## Abstract

Recent studies have shown the emerging role of noncoding RNA (ncRNAs) such as piRNA in stress response and transgenerational inheritance. Zebrafish were placed under conditions mimicking global climate change stress: elevated temperature, decreased pH and reduced nutrient availability. The total RNA of zebrafish were then isolated and purified. This will be used to identify any steady-state level changes of the piRNA expressed from the genome in response to stress. Here we report the quality of the purified total RNA from the gonads of the zebrafishes.

## Introduction

### Significance

PIWI-interacting RNAs (**piRNAs**) belong to a broad class of regulatory RNA called non-coding RNA (ncRNA). piRNA are known to have transgenerational effects on gene expression, since they operate in sex cells, allowing epigenetic inheritance thru multiple generations.<sup>(1)</sup> It is anticipated that when an organism undergoes climatic stress such as heat shock, acidity, or nutrient deprivation, the piRNAs within the organism alters the expression of genes in sex cells and thus may lead to transgenerational genetic change. Intriguingly, zebrafish also rely on piRNAs for sexual determination and sex cell production. <sup>(2)</sup>

### Question

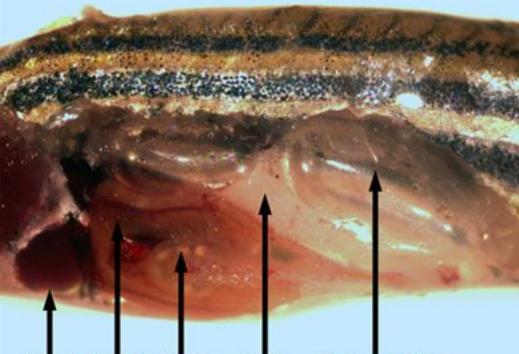
Do piRNAs play a significant role in response to adaptive stressors mimicking global climate change?

### Hypothesis

We hypothesize that piRNA expression has a role in regulating transgenerational adaptive stress response in Zebrafish.



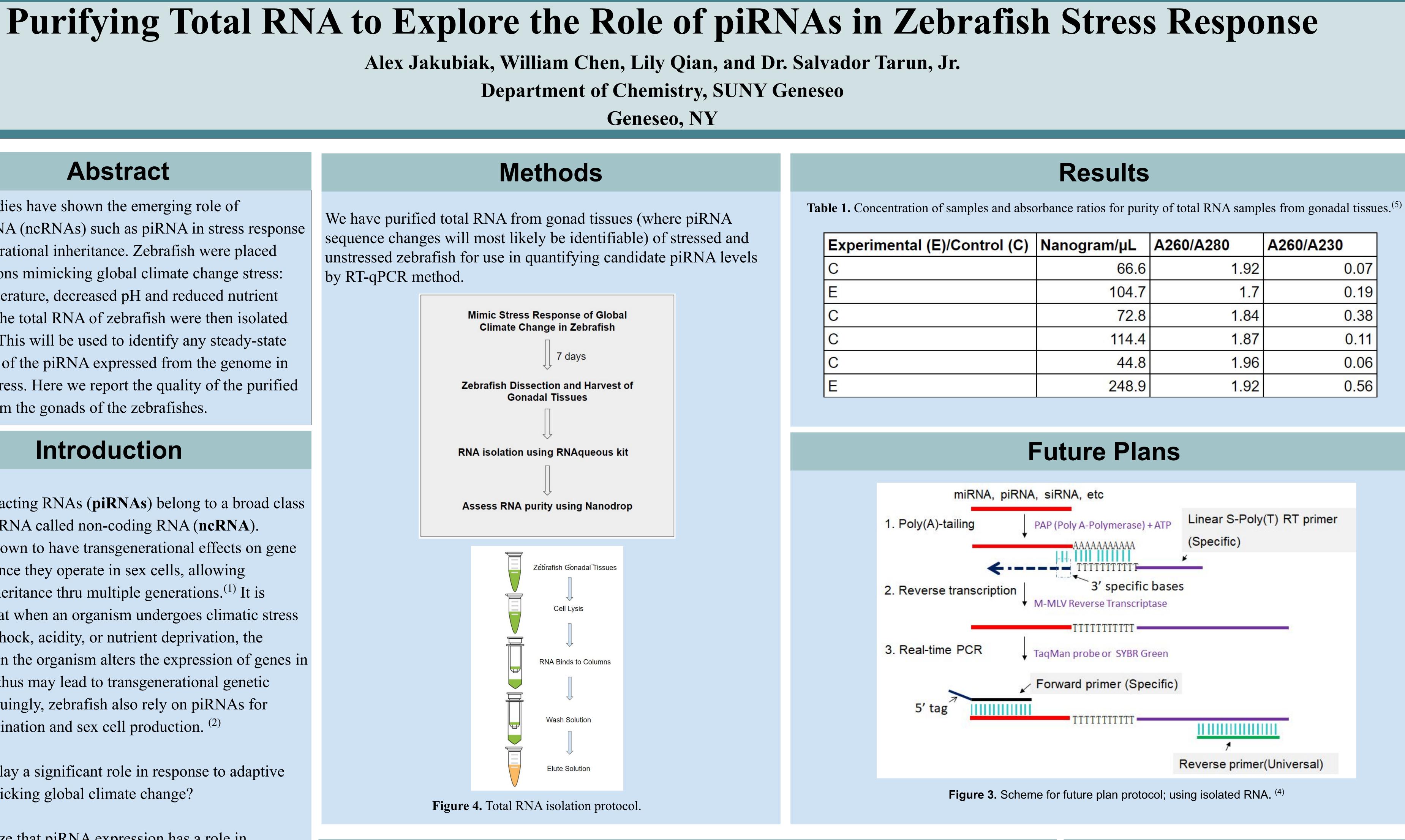
Figure 1. Experimental tank after 7 days



Swim bladder Figure 2. Example of a Zebrafish Dissection.<sup>(3)</sup>

# Acknowledgements

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## **Discussion/Conclusions**

### **Raising and Harvesting Zebrafish**

Maintaining stressed conditions (in particular, pH) in the experimental tank proved to be difficult, as a result of the pH solution being diluted by the large volume of the tank. However, we maintained conditions of temperature and nutrient stress consistently throughout the experiment.

### **RNA Purity and Contamination**

It seems that the quality and amount of RNA isolated was sufficient in some samples. Since the goal was to obtain A260/280 values of approximately 2.0, most of our samples came close.<sup>(6)</sup> In others, we see significantly less than 2.0, indicating a contamination of DNA in our RNA samples. This was most likely a result of ineffective DNAse treatment. However, our issue lie in the contamination of our RNA. Our goal was to have a A260/230 value around 1.8-2.2, and all of our samples reveal a significant deviation from this value, suggesting high contamination of our samples. This is most likely due to residual reagents involved in the purification process. This requires us to further improve our method of purification. **Future Plans** 

To remedy the contamination issue, we plan to redo the experiment and wash our RNA more extensively before the elution step.<sup>(5)</sup> Our future plan is to follow this up with RT-qPCR of our isolated RNA. By doing this, we plan to quantify candidate piRNA levels that we have already identified by bioinformatic analyses.

Experimental (E)/Control (C)	Nanogram/µL	A260/A280
С	66.6	1.92
E	104.7	1.7
С	72.8	1. <mark>8</mark> 4
С	114.4	1.87
С	44.8	1.96
E	248.9	1.92

## References

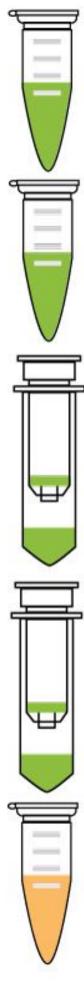
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## A260/A230 0.07 0.19 0.38 0.11 0.06 0.56

Mimic Stress Response of Global Climate Change in Zebrafish	
↓ 7 days	
Zebrafish Dissection and Harvest of Gonadal Tissues	
RNA isolation using RNAqueous kit	
Assess RNA purity using Nanodrop	

Simulate Global Climate Change  $\rightarrow$  Zebrafish Dissection  $\rightarrow$  RNA isolation using RNAqueous kit  $\rightarrow$  Assess RNA purity using Nanodrop



Mammalian cells or tissues



Cell lysis



RNA binds to the columns



Wash



Elute RNA