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Scriptwriter Name: Pallavi Sharma

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Title: Establishment and Quantification of *De Novo* Lytic Infection by Cell-Free Kaposi's Sarcoma-Associated Herpesvirus

Authors and Affiliations:

Qing Zhu¹, Janvhi Suresh Machhar¹, Pinghui Feng², Chengyu Liang¹

¹Program in Molecular and Cellular Oncogenesis, The Wistar Institute ²Section of Infection and Immunity, Herman Ostrow School of Dentistry, University of Southern California

Corresponding Authors:

Qing Zhu (qzhu@wistar.org)
Chengyu Liang (cliang@wistar.org)

Email Addresses for All Authors:

Janvhi Suresh Machhar

Pinghui Feng

Qing Zhu

Chengyu Liang

(imachhar@wistar.org)

(pinghuif@usc.edu)

(qzhu@wistar.org)

(cliang@wistar.org)



Author Questionnaire

- Microscopy: Does your protocol require the use of a dissecting or stereomicroscope for performing a complex dissection, microinjection technique, or something similar? Yes, all done
- **2. Software:** Does the part of your protocol being filmed include step-by-step descriptions of software usage? **No**
- **3. Filming location:** Will the filming need to take place in multiple locations? **Yes**They are on the same floor, about 50 meters apart.

Current Protocol Length

Number of Steps: 18 Number of Shots: 35



Introduction

Videographer: Obtain headshots for all authors available at the filming location.

- 1.1. <u>Janvhi Machhar:</u> Tractable and reproducible systems to model KSHV primary lytic infection are limited. We established a reliable and robust *in vitro* platform for modeling KSHV de *novo* lytic infection.
 - 1.1.1. INTERVIEW: Named talent says the statement above in an interview-style shot, looking slightly off-camera. Suggested B roll: Figure 1

What are the current experimental challenges?

- 1.2. <u>Janvhi Machhar:</u> It's challenging to model KSHV lytic infection because of its strong latency bias, and the few primary cells that support it are delicate and have a limited lifespan, which severely limits reproducibility and scalability.
 - 1.2.1. INTERVIEW: Named talent says the statement above in an interview-style shot, looking slightly off-camera.

What significant findings have you established in your field?

- 1.3. <u>Janvhi Machhar:</u> We developed a tractable and reproducible model in HCT116 cells that captures efficient KSHV lytic replication during primary infection, offering a unique platform to study early events in herpesvirus replication.
 - 1.3.1. INTERVIEW: Named talent says the statement above in an interview-style shot, looking slightly off-camera. Suggested B roll: Figure 2

Videographer: Obtain headshots for all authors available at the filming location.



Protocol

NOTE: LAB MEDIA/SCOPE timestamps for protocol were added at the postshoot stage. Please contact the postshoot note integrator (Sulakshana Karkala) for queries regarding lab media

2. Production of Cell-Free BAC16-Derived KSHV

Demonstrator: Qing Zhu

- 2.1. To begin, seed five T75 (*T-Seventy*) flasks with Dox-inducible iSLK (*I-S-L-K*) cells containing wild-type KSHV (*K-S-H-V*) BAC16 (*Bac-Sixteen*) at approximately 80 percent confluency [1]. Incubate the flasks at 37 degrees Celsius in a humidified incubator with 5 percent carbon dioxide for 24 hours [2].
 - 2.1.1. WIDE: Talent seeding T75 flasks with Dox-inducible iSLK cells using a pipette.
 - 2.1.2. Talent placing the flasks in a 37 degrees Celsius incubator set at 5 percent carbon dioxide.
- 2.2. Add 1 microgram per milliliter of doxycycline and 1 millimolar sodium butyrate to each flask to induce lytic reactivation [1-TXT].
 - 2.2.1. Talent pipetting doxycycline and sodium butyrate into each flask. **TXT: Incubate** for 76 h and 96 h
- 2.3. Using phase-contrast microscopy, monitor the cells daily [1]. Collect the supernatant once more than 90 percent of cells show rounding, detachment from the flask surface, or visible lysis [4].
 - 2.3.1. LAB MEDIA: 2.3.1-shot-of-the-cell.tif.
 - 2.3.2. Talent collecting the supernatant from the flasks after confirming cell rounding and lysis.
- 2.4. Then, centrifuge the collected supernatants at 1,500 g for 10 minutes at 4 degrees Celsius to remove cellular debris [1]. Filter the resulting supernatant through a 0.45 micrometer polyethersulfone membrane filter without disturbing the pellet [2].
 - 2.4.1. Talent placing the supernatant tubes into a centrifuge set at 1,500 g and 4 degrees Celsius.
 - 2.4.2. Talent filtering the supernatant through a 0.45 micrometer polyethersulfone



membrane filter.

- 2.5. Transfer the filtered supernatants to ultracentrifuge tubes [1]. Centrifuge the tubes at 25,000 g for 3 hours at 4 degrees Celsius using an SW28 (S-W-Twenty-Eight) rotor to pellet the viral particles [2].
 - 2.5.1. Talent transferring filtered supernatant into ultracentrifuge tubes.
 - 2.5.2. Talent loading the tubes into an ultracentrifuge with an SW28 rotor and setting it to 25,000 g at 4 degrees Celsius.Videographer's Note: Settings on the centrifuge were incorrect for this shot.

Please blur or crop

- 2.6. Carefully discard the supernatant into a container with 10 percent bleach [1]. Resuspend the viral pellet in 1 milliliter of serum-free DMEM (*D-M-E-M*) [2].
 - 2.6.1. Talent pouring off the supernatant into a bleach-containing waste container.
 - 2.6.2. Talent resuspending the viral pellet in 1 milliliter of serum-free DMEM using a pipette.

3. De Novo Infection of HCT116 Cells

- 3.1. Seed 1×10^5 HCT116 (*H-C-T-One-One-Six*) cells per well in a 12-well tissue culture plate [1]. Incubate the plate at 37 degrees Celsius in a humidified incubator with 5 percent carbon dioxide for 24 hours [2].
 - 3.1.1. Talent seeding 1×10^5 HCT116 cells in each well of a 12-well plate using a pipette.
 - 3.1.2. Talent placing the culture plate inside a 37 degrees Celsius, 5 percent carbon dioxide incubator.
- 3.2. Dilute the concentrated virus stock in 250 microliters of pre-warmed, serum-free DMEM to achieve a multiplicity of infection of 10 [1].
 - 3.2.1. Talent pipetting virus stock into a microcentrifuge tube containing 250 microliters of pre-warmed, serum-free DMEM.
- 3.3. Aspirate the culture medium from each well [1] and add 250 microliters of the diluted virus directly onto the cells [2].
 - 3.3.1. Talent aspirating the culture medium from each well using a vacuum aspirator.
 - 3.3.2. Talent pipetting 250 microliters of the virus dilution into each well.
- 3.4. Centrifuge the plate at 1,500 g for 1 hour at 30 degrees Celsius to synchronize infection



- [1]. Immediately transfer the plate to a 37 degrees Celsius incubator and continue incubation for 1 hour [2].
- 3.4.1. Talent placing the plate into a centrifuge and setting it to 1,500 g and 30 degrees Celsius for 1 hour.
- 3.4.2. Talent moving the plate from the centrifuge into a 37 degrees Celsius incubator.
- 3.5. Aspirate the viral inoculum from each well [1]. Wash the cells three times with prewarmed PBS to remove residual virus [2]. Add 1 milliliter of complete DMEM to each well [3].
 - 3.5.1. Talent aspirating the inoculum from the wells.
 - 3.5.2. Talent washing each well three times with pre-warmed phosphate-buffered saline using a pipette.
 - 3.5.3. Talent adding 1 milliliter of complete DMEM to each well. **TXT: Monitor GFP** expression at 36 h, 60 h, and 96 h post-infection
- 3.6. Then, collect 1 milliliter of culture supernatant from each well [2].
 - 3.6.1. Talent pipetting 1 milliliter of culture supernatant from each well into labeled collection tubes.

4. Infectious Units Assay (IU assay)

- 4.1. Seed 1×10^4 SLK cells per well in a 96-well plate 24 hours before infection [1-TXT].
 - 4.1.1. WIDE: Talent seeding SLK cells into a 96-well plate using a multichannel pipette. **TXT: Seed cells to reach ~60–70% confluence on infection day**
- 4.2. Label nine 1.5-milliliter microcentrifuge tubes for a serial two-fold dilution series [1].
 - 4.2.1. Talent labeling each microcentrifuge tube with dilution ratios using a permanent marker.
- 4.3. Add 200 microliters of virus supernatant to Tube 1 [1]. Add 100 microliters of prewarmed, serum-free DMEM to each of Tubes 2 through 9 [2]. Begin the serial dilution by transferring 100 microliters from Tube 1 to Tube 2 and mixing thoroughly by pipetting [3-TXT].
 - 4.3.1. Talent pipetting virus supernatant into Tube 1.
 - 4.3.2. Talent adding DMEM into Tubes 2 through 9.
 - 4.3.3. Talent transferring 100 microliters from Tube 1 to Tube 2 and mixing. **TXT**: Repeat the process sequentially through tube 9



Videographer's Note: Take 2 is better shot for 4.3.3

- 4.4. Now, aspirate the culture medium from SLK cells [1]. Add 100 microliters of each virus dilution to the designated wells, using 27 wells in total [2].
 - 4.4.1. Talent aspirating media from the 96-well plate using a multichannel aspirator.
 - 4.4.2. Talent dispensing 100 microliters of each dilution into designated triplicate wells.
- 4.5. Centrifuge the 96-well plate at 1,500 *g* for 1 hour at 30 degrees Celsius to promote viral adsorption [1]. Transfer the plate to a 37 degrees Celsius incubator with 5 percent carbon dioxide and incubate for an additional hour [2].
 - 4.5.1. Talent placing the 96-well plate in a centrifuge set at 1,500 g and 30 degrees Celsius.
 - 4.5.2. Talent transferring the plate from the centrifuge to a 37 degrees Celsius carbon dioxide incubator.
- 4.6. Remove the inoculum from each well [1]. Then, gently add 100 microliters of complete DMEM to each well [2]. Incubate the plate for 24 hours at 37 degrees Celsius [3]. Then, analyse the GFP positive cells [4].
 - 4.6.1. Talent aspirating the inoculum from all wells in the 96-well plate.
 - 4.6.2. Talent adding 100 microliters of complete DMEM to each well using a multichannel pipette.
 - 4.6.3. Talent placing the plate back in the 37 degrees Celsius incubator.
 - 4.6.4. LAB MEDIA: 4.6.4-shot-of-the-GFP-pisitive-cells.tif



Results

5. Results

- 5.1. A two-fold serial dilution of BAC16-derived virus stock showed a progressive decrease in the number of GFP-positive SLK cells, confirming a measurable and dilution-sensitive infectious titer [1]. The calculated infectious titer of the BAC16-derived virus stock was approximately 1.3 million infectious units per milliliter [2].
 - 5.1.1. LAB MEDIA: Figure 2A. *Video editor: Highlight the downward trend in the bar graph, moving left to right across the dilution factors.*
 - 5.1.2. LAB MEDIA: Figure 2B.
- 5.2. GFP expression in HCT116 cells increased over time following de novo infection, with the most intense signal observed at 60 hours post-infection [1], indicating robust viral replication at that timepoint [2].
 - 5.2.1. LAB MEDIA: Figure 2C. Video editor: Highlight the three green micrographs under the 36 h, 60 h, and 96 h timepoints, emphasizing the brightest image at 60 h.
 - 5.2.2. LAB MEDIA: Figure 2D. Video editor: Highlight the corresponding three green-stained SLK cell images in the right column labeled 36 h, 60 h, and 96 h, emphasizing the most densely green image at 60 h.
- 5.3. Flow cytometry analysis confirmed a peak in GFP–positive SLK cells at 60 hours post-infection, reaching 74.4% positivity [1], followed by a decrease to 62.7% at 96 hours [2].
 - 5.3.1. LAB MEDIA: Figure 2E. Video editor: Highlight the orange peak labeled "60 h" with the 74.4% annotation.
 - 5.3.2. LAB MEDIA: Figure 2E. Video editor: Highlight the green peak labeled "96 h" with the 62.7% annotation.
- 5.4. Quantitative PCR of viral DNA in supernatants showed a sharp increase in viral genome copies, peaking at 60 hours post-infection and declining thereafter [1].
 - 5.4.1. LAB MEDIA: Figure 2F.



Pronunciation Guide:

1. Kaposi's sarcoma-associated herpesvirus

- Pronunciation link: Cambridge Dictionary provides the pronunciation for Kaposi's sarcoma; the rest (herpesvirus) follows standard pronunciation patterns YouTube+11Cambridge Dictionary+11Wikipedia+11
- o IPA (American): /kəˌpoʊ.siːz sɑːrˈkoʊ.mə-əˈsoʊ.siˌeɪ.tɪd ˈhɜr.pizˌvɜr.əs/
- Phonetic Spelling: kuh-POH-seez sar-KOH-muh-uh-soh-see-ay-tud HER-peez-vur-us

2. KSHV (Kaposi's sarcoma-associated herpesvirus)

- Pronunciation link: Defined by its abbreviation; "K-S-H-V" pronounced as individual letters, matching the full term <u>Cambridge Dictionary+1How To</u> <u>Pronounce+3Wikipedia+3Wikipedia+3</u>
- IPA: /keɪ-εs-eɪtʃ-viː/
- o **Phonetic Spelling:** kay-ess-aych-vee

3. BAC16 (BAC Sixteen derived KSHV)

- Pronunciation link: Cambridge Dictionary for "BAC" covers the abbreviation Wikipedia+9Cambridge Dictionary+9YouTube+9How To PronouncePronounce Names+6DigiKey+6L-com+6Cambridge Dictionary
- o IPA: /biː.eɪ.siː 'sɪkˌstiːn/
- Phonetic Spelling: bee-ay-see SIX-teen

4. Dox-inducible

- Pronunciation link: While not listed in dictionaries, derived from "doxycycline-inducible"—standard pronunciation rules apply
- o IPA: /dpk.sɪnˈdju.sə.bəl/ (American: /dak.sɪnˈdju.sə.bəl/)
- Phonetic Spelling: dok-sin-DYOO-suh-buhl

5. Ultracentrifuge (Ultracentrifugation)

- Pronunciation link: The base word ultracentrifuge is pronounced per standard patterns
- IPA: /ˌʌl.trəˌsɛn.trəˈfjuːdʒ/
- o Phonetic Spelling: uhl-truh-sen-truh-FYOODGE

6. iSLK (cell line name)

- o **Pronunciation link:** Treated as separate letters in common lab usage
- IPA: /aι-εs-εl-keι/
- Phonetic Spelling: eye-ess-el-kay

7. HCT116 (cell line name)

- Pronunciation link: Standard scientific naming; pronounced as letters and numbers
- o IPA: /eɪt∫-si-ti-wʌn-wʌn-sɪks/
- o **Phonetic Spelling:** H-C-T-one-one-six

8. Multiplicity of Infection (MOI)

- o **Pronunciation link:** Standard scientific phrase
- IPA: / mʌl.təˈplɪs.ɪ.ti əv ɪnˈfɛk.ʃən/
- o **Phonetic Spelling:** mul-tuh-PLIS-ih-tee of in-FEK-shun



9. Phase-contrast microscopy

- o **Pronunciation link:** Standard compound term in microscopy
- o **IPA:** /feɪz-ˈkɑn.træst maɪˈkrɑs.kə.pi/
- o **Phonetic Spelling:** faz-KON-trast my-KRAS-kuh-pee

10. Serial two-fold dilution

- o **Pronunciation link:** Common lab phrase
- o **IPA:** /ˈsɪə.ri.əl tuː-foʊld daɪˈluː.ʃən/
- o **Phonetic Spelling:** SEER-ee-uhl too-fold dy-LOO-shun