

## Introduction to Generative AI with AWS

### Project Documentation Report

QUESTIONS	ANSWERS:
<p><b>Step 2: Domain Choice</b></p> <p>What domain did you choose to fine-tune the Meta Llama 2 7B model on?</p> <p>Choices:</p> <ol style="list-style-type: none"><li>1. Financial</li><li>2. Healthcare</li><li>3. IT</li></ol>	<p>2. Healthcare</p>
<p><b>Step 3: Model Evaluation Section</b></p> <p>What was the response of the model to your domain-specific input in the <b>model_evaluation.ipynb</b> file?</p>	<p>Certain germline disorders may be associated with &gt; an increased risk of developing breast cancer. Certain mutations in the BRCA1 and BRCA2 genes increase the risk of developing breast cancer. These mutations can be passed down from a parent to a child. Inheriting a mutation in BRCA1 or BRCA2 can increase the</p> <p>=====</p>
<p><b>Step 4: Fine-Tuning Section</b></p> <p>After fine-tuning the model, what was the response of the model to your domain-specific input in the <b>model_finetuning.ipynb</b> file?</p>	<p>Genomic characterization is essential for &gt; [{'generated_text': ' the development of personalized therapies in cancer. Genomic characterization of tumor samples is typically achieved by next-generation sequencing (NGS) of exons, which is expensive and time-consuming. We propose a novel method, called DNase-Seq, to characterize genomic'}]</p> <p>=====</p>