

## Gene Expression Profile in BCBL-1 Cell Line

Samantha Haines<sup>1</sup>, Sabrina Bishop, Devin M. Drown, Andrej Podlutsky<sup>1,2</sup>

<sup>1</sup>Department of Wildlife and Biology, <sup>2</sup>Institute of Arctic Biology, University of Alaska Fairbanks



### **OBJECTIVES**

Obtain gene expression profile of BCBL-1 cells using Oxford's Nanopore MinION sequencer

PROGRAM

CONSORTHIM

Determine differences in gene expression between latent and lytic BCBL-I cells following X-ray exposure

Identify DNA repair-associated gene expression in each treatment

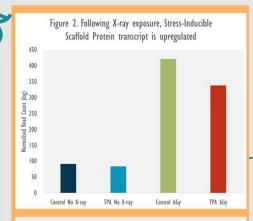
### **BACKGROUND**

- Faulty DNA repair leads to cancer formation<sup>1</sup>
- We previously found viral activity affects DNA repair
- Exact viral influence on DNA repair is unknown<sup>2</sup>
- BCBL-I cells are infected with cancercausing virus<sup>3</sup>
- Nanopore technology generates full length reads of viral transcripts<sup>4-6</sup>



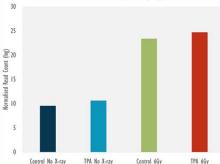
Figure 1. Computer generated image of Kaposi's Sarcoma Herpes Virus

### RESULTS



- SQSTM1—stress-inducible scaffold protein
- Regulates endosomal trafficking, apoptosis, and inflammation
- Known protein diagnostic and prognostic marker for many cancers<sup>7</sup>

### Figure 3. Following X-ray exposure, Cyclin-Dependent Kinase Inhibitor transcript is upregulated



- CDKNIA—cyclin-dependent kinase inhibitor
- Arrests cell at G<sub>1</sub> stage to repair damage or induce apoptosis<sup>8</sup>

# Figure 4. Following viral induction by TPA, Galectin-I transcript is upregulated 400 350 300 (80 1) 100 100 50

- LGALS I Galectin-I
- Modulates cell-cell and cell-matrix interactions

TPA No X-ray

Control 6Gy

TPA 6Gv

Regulates cell proliferation<sup>9</sup>

Control No X-ray

### **INTERPRETATIONS**

- Proof of principal concept demonstrated transcriptomic sequencing ability
- SQSTM1 expression may be due to its role in the apoptotic pathway in response to damage<sup>7</sup>
- CDKNIA expression likely related to cell cycle arrest at G<sub>1</sub> in order to repair damaged DNA or lead to apoptosis<sup>8</sup>
- Viral induction upregulates LGALS1 perhaps due to protein's ability to inhibit infectivity and viral production<sup>9</sup>

### **FUTURE DIRECTIONS**

- Analyze and interpret remainder of 16 genes identified
- Improve validity through separate replication of each time sample for treatments
- Examine isoforms of genes through analysis of long reads of mRNA from Nanopore
- Induce DNA damage with UV-C radiation to seek out repair pathway differences between UV and X-ray

### REFERENCES AND ACKNOWLEDGEMENTS

Authors would like to acknowledge support from the Biomedical Learning and Student Training (BLAST) Program at UAF. Research reported in the publication was supported by an institutional Development-Averal (DLAs) from the Nazional Institute of General Medical Sciences of the Nazional Institute of Health under grant number 2920GH103795. The content is solely the responsibility of the suplices and does occasiately related the efficial select of the DHI.

BLST is supported by the NH Common Fund, through the Office of Strategic Coordination, Office of the NHI Director with the Inited swards 174.CMH 189974, LDV (2011 189974, LD) COUNT 18991 The content is solely be re-proprietally of the subtrons and does not necessaryly represent the official views of the National Institutes of Health, LIAF is an affirmative action / equal employment opportunity employer and educational institutions www.salaksa.cd/involved.com/scrimination.

Fig. 1 countery of Science Photo Library/Allary Stock Photo. All corn courtery of Microsoft.

1. Vig. 1 Somatic mustaions, groome mosalicism, cancer and aging, Curr Opin Genet Dev. 2014;(26):141-149.doi:10.1016/j.gde.2014.04.002.

2. Moore PS. Chang (Why do visues cause cancer! Highlights of the first century of human tumour virology. Nat Rev Cancer.2010;10(12):8784

- doi:10.1088/mc2961

  3. Yu.Y. Black JB, Goldenith CS, Browning PJ, Bhalla K, Offerman MK. Induction of human horpesvirus-8-DNA replication and transcription by butyriste and TPA in BCB-1 cells. J Cen Virel. 1999;(80):83-90.
- a survige y notation L. Amount D. G. in Certify Preferency in Amorphism and processors expending occurring growing references complexity in clinical samples. PLoS One. 2018; 1(3), doi:10.1371/journal/poor.191406
   5. Kilanskis A. Hass JR. Corrivosa Et. et al. Becterial and viral infendication and differentiation by amplicon sequencing on the MinION nanopore sequencing configuration.
- requestors agreement and 17-31 (17.2 doi:10.11.10941) //26/3-30-12.40-12

  6. Bistavilla, I.J. puri SE. Rodonie SE, sowbridge TL, Cogan NO. Metagenomic arbovirus detection using MinION nanopore sequencing J Virol Method 2017;24979-84 doi:10.1016/j.jviromet.2017.08.019

  7. Sinches-Mann R Konastu M. AG/SOSTMI Severies the cell through health and disease. [Cell Sci. 2018;131/21]. doi:10.1242/sci.222836.
- 2020. Accessed April 10, 2020.

  9. Chen Y, Zhou J, Cheng Z, et all. Functional variants regulating LGALS1 (Galectin 1) expression affect human susceptibility to influenza A(H7N9).

  Rev. 2015; Vil. 1: 10. doi: 10.1018/seve08517

### **METHODS**

- Cultured BCBL-I cells
- Activated viral production with
   12-O-tetradecanoylphorbol-13-acetate (TPA)
- Treated half the flasks with 6Gy X-ray to induce DNA damage
- Extracted RNA 4 hours following X-Ray treatment
- Sequenced transcriptome using the Nanopore MinION Sequencer (PCR-cDNA Barcoding kit)
- Used DESeq2 for differential expression analysis