

Supplementary Material

## 1. Python Script – Lipid Droplet Analysis

```

import os
import datetime
import numpy as np
from skimage import io, color, exposure, feature
import cv2
from PIL import Image
import multiprocessing

input_folder = "/ImagesFolderPath/Images/"
output_folder = os.path.join(os.path.dirname(input_folder),
"Image_Outputs")
timestamp = datetime.datetime.now().strftime("%Y%m%d%H%M%S")
output_folder_timestamped = os.path.join(output_folder, timestamp)
os.makedirs(output_folder_timestamped, exist_ok=True)

def process_image(image_path):
    img = io.imread(image_path)
    img = exposure.rescale_intensity(img, in_range='image',
out_range=(0, 1))
    img = exposure.rescale_intensity(img, in_range=(0.2, 0.8),
out_range=(0, 1))

    img_gray = color.rgb2gray(img)
    img_contrast = exposure.equalize_adapthist(img_gray,
clip_limit=0.03)

    blobs = feature.blob_log(img_contrast, max_sigma=5,
threshold=0.01)
    black_background = np.zeros_like(img_gray)

    particle_count = 0
    sum_diameter = 0.0

    for blob in blobs:
        y, x, r = blob
        if r > 2:
            cv2.circle(black_background, (int(x), int(y)), int(r),
(1, 1, 1), -1)
            particle_count += 1
            sum_diameter += 2 * r

    output_path = os.path.join(output_folder_timestamped,
os.path.basename(image_path))
    Image.fromarray((black_background *
255).astype(np.uint8)).save(output_path, format="PNG")

    return os.path.basename(image_path), particle_count,
sum_diameter

if __name__ == '__main__':
    num_processes = os.cpu_count()

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with multiprocessing.Pool(processes=num_processes) as pool:
    image_paths = [os.path.join(input_folder, filename) for
filename in os.listdir(input_folder) if
                filename.endswith(".jpg") or
filename.endswith(".png")]

    results = pool imap_unordered(process_image, image_paths)

    for filename, particle_count, diameter_sum in results:
        if particle_count > 0:
            average_diameter = diameter_sum / particle_count
        else:
            average_diameter = 0.0
        print(f"Image: {filename}, Particles: {particle_count}, "
Average Diameter: {average_diameter:.2f} pixels")

    print("Image processing completed and saved in:",
output_folder_timestamped)

```

## 2. Python Script – Cell Count

```

3. import cv2
import numpy as np
import os
import openpyxl

def count_cells(image_path):
    image = cv2.imread(image_path)

    if image is None:
        print(f"Error: Unable to load the image from
'{image_path}'")
        return 0

    gray = cv2.cvtColor(image, cv2.COLOR_BGR2GRAY)
    blurred = cv2.GaussianBlur(gray, (5, 5), 0)
    _, thresh = cv2.threshold(blurred, 0, 255, cv2.THRESH_BINARY
+ cv2.THRESH_OTSU)
    contours, _ = cv2.findContours(thresh, cv2.RETR_EXTERNAL,
cv2.CHAIN_APPROX_SIMPLE)
    marked_image = image.copy()

    cell_count = 0
    for contour in contours:
        area = cv2.contourArea(contour)

        if 100 < area < 25000:
            x, y, w, h = cv2.boundingRect(contour)
            cv2.rectangle(marked_image, (x, y), (x + w, y + h),
(0, 255, 0), 2)
            cell_count += 1

    return cell_count, marked_image

def process_images_in_folder(folder_path):

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results = []
excel_filename = "cell_counts.xlsx"
output_folder = os.path.join(folder_path, "cell_outputs")
os.makedirs(output_folder, exist_ok=True)

wb = openpyxl.Workbook()
ws = wb.active
ws.append(["Image Name", "Cell Count"])

for filename in os.listdir(folder_path):
    if filename.endswith(('.jpg', '.jpeg', '.png', '.bmp')):
        image_path = os.path.join(folder_path, filename)
        cell_count, marked_image = count_cells(image_path)
        results.append((filename, cell_count))
        marked_image_path = os.path.join(output_folder,
f"marked_{filename}")
        cv2.imwrite(marked_image_path, marked_image)
        ws.append([filename, cell_count])

wb.save(os.path.join(output_folder, excel_filename))
return results

if __name__ == "__main__":
    folder_path = "/ImagesFolderPath/Images/"
    if not os.path.exists(folder_path):
        print(f"Error: Folder '{folder_path}' does not exist.")
    else:
        results = process_images_in_folder(folder_path)
        print("Analysis complete. Results saved to 'cell_outputs' folder.")
```