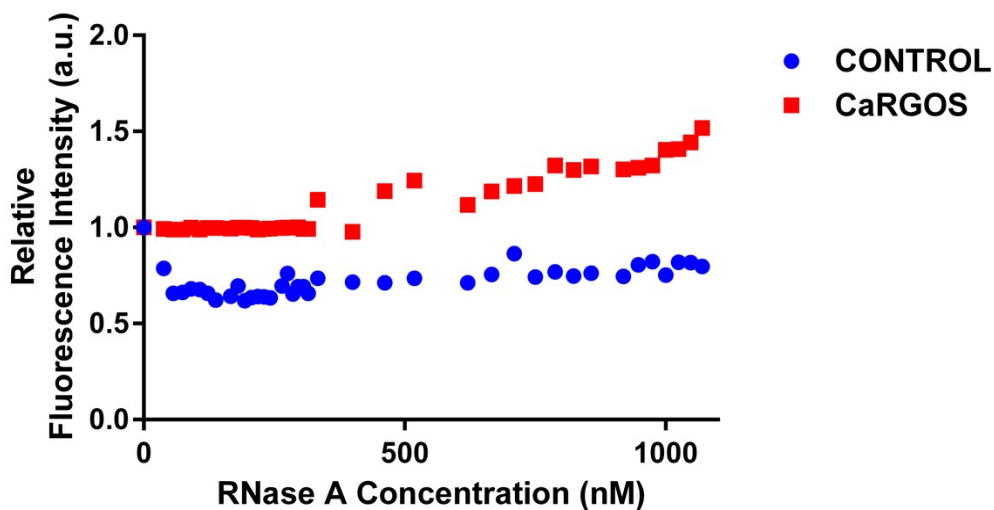


Figure. S4. A plot of relative fluorescence intensity of Ethidium bromide against RNase A concentrations in (0-1200) nM range: An increase in relative fluorescence emission intensities of EtBr was observed in (320-1200) nM RNase A concentrations range.

**Table S1**

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**Theoretical and calculated molar methanol yield : Efficiency of the hydrolysis was computed utilizing the Raman peak of methanol aqueous solutions.**

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<b>TMOS (v/v %)</b>	0.5	1.0	5	10
<b>Theoretical Methanol (mol/L)</b>	0.1353	0.2707	1.3533	2.7066
<b>Calculated Methanol (mol/L)</b>	0.1311	0.2250	1.2519	2.5707
<b>Hydrolysis Efficiency (%)</b>	98.4	83.1	92.5	95.0

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**Table S2**

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**Methanol content of hydrolyzed TMOS formulations : A significant increase in methanol concentrations is demonstrated with the increase in TMOS concentrations.**

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<b>TMOS (v/v %)</b>	<b>Methanol Peak (Counts)</b>	<b>Methanol Content (v/v %)</b>
<b>0.5</b>	191.0	0.74
<b>1.0</b>	235.3	0.91
<b>5.0</b>	1288.2	5.06
<b>10.0</b>	2640.4	10.40

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**Table S3**

**Size and stability characterization of CaRGOS formulations : Hydrodynamic Size (DLS), Polydispersity Index (PDI) and Stability [zeta potential ( $\zeta$ )] characterization of CaRGOS.**

<b>Sample</b>	<b>DLS (nm)</b>	<b>PDI</b>	<b>Zeta Potential (mV)</b>
<b>CaRGOS without buffer (1.25 v/v %)</b>	$*0.79 \pm 0.11$	$0.983 \pm 0.026$	$-22.07 \pm 1.01$
<b>CaRGOS without buffer (0.5 v/v %)</b>	**	**	$-26.58 \pm 7.69$
<b>CaRGOS with Buffer (0.5 v/v %)</b>	$67.22 \pm 1.65$	$0.248 \pm 0.006$	$-10.50 \pm 1.66$
<b>CaRGOS with Buffer (0.5 v/v %) and miRNA 21</b>	$69.95 \pm 0.47$	$0.308 \pm 0.004$	$-20.04 \pm 1.26$
<b>CaRGOS with Buffer (0.5 v/v %) without miRNA 21</b>	$70.02 \pm 2.09$	$0.338 \pm 0.035$	$-22.07 \pm 1.01$

*\*A (~1 nm) Hydrodynamic size is insignificant*

*\*\*Count-rate too low for measurement*

**Table S4:**

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**Reverse Transcription (RT) reaction mixture for a 15  $\mu$ L reaction : 15  $\mu$ L reaction consists of 7  $\mu$ L master mix, 3  $\mu$ L of 5X primer and 5  $\mu$ L miRNA 21 sample (with or w/o CaRGOS).**

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<b>Component</b>	<b>Volume (<math>\mu</math>L) per 15- <math>\mu</math>L reaction</b>
<b>100 mM dNTPs</b>	0.15
<b>MultiScribe Reverse Transcriptase, 50 U/ <math>\mu</math>L</b>	1.00
<b>10X Reverse Transcription Buffer</b>	1.50
<b>RNase Inhibitor 20 U/<math>\mu</math>L</b>	0.19
<b>Nuclease-free water</b>	4.16
<b>5X miRNA</b>	<u>3.00</u>
<b>Total volume(<math>\mu</math>L)</b>	10.00

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**Table S5**

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**PCR reaction mixture for a 10  $\mu$ L reaction: Each 10  $\mu$ L reaction consists of 5  $\mu$ L master mix, 0.5  $\mu$ L of 20X primer, 3.17 $\mu$ L of nuclease-free water and 1.33 $\mu$ L of cDNA (RT product).**

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<b>Component</b>	<b>Volume (<math>\mu</math>L) per 10- <math>\mu</math>L reaction</b>
<b>20X miRNA Primer</b>	0.5
<b>Universal Master Mix</b>	5.00
<b>Nuclease-free water</b>	<u>3.17</u>
<b>Total Volume</b>	8.67

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