

Differential Plasma Exosomal Long Non-Coding RNAs Expression Profiles and Their Emerging Role in Smokers and Vapers

Gagandeep Kaur, PhD¹, Kameshwar Singh, DMV¹, Krishna Maremanda, PhD¹, Dongmei Li, PhD², Hitendra Chand, PhD³ and Irfan Rahman, PhD¹

¹Environmental Medicine, University of Rochester, NY, ²Clinical and Translational Science Institute, University of Rochester Medical Center, Rochester, NY, ³ Department of Immunology and Nanomedicine, Florida International University, Miami, FL.

INTRODUCTION

- •Long non-coding RNA (lncRNA) are diverse se transcripts that play a critical role in biological processes regulation, transcription, post-transcrip gene modification, and chromatin remodeling.
- •Studies have reported the presence of **lncRNA** in the exosomes.
- •Exosomes are the smallest subtype of extracellular vesicles (30-150 nm in size) surrounded by phospholipid bilayer.
- •They are known to be involved in regulating **cell-to-cell** • Exosomal RNA extraction: Total RNA from exosomes was isolated using Exosomal RNA isolation kit (Norgen Bioteck communication in lung pathologies including lung cancer, Corporation, Cat# 58000) per the manufacturer's protocol. COPD, asthma and IPF.
- •Importantly, exosomal lncRNAs are being extensively investigated as potential diagnostic biomarkers, especially for cancers.
- •**Cigarette smoking** has been found to alter lncRNA associated with activation of crucial metabolic pathway. Thus, It is likely that tobacco smoke exposure can alter the lncRNA profile within the lung tissue and can be involved in the pathogenesis of lung diseases.
- •Smoking habits amongst adults and adolescents are not confined to combustible tobacco cigarettes only. In the last few decades, the popularity of e-cigarette and related **products** is on a rise amongst young adults in the US.
- •We thus were interested in comparing how the lncRNA profiles vary amongst non-smokers, cigarette smokers and ecig users.
- Successful completion of this project will characterize the **IncRNA content of the plasma-derived exosomes** from various smoking populations to identify biomarkers relevant to pulmonary pathologies like COPD, asthma or IPF.

HYPOTHESIS

Plasma derived exosomal lncRNA profile is distinct amongst smokers, vapers and non-smokers and are responsible in regulating the inflammatory responses eventually leading to lung-related pathologies amongst these populations.

METHODS

et	of
es	like
oti	onal

- Ethics approval and consent to participate: All the protocols and procedures were approved by the Institutional Review Board (IRB)/Research Subject Review Board (RSRB) committee at the University of Rochester Medical Center, Rochester, NY with an approval number CR00002635. All the study participants recruited for the study signed an informed written consent form before recruitment and sample collection.
- Study design and subjects: 20-25 ml of venous blood was collected by venipuncture from smokers (CS), vapers (E-cig), and non smokers (NS). In brief, n=6-8 subjects were selected in each group.
- Plasma exosome isolation: We employed commercially available kit from Norgen Biotek (Cat# 57400; Ontario, Canada) to isolate exosomes from human plasma. Plasma exosomes were isolated as per manufacturer's protocol. Transmission electron microscopy (TEM) was used to visualize and characterize the isolated exosomes and nanoparticle tracking analysis was performed to analyze the particle size and concentration.
- Library preparation and Microarray analysis: GeneChip™ WT Pico kit (Cat# 902622 and 902623, Applied Biosystems, Foster city, CA) was employed for transcriptome profiling of the RNA isolated from various experimental groups. The library preparation and microarray hybridization were outsourced to the Microarray & NextGen Sequencing Core at the Center for Functional Genomics at SUNY Albany.
- Statistical analysis: The long non-coding RNA (lncRNA) data from two batches were first normalized and they analyzed in the R/Bioconductor. Volcano plots were generated to highlight significant lncRNAs with significant fold changes (> 2-fold changes) and raw P-values (less than 0.0001). Heatmaps were generated to show the lncRNA expressions in different groups. To examine the overlaps in identified lncRNAs from related group comparisons, venn.diagram function in R/Bioconductor was used.



Hierarchical cluster analyses of differentially expressed

Heat map showing differentially expressed lncRNAs that are significantly varies between (A) Non-smokers and cigarette smokers, (B) Non-smokers and E-cigarette users, and (C) Cigarette smokers and E-cig users. These lncRNAs were identified based on individual individual lncRNA, and each column represents individual sample. The relative lncRNA expression is depicted according to the color scale as shown at the right side of the figure.

gene targets on pairwise

Significant differentially expressed lncRNAs on comparing E-cig users with non-smokers and their significance.

PROB

E-cig vs NS **TC0400010170** TC0500007633

TC0600013228

TC0600014328

TC070000961

TC0700010504

TC0700011130

TC0800007080

TC0800007593

TC1300009105 TC1400006773 TC1700012104

TC1900007497

- observed.

This study was supported by the National Institutes of Health (NIH) 1R01HL135613 and U54 CA228110.

Authors would like to thank **Ms Janice Gerloff** and **Dr. Naushad** Ahmad Khan for initial recruitment of subjects and technical assistance.







RESULTS

CID	GENENAME	log2 Fold	P. Value	Description
		change		
.hg.1	skusworbu	-0.854161017	3.15E-05	Putative Protein
.hg.1	lozorby	1.644179344	2.96E-05	RNA-binding protein like family member.
.hg.1	SLC2A12	-0.792272819	3.95E-05	Solute carrier family 2 member 12
.hg.1	OSTM1	0.764009224	1.32E-05	Osteoclastogenesis associated transmembrane protein 1
.hg.1	Transfer RNA-Cys (GCA) 1-1	1.115171958	4.41E-05	tRNA; Genhancer regulatory region
.hg.1	OSBPL3	0.721080223	6.85E-05	Oxysterol binding protein like 3
.hg.1	T324413 (miTranscriptome)	-0.862064073	2.30E-05	Unannotated
.hg.1	BNIP3L	3.736036439	2.80E-05	Bcl2 interacting protein
.hg.1	zoyberby	1.81372327	4.40E-05	Endonuclease reverse transcriptase family member
.hg.1	Inc-PCDH20-13	1.325335976	3.19E-05	RNA gene
.hg.1	miR4307	-0.779299012	3.36E-05	micro RNA
.hg.1	T156883 (miTranscriptome)	-1.361512953	4.10E-05	Unannotated
.hg.1	flarchoy	1.258862445	7.32E-05	Endonuclease reverse transcriptase

SUMMARY

13 lncRNA gene loci showed altered expression among e-cig users as compared to NS.

A **four-fold increase** in the TC0800007080.hg.1 gene locus that encodes Bcl2 interacting protein 3 like-protein (BNIP3L) was

Gene enrichment analyses revealed a significant change (p=0.049) in the expression of **lncRNAs involved in steroid binding** in the plasma-derived exosomes from e-cig users as compared to their non-smoking controls.

ACKNOWLEDGEMENTS