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14. ABSTRACT Early detection and prevention of respiratory infections is a critical mission of military and civilian medical facilities. For example, adenoviruses are a major cause of acute respiratory illness in the military. The hypothesis that different types of cells produce unique profiles of respiratory chemicals (gaseous biomarkers) is substantiated by apparent differences in cell-specific metabolic biochemistry. The research team comprised of Drs. Ivan Medvedev (WSU-Physics), Katherine Excoffon (WSU-Biology), and Jennifer Martin (AFRL) is studying the variability of respiratory chemical profiles for a variety of cellular species, as well as a range of metabolic, homeostatic, and infectious conditions. The analysis is done using THz chemical sensing facility at WSU and Gas Chromatography-Mass Spectrometry (GC-MS) facility at AFRL. We are studying variability of respiration associated with viral infections (such as adenovirus). We are studying the effects of variability of cellular metabolism in response to varying concentrations of glucose, insulin, melatonin, and cortisol intended to simulate a range of cellular stress conditions. An exhaustive range of respiratory biomarkers from cellular models will be collected to create biomarker libraries and investigate the feasibility of using this information for pre-symptomatic detection of related pathologies, unadulterated by competing bio-processes in human body.					
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1. **INTRODUCTION:**

Early detection and prevention of respiratory infections is a critical mission of military and civilian medical facilities. For example, adenoviruses are a major cause of acute respiratory illness in the military. The hypothesis that different types of cells produce unique profiles of respiratory chemicals (gaseous biomarkers) is substantiated by apparent differences in cell-specific metabolic biochemistry. We have recently demonstrated the use of THz chemical sensors in detecting chemicals produced by cell cultures in a well-controlled laboratory environment, thus, providing the experimental basis for investigation of this hypothesis. The research team comprised of Drs. Ivan Medvedev (WSU-Physics), Katherine Excoffon (WSU-Biology), and Jennifer Martin (AFRL) is studying the variability of respiratory chemical profiles for a variety of cellular species, as well as a range of metabolic, homeostatic, and infectious conditions. The analysis is done using THz chemical sensing facility at WSU and Gas Chromatography-Mass Spectrometry (GC-MS) facility at AFRL. To identify tissue specific respiratory bio-markers we are studying cellular respiration of the following normal human cell types: lung, liver. Liver cells will be used for baseline comparison. We are studying variability of respiration associated with viral infections (such as adenovirus). We are studying the effects of variability of cellular metabolism in response to varying concentrations of glucose, insulin, melatonin, and cortisol intended to simulate a range of cellular stress conditions. An exhaustive range of respiratory biomarkers from cellular models will be collected to create biomarker libraries and investigate the feasibility of using this information for pre-symptomatic detection of related pathologies, unadulterated by competing biochemical processes in the human body.

2. **KEYWORDS:** Adenovirus, Virus, Infection, Cellular, Metabolism, Bio-markers, Terahertz, Spectroscopy, Breath, Analysis

3. **ACCOMPLISHMENTS:**

▪ **What were the major goals of the project?**

The main objective of this research program is to determine if Adenovirus infections cause detectable changes in gaseous metabolites produced by in-vitro cell cultures. The quantitative analysis of gaseous metabolites is performed using complementary Terahertz and GC-MS sensing approaches.

This objective is being reached through the following three major tasks.

Major Task 1: To grow primary airway epithelia and liver cell and collection of related Respiratory Volatile Organic Compounds

Major Task 2: Collect all relevant Terahertz spectral libraries

Major Task 3: To infect airway epithelia with adenovirus and identify unique VOC bio-signature

▪ **What was accomplished under these goals?**

The list of major achievements during the second year (13-24 months) of the research program:

1. Seven runs of human primary epithelial cell growth (each lasting 4-6 weeks) have been conducted. Three runs had to be terminated due to bacterial contamination. Four runs were successful and allowed us to study gaseous metabolic biomarkers of adenovirus infections with THz and GC-MS methods.
2. In Year 1 we tentatively determined that primary epithelial cell gaseous metabolites manifested systematic trends in ethanol and acetaldehyde concentrations associated with adenovirus infection. Year 1 results shown in Figure 4 exhibit rather high concentration of ethanol (~1 ppm), which was traced in Year 2 to a contamination associated with the use of ethanol as a cleaning agent in the incubation facility. In year 2 we took extra precautions to minimize this contamination. All cell cultures were handled in a dedicated fume hood with hydrogen peroxide used as a cleaning agent.
3. Four successfully grown cell cultures we infected with adenovirus. These are referred to as Run 1, 2, 3, and 4 in what follows. The multiplicity of infection used in Year 2 runs was increased to 100 from 10 used in Year 1.
4. To better differentiate gaseous metabolites associated with viral infections we concurrently sampled infected and control cells to look for biomarkers of adenovirus infections.
5. Gaseous metabolites of Runs 1-4 were studied with THz sensing.
6. Gaseous metabolites of Run 2 for infected and control cells at 48 hours were analyzed with GC-MS instruments at AFRL chemical sensing facility (Table 1, Figures 11-14).
7. Gaseous metabolites of Run 4 for infected cells were analyzed with GC-MS instruments at AFRL chemical sensing facility. Six samples were studied: 1. Pre-infected cells; 2. Pre-infected media; 3. 4-hours after infection; 4. 24-hours after infection; 5. 48-hours after infection; 6. 72-hours after infection.
8. THz analyses of gaseous metabolites reveal systematic decrease of ethanol and acetaldehyde concentrations along the timeline of the experiment. Both control and infected cells manifest this effect. The initial concentration of ethanol and acetaldehyde is thought to be present in the cell nutritional media. The concentration of ethanol and acetaldehyde is gradually decreasing (Figure 8) due to evaporation and possible metabolic consumption/production by cells. We detected a slight difference between concentration decay time constant for ethanol and acetaldehyde between control and infected cells (Figure 10). The small margin of variance can be due to

metabolic differences between control and infected cells, but the magnitude of the difference does not allow us to state with confidence that ethanol and acetaldehyde can be used as biomarkers of viral infection.

9. GC-MS analyses for Runs 2 and 4 revealed several chemicals whose concentrations manifest patterns systematic with adenovirus infection (Figures 11-14, Table 1).
10. We are currently preparing to conduct several more cell growth sessions followed by infection with adenovirus to further validate the findings presented in this report.

Goals not met in the during the second year (13-24 months) period:

1. Bacterial contaminations during cell growth has limited us to 4 useful cell growths runs in Year 2. We are currently working on conducting several more cell growth sessions.
2. The GC-MS results presented in this report make us optimistic about finding gaseous biomarkers of viral infections and we are looking to further validate these findings.
3. We will work to develop additional THz spectral libraries for the chemicals deemed

A detailed account of our achievements and approaches follows.

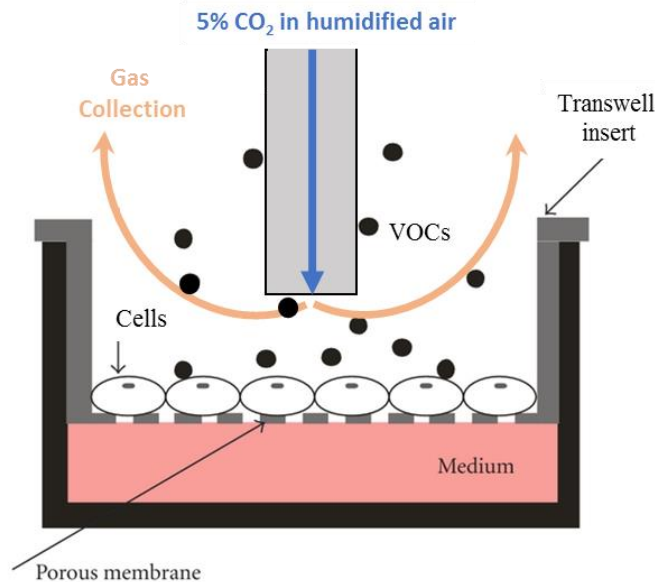


Figure 1. In-vitro THz identification of bio-markers of adenovirus. Organotypic cell cultures grown on a semi-permeable membrane inside a transwell insert and placed at the air-liquid interface (ALI).

Figure 1 shows our approaches to collection of metabolic VOCs from in-vitro cell cultures. In this project we focused on the development of an experimental system capable of collecting metabolic VOCs from

organotypic cell cultures grown at the air-liquid interface (ALI) shown in Figure 1. Cells are grown on top of semi-permeable membrane located at the bottom of a transwell. The transwell is placed on top of the medium. Nutrients from the medium are delivered to cell cultures through the membrane, but the medium surface is not directly exposed to the flow of buffer gas (in our case we use 5% CO₂ in humidified air). When cell cultures are fully polarized permeation of VOCs dissolved in medium is further reduced. This facilitates selective collection of metabolic VOCs from cell cultures. Buffer gas has two roles: 1) it creates the optimum environment for cell wellbeing (for epithelial lung cells it mimics gas composition in human lungs); 2) it transports metabolic VOCs to gas collection hardware. In our case we use Tedlar bags for gas sampling, which are certified by EPA for applications related to sampling and analysis of environmental gas mixtures. Tedlar bags are routinely used in PI's laboratory for collection and THz analysis of human breath, and have proven as a robust method of breath storage, with reliable sample retention up to 72 hours after sample collection.

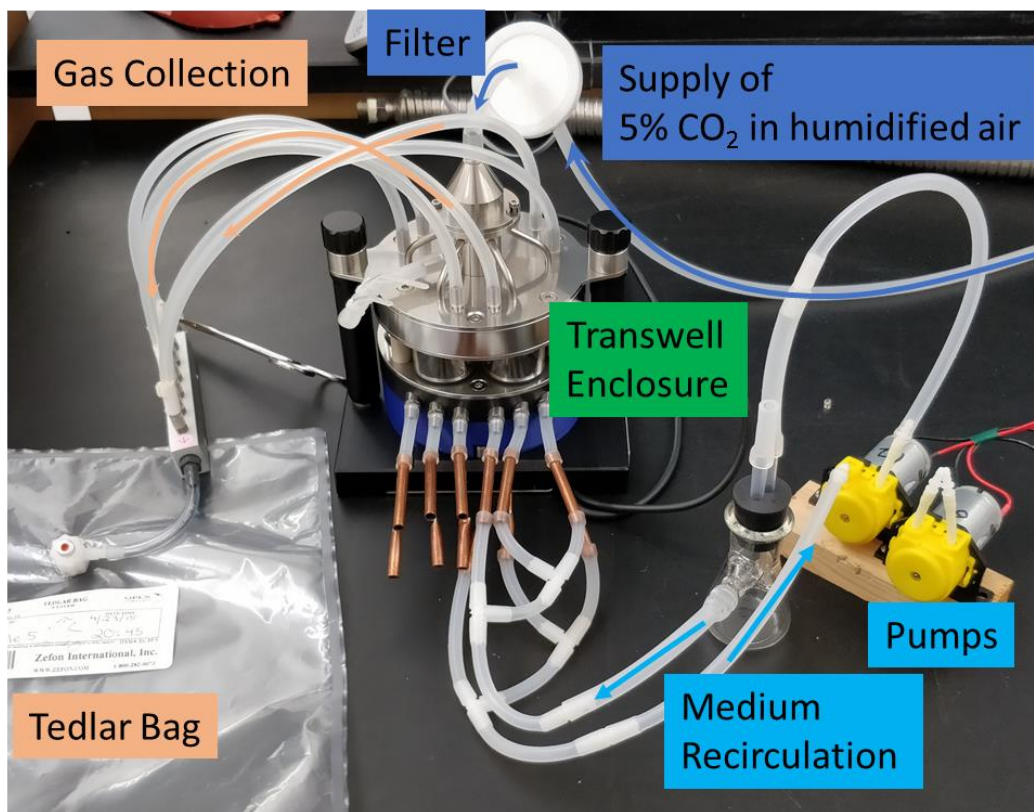


Figure 2. Cell exposure and gas collection system. Cultex exposure system shown in the center, is capable of housing 6 transwell insert. In Blue – A mixture of 5% CO₂ in humidified air flows through Cultex from a pressurized tank at a rate of 10 cc/min controlled by a flow mass controller (not shown in figure). In Orange – Buffer gas enriched with metabolic VOCs is collected into a Tedlar bag. In Cyan - Nutritional

medium is recirculated by a water pump. The set-up shown provides medium recirculation to 3 out of 6 enclosures.

Figure 2 shows the modified Cultex cell exposure system procured during this research. Figure 2 shows the custom media recirculation system comprised of 2 peristaltic pumps (shown in yellow), a Buchner flask for medium storage, and custom copper adapters to implement nested tubing for supply and return of media to individual chambers through a single inlet port. A custom base for Cultex system was 3-D printed (blue bottom of Cultex visible in Figures 2 and 3) to accommodate the additional tubing for the recirculation system. Cultex system has proven to be hard to clean and has resulted in bacterial contamination in some of the cell growth/infection runs. Cultex system was used in Run 4 with minimal contamination.

The results presented in this report for Runs 1-3 have been obtained using a custom-built exposure system shown in Figure 3. The system shown in Figure 3 allows us to grow cell cultures with greater surface area. The system consists of an aluminum base with a 4 inch opening for dishes with cell cultures. An air-tight lid has ports for buffer gas supply and return. The system is temperature controlled at 37 degrees Celsius by means of two heaters controlled by two Omega temperature controllers, with temperature readout provided by two thermocouples. The heaters and thermocouples are placed on two sides of the well for better temperature uniformity.

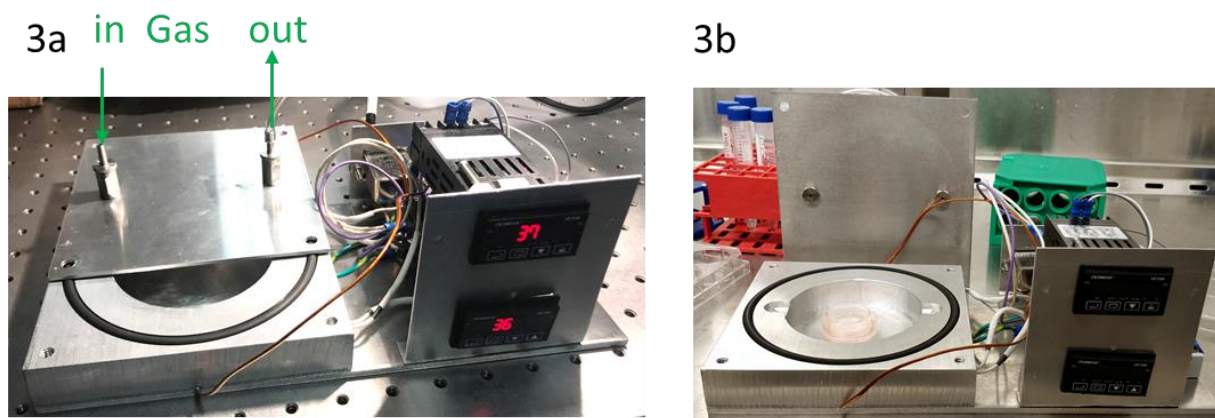


Figure 3. Custom bio-containment/exposure system designed to collect gaseous VOCs from cell cultures. 3a - Bio-containment/exposure system is temperature stabilized at 37°C. 3b - shows dish containing a 2" transwell with medium.

The results presented below for Runs 1-3 were obtained with cells placed into a 2" transwell inside the custom exposure system shown in Figure 3. The cells were grown on a semi-permeable membrane which provided facilitated medium isolation from the buffer gas stream.

Gas chromatography mass spectrometry (GC-MS)

All sorbent tubes were thermally desorbed on a Markes International TD-100 thermal desorber and analyzed on a Trace Ultra-ISQ gas chromatograph in line with a single quadrupole mass spectrometer (Thermo Scientific, Waltham, MA). Thermal desorption and GC-MS analysis of sample tubes were conducted as described previously¹. Briefly, thermal desorption was carried out at 310 °C over 10 min. Trap settings were as follows: flow path temperature 160 °C, trap flow rate 50 ml min⁻¹, trap purge time 1 min, trap low temperature of 25 °C, trap high trap temperature 315 °C, trap heating rate 40 °C s⁻¹ and post trap split 3.5 : 1. TO-14 A internal standards, bromochloromethane, 4-bromofluorobenzene, chlorobenzene-d5, and 1,4-difluorobenzene, were applied automatically to the sorbent tube by the Markes TD-100 prior to thermal desorption (25 ppm, Linde Gas North America, Stewartville, NJ). GC separations were carried out on a Restek Rxi-624Sil GC column (Bellefonte, PA, 60 m × 0.32 mm ID × 1.80 μm df), with a constant flow of helium (2 ml min⁻¹), over a temperature range of 40 to 240 °C at an increase of 10 °C min⁻¹ with a 20 min hold time at the maximum temperature. Mass spectral analysis was carried out via electron impact ionization at 70 eV with an ion source of 275 °C while scanning over a 35–300 m/z range every 0.154 s. Data was acquired using the Thermo Scientific Trace finder EFS software package (v. 3.0). Raw data file conversion from .RAW to .CDF occurred using the File Conversion tool as part of the Xcalibur software package (v. 3.0, Thermo Scientific).

A novel Terahertz gas sensor utilized in this program for chemical analysis has been recently developed in the PI's laboratory with funding from AFOSR². It can detect a wide range of gaseous chemicals at a part per billion to part per trillion level of dilution. The distinct advantage of this technology is the near 'absolute' specificity (extremely low probability of a false alarm) stemming from the highly redundant spectroscopic signatures being used for chemical detection.

¹ Harshman, S.W., Geier, B.A., Fan, M., Rinehardt, S., Watts, B.S., Drummond, L.A., Preti, G., Phillips, J.B., Ott, D.K., and Grigsby, C.C. (2015). "The identification of hypoxia biomarkers from exhaled breath under normobaric conditions". *Journal of Breath Research* 9, 047103.

² "Terahertz Spectroscopic Chemical Sensor for Analysis of Fatigued Human Breath", STTR Phase II, Air Force Research Laboratory, FA8650-16-C-6693

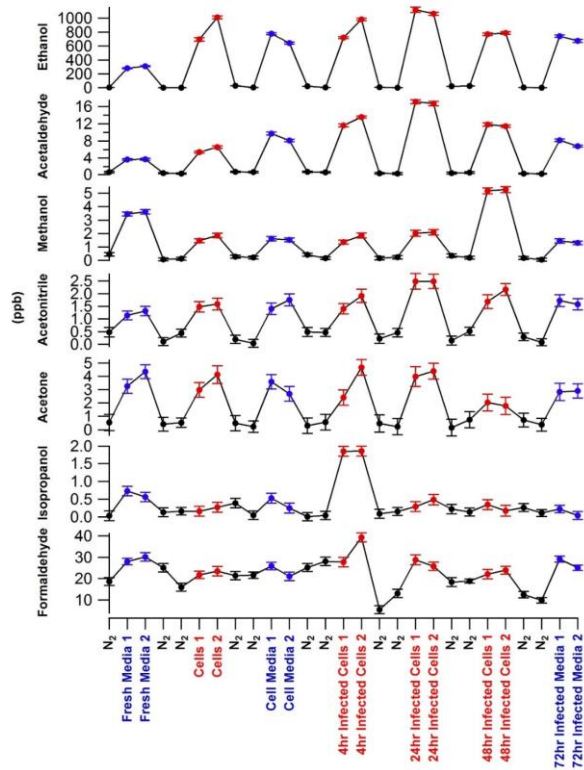


Figure 4. THz quantitative analyses of gaseous samples collected from a sample of healthy Primary human epithelial cell cultures in Year 1. Blue markers correspond to composition of media. Black markers correspond to composition of Tedlar bags filled with pure nitrogen. Red markers correspond to VOCs detected in presence of cell cultures. Consecutive unicolor markers shown correspond to sampling from the same Tedlar bag. All concentrations are measured in parts per billion.

Figure 4 shows Year 1 results from the Year 1 report. Concentrations of gaseous samples produced by primary epithelial cells were determined with THz sensing at WSU. The concentrations of ethanol and acetaldehyde appear to increase and peak at 24h after infection by adenovirus. The high concentration of ethanol (~1 ppm) exceeds the usual abundance of ethanol in human breath, which made us think that the most likely source of ethanol is an external contamination, as ethanol is used as a cleaning agent in Dr. Excoffon's laboratory. In Year 2 we took extra precautions to minimize this contamination. We moved all cell growth and manipulation hardware to a separate hood and started to use hydrogen peroxide as a cleaning agent. This yielded good results, as detected ethanol and acetaldehyde levels dropped significantly. Figures 5,6,7 show much lower levels of ethanol.

In Year 2 we conducted 4 successful cycles of primary human epithelial cell growth and infection referred to as Runs 1-4 in what follows. For Runs 1-3 gaseous metabolites were collected using custom

biocontainment/exposure system shown in Figure 3. For Run 4 gaseous metabolites were collected using Cultex system shown in Figure 4.

The growth of primary human airway requires feeder fibroblast irradiated cells and a unique cocktail of nutrients. The production of sufficient number of feeder fibroblasts and primary human airway epithelial cells requires 4-6 weeks for each run. At that time primary cells can be seeded in transwells as described below and polarized for a minimum of 7 days prior to experiment.

For Runs 1-3 three million primary human airway epithelial cells from healthy donors were seeded onto transwells (3 cm diameter, 0.4 μ m pore size). Cells were allowed to polarize over 7-14 days by incubation with polarization medium (1:1 mix of Dulbecco's modified Eagle's medium (DMEM) and F12 medium, 2% Ultrosor G serum substitute, 1% penicillin-streptomycin) on the basolateral surface. Medium was changed every 2-3 days. This resulted in polarized epithelia containing roughly 1.4 million cells. Epithelia were infected with wild human adenovirus (AdV5), suspended in phosphate buffered saline, at multiplicity of infection of 100 plaque forming units per cell for 2 hours at the apical and basolateral surfaces of the polarized epithelium. During this infection period, baseline gaseous sampling of the medium was performed in order to obtain a profile of the medium for background subtraction. Epithelia were then returned to the medium and gaseous samples were collected from uninfected control epithelia and infected epithelia at 4, 24, 48, and 72 hours after the end of mock (control) or viral exposure. Gaseous sampling of the conditioned media was performed along the timeline of the experiment in order to obtain a profile of the medium post-infection for background subtraction. THz analyses of gaseous samples for control and infected cell/media are shown in Figures 5,6, and 7.

Figures 5,6 and 7 show concentrations ethanol, acetaldehyde, acetone, methanol, methyl cyanide (also known as acetonitrile), isopropanol, n-propanol, and formaldehyde determined with THz sensing. For each time point we measured gaseous dilutions for cells and media in which cell resided (conditioned media). For conditioned media we removed the cells and collected gaseous samples evaporated by the media itself in absence of cell cultures.

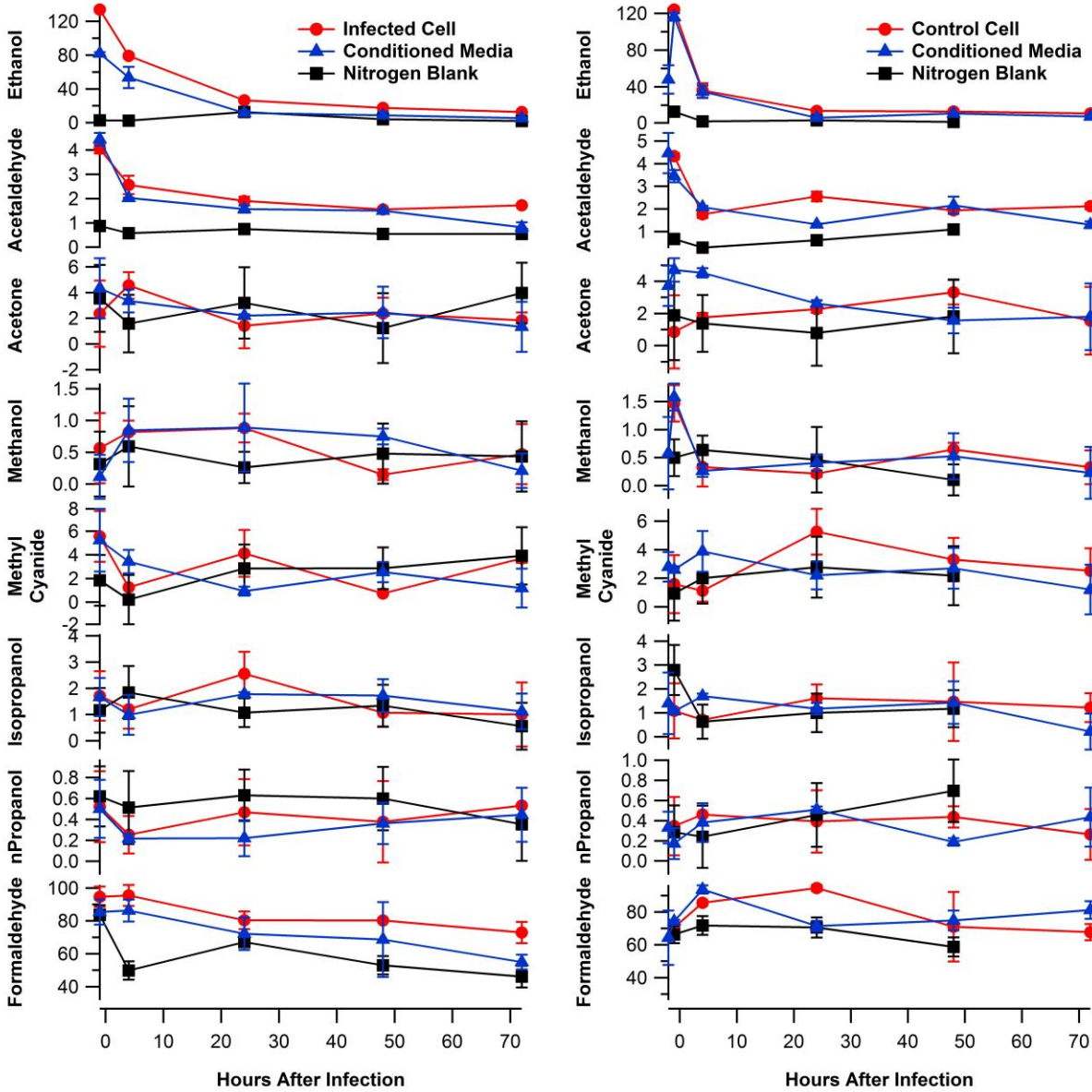


Figure 5. Gaseous metabolite concentrations determined with THz sensing during Run1. Gaseous sample were collected using custom bio-containment/exposure system shown in Figure 3. Gaseous dilutions for Control (right) and Infected (left) cell, Conditioned Media (blue) and Pure Nitrogen Without cells or media (black) are shown in part per billion. Ethanol and Acetaldehyde initial concentrations show gradual reduction due to either evaporation or metabolic consumption. Other chemicals do not show any significant pattern.

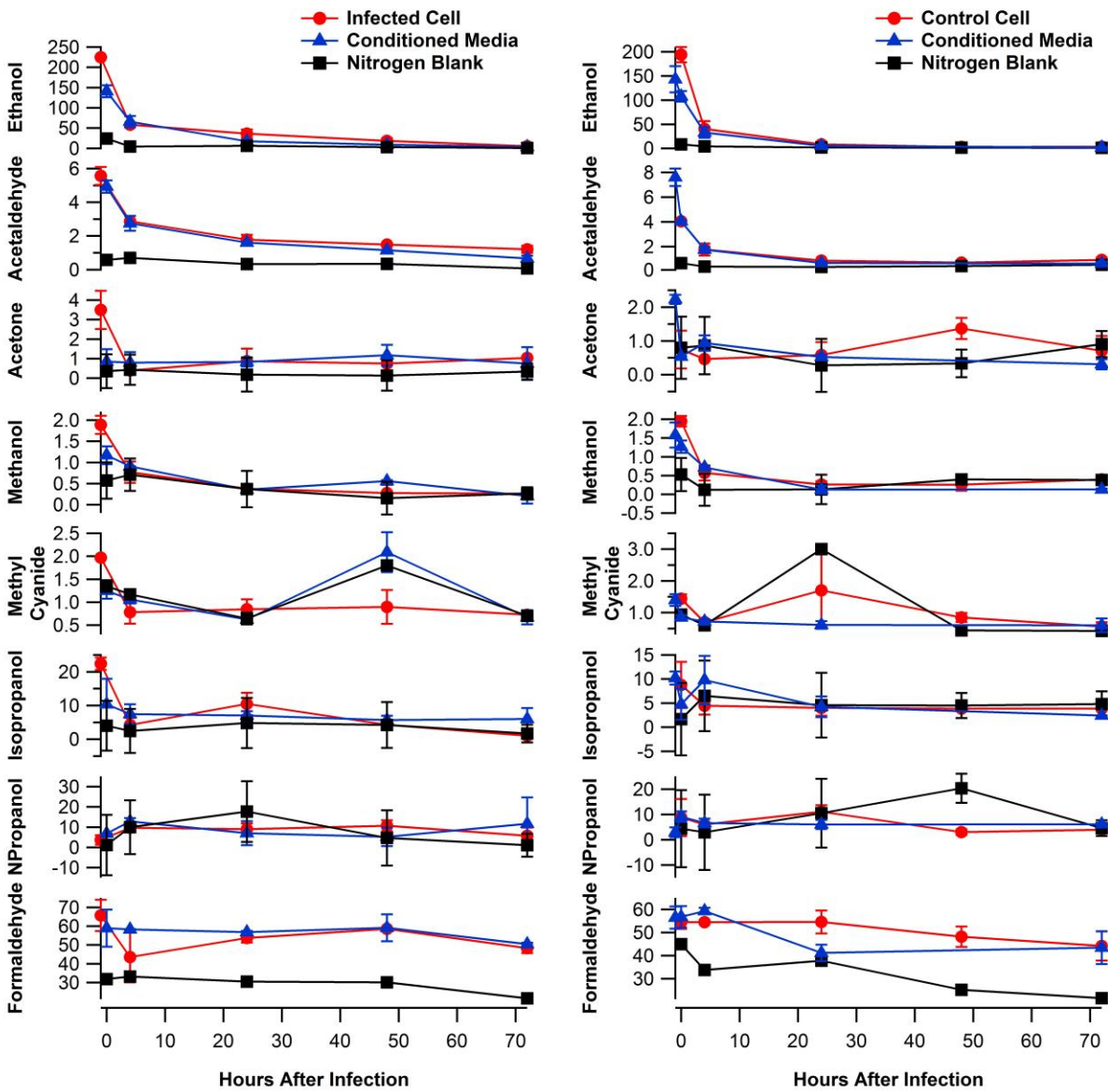


Figure 6. Gaseous metabolite concentrations determined with THz sensing during Run 2. Gaseous sample were collected using custom bio-containment/exposure system shown in Figure 3. Gaseous dilutions for Control (right) and Infected (left) cell, Conditioned Media (blue) and Pure Nitrogen Without cells or media (black) are shown in part per billion. Ethanol and Acetaldehyde initial concentrations show gradual reduction due to either evaporation or metabolic consumption. Other chemicals do not show any significant pattern.

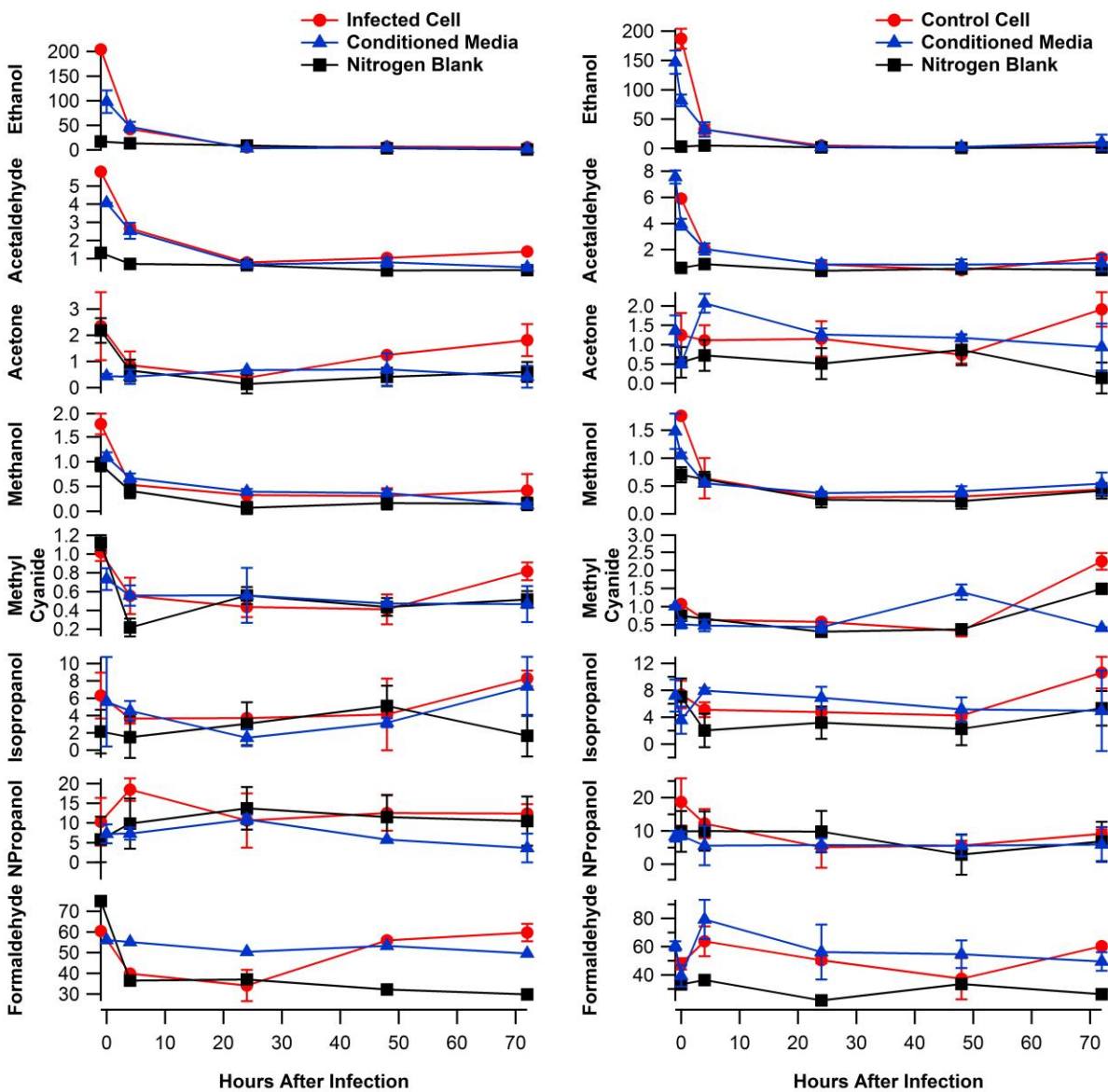


Figure 7. Gaseous metabolite concentrations determined with THz sensing during Run 3. Gaseous sample were collected using custom bio-containment/exposure system shown in Figure 3. Gaseous dilutions for Control (right) and Infected (left) cell, Conditioned Media (blue) and Pure Nitrogen Without cells or media (black) are shown in part per billion. Ethanol and Acetaldehyde initial concentrations show gradual reduction due to either evaporation or metabolic consumption. Other chemicals do not show any significant pattern.

In Runs 1-3 ethanol and acetaldehyde show systematic reduction in dilution along the timeline of the experiment. The initial abundances of these chemicals are most likely due to their presence in nutritional media in trace amounts. Other chemicals show no prominent patterns. To further investigate the patterns of ethanol and acetaldehyde we fitted their concentration to exponential decay equation ($A e^{-t/\tau}$, where τ is a decay constant). Excessive scatter in acetaldehyde Run-1 control set did not allow us to get a good fit

of this data. Figure 8 shows the fitting results while Figure 9 show the graphs of decay constants

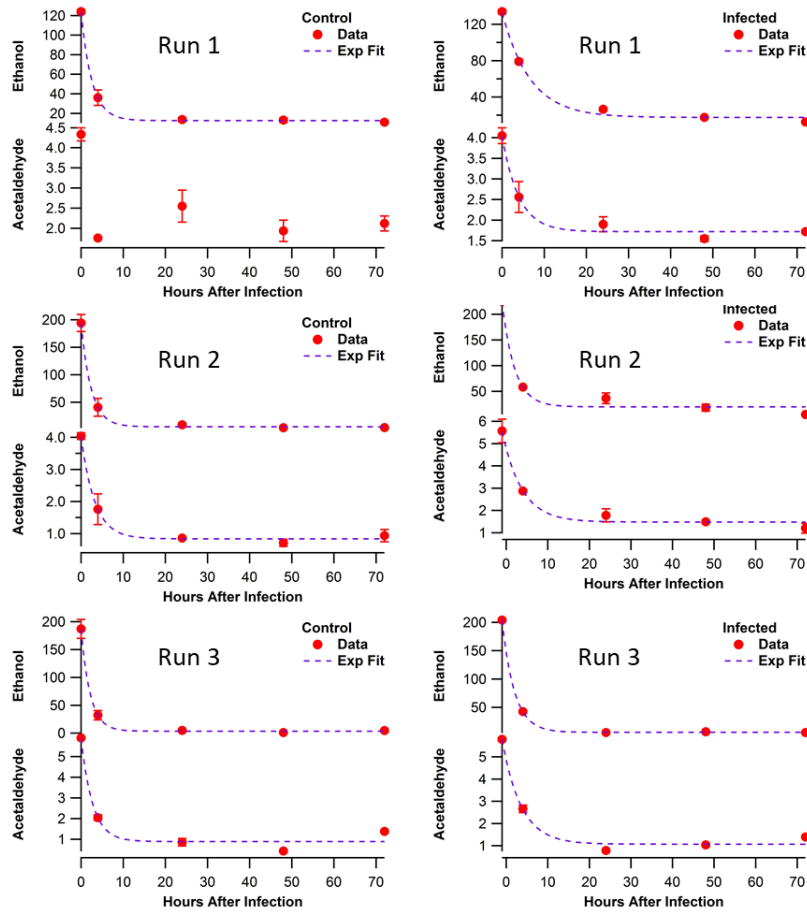


Figure 8. Fitting of ethanol and acetaldehyde concentrations (red) for Runs 1-3 to exponential decay equation. The fitted curves are shown in blue dashed lines. Infected cells exhibit consistently high decay constants.

for ethanol and acetaldehyde. The control (red trace in Figure 9) shown systematically lower decay constant than infected cells (green trace in Figure 9). While the decay itself is most likely due to evaporation, the systematic differences in decay constant shown in Figure 9 can be an indication of metabolic differences between infected and control cells and deserve further investigation.

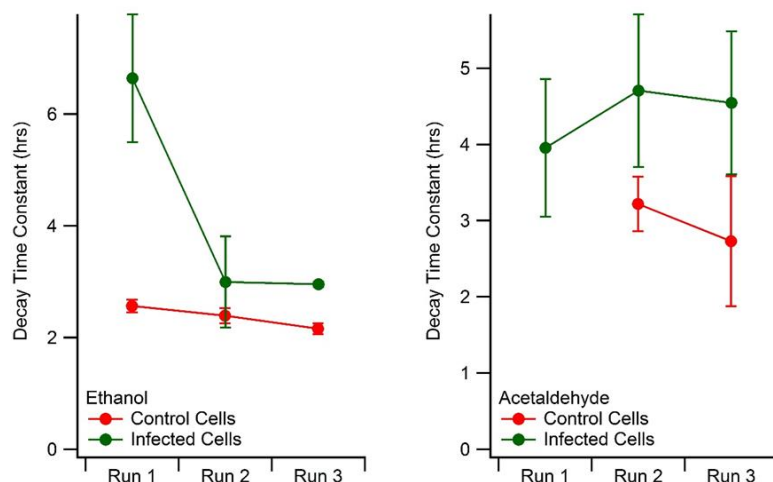


Figure 9. Decay time constants (τ) for ethanol (left) and acetaldehyde (right) dilutions determined with THz sensing. Control cells (red) exhibits systematically lower decay constant than infected cells (green).

For Run 4 one million primary human airway epithelial cells from healthy donors were seeded onto transwells (0.6 cm diameter, 0.4 μ m pore size). Cells were allowed to polarize over 7 days by incubation with polarization medium (1:1 mix of Dulbecco's modified Eagle's medium (DMEM) and F12 medium, 2% Ultrosor G serum substitute, 1% penicillin-streptomycin) on the basolateral surface. Medium was changed every 2-3 days. This resulted in polarized epithelia containing roughly .8 million cells. Epithelia were infected with wild human adenovirus (AdV5), suspended in phosphate buffered saline, at multiplicity of infection of 100 plaque forming units per cell for 2 hours at the apical and basolateral surfaces of the polarized epithelium. During this infection period, baseline gaseous sampling of the medium was performed in order to obtain a profile of the medium for background subtraction. Epithelia were then returned to the medium and gaseous samples were collected from uninfected control epithelia and infected epithelia 4, 24, and 48 hours after the end of mock (control) or viral exposure. Gaseous sampling of the conditioned media was performed along the timeline of the experiment in order to obtain a profile of the medium post-infection for background subtraction. THz analyses of gaseous samples for control and infected cell/media are shown in Figure 10.

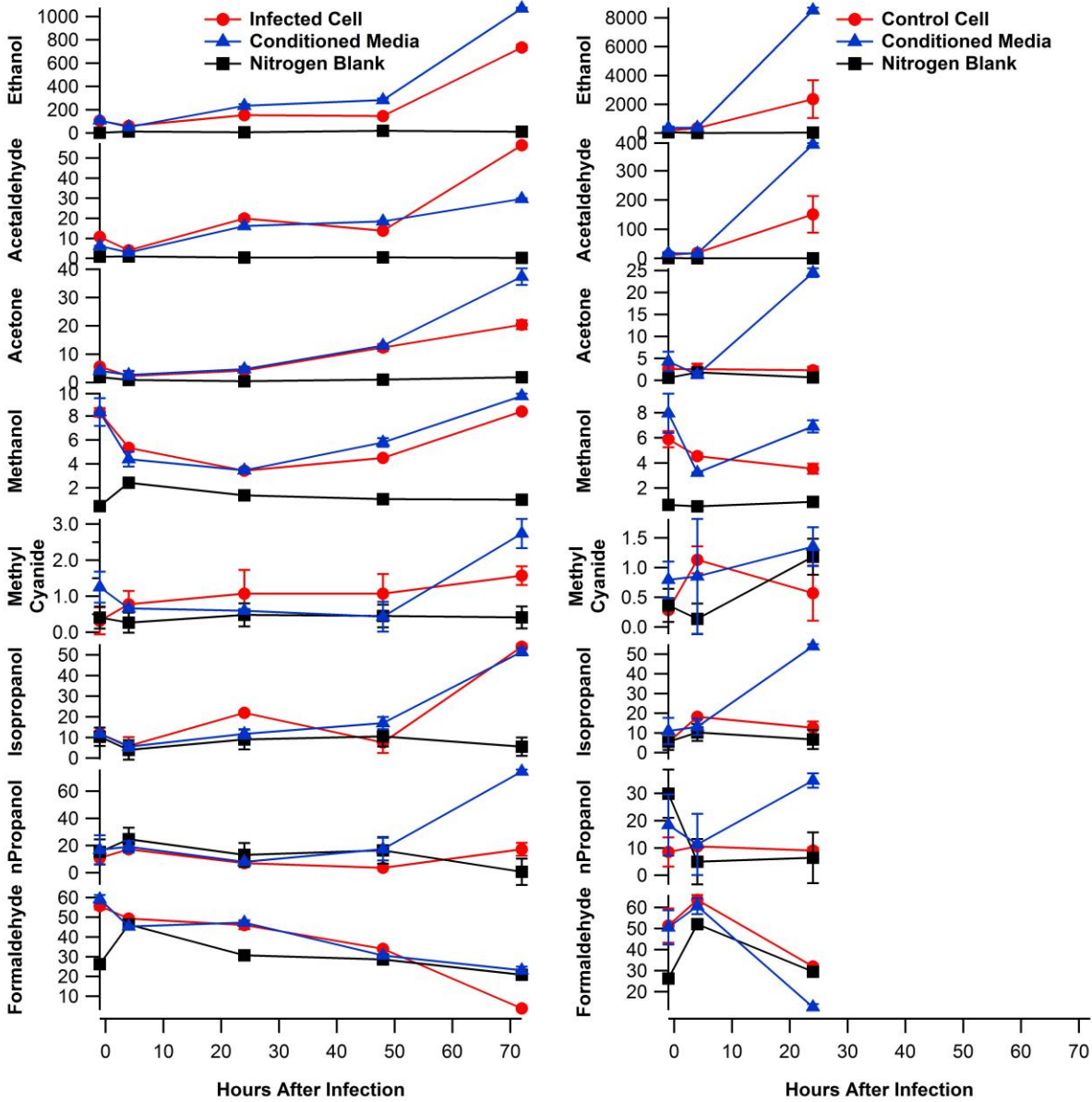


Figure 10. Gaseous metabolite concentrations determined with THz sensing during Run 3. Gaseous sample were collected using Cultex bio-containment/exposure system shown in Figure 2. Gaseous dilutions for Control (right) and Infected (left) cell, Conditioned Media (blue) and Pure Nitrogen Without cells or media (black) are shown in part per billion.

Figure 10 shows THz analyses of gaseous samples collected during Run 4, which was done on Cultex exposure system. Notably the concentrations of ethanol, acetaldehyde, acetone, methanol, methyl cyanide, and iso-propanol all appear to increase during infected cell run. No noticeable contamination was noticed during infected run, but the control cell run had to be abandoned after 24 hours due to obvious bacterial contamination of the control cell culture. The bacterial contamination most likely started during the infected

cell run, which was done first, and got rather prominent during the control cell run despite our efforts to clean the Cultex system with hydrogen peroxide. Despite this we think the GC-MS data collected during Run 4 (see below) provides valuable insight into the gaseous metabolite of viral infection, as the bacterial contamination did not manifest itself up until 48 hours after the infection.

The GC-MS analyses were conducted at AFRL using the state-of-the-art hardware described earlier. Instead of Tedlar bags, the gaseous metabolites were directly loaded onto sorbent tubes containing Tenax sorbent material. A total of 0.5L of gas was drawn through sorbent tubes for each collection. For Run 2 we sampled gas at 48 hour mark for control and infected (labeled T1 and T2 in Table 1). For Run 4 we collected 6 samples: T1 – pre-infected cells (right before infection), T2 – pre-infected media (media conditioned by cell right before infection), T3 – 4 hours after infection, T4 – 24 hour after infection, T5 – 48 hours after infection, T6 – 72 hours after infection. Table 1 summarizes the results of GC-MS analyses and provides chemical name, retention time in minutes (time of sample evolution from the gas-chromatography column), chemical dilutions and associated probabilities of identification. GC-MS analyses were run in a tentative identification mode, when chemicals are identified by fits against libraries. The software determines the match probability which needs to be used as guidance in determining the correctness of the measurement. We present Table 1 in its entirety, but highlight few chemicals (Figures 11, 12, 13, 14 and bold green font in Table 1) that exhibit prominent patterns that can be caused by the introduction of viral infection.

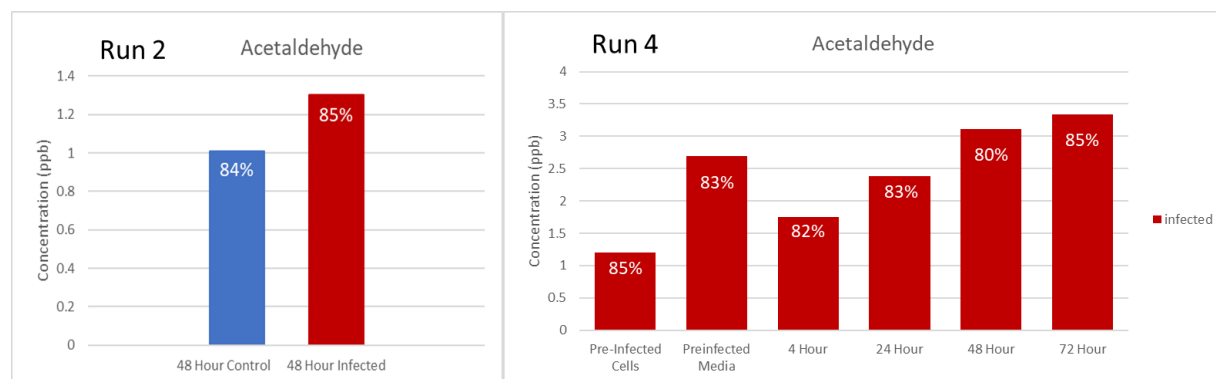


Figure 11. Detection of acetaldehyde with GC-MS. Run 2 measurements of control and infected metabolites at 48 hours are presented in the left pane, while Run 4 data represented on the right. Each bar is labeled with corresponding “Match Probability”.

Figure 11 shows concentrations of acetaldehyde detected in Runs 2 and 4. In both cases THz sensing returns a higher value of acetaldehyde dilution, which speaks for better retention of the analyte in THz

measurement. Notably, ethanol was only detected only in T6 for Run 4 with GC-MS (Table 1), which is emblematic of THz sensor ability to detect light volatiles such as ethanol. Concentration of acetaldehyde increases through Run 4 which tracks well the THz measurements shown in Figure 10. The rise of acetaldehyde (not detected in Runs 1-3) is most likely due to the onset of bacterial infection in the Cultex system, that became prominent during the control run (Figure 10).

Through analysis of data presented in Table 1 we identified several chemicals that could be linked to the onset of viral infections.

Figure 12 shows a rise in concentration of Caprolactam detected with GC-MS in Run 4. No Caprolactam was detected in Run 2. Notably Caprolactam was identified as a possible biomarker of lung cancer in a publication by Wang et al.³. Caprolactam “(CH₂)₅C(O)NH” is a good candidate for THz detection, with detection at a part per billion level of dilution.

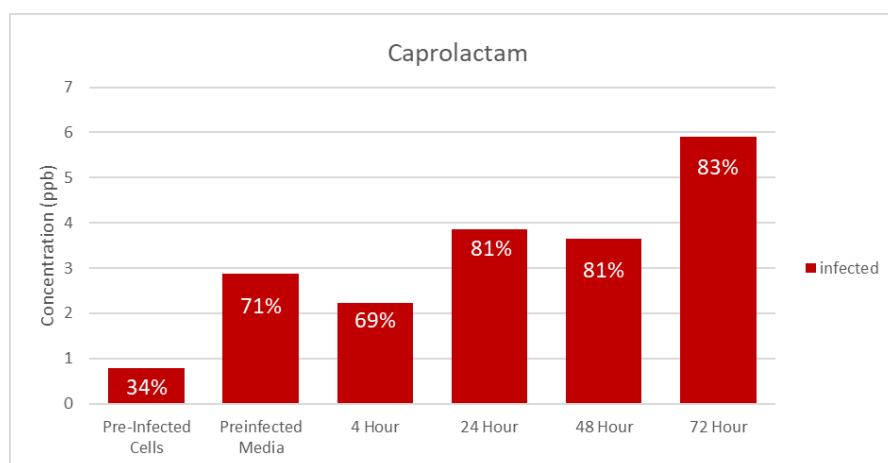


Figure 12. Detection of Caprolactam with GC-MS in Run 4. Each bar is labeled with corresponding “Match Probability”.

Figure 13 shows a rise in concentration of 1,2-Benzisothiazole detected with GC-MS in Run 4. No 1,2-Benzisothiazole was detected in Run 2. 1,2-Benzisothiazole “C₇H₅NS” is a good candidate for THz detection, with detection at a part per billion level of dilution.

Figure 14 shows a rise in concentration of acetophenone detected with GC-MS in Run 4. In Run 2 acetophenone was detected at slightly higher concentration in infected cells a 48h mark. The fact that it was present in 48-hour control in Run 2 makes it a less likely candidate as a biomarker of viral infection. Acetophenone “C₆H₅CCH₃” is a good candidate for THz detection, with detection at a part per billion level

³ Wang, C., Dong, R., Wang, X., Lian, A., Chi, C., Ke, C., Guo, L., Liu, S., Zhao, W., Xu, G., and Li, E. (2014). "Exhaled volatile organic compounds as lung cancer biomarkers during one-lung ventilation". Scientific reports 4, 7312.

of dilution.

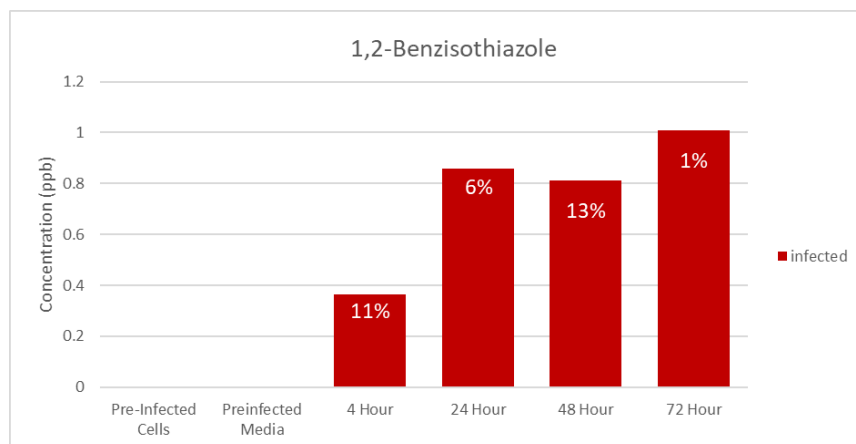


Figure 13. Detection of 1,2-Benzisothiazole with GC-MS in Run 4. Each bar is labeled with corresponding “Match Probability”.

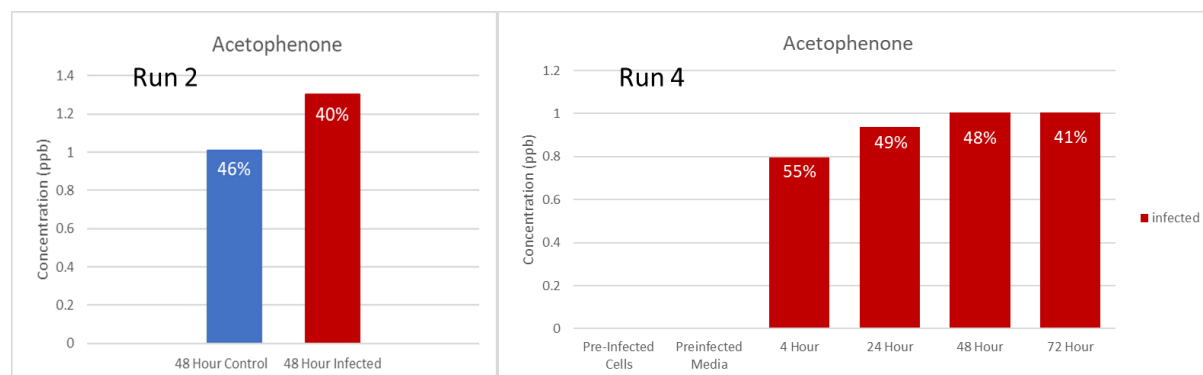


Figure 14. Detection of Acetophenone with GC-MS. Run 2 measurements of control and infected metabolites at 48 hours are presented in the left pane, while Run 4 data represented on the right. Each bar is labeled with corresponding “Match Probability”.

2-Butene was detected by GC-MS in 48-hour and 72-hour samples. 2-Butene “C₄H₈” exists in two conformations (cis and trans) and is a very good candidate for detection at a part per billion level of dilution. Ethanol, acetonitrile, and 1 -propanol were detected in 72-hour sample and are also detected by THz in all of Run 4 samples. We hypothesize that increases in concentrations for these chemicals are most likely caused by the onset of the bacterial contamination during Run 4.

2-methyl-propanal and 2-methyl-1-butanol were detected in 48-hour and 72-hour samples of Run 4. Both are excellent candidates for THz detection and will be validated in future experiments.

Spiro-1-(cyclohex-2-ene)-2-(5oxabicyclo[2,1,0]pentane) 1,4,2,6,6-pentamethyl were detected in 48-hour and 72-hour samples of Run 4. This molecule is an unlikely candidate for THz detection, but we will monitor

its presence in future experiments.

Table 1. Chemical analyses of gaseous samples from Runs 2 and 4 conducted with GC-MS.

Chemical Name	Retention Time (min)	Concentration (ppb)															
		Run 2				Run 4											
		T1		T2		T1		T2		T3		T4		T5		T6	
		Control 48 h	Match Probability	Infected 48 h	Match Probability	Pre-Infected Cells	Match Probability	Pre-Infected Media	Match Probability	4 h	Match Probability	24 h	Match Probability	48 h	Match Probability	72 h	Match Probability
Carbamic acid, monoammonium salt	3.31	0.42	60%	0.334	57%	0.473	33%	0.739	59%	0.657	59%	0.894	57%		0.565	59%	
Carbon dioxide	3.31	0.42	32%	0.334	31%	0.473	57%	0.739	32%	0.657	32%	0.894	31%		0.565	32%	
Aminomethanesulfonic Acid	3.79					0.67	4%	1.123	5%	0.939	5%	0.926	5%		1.154	5%	
Sulfur dioxide	3.79			0.201	22%	0.67	94%	1.123	89%	0.939	93%	0.926	90%		1.154	92%	
L-Alanine, 3-sulfo	3.79			0.201	66%												
1,3-Difluoro-2-propanol	3.8													1.914	12%		
Phosgene	3.8													1.914	2%		
1-Propene, 2- methyl	4.05													0.544	10%	0.52	
2-Butene	4.05													0.544	47%	0.524	
Acetaldehyde	4.22	1.223	84%	1.409	85%	1.198	85%	2.696	83%	1.745	82%	2.382	83%	3.106	80%	3.334	
Ethyne,fluoro-	4.22					1.198	0%			1.745	0%						
Ethylene Oxide	4.22	1.223	4%	1.409	5%			2.696	4%			2.382	4%		3.334	4%	
Oxiranemethanol (R)	4.22													3.106	3%		
Pentane	5.1			2.932	69%					6.842	73%						
Propane, 1-chloro-2-methyl	5.1																

Disulfide, bis[1-(methylthio)ethyl]	7.08	7.08	0.138	0.138	1%					0.814	0%	0.655	1%	0.59	0.59	1%	0.932	0.932	1%	0.652	0.652	1%			
Silanol, trimethyl-	7.08	7.08	0.138	0.138	41%							0.655	65%	0.59	0.59	70%	0.932	0.932	73%	0.652	0.652	70%	0.577	0.577	70%
tert-butyldimethylsilanol	7.08	7.08																					0.577	0.577	5%
Butanal, 2-ethyl-	7.21	7.21										0.468	2%												
Oxirane, 2,3-dimethyl-, trans-	7.21	7.21								0.278	9%														
1,3-Butanediol (S)-	7.21	7.21												0.371	0.371	20%						0.375	0.375	12%	
Butane, 1-(ethenyl)-3-methyl-	7.21	7.21												0.371	0.371	4%									
Cyclobutanol	7.21	7.21	0.094	0.094	13%																0.375	0.375	9%		
2-Propanone, 1-cyclopropyl-	7.2	7.2																					0.484	0.484	0%
Formic Acid ethenyl ester	7.2	7.2																					0.484	0.484	1%
2-propen-1-ol-2-methyl. Acetate	7.21	7.21								0.278	3%														
4-Methyl-1,3,2-dioxathiene 2-oxide	7.22	7.22	0.094	0.094	4%							0.468	11%												
Hexane, 3,3,4,4-tetrafluoro-	7.27	7.27	0.225	0.225	6%							0.415	2%	0.327	0.327	2%									
Methanesulfonyl azide	7.27	7.27	0.225	0.225	15%							0.415	11%	0.327	0.327	23%									
2(3H)-Furanone, dihydro-3,4-dimethyl	7.27	7.27									0.366	0%													
Isoxazolidine, 5-ethyl-2,4-dimethyl	7.27	7.27									0.366	0%													
5-[2-(Trimethylsilyl)cycloprop-1-ylidene]cyclopenta-1,3-diene	7.46	7.46								0.609	0%														
N-Acetylenediamine	7.46	7.46								0.609	0%														
Acetic acid	7.87	7.87	0.527	0.527	67%	0.724	32%	2.884	2.884	71%	3.062	18%	1.946	1.946	77%	2.802	2.802	77%	1.604	1.604	79%	2.241	2.241	75%	
Ammonium Acetate	7.87	7.87						2.884	2.884	23%	3.062	18%	1.946	1.946	17%	2.802	2.802	18%	1.604	1.604	15%	2.241	2.241	19%	
Acetic Acid anhydride with formic acid	7.87	7.87	0.5274	0.5274	16%	0.724	32%																		
Propanoic Acid, 2- (aminooxy)	8.09	8.09	0.134	0.134	1%																				

What opportunities for training and professional development has the project provided?

"Nothing to Report."

How were the results disseminated to communities of interest?

"Nothing to Report."

What do you plan to do during the next reporting period to accomplish the goals?

We will conduct several more cell growth/infection runs to further validate/discover gaseous biomarkers of viral infection. We will conduct GC-MS and THz analyses of gaseous samples collected from cell cultures.

4. IMPACT:

What was the impact on the development of the principal discipline(s) of the project?

Our preliminary results, outlined in this report, indicate the possible importance of caprolactam, acetophenone, 1,2 benzisothiazole, 2-butene, 2-methyl-1-butanol, and 2-methyl-propanal as a bio-marker of viral infections. We will further validate this funding. If confirmed the finding will contribute to scientific understanding of viral bio-chemistry. THz sensing technology used in this project will provide capability to track the onset of viral infection.

What was the impact on technology transfer?

"Nothing to Report."

What was the impact on society beyond science and technology?

"Nothing to Report."

5. CHANGES/PROBLEMS:

"Nothing to Report"

Changes in approach and reasons for change

"Nothing to Report"

Actual or anticipated problems or delays and actions or plans to resolve them

In Year 2 several cell growth experiments were aborted due to bacterial infection during incubation. We will be taking extra precautions in future experiments to avoid this problem.

Changes that had a significant impact on expenditures

Nothing to report.

Significant changes in use or care of human subjects, vertebrate animals, biohazards, and/or select agents

"Nothing to Report"

Significant changes in use or care of vertebrate animals.

"Nothing to Report"

Significant changes in use of biohazards and/or select agents

"Nothing to Report"

6. PRODUCTS:

"Nothing to Report."

7. PARTICIPANTS & OTHER COLLABORATING ORGANIZATIONS

What individuals have worked on the project?

- *Provide the following information for: (1) PDs/PIs; and (2) each person who has worked at least one person month per year on the project during the reporting period, regardless of the source of compensation (a person month equals approximately 160 hours of effort). If information is unchanged from a previous submission, provide the name only and indicate "no change."*

Name:	<i>Ivan Medvedev</i>
Project Role:	<i>PI</i>
Nearest person month worked:	<i>2</i>
Contribution to Project:	<i>Dr. Medvedev's laboratory performs THz analysis of the mixture, lead experimental design for the project. Dr. Medvedev performs overall coordination of the project.</i>
Funding Support:	

Name:	<i>Kate Excoffon</i>
Project Role:	<i>Co-PI</i>
Nearest person month worked:	<i>2</i>

Contribution to Project:	<i>Dr. Excoffon's laboratory is responsible for grows of cell cultures, viral exposure and sample collection. Dr. Excoffon assists Dr. Medvedev in project coordination.</i>
Funding Support:	

Name:	<i>Jennifer Martin</i>
Project Role:	<i>Co-PI</i>
Nearest person month worked:	<i>0</i>
Contribution to Project:	<i>Dr. Martin performs GC-MS analyses of gaseous cellular samples.</i>
Funding Support:	

Name:	<i>Priyanka Sharma</i>
Project Role:	<i>Research Scientist</i>
Nearest person month worked:	5
Contribution to Project:	<i>Dr. Sharma is responsible for growth of cell cultures.</i>
Funding Support:	

Name:	<i>Daniel Tyree</i>
Project Role:	<i>Graduate Student</i>
Nearest person month worked:	12
Contribution to Project:	<i>Mr. Tyree is responsible for design of experiments. He is responsible for THz sensing of cellular samples. In coordination with the PI he leads data analysis.</i>
Funding Support:	

Name:	<i>James Readler</i>
Project Role:	<i>Graduate Student</i>
Nearest person month worked:	3
Contribution to Project:	<i>Mr. Readler is responsible for growth and cell cultures. He assists the PI and Mr. Tyree in design of experimental protocols.</i>
Funding Support:	NIH/NIAID 1R01AI127816-01A1

Name:	<i>Hannah Benston</i>
Project Role:	<i>Undergraduate Student</i>
Nearest person month worked:	1
Contribution to Project:	<i>Ms. Benston is assisting Mr. Tyree in performing THz gas sensing and associated data analysis</i>
Funding Support:	<i>STTR Phase II, AFOSR FA8650-16-C-6693</i>

Has there been a change in the active other support of the PD/PI(s) or senior/key personnel since the last reporting period?

NIH/NIAID 1R01AI127816-01A1 (NIH) Excoffon (PI)

9/25/17-8/31/22

“Prevention of adenovirus pathogenesis through downregulation of the apical adenovirus receptor”; The molecular mechanism and anti-adenovirus efficacy of novel molecules that downregulate the expression of the apical adenovirus receptor and adenovirus infection will be investigated in epithelia in vitro and in the cotton rat model for adenovirus pathogenesis in vivo.

Role: PI

No overlap

What other organizations were involved as partners?

"Nothing to Report."