

Q/A:-

- 1. What specific types of cancer cells are you looking to identify and classify?
 - o Answer: Breast cancer cells, lung cancer cells, and melanoma.
- 2. What type of medical images will we be using (e.g., histopathology slides, MRI, CT scans)?
 - Answer: Histopathology slides and CT scans.
- 3. Are there specific imaging modalities preferred for this project?
 - Answer: We prefer histopathology slides due to their detail.
- 4. What is the desired accuracy level for the segmentation algorithm?
 - o Answer: We aim for at least 90% accuracy in segmentation.

- 5. Do you have a dataset available for training the algorithm? If so, what is its size and format?
 - Answer: Yes, we have a dataset of 5,000 annotated histopathology images in TIFF format.
- 6. How diverse is the dataset in terms of different cancer types and stages?
 - Answer: The dataset includes samples from early to late stages across various cancer types.
- 7. Are there any existing annotated datasets you would like us to use?
 - Answer: We would like to use the CAMELYON dataset as a supplementary resource.
- 8. What preprocessing steps do you envision are necessary for the images?
 - o Answer: Normalization, resizing, and augmentation to enhance the dataset.
- 9. Do you have specific software or tools in mind for developing the algorithm?
 - o Answer: We prefer using TensorFlow and Keras for model development.
- 10. What programming languages are preferred for this project?
 - o Answer: Python is the preferred language due to its extensive libraries.