

SCHOOL OF COMPUTER SCIENCE AND ARTIFICIAL INTELLIGENCE			DEPARTMENT OF COMPUTER SCIENCE ENGINEERING	
Program Name: B. Tech		Assignment Type: Lab		Academic Year: 2025-26
Course Coordinator Name		Dr.Vairachilai Shenbagavel		
Instructor(s) Name		Srinivas Komakula		
Course Code	23CA201SE402	Course Title	Explainable AI (P)	
Year/Sem	III/V	Regulation	R24	
Date and Day of Assignment	28-08-2025	Time(s)	09:00AM -05:00PM	
Duration	2 Hours	Applicable to Batch	23CSBTB32	
Assignment Number: 04				
Q. No.	Question			Expected Time to complete
1	Assignment 2 — Breast Cancer (Binary Classification)			
<p><b>Objectives:</b></p> <ul style="list-style-type: none"><li>• Use <b>Permutation Importance</b> for global feature contributions.</li><li>• Apply <b>SHAP</b> to visualize why some patients are predicted malignant/benign.</li><li>• Use <b>LIME</b> to generate local explanations for two patients.</li><li>• Compare alignment and differences between methods.</li></ul> <p><b>Assignment Details:</b></p> <ul style="list-style-type: none"><li>• <b>Goal:</b> Interpret how features influence breast cancer classification across multiple explanation techniques.</li><li>• <b>Data:</b> <code>sklearn.datasets.load_breast_cancer()</code></li><li>• <b>Model:</b> <code>GradientBoostingClassifier</code></li><li>• <b>Steps:</b><ol style="list-style-type: none"><li>1. Train <code>GradientBoostingClassifier</code>.</li><li>2. <b>Permutation Importance:</b> Rank features by prediction impact.</li><li>3. <b>SHAP:</b> Create global summary plot and local force plots for one malignant and one benign case.</li><li>4. <b>LIME:</b> Generate local explanations for the same two cases.</li><li>5. Compare results across methods.</li></ol></li><li>• <b>Deliverables:</b><ul style="list-style-type: none"><li>◦ Permutation Importance plot.</li><li>◦ SHAP summary + 2 force plots.</li><li>◦ LIME explanations for 2 cases.</li><li>◦ Comparative analysis across PI, SHAP, and LIME.</li></ul></li></ul> <p><b>Submission Requirements:</b></p> <ul style="list-style-type: none"><li>• Short methods summary (3–5 lines).</li><li>• Clean, runnable code/notebook.</li><li>• All required plots (PI, SHAP global + local, LIME local).</li><li>• 5–10 bullet insights highlighting consistencies and differences.</li></ul>				