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## Use of an in vitro scratch assay to demonstrate the effects of arsenic on skin cell migration --Manuscript Draft--

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To Whom It May Concern:

Enclosed, please find our manuscript entitled "**Use of an *in vitro* scratch assay to demonstrate the effects of arsenic on skin cell migration**".

This enclosed manuscript is truthful and original work and represents a useful in-vitro assay that will be of interest to your readership. The submitting author has the written consent from all authors to submit the manuscript and all authors accept complete responsibility for the contents of the manuscript.

The manuscript has not been previously published, and is not currently under consideration elsewhere. The work reported will not be submitted for publication elsewhere until a final decision has been made as to its acceptability by **JoVE**.

Please address all future correspondence regarding this manuscript to myself. Thank you for your help and consideration.

Yours truly,

A handwritten signature in cursive script, appearing to read 'R. Kellar'.

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**TITLE:**

*In Vitro* Scratch Assay to Demonstrate Effects of Arsenic on Skin Cell Migration

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**KEYWORDS:**

Scratch assay, cellular migration, wound healing, arsenic, environmental contaminants, human dermal fibroblasts

**SUMMARY:**

This study focuses on an *in vitro* model of wound healing (scratch assay) as a mechanism for determining how environmental contaminants such as arsenic influence cellular migration. The results demonstrate that this *in vitro* assay provides rapid and early indications of changes to cellular migration prior to *in vivo* experimentation.

**ABSTRACT:**

Understanding the physiologic mechanisms of wound healing has been the focus of ongoing research for many years. This research directly translates into changes in clinical standards used for treating wounds and decreasing morbidity and mortality for patients. Wound healing is a complex process that requires strategic cell and tissue interaction and function. One of the many critically important functions of wound healing is individual and collective cellular migration. Upon injury, various cells from the blood, surrounding connective, and epithelial tissues rapidly migrate to the wound site by way of chemical and/or physical stimuli. This migration response can largely dictate the outcomes and success of a healing wound. Understanding this specific cellular function is important for translational medicine that can lead to improved wound healing outcomes. Here, we describe a protocol used to better understand cellular migration as it pertains to wound healing, and how changes to the cellular environment can significantly alter

this process. In this example study, dermal fibroblasts were grown in media supplemented with fetal bovine serum (FBS) as monolayer cultures in tissue culture flasks. Cells were aseptically transferred into tissue culture treated 12-well plates and grown to 100% confluence. Upon reaching confluence, the cells in the monolayer were vertically scratched using a p200 pipet tip. Arsenic diluted in culture media supplemented with FBS was added to individual wells at environmentally relevant doses ranging 0.1–10  $\mu$ M. Images were captured every 4 hours (h) over a 24 h period using an inverted light microscope to observe cellular migration (wound closure). Images were individually analyzed using image analysis software, and percent wound closure was calculated. Results demonstrate that arsenic slows down wound healing. This technique provides a rapid and inexpensive first screen for evaluation of the effects of contaminants on wound healing.

## INTRODUCTION:

Wound healing consists of a complex series of overlapping steps that involve various cell types, signaling molecules, and extracellular matrix components<sup>1</sup>. Together, these components work in concert to achieve wound resolution or repair, which ultimately leads to the formation of a protective fibrotic scar depending on the degree of trauma<sup>1</sup>. To capture the individual processes and functions of wound healing in one experiment is difficult; individual steps within the wound healing process need to be identified and tested separately. Among the many steps of wound healing, the process of cellular migration has been considered critical when evaluating healing rates<sup>2</sup>. Cellular migration can be observed in all phases of wound healing, and thus necessitates further understanding of how alterations to cell migration can impact wound closure. For example, cellular migration is seen in both early and late phase inflammation, with blood coagulation and platelet aggregation at the wound site<sup>3</sup>. These events prevent excessive blood loss by forming a temporary blockage to fill wounded areas devoid of tissue<sup>3</sup>. Platelets in circulation will synthesize and secrete vasoactive mediators along with chemotactic factors, which together signal for inflammatory leukocyte migration (white blood cells) to the wounded tissue for repair<sup>4</sup>. Re-epithelialization occurs within hours of tissue injury and involves covering the wound site through activity and migration of epithelial keratinocytes to prevent further contamination or invasion of foreign particles to the wound site<sup>2</sup>. These examples capture only a few of the many cell migration functions associated with wound healing.

The scratch assay has been described and used to better understand cell migration and its many roles in wound healing<sup>5,6</sup>. This assay is considered simplistic, provides high-throughput, and enables the investigator to collect representative cell migration data. Migration assays have successfully demonstrated that certain compounds may accelerate or slow down the rate of cellular migration<sup>6</sup>. Furthermore, these assay results provide translational physiologic information that can be used to formulate target treatments, dosing concentrations, and genes of interest prior to *in vivo* experimentation. The assay requires basic cell culture techniques and supplies, a cell line of choice, and imaging technology to capture movement over time or movement in response to treatments.

The assay can be readily manipulated or tailored to suit the needs of the investigator. However, there are four basic questions to consider prior to experimentation. What general or specific



question(s) is/are the investigator(s) trying to answer? This holds true for most hypotheses driven research; however, it is critical to this assay because it will determine many of the parameters needed to accurately and completely answer the question(s). What cell line or cell lines (if multiple) should be used to represent the physiologic system being studied? For example, in a cutaneous wound healing study, a dermal fibroblast<sup>5,6,7</sup>, stem cell<sup>8</sup>, or epidermal keratinocyte<sup>9</sup> might be considered to represent cell populations that are normally found in abundance in skin tissue<sup>10</sup>. Alternatively, neutrophils, macrophages or other inflammatory cells could be evaluated in this system to represent cell migration of other components of wound healing within skin tissue<sup>10</sup>. Additionally, identifying the specific cell line being evaluated will help to identify which reagents are optimal to use to promote cell metabolic activity. How will cell migration be tracked/imaged? There are a number of techniques used to observe cell migration within a plate or flask. For example, dyeing or fluorescently tagging cells and using fluorescence or confocal imaging to track cells provides a strong contrast, which allows the investigator to clearly define cellular vs. non-cellular locations and cellular movement<sup>11</sup>. Another common technique, described in this study, is the use of inverted light microscopy with phase contrast<sup>6,7</sup>. This alternative to cell dyes and fluorescence allows the user to live track cells without having to terminally fix cells prior to imaging and also allows for capturing images of the same individual wells containing wounded monolayers of cells across time points. What outcome measures will be used for analysis? Using the images gathered throughout the study, data can be generated in which changes to cellular migration rates can be understood. In our lab, microscopic images captured on the inverted light microscope are uploaded to image analysis software and the percent wound closure and summed area under the curve (AUC) are calculated.

We will highlight the specific use of this assay to elucidate changes to dermal fibroblast migration in the presence of a known environmental contaminant, arsenic. Arsenic is a naturally occurring metalloid found within the earth's crust. Various locations have been found with elevated levels of arsenic, which poses a risk to individuals who live near these locations. As a result, food and water sources have been shown to have increased levels of arsenic contamination, which increases the likelihood for individuals to become exposed to arsenic. Environmental arsenic exposure has been shown to have long-term health consequences in human populations above or even below the current United States Environmental Protection Agency (USEPA) Max Contaminant Level (MCL) of 10 ppb (10 µg/L)<sup>12,13</sup>. Although the USEPA has set this MCL and deemed it safe for drinking limits, arsenic concentrations that exceed this limit are not uncommon in water sources worldwide. In the Southwestern United States, numerous groundwater, well water, and springs have been documented to contain concerning levels of arsenic<sup>14</sup>. For example, in Verde Valley, AZ, Montezuma Well contains 210 µg/L of arsenic<sup>15</sup>. Groundwater around the Verde River, AZ contains 16 µg/L of arsenic on average, with peak values reaching 1.3 mg/L<sup>15,16</sup>. Notably, arsenic concentrations in many wells in the Verde River water shed exceed 50 µg/L<sup>15,16</sup>. With this knowledge, environmentally relevant arsenic concentrations were selected that range from below the MCL at 0.1 µM (7.5 ppb), to above the MCL at 10 µM (750 ppb) in the current study<sup>14-20</sup>.

The information gathered from these experiments will be used as an early indicator of potential changes to cellular migration rates in an *in vivo* wound contaminated with arsenic. Afterward,

signaling molecules can be selected based on their role in facilitating wound healing and cellular migration, and future quantitative polymerase chain reaction (qPCR) based gene expression experiments may be set up upon completion of the current studies. If migration rates in this assay change in the presence of arsenic, specific gene targets involved in cellular migration may be the target for deleterious effects of arsenic, and thus may be the mechanism of disruption.

## **PROTOCOL:**

### **1) Preparation of a Biosafety II Cabinet (BSC)**

NOTE: The methods described in this protocol should be conducted with personal protective equipment, while using good laboratory practices to achieve aseptic technique.

1.1) Lift the BSC protective glass to operating height and turn on laminar flow fans.

1.1.1) Use 70% isopropanol aqueous solution to perform a sanitation wipe of the working area. Wipe BSC from the back wall and side walls and work down to the base plate.

1.1.2) Wipe all materials that will be used in the assay with 70% alcohol (isopropanol or EtOH) and place them inside the BSC.

### **2) Human Dermal Fibroblasts Seeding in a T75 Flask**

2.1) Obtain vials with the desired cryopreserved cell line (neonatal human dermal fibroblasts [hDFn], at cell concentration of 500,000 cells/vial, and at passage p3–p5).

2.1.1) Place a vial with cells in a 1.0-cm-deep water bath at 37 °C to allow thawing. Thaw vial until approximately 70% of cell solution is liquid. Wipe down vial with 70% alcohol and place vial in vial rack inside BSC.

NOTE: Ensure water does not come in contact with threads or cap of vial to prevent potential contamination during cell seeding. Prevent complete thawing of cells in water bath as failure to do so may cause unintentional cell death.

2.2) Draw out 10 mL of media supplemented with 10% Fetal Bovine Serum (FBS). Use an automatic pipettor and 10 mL pipette tip to aspirate into T75 tissue culture flask. Allow media to fully saturate the base of the flask by gently rocking the flask back and forth.

2.2.1) Open vial with cell stock and aspirate solution from the vial, being mindful of pipetting speed, as cell membranes will be vulnerable from cryopreservation.

2.2.2) Transfer cells to a T75 tissue flask with media. Eject thawed cell into tissue flask, again being mindful of pipetting speed.

NOTE: Seeding density of cells in tissue flask is approximated to be 6,600 cells/cm<sup>2</sup>. Seeding density can be altered to accommodate the user and desired timing of experimentation.

2.2.3) Measure out 1 mL of media from T75 tissue flask and transfer volume back into the vial. Transfer volume from the vial back into the T75 flask.

NOTE: Performing this step helps maximize cell yield from the vial.

2.2.4) Tighten the tissue culture flask cap and gently rock to uniformly disperse cells in suspension across the entire surface. Check for even distribution of cells using an inverted microscope.

2.3) Label flask with cell type, date, initials, lineage, and purpose (*e.g.*, hDFn, 07/11/2018, NDC, Passage 4, Arsenic Scratch Assay). Place tissue culture flask in a humidified incubator at 37 °C and 5.0% CO<sub>2</sub> for 72 h.

NOTE: Growth medium should be changed 12-24 h after initial seeding to remove trace amounts of dimethyl sulfoxide (DMSO) cryopreservative, and then changed every 48 h after that to ensure cells are properly nourished. The incubation and growth period will depend on the calculated number of cells needed for the scratch assay. The doubling time for hDFn cells is typically 16-24 h under normal conditions. However, doubling rate may change due to elevation, pH, temperature, humidity, medium type, *etc.*, and must be accounted for when planning experiments.

### 3) Cell Counting and Subculture into 12-Well Tissue Culture Plate

3.1) Retrieve the T75 tissue culture flask from incubator. Observe adhered cells using an inverted microscope at 10X objective magnification (100X total magnification) and determine the approximate cell confluence.

3.1.1) Scan the majority of the tissue culture flask when observing confluence to ensure the most representative percentage is approximated. Evaluate individual cell morphology for any abnormalities, or for bacterial and/or fungal contamination.

3.2) Assemble the automatic pipettor with a 10 mL pipette tip. Use pipettor to aspirate the full volume of media from the T75 and transfer solution to a waste container. Prevent the pipet tip from contacting the cell adhering surface. Discard pipette tip in a biohazard bin.

3.2.1) Assemble the automatic pipettor with a 5 mL pipette tip. Aspirate 5 mL of the balanced salt solution, dispense in the flask, and gently rock the flask to rinse excess media, dead cells, and waste product.

3.2.2) Draw the volume from the T75 and dispense into the waste container. Discard the pipette tip into the biohazard bin.

221 3.2.3) Assemble the automatic pipettor with a 5 mL pipette adapter. Aspirate 2 mL of trypsin  
222 from stock, dispense in the flask, and gently rock the flask to disperse the enzyme over the  
223 adhered cells. Ensure complete saturation of the adherent surface. Place the tissue flask in the  
224 incubator for 2-3 minutes (min).

225  
226 NOTE: The maximum enzyme-substrate reaction should not exceed 10 min.

227  
228 3.2.4) Observe the tissue flask under an inverted microscope at a 10X objective lens magnification  
229 (100X total magnification). Apply gentle vibrations to facilitate cell detachment.

230  
231 3.2.5) Wipe the T75 flask with 70% alcohol and place inside the BSC.

232  
233 3.2.6) Assemble the automatic pipettor with a 5 mL pipette tip. Aspirate 5 mL of media from the  
234 stock and add to the T75 tissue flask to stop the enzyme-substrate reaction. The media:trypsin  
235 ratio should be 3:1. Pipet up and down the base of the flask 2-3 times to rinse and ensure that  
236 the reaction has stopped. Aspirate the complete volume of fluid from the flask and transfer into  
237 a sterile 15 mL conical tube.

238  
239 3.2.7) Fill a second 15 mL conical tube with an identical volume of media.

240  
241 3.2.8) Place both 15 mL conical tubes in an antipodal position to one another. Apply 200 g of  
242 centrifugal force for 5 min at 25° C by setting equipment's running parameters.

243  
244 3.2.9) Wipe the 15 mL conical tube with pelleted cells with 70% alcohol and place inside the BSC.

245  
246 3.2.10) Use the automatic pipettor with a 5 mL attachment to pipet off media, leaving behind  
247 approximately 250  $\mu$ L of fluid and cell pellet. Pay attention to the location of the pipet tip, as the  
248 cell pellet should remain undisturbed. Dispense the volume collected into a waste container and  
249 discard the tip in a biohazard bin.

250  
251 3.2.11) Use a microcentrifuge tube rack to run the bottom of the conical tube across the vial slots,  
252 creating a swift vibration, to disturb and re-suspend the cell pellet (sometimes called "rack  
253 raking").

254  
255 NOTE: Avoid using a vortex mixer to perform this step. Gravity forces created by vortex mixers  
256 exceed cell g-force thresholds and can cause lysing. When completed correctly, the new cell  
257 solution will appear as a cloudy mixture with no remaining pellet. Diligence performing this step  
258 will oftentimes create a single cell suspension, leading to increased accuracy when counting cells.

259  
260 3.2.12) Add 2 mL of media to the cell stock or resuspension. Pipet the cells gently down the side  
261 of the tube 20 times to break up any clumps. Record the total cell suspension volume and set  
262 aside briefly.

3.3) Pipet 20  $\mu$ L of Trypan Blue stock (0.4%) into an empty well of a 96-well plate using a sterile pipet.

3.3.1) Mix the stock cell solution to generate a uniform cell suspension before transfer. Use a sterile pipet tip to remove 20  $\mu$ L of cell stock (avoiding touching the tube walls). Add to the 20  $\mu$ L of Trypan Blue solution in the 96-well plate.

3.3.2) Pipet gently to mix contents. Pipet 10  $\mu$ L of the mixture between the coverslip and the counting chamber on the hemocytometer.

3.3.3) Observe the cells and the grid under a 10X objective. Observe nine large squares within the field of view. Count cells only in the center and 4 corner squares (total of 5 squares).

NOTE: Look for a uniform distribution of cells across the entire grid to ensure an accurate count.

3.3.4) Tally all cells (transparent and blue) in the 5 identified squares of the grid. Separately, tally only the nonviable (blue) cells in the same 5 squares.

3.3.5) Calculate viable cell count using:

$$x = a - b$$

where  $x$  = viable cell count,  $a$  = total cells,  $b$  = non-viable cells

3.3.6) Calculate the viable cell density using:

$$x = \left(\frac{y}{5}\right) \times \left(\frac{1.0 \times 10^4}{1 \text{ mL}}\right) \times 2$$

where  $x$  = viable cell density and  $y$  = total sum of viable cells.

3.3.7) Calculate total viable cells:

$$x = y \times f$$

where  $x$  = total viable cells,  $y$  = viable cell density,  $f$  = cell suspension volume (mL).

3.3.8) Calculate %Cell viability:

$$x = \left(\frac{y}{f}\right) \times 100$$

where  $x$  = Percent cell viability,  $y$  = viable cells counted on grid,  $f$  = total cells counted on grid.

3.4) Mark a horizontal line across the bottom of the 12-well plate to serve as reference marks during imaging.

3.4.1) Use the viable cell density (calculated in 3.3.7) to calculate the volume of media to dilute cell stock in for seeding a 12-well tissue culture plate.

NOTE: The optimal cell seeding density for hDFn cells is 5,000 cells/cm<sup>2</sup>.

3.4.2) Dilute cell stock to 19,000 cells/mL using media and seed every well with 1 mL of cell stock. Mix cell stock periodically to ensure even cell seeding across all 12 wells.

3.4.3) Label each plate with the cell type, the lineage, the user initials, and the purpose. Place plates in incubator set to 37 °C and 5.0% CO<sub>2</sub>. Allow cells to grow until 100% confluence is achieved for scratch assay.

#### 4) Arsenic (As) Stock Solutions and Calculations

4.1) Dilute sodium arsenite (NaAsO<sub>2</sub>) directly into cell culture media with 10% FBS at working concentrations of 10, 1.0, 0.1, and 0.01 µM As.

4.2) For this, make an initial 1 mM stock As solution by diluting 3.9 mg of sodium arsenite into 30 mL of media. Serially dilute the 1 mM stock for the remaining As solutions.

[Place **Table 1** here]

#### 5) Scratch Assay Technique and Arsenic Contamination

5.1) Obtain a 12-well plate with cells from the incubator. View cells at a 10X objective magnification (100X total magnification). Determine cell confluence within each well.

**NOTE:** Each individual well **must** reach 100% confluence prior to scratching. This critical step ensures consistency within the assay and helps standardize assay protocols across experiments. If any wells have not reached 100% confluence, allow cells to incubate for additional time until **all** wells have reached 100% confluence. Additional growth time in wells that have already reached 100% confluence will not be affected. Confluent cells will typically become growth arrested and remain as a monolayer culture, while allowing sub-confluent wells to reach 100% confluence.

5.2) Assemble an automatic pipettor with a 10 mL pipette tip. Aspirate the growth media out of all wells. Dispose the volume in liquid waste and the pipette tip in the biohazards bin.

5.2.1) Assemble a p200 pipettor with a 1-200 µL sterile tip. In each well, produce a scratch “mock wound” by gliding the tip across cell surface from a 12-o’clock to 6-o’clock location.

**NOTE:** Three factors that will greatly affect consistency are speed, pressure, and tip angle. The scratch should be performed swiftly, with a constant pressure, while being mindful of the tip angle staying perpendicular to the base of the plate. Scratching too slowly will cause crooked lines, excess pressure can permanently score cell adhering surfaces, and a tip angle less than 90° will narrow the scratch width. Any of these common mistakes can result in altered migration rates and patterns.

[Place **Figure 1** here]

5.2.2) Clear away excess debris and cell clumps by rinsing with 1 mL of balanced salt solution per well. Rock the plate side-to-side to facilitate washing. Remove the balanced salt solution from wells and dispose in a waste container. Dispose of the 5 mL pipette tip into the biohazards bin.

5.3) Use a p1000 pipettor with a 1000  $\mu$ L tip to pipet 1000  $\mu$ L from arsenic stocks into wells that have been randomly assigned with a treatment group. Use a new pipet tip for each well to prevent cross contamination. Record the time when treatments are added. Place 12-well plates in an incubator set at 37 °C and 5.0% CO<sub>2</sub>.

## 6) Imaging

6.1) Capture images of each well at 10X objective magnification every 4 h over a 24 h period (0, 4, 8, 12, 16, 20, and 24 h). Reference the line previously drawn on the bottom of the plate to image the same location along the scratch within each well at subsequent time points.

NOTE: It is imperative that the **same locations** are photographed at each time point to allow for accurate image analysis and to obtain representative data.

## 7) Image Analysis

7.1) Upload images captured on the inverted light microscope to a computer and save as a JPEG image with a detailed file name.

7.2) Open ImageJ.

7.3) Upload an image of a stage micrometer with known distances to the software to convert pixels to millimeters (mm) when taking image measurements by dragging and dropping the file into the software window.

NOTE: The micrometer should have lines demarcating a known 1 mm length and the image must be taken at the same magnification used to capture images of cells. For example, in this study images were taken at 100X total magnification, thus the image of the stage micrometer was taken at 100X total magnification.

7.3.1) Click the **Straight**, **Segmented**, or **Freehand Lines** options.

7.3.2) Draw a line of known distance on the micrometer image by using the tick-marks with a length associated to each tick marks. To draw a straight line: click mouse, hold, drag the cursor to desired location, then let go of mouse.

7.3.3) Select **Analyze** then select **Set Scale**.

7.3.4) Set **Known Distance** to the known length of the line drawn in previous step.

394  
395 7.3.5) Set **Unit of Length** to **mm**.  
396  
397 7.3.6) Check the global box.  
398  
399 7.3.7) Select **OK** to exit out of the menu.  
400  
401 7.3.8) Close out of the stage micrometer image.  
402  
403 7.4) Upload one scratch assay image into ImageJ by dragging and dropping image file into the  
404 window.  
405  
406 7.4.1) Draw a straight line across the width of the scratch (from left leading edge to right leading  
407 edge).  
408  
409 7.4.2) Press the letter **M** on keyboard to record the measurement of the straight line drawn. A  
410 small **Results** window will immediately appear after typing the letter **M**. This is where the width  
411 measurement will be.  
412  
413 7.5) Repeat step 7.4 a total of 10 times to obtain 10 measurements along the length of the  
414 scratch.  
415  
416 NOTE: It is important to generate the most representational width values down the entire length  
417 of the scratch. Make sure to space out the 10 width measurements equally along the length of  
418 the scratch from top to bottom.  
419  
420 7.6) Create a scratch assay spreadsheet file to copy and paste the measurements.  
421  
422 7.6.1) Copy and paste the 10 width measurements from the **Results** window into appropriate  
423 column in a spreadsheet file (**Table 2**).  
424  
425 [Place table 2 here]  
426  
427 NOTE: When the measurements are copied from the **Results** window and pasted into the  
428 spreadsheet, there will be extraneous columns of values; these values should be deleted,  
429 preserving only the 10 width measurements.  
430  
431 7.7) Repeat the above step for the remainder of the scratch assay images.  
432  
433 7.8) Calculate the average of the 10 measurements at each time point (0-24 h) for every well.  
434  
435 7.9) Determine percent closure calculations using values from step 7.8.  
436  
437 7.9.1) Create a table at bottom of spreadsheet titled “width measurement averages” (**Table 3**).



[Place table 3 here]

7.9.2) Copy and paste the **Average of 10 Measurements** row calculated in step 7.8 into the “width measurement averages” table.

7.9.3) Create another table next to the **Width Measurement Averages** table. Title this table **Percent Wound Closure (Table 4)**.

[Place table 4 here]

NOTE: The value in the 0 h column for each well should be 0 because the scratch is 0% closed at the initial 0 h photo for all wells.

7.9.4) Navigate to the next column over (4 h).

7.9.5) Calculate the percent closure for 4, 8, 12, 16, 20, and 24 h time points:

$$x = \left( \frac{i - f}{i} \right) \times 100$$

where,  $x$  = percent closure,  $i$  = initial scratch width (0 h width),  $f$  = final scratch width (4, 8, 12, 16, 20, or 24 h width).

7.10) Create another table next to the **Percent Wound Closure** table. Title this table **AUC (Table 5)**.

[Place table 5 here]

7.10.1) Calculate the 0-4 h area under the curve (AUC) value in the 0-4 h column for each well:

$$x = \left( \frac{a + b}{2} \right) \times h$$

where  $x$  = 0-4 h AUC value,  $a$  = % closure value for 0 h,  $b$  = % closure value for 4 h,  $h$  = the time between the two measures (4 h in this study).

7.10.2) Calculate the 4-8 h area under the curve (AUC) value in the 4-8 h column for each well:

$$x = \left( \frac{a + b}{2} \right) \times h$$

Where  $x$  = 4-8 h AUC value,  $a$  = % closure value for 4 h,  $b$  = % closure value for 8 h,  $h$  = the time between the two measures (4 h in this study).

NOTE: One AUC value should be calculated for each 4 h time interval over the 24 h period for each well.

7.10.3) Sum all of the AUC values from 0-24 h (calculated in steps 7.10.1-7.10.2) to get a summed AUC value for each well. Ensure these values are properly saved for statistical analysis.

## 8) Statistical Analysis

8.1) Determine the statistical analysis method that best fits both the experimental design and summed AUC values obtained during the assay.

### REPRESENTATIVE RESULTS:

Sample sizes ( $n$ ) for all treatment groups was  $n = 28$ . Cells were successfully cultured following aseptic techniques and lab protocols (**Figure 1**). Uniform scratches were observed across all treatment groups with the mean scratch width measuring  $0.8 \text{ mm} \pm 0.4 \text{ mm}$  (**Figure 2**). Control groups had an average summed area under the curve value of 1,355.83;  $0.01 \text{ } \mu\text{M}$  As treatment groups had an average summed area under the curve value of 1,366.21;  $0.1 \text{ } \mu\text{M}$  As treatment groups had an average summed area under the curve value of 1,326.24,  $1.0 \text{ } \mu\text{M}$  As treatment group had an average summed area under the curve value of 1,295.10; and finally the  $10 \text{ } \mu\text{M}$  As treatment group had an average summed area under the curve value of 535.12, which was statistically lower compared to all other groups ( $p < 0.05$ ) (**Figure 3**). R-program for statistical analysis was used to run a one-way ANOVA with a Tukey *post hoc* adjustment to determine statistical differences among treatment groups ( $p < 0.05$ ).

[Place figures 2 and 3 here]

### FIGURE AND TABLE LEGENDS:

**Figure 1. Schematic representation of the scratch assay.** All work is done within an aseptic field to decrease the chance of contamination. Cells are seeded at a desired density in tissue culture flasks and grown in an incubator set to human physiologic conditions (5%  $\text{CO}_2$ ,  $37^\circ\text{C}$ ). Upon reaching a desired confluence, cells are aseptically sub-cultured into 12-well tissue culture plates at a desired seeding density. Cells are grown to 100% confluence in each well of the 12-well plate and scratched using a sterile pipet tip. This “scratch” creates an *in vitro* mock wound (denoted by double headed arrow), in which cells naturally migrate to close the open area. Each well can be uniquely conditioned and cellular migration rates can be observed and quantified in response to different treatment conditions.

**Figure 2. Representative scratch assay images over a 20 h period.** Images A-D represent cellular migration of control cells; images E-H represent cellular migration of cells contaminated with  $1.0 \text{ } \mu\text{M}$  As; images I-L represent cellular migration of cells contaminated with  $10 \text{ } \mu\text{M}$  As. From these images, it is clear that cellular migration is slowed in the presence of arsenic. The control images show a completely “healed” scratch after 20 h, while the other two As-treatment groups were not “healed”. The  $10 \text{ } \mu\text{M}$  As treatment inhibited complete closure altogether after 24 h.

**Figure 3. Arsenic slows cellular migration in the scratch assay.** Images were captured over a 24 h period to observe changes to cellular migration rates in the presence of varying concentrations

of arsenic. Raw images were analyzed using an automated software (e.g., in MATLAB), which initially measured wound width, then calculated percent closure and finally area under the curve (AUC). Error bars represent standard error of the mean. The 10  $\mu$ M As treatment group statistically slowed cellular migration of human dermal fibroblasts neonatal (hDFn), shown by a statistically lower AUC value, compared to all other treatment groups using a one-way ANOVA with a Tukey *post hoc* adjustment ( $p < 0.05$ ). Statistical differences are represented by the letters “A” and “B”, which were obtained through using the Compact Letter Display method available in R-program. Treatments that do not share a common letter are statistically significant from each other ( $p < 0.05$ ).

**Table 1. Arsenic stocks and serial dilutions.** This table explains the calculations used to create the As stock solutions and working stock solutions for treatment groups.

**Table 2. Raw width measurements table format.** This table shows the organizational structure for the raw measurements obtained.

**Table 3. Width measurement averages table format.** This table shows the organizational structure for the width measurement averages calculated using the raw width measurements in **Table 2**.

**Table 4. Percent wound closure table format.** This table shows the organizational structure for the percent wound closure values calculated using the width measurement averages in **Table 3**.

**Table 5. AUC table format.** This table shows the organizational structure for the AUC values calculated using the percent wound closure values in **Table 4**.

## DISCUSSION:

Application of this assay provides a high-throughput and timely evaluation of cellular migration and may directly translate to complex physiological systems<sup>5-7</sup>. For this reason, the proposed bench-top model has been tuned and practiced to evaluate the effects of various therapeutic agents and environmental toxins on wound healing<sup>6</sup>. The concentrations of arsenic used in the current study were deemed environmentally relevant through extensive literature search of known exposure levels, and from retrospective review of various data bases<sup>14-21</sup>. Use of this *in vitro* scratch assay demonstrated that exposure to an arsenic concentration within the high, but environmentally relevant range, delayed wound closure (cellular migration).

This bench top model was also employed to isolate and study an individual cell line (fibroblast) to further understand how fibroblast migration contributes to wound healing outcomes. Furthermore, it is difficult to study the functions of an individual cell line in complex *in vivo* systems with many other cells and tissues present that can interfere with the analysis. Additionally, the assay provides a means of achieving semi-quantitative percent wound closure data that can be used to target specific cellular mechanisms through which compounds may influence wound healing. For example, cells used in the scratch assay can be collected and stored in a desired reagent and used in gene expression (signal mRNA) or protein expression assays<sup>22</sup>.

This information can be useful to the investigator if they seek to understand what signals or proteins may be largely contributing to changes in cellular migration in the presence of varying treatments (*i.e.*, arsenic)<sup>22</sup>.

Human neonatal dermal fibroblasts were selected for this work based on their prominent role in wound healing. However, this scratch assay protocol has been implemented using various cell types and for a range of research purposes. For example, the scratch assay has been adopted into the field of oncology for studying migration inhibition of chemotherapy drugs<sup>23</sup>; for skeletal muscle cell migration, proliferation, and diffusion<sup>24</sup>; to study the effect of plant extracts on cellular migration<sup>25</sup>; and to understand how keratinocyte migration and proliferation contribute to wound healing<sup>9</sup>. Additionally, a previous paper by Pinto *et al.* evaluated the effects of uranium (U) and platelet-rich plasma (PRP) on hDFn migration using the scratch assay<sup>6</sup>. The purpose of this work was to understand the extent to which this assay could test the effect of an agent thought to slow cellular migration (U), as well as an agent thought to speed up cellular migration (PRP). As can be seen in the aforementioned work and research by other investigators, application of the scratch assay has high potential for use in various experimental models<sup>26</sup>. In spite of the detail provided within the methods, variance should be expected between investigators and experiments. For example, cellular migration rates will likely fluctuate based on the cell line in use, in addition to the required micronutrients needed for unique cell types. Many of the important observations to aid in cell yield and viability are listed as **NOTES** throughout this protocol. However, it is strongly recommended that investigators follow manufacturer's recommendations for optimal growth conditions, and to use this method as supplementary instruction for general cell culture work.

Although the scratch assay is extremely versatile for study of cellular activity; unforeseen biases may arise in measuring techniques. For example, when using ImageJ to manually trace cell migration fronts, variation may exist among users, which can limit accuracy and representative data collection. It should also be noted that migration fronts should be read blind to the treatment to minimize user bias. Furthermore, manually identifying migration fronts may result in miscalculation of average distance traveled due to pseudopod formation by cells and their ability to reach out to form focal adhesions, which can be visually misleading. In like manner, the use of fluorescent "cell tracker" dyes on cell surface or nuclear proteins to detect migration may unintentionally fix cells resulting in inaccuracies when measuring cellular migration over time. It is recommended that appropriate positive and/or negative controls be enrolled within the experiment to completely assess the effect of treatment on the cell lines. Practice of this assay must be completed in recognition of its limitations. This assay does not guarantee that cellular migration results obtained from *in vitro* experimentation will directly translate to *in vivo* outcomes. Wound healing and cellular migration in a complex living system involves a multitude of cell types and molecular signals; each of which has the potential to influence healing responses.

In summary, the scratch assay has been found to provide high-throughput screening of cellular migration in response to changing environments<sup>6,25</sup>. We specifically utilized this protocol to successfully detect changes in cellular migration as a result of arsenic exposure at various

environmentally relevant doses. These data have provided justification to use the scratch assay to further study and evaluate the mechanistic pathways of cellular migration in this assay and how these pathways are altered by arsenic exposure.

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#### DISCLOSURES:

The authors have nothing to disclose.

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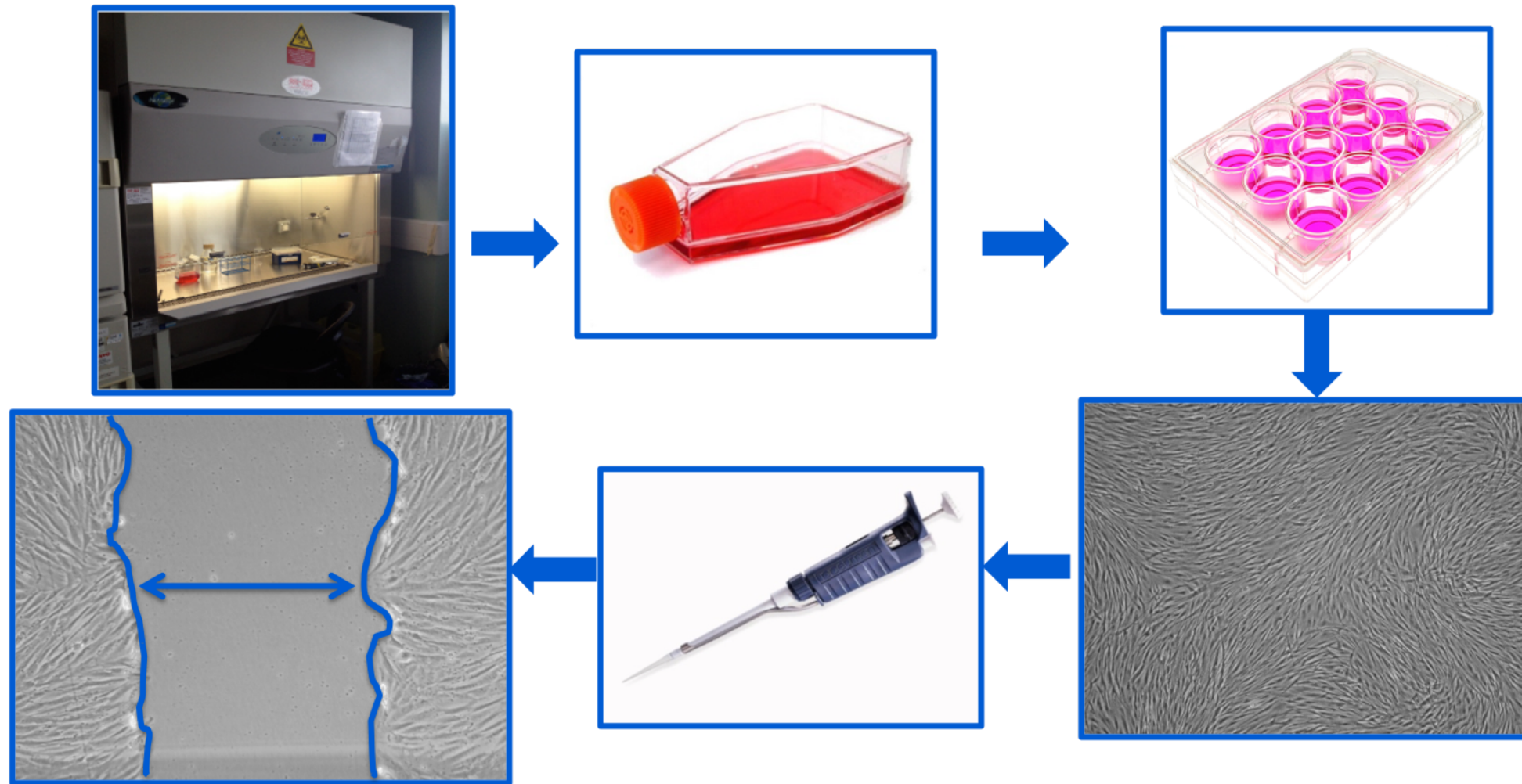




Figure 2

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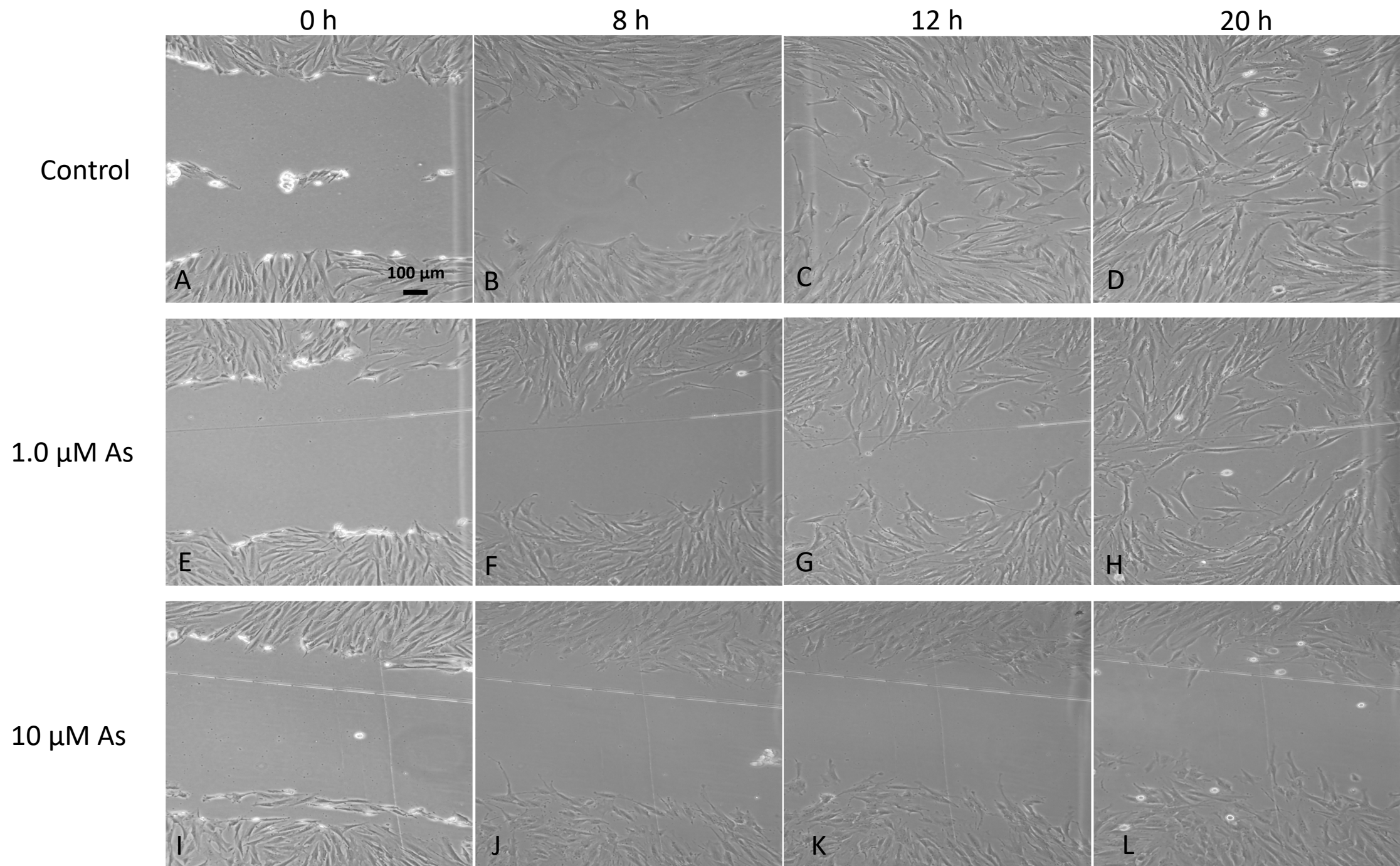
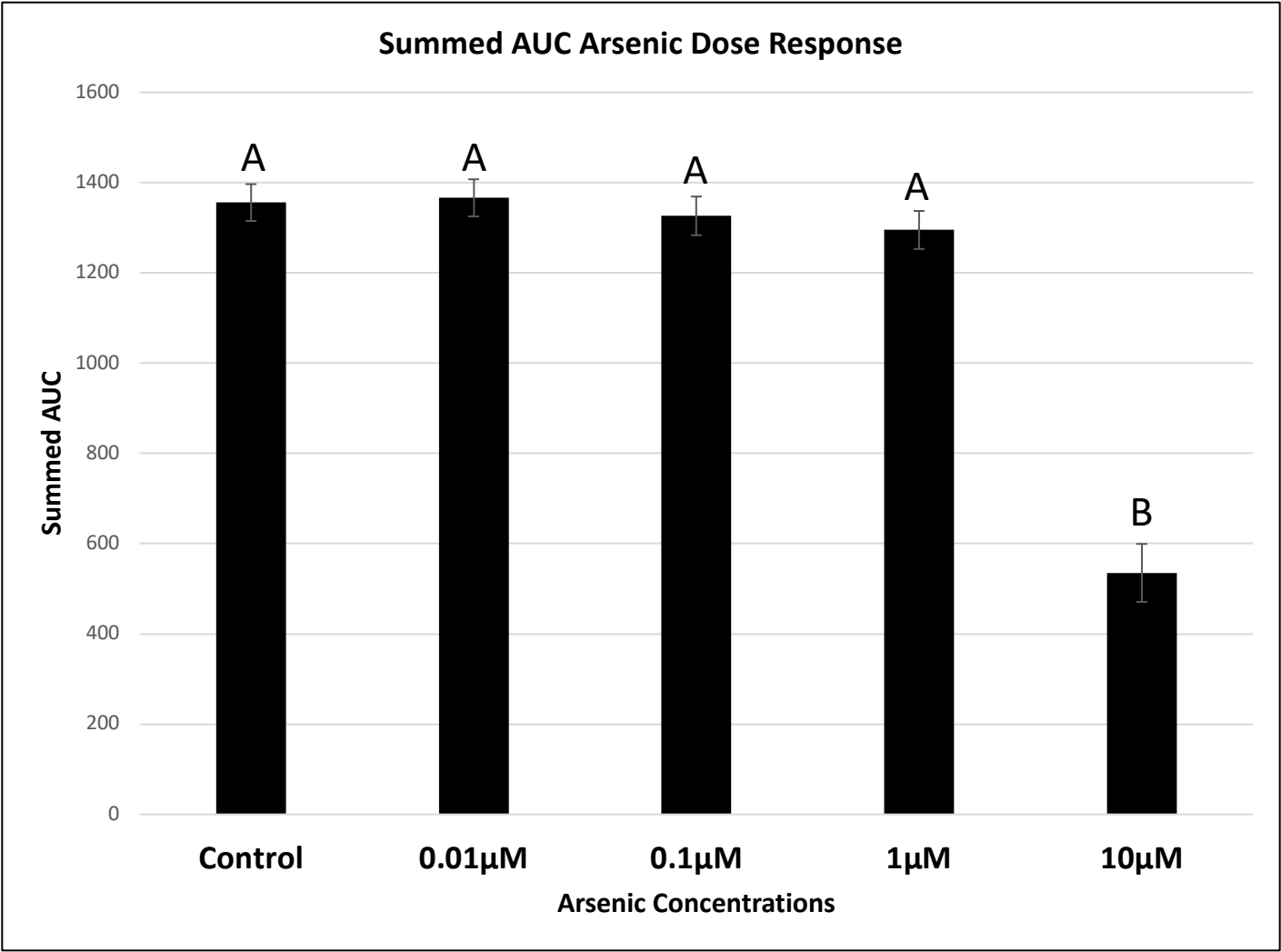




Figure 3



1000 $\mu\text{M}$ As Stock Calculations	Arsenic Working Stock Concentrations	$M_1V_1=M_2V_2$
NaAsO <sub>2</sub> molecular weight: 129.9 g/mol	10 $\mu\text{M}$ As	$(x \text{ mL})(1000 \mu\text{M}) =$ $(30 \text{ mL})(10 \mu\text{M})$ $x =$ 0.3 mL
.03 L x .001 mol/L x 130 g/mol = 3.9 mg of Arsenic	1.0 $\mu\text{M}$ As	$(x \text{ mL})(10 \mu\text{M}) =$ $(30 \text{ mL})(1 \mu\text{M})$ $x$ = 3 mL
Dilute 3.9 mg of sodium arsenite into 30 mL media	0.1 $\mu\text{M}$ As	$(x \text{ mL})(1 \mu\text{M}) = (30$ $\text{mL})(0.1 \mu\text{M})$ $x = 3$ mL
	0.01 $\mu\text{M}$ As	$(x \text{ mL})(0.1 \mu\text{M}) =$ $(30 \text{ mL})(0.01 \mu\text{M})$ $x$ = 3 mL

Volume of Previous Arsenic Solution	Volume of Media	Total Volume of Arsenic Working Stock
0.3 mL = 300 $\mu$ L of 1mM As solution	29.7 mL media	30
3 mL of 10 $\mu$ M As solution	27 mL of media	30
3 mL of 1 $\mu$ M As	27 mL of media	30
3 mL of 0.1 $\mu$ M As	27 mL of media	30

10 random width measurements	0 h	4 h	8 h	12 h	16 h	20 h
1						
2						
3						
4						
5						
6						
7						
8						
9						
10						
Average of 10 measurements						

[illegible]

Well width averages	0 h	4 h	8 h	12 h	16 h	20 h
1a						
2a						
3a						
4a						
1b						
2b						
3b						
4b						
1c						
2c						
3c						
4c						

[illegible]

Well	0 h	4 h	8 h	12 h	16 h	20 h
1a	0					
2a	0					
3a	0					
4a	0					
1b	0					
2b	0					
3b	0					
4b	0					
1c	0					
2c	0					
3c	0					
4c	0					



[illegible]

Well	0-4 h	4-8 h	8-12 h	12-16 h	16-20 h	20-24 h
1a						
2a						
3a						
4a						
1b						
2b						
3b						
4b						
1c						
2c						
3c						
4c						

[illegible]

Name of Material/ Equipment	Company	Catalog Number	Comments/Description
Class II A/B3 Biological Safety Cabinet	Forma Scientific	15201-816	
Nitrile Gloves	Kimberly-Clark	55082	
Water Bath	Precision	Model 182	
Inverted Microscope	Leica	Q080318	
CPR Centrifuge	Beckman	349702	
		I093	
Analytical Scale	Ohaus	1125062111P	
Weighting paper	VWR	H351125781212A	
Biohazard bags	Fisherbra n	01-830A	
Water Jacketed CO <sub>2</sub> Incubator	Thermo Scientific	309739-31053	
Hanks Balanced Salt Solution	Gibco	14175-095	
Dulbecco's Modified Eagle Medium	Gibco	10566-016	
Fetal Bovine Serum	Gibco	26140-079	
	Drummon		
Automatic Pipettor		140685	
p200 Pipette	Gilson		
p20 Pipette	Gilson		
	Fisher		
Denominator	Scientific	D59707	
Trypan Blue Stain	Gibco	15250-061	
T75 tissue flask	Corning	430725U	
	Fisherbra		
15 mL Conical Tube	nd	05-539-5	
50 mL Conical Tube	VWR	21008-178	

5 mL Pipette	Fisherbrand	13-678-11D
10 mL Pipette	Fisherbrand	13-678-11E
Tissue Marker	Securline	92167
Wipes	Kintech	34155
70% Ethanol	Fisher	63-67-0
Non-Filter Pipet Tips	Fisherbrand	16160210
Bleach	Great Value	220-04
96-well Cryovial Rack	Falcon	353915
Hemocytometer	Nalgene	5030-0505
TrypLE Express	Gizmo Supply	B00SCOGY56
Conical Tube Rack	Gibco	12605-028
12-well plate	n/a	n/a
Waste Beaker	Corning	3598
1 mL Pipette	n/a	n/a
Straight Edge Ruler	Fisherbrand	13-678-11B
Human neonatal dermal fibroblasts (hDFn)	lot: 1734, 2902	106-05n
Rstudio Statistical Software		ver: 3.5.1
Image J	NIH	ver: 1.8.0_112



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Date:

7-24-18

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NORTHERN ARIZONA  
UNIVERSITY  
*College of Engineering, Forestry & Natural Sciences*  
**Center for Bioengineering Innovation**

September 18, 2018

*Journal of Visualized Experiments*  
1 Alewife Center, Suite 200  
Cambridge, MA 02140

To Whom It May Concern:

Enclosed please find our point-by-point response (in red ink) to the Reviewers' comments to our manuscript entitled "**Use of an *in vitro* scratch assay to demonstrate the effects of arsenic on skin cell migration**". We have responded to the reviewers' requests and we believe these changes have improved the manuscript. Please find the changes highlighted as "track changes" in the newest uploaded document.

This enclosed manuscript is truthful and original work and represents a useful in-vitro assay that will be of interest to your readership. The submitting author has the written consent from all authors to submit the manuscript and all authors accept complete responsibility for the contents of the manuscript.

The manuscript has not been previously published, and is not currently under consideration elsewhere. The work reported will not be submitted for publication elsewhere until a final decision has been made as to its acceptability by **JoVE**.

Please address all future correspondence regarding this manuscript to myself. Thank you for your help and consideration.

Yours truly,



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## **Editorial Comments:**

- Please take this opportunity to thoroughly proofread the manuscript to ensure that there are no spelling or grammatical errors.

### **• Protocol Language:**

1) Please ensure that all text in the protocol section is written in the imperative voice/tense as if you are telling someone how to do the technique (i.e. “Do this”, “Measure that” etc.) Any text that cannot be written in the imperative tense may be added as a “Note”, however, notes should be used sparingly and actions should be described in the imperative tense wherever possible. Examples NOT in the imperative voice: 7.1, 7.1.2, etc.

-We have made the required changes.

2) Lines 315- 331 should be made into a table for clarity and conciseness of the protocol.

-We have included Table 1 that details these calculations.

• **Protocol Detail:** Please note that your protocol will be used to generate the script for the video, and must contain everything that you would like shown in the video. **Please add more specific details (e.g. button clicks for software actions, numerical values for settings, etc) to your protocol steps.** There should be enough detail in each step to supplement the actions seen in the video so that viewers can easily replicate the protocol.

-We have made numerous changes to the protocol.

• **Protocol Numbering:** Please adjust the numbering of your protocol section to follow JoVE’s instructions for authors, 1. should be followed by 1.1. and then 1.1.1. Some numbering errors are present. **–Corrections have been made.**

• **Protocol Highlight:** Please highlight ~2.5 pages or less of text (which includes headings and spaces) in yellow, to identify which steps should be visualized to tell the most cohesive story of your protocol steps.

1) The highlighting must include all relevant details that are required to perform the step. For example, if step 2.5 is highlighted for filming and the details of how to perform the step are given in steps 2.5.1 and 2.5.2, then the sub-steps where the details are provided must be included in the highlighting.

2) The highlighted steps should form a cohesive narrative, that is, there must be a logical flow from one highlighted step to the next.

3) Please highlight complete sentences (not parts of sentences). Include sub-headings and spaces when calculating the final highlighted length.

4) Notes cannot be filmed and should be excluded from highlighting.

5) Please bear in mind that software steps without a graphical user interface/calculations (e.g 3.3.5-3.3.9) cannot be filmed.

-We have updated the highlighted sections.

- **Results:** Please mention the statistical tests performed and report sample sizes.

-We have made the requested changes. See “representative results” section.

- **Discussion:** JoVE articles are focused on the methods and the protocol, thus the discussion should be similarly focused. Please ensure that the discussion covers the following in detail and in paragraph form (3-6 paragraphs): 1) modifications and troubleshooting, 2) limitations of the technique, 3) significance with respect to existing methods, 4) future applications and 5) critical steps within the protocol.

-We have edited the discussion to include the relevant information mentioned above.

- **Figures:**

1) Fig 2: Please use a different panel labeling scheme. We suggest A-L.

-We have made the suggested change.

- **Figure/Table Legends:**

1) Fig 3: Define error bars and A,B.

-We have made the required changes.

- **Commercial Language:** JoVE is unable to publish manuscripts containing commercial sounding language, including trademark or registered trademark symbols (TM/R) and the mention of company brand names before an instrument or reagent. Examples of commercial sounding language in your manuscript are cryovial, TrypLE Express, MatLab

1) Please use MS Word’s find function (Ctrl+F), to locate and replace all commercial sounding language in your manuscript with generic names that are not company-specific. All commercial products should be sufficiently referenced in the table of materials/reagents. You may use the generic term followed by “(see table of materials)” to draw the readers’ attention to specific commercial names.

-We have made the required changes.

- **Table of Materials:** Please revise the table of the essential supplies, reagents, and equipment. The table should include the name, company, and catalog number of all relevant materials/software in separate columns in an xls/xlsx file. Please include items such as cell lines, software, microscope, etc.

-We have updated the Table of Materials.

- Please define all abbreviations at first use.

- Please use standard abbreviations and symbols for SI Units such as  $\mu\text{L}$ , mL, L, etc., and abbreviations for non-SI units such as h, min, s for time units. Please use a single space between the numerical value and unit.

-We have made the required changes.

- If your figures and tables are original and not published previously or you have already

obtained figure permissions, please ignore this comment. If you are re-using figures from a previous publication, you must obtain explicit permission to re-use the figure from the previous publisher (this can be in the form of a letter from an editor or a link to the editorial policies that allows you to re-publish the figure). Please upload the text of the re-print permission (may be copied and pasted from an email/website) as a Word document to the Editorial Manager site in the "Supplemental files (as requested by JoVE)" section. Please also cite the figure appropriately in the figure legend, i.e. "This figure has been modified from [citation]."

### **Reviewer #1:**

#### Manuscript Summary:

Here the authors present a modification to a very well established protocol to investigate the impact of arsenic contamination on skin cell migration.

#### Major Concerns:

1. The steps in the procedure are clearly written and would result in the desired outcome. The critical steps are highlighted and no important information is missing. Anticipated results are reasonable and useful to the reader if more context is provided. However, due to the simple nature of this protocol, the incremental modification and the amount of papers that have been published previously describing this protocol, it is recommended that this publication not be accepted for publication in the current form. Some modifications that could improve the manuscript include:

1. Discussing the broad application of this method/protocol for the study of many types of environmental contaminants, not just arsenic.

-We have included highlights and references from our previous work with the scratch assay and the influence uranium (environmental contaminant) and platelet-rich plasma (therapeutic) independently have on *in vitro* cellular migration. This ref demonstrates that the scratch assay can be used to study compounds that both speed up- and slow down, cellular migration. This proves its value for broad applications.

2. Providing more background on how people are exposed to these contaminants and in what concentrations to make the assay relevant in the context of healthcare.

-Agreed. We have included an additional paragraph in the "introduction" section that specifically addresses arsenic exposure routes and justification for our selected arsenic concentrations, which come from the literature.

3. Including the procedure for creating and using the automated algorithm to calculate the decrease in open wound area would also advance this protocol beyond what is currently available in the literature.

-We have chosen to remove the use of the automated algorithm because we plan to submit the algorithm work as a separate and unique publication. Additionally, we have updated the "protocol" section, which now describes manual image analysis that utilizes NIH image-j software.

4. Summed area under the curve is used as an output but is not described in detail. Please add more information on how this value is calculated.

-We have addressed this comment and describe, in detail, the mathematical equation for calculating area under the curve using percent wound closure values.

Minor Concerns:

There are a few spelling errors, please review for grammar.

-All errors have been corrected to the best of our knowledge.

## **Reviewer #2:**

Manuscript Summary:

The manuscript by Pinto et al. describes a protocol to evaluate cell migration of human dermal fibroblasts in the presence of arsenic using the in vitro scratch assay. The in vitro scratch assay has been extensively used in particular for screening of compounds which potentially modulate the process of cell migration. Accordingly, several protocols have been published during the last years. Although the paper is well written, the scope of the manuscript is somewhat limited focussing on human dermal fibroblasts and arsenic only. Importantly, there are major issues with the methodology that need to be addressed before this work is publishable.

Major Concerns:

1. The authors state that confluent cells will typically become growth arrested. However, after scratching the monolayer cells may proliferate again. Thus, it is mandatory to use proper controls (such as mitomycin C) to discard proliferation instead of migration. In addition, migration assays may be performed under serum-free conditions.

-Thank you, that is a good point and something to consider when conducting cellular migration assays. However, upon scratching, we allow cellular migration to take place for 24 hours, at which point we terminate the experiment. Experimentation in our lab has determined that human dermal fibroblasts have an approximate doubling time of 24-28 hours at 7,000 feet elevation where our lab in Flagstaff, AZ is located. This would not give the majority of cells enough time to fully double (proliferate) and significantly change the closure rates that we observe within the 24 hour migration period. With this knowledge, we believe that the majority of the “wound closure” observed is from cellular migration alone, with little effect from proliferation.

2. Since dermal fibroblasts are extensively embedded in ECM, the cells may behave quite differently in an artificial monolayer setting. To mimic the situation in vivo more closely, a protocol (incl. results) should be presented where cells are also seeded on plates coated with proper ECM substrates.

-Yes, we agree that the 3-dimensional *in vivo* environment where dermal fibroblasts reside is much different than that of an *in vitro* monolayer culture. We understand that there are many changes/additions that can be made to this well described assay, which may include coating the growth surface with proteins such as collagen, fibronectin, elastin and hyaluronic acid. However, coating may not be applicable to all scratch assays intended to assess migration. This methods

paper is intended to provide a detailed technique to investigators such that it can be readily manipulated and tailored to fit a broad range of projects (i.e. coating). Another piece to consider is that much of any coating added to the wells would likely be removed during the time of scratching, leaving little to no substrate at the scratch site for the cells to adhere to and migrate on; thus not having a significant effect on migration rates. Lastly, we believe that the addition of an ECM coating would be outside the scope of this specific work, and would be a great question to address in future work.

#### Minor Concerns:

1. Appropriate positive controls (besides arsenic) which inhibit cell migration should be added in the protocol and results.

-The purpose of this paper was to describe the scratch assay as a model, therefore describing this method we've chosen to share representative data sets. We agree with the reviewer that during experimental design, positive and negative controls should be selected based on the study design the investigator is pursuing. We have added a statement to this effect in the body of the manuscript "It is recommended that appropriate positive and/or negative controls be enrolled within the experiment to completely assess the effect of treatment on the cell lines."

2. Although the manuscript focusses on arsenic, the authors should add details for the protocol when evaluating other environmental contaminants for the modulation of cell migration.

-We have included highlights and refs from our previous work with the scratch assay and the influence Uranium (environmental contaminant) and platelet-rich plasma (therapeutic) independently have on *in vitro* cellular migration. This ref includes a detailed protocol for both compounds, and gives insight on how this simple assay can be readily manipulated for various projects.

3. The authors may want to include discussion on the required modifications of the protocol when using other cell types (e.g. keratinocytes, endothelial cells).

-We have included references in the introduction that describe, in detail, migration assays using the cell lines mentioned in comment #3. Additionally, we describe important details to consider when designing this assay in paragraph 3 of the introduction.

4. What is known about the mechanism of action of arsenic regarding the inhibition of fibroblast cell migration. This should be added in the manuscript.

-Arsenic was used as "representative data" to primarily provide information on how this assay can help investigators understand cellular migration as it pertains to specific questions/hypotheses, such as environmental contaminants in this manuscript. In this journal, we were encouraged to limit discussion regarding arsenic and its toxic effects and focus on the methods. We have added references in the discussion that discuss mechanism of action regarding the inhibition of fibroblast cell migration. Additionally, please see below the instructions for authors regarding the "Discussion" section. Lastly, we have NIH funding in place to explore the mechanism behind arsenic disruption of wound healing (Southwest Health Equity Research Collaborative - NIH RCMI, 1 U54 MD012388-01). Our specific aims involve; 1) elucidating how arsenic impacts dermal fibroblasts *in vitro*; 2) elucidating how estrogen may

reverse the negative effects of arsenic *in vitro*; and 3) translating the findings from aims 1 and 2 into a full thickness *in vivo* wound healing model (murine).

**“...DISCUSSION:** (3-6 paragraphs)

JoVE is a methods-based journal. Thus, the Discussion of the article should be focused on the protocol and not the representative results.

This section should discuss the following with citations:

- Critical steps in the protocol.
- Modifications and troubleshooting of the method.
- Limitations of the method.
- The significance of the method with respect to existing/alternative methods.
- Future applications or directions of the method.”