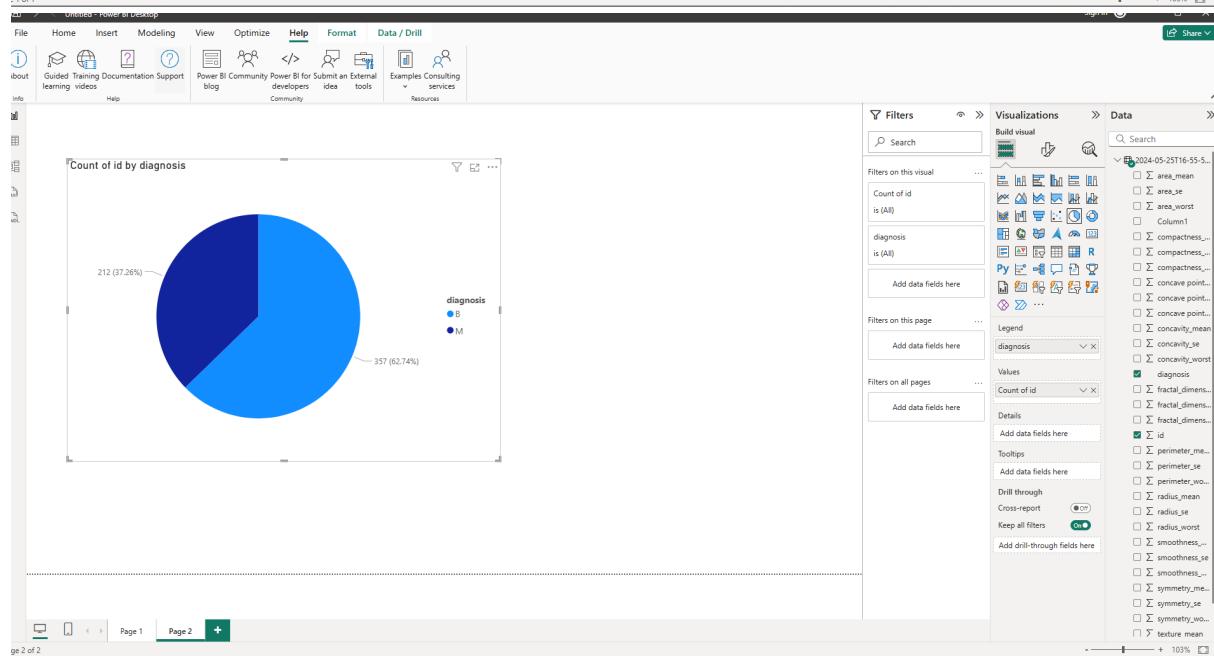
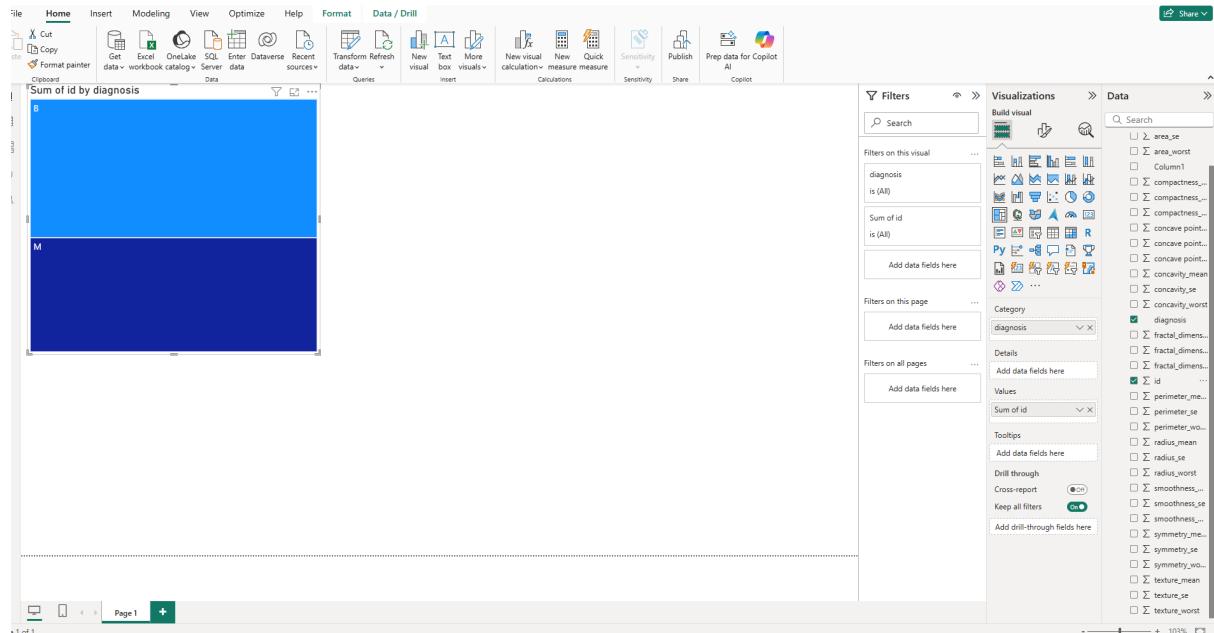
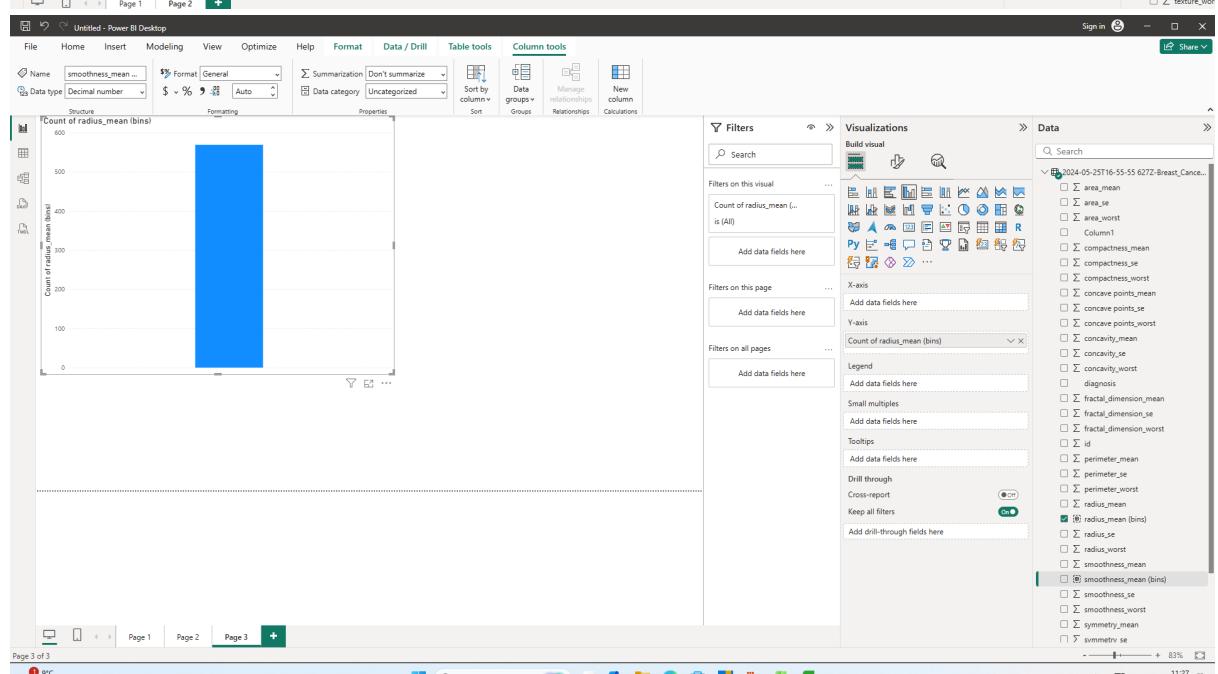
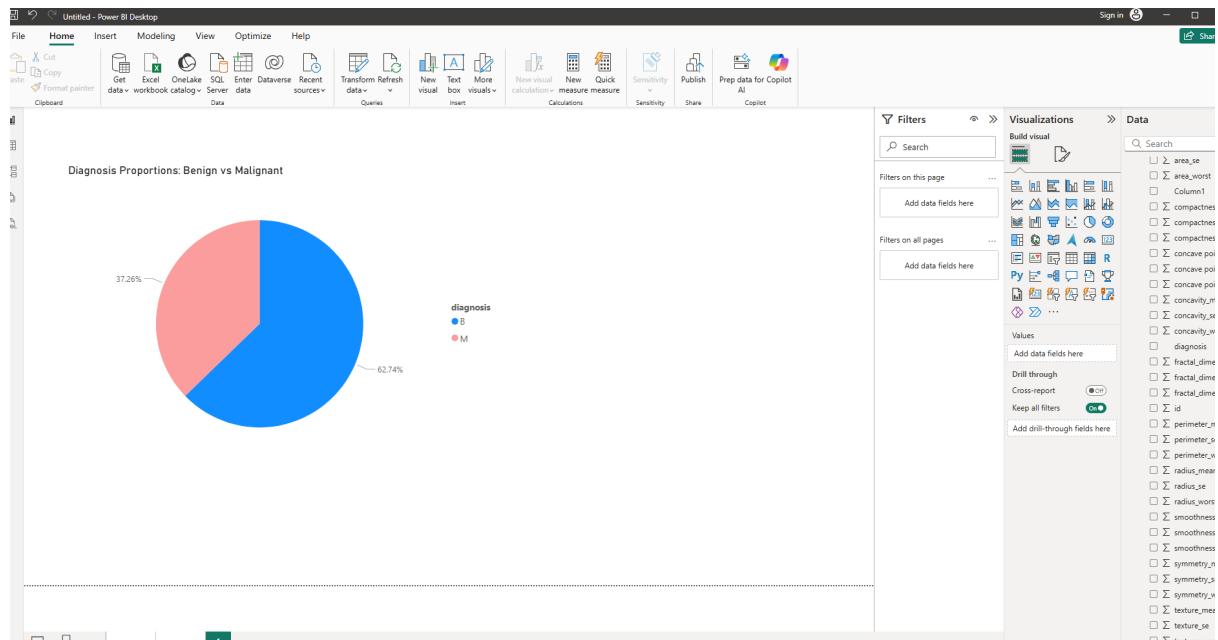
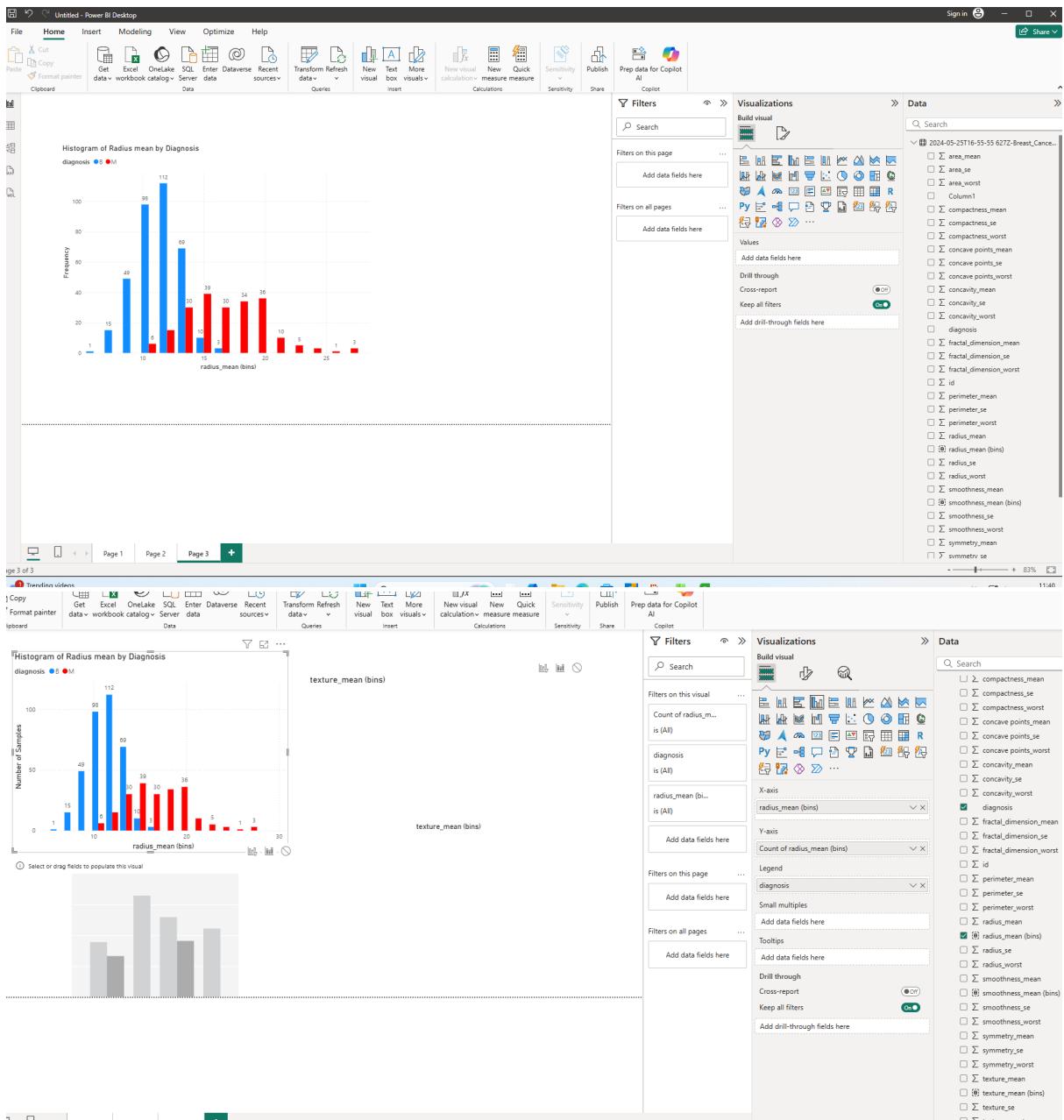


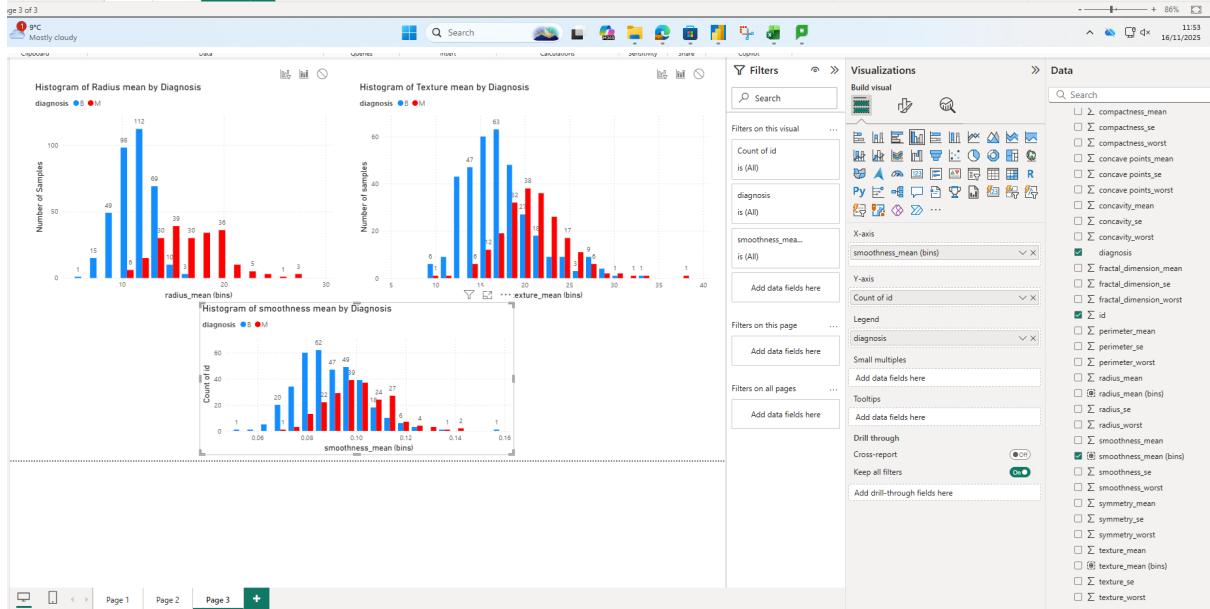
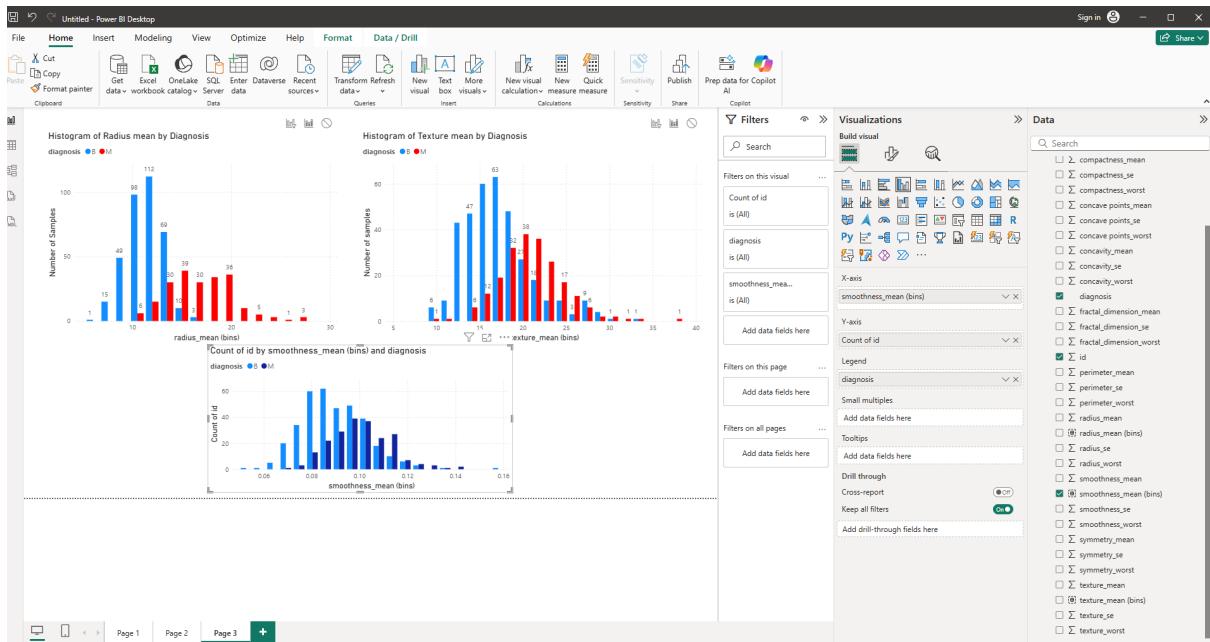
Supporting Info Upload – Group Project

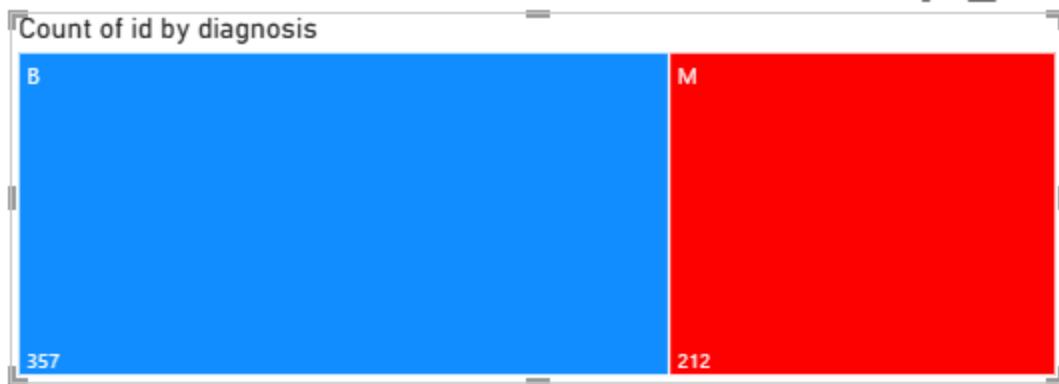
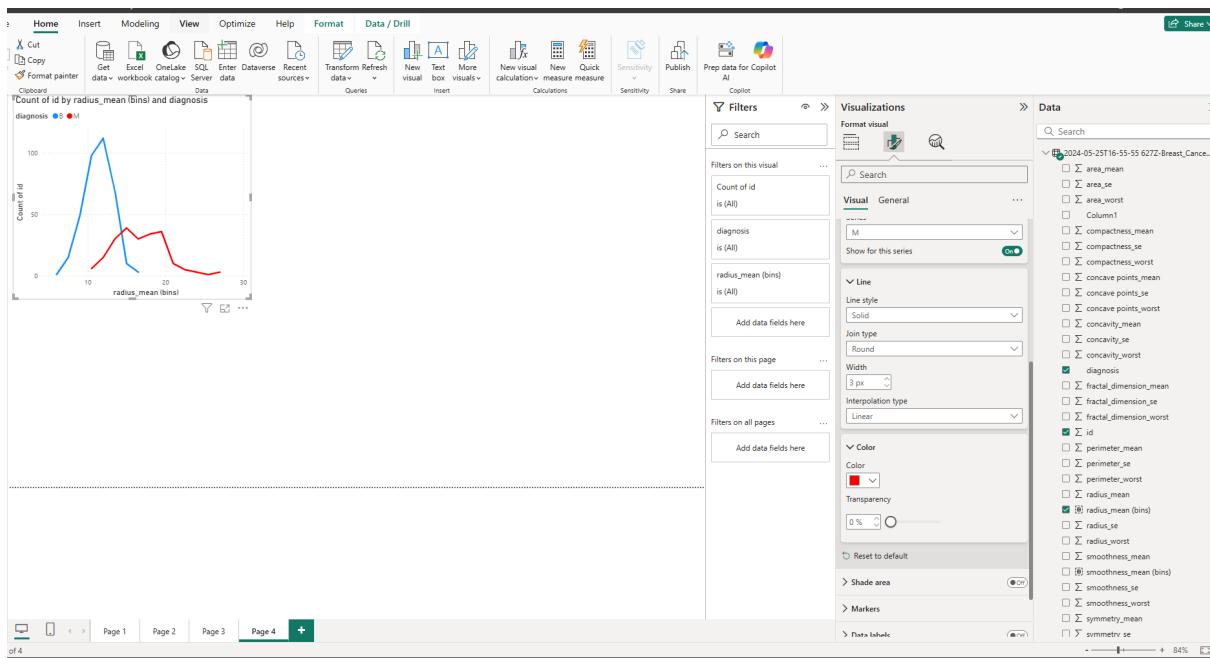
These are the images of the progress of few Visualizations.



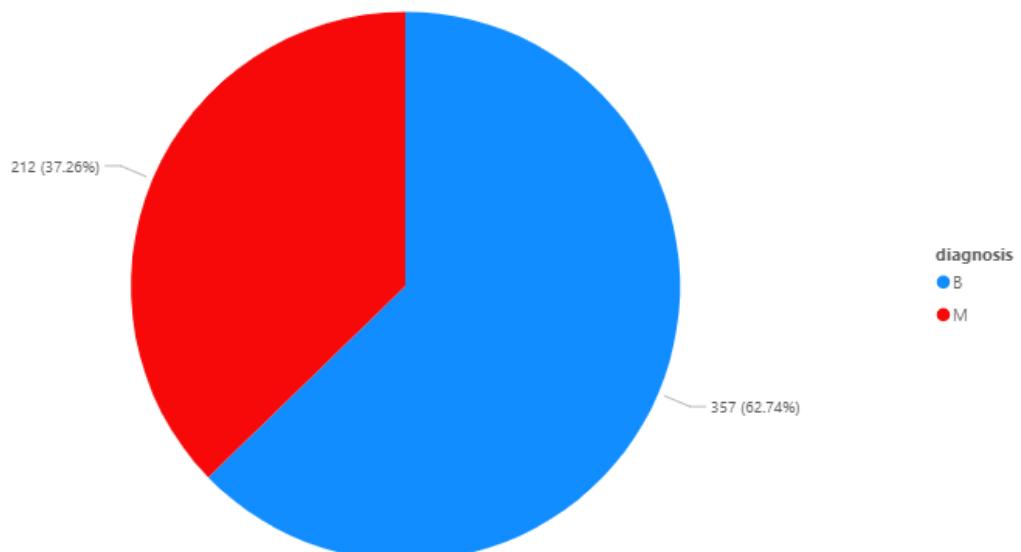






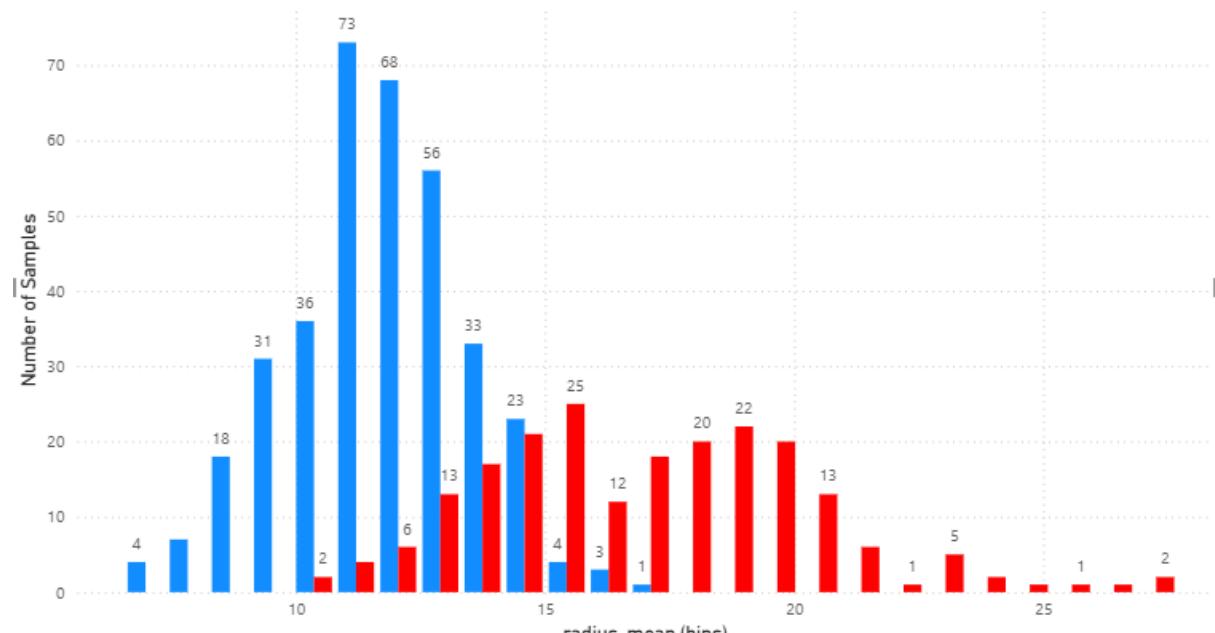


Diagnosis Proportions: Benign vs Malignant



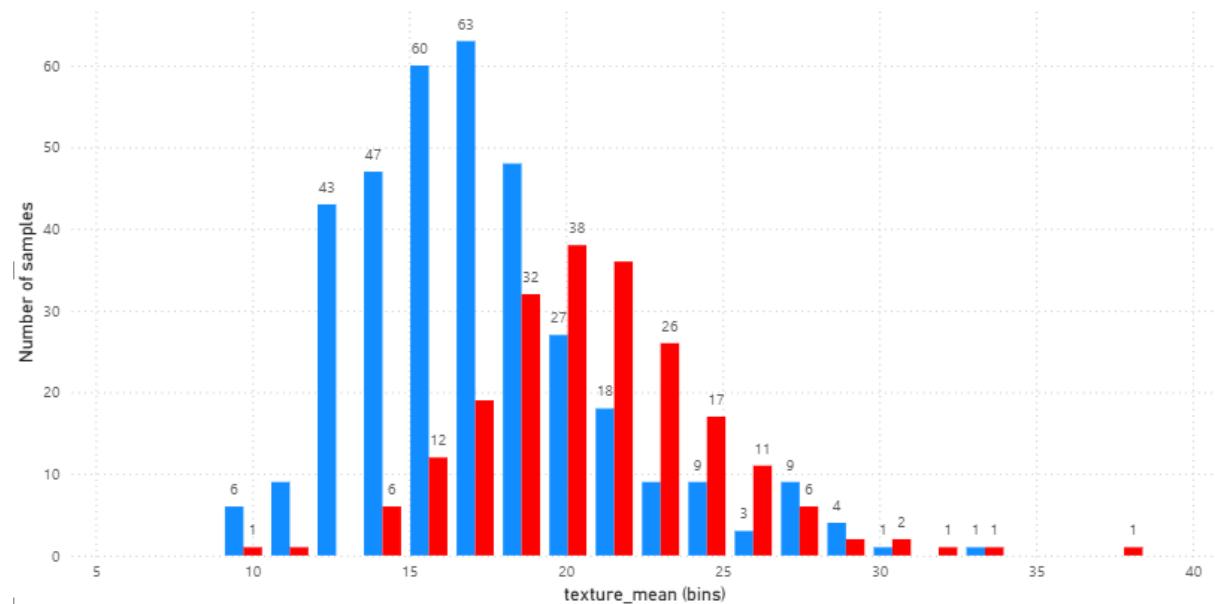
Histogram of Radius mean by Diagnosis

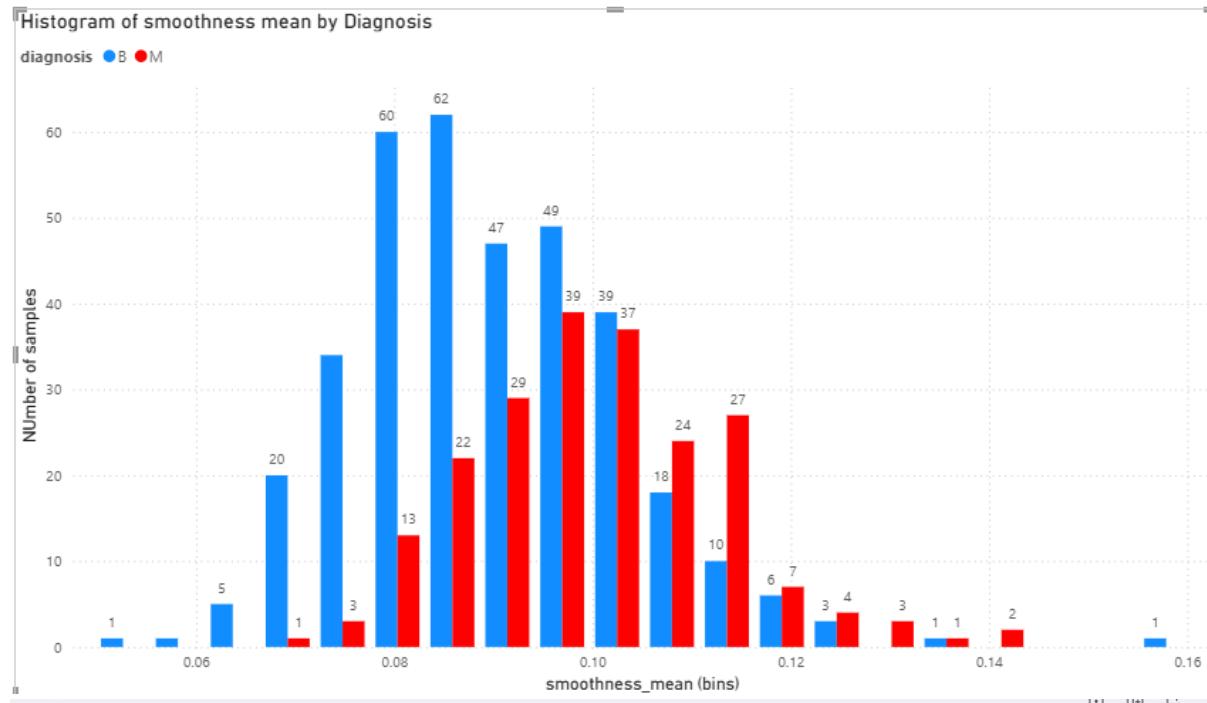
diagnosis ● B ● M



Histogram of Texture mean by Diagnosis

diagnosis ● B ● M





Comparative Analysis of Cell Radius, Texture, and Smoothness in Malignant vs Benign Breast Tumors

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Prifysgol
De Cymru

Introduction

Breast cancer remains one of the leading causes of mortality among women worldwide. Early diagnosis is critical for improving outcomes, and data-driven approaches can support clinical decision-making. We analyzed over 569 dataset which is created by William H. Wolberg. We researched over this dataset whether a tumor is benign or malignant (Cancer) using total 569 records of patients. Specifically, from cell nucleus features extracted from digitized images. Our goal is to

Dataset

The dataset contains 569 patient records, each identified by a unique ID. It includes 30 numerical features extracted from digitized images of cell nuclei, with 357 benign and 212 malignant cases.

Diagnosis	N (No.)	Variable	Mean	Std Dev	Minimum	Maximum	Median	N (No.)	
B	357	radius_mean	12.845538	1.7950118	9.91	20.000000	12.200000	357	0
B	357	radius_mean	12.845538	1.7950118	9.91	20.000000	12.200000	357	0

Hypothesis

H₀: There is no significant difference in cell radius, texture, or smoothness between malignant and benign tumors.
H₁: At least one of these features differs significantly between the two groups.

Methodology 1

Fitted Normal Distribution for radius_mean			
Goodness-of-Fit Test for Normal Distribution			
Test	Anderson-Darling	Shapiro-Wilk	Kolmogorov-Smirnov
Anderson-Darling	0.00000000	0.00000000	0.00000000
Shapiro-Wilk	0.00000000	0.00000000	0.00000000
Kolmogorov-Smirnov	0.00000000	0.00000000	0.00000000
Crps vs Mean	0.00000000	0.00000000	0.00000000
Anderson-Darling	0.00000000	0.00000000	0.00000000

Normality Test

Fitted Normal Distribution for radius_mean			
Goodness-of-Fit Test for Normal Distribution			
Test	Anderson-Darling	Shapiro-Wilk	Kolmogorov-Smirnov
Anderson-Darling	0.00000000	0.00000000	0.00000000
Shapiro-Wilk	0.00000000	0.00000000	0.00000000
Kolmogorov-Smirnov	0.00000000	0.00000000	0.00000000
Crps vs Mean	0.00000000	0.00000000	0.00000000
Anderson-Darling	0.00000000	0.00000000	0.00000000

All three variables (radius_mean, texture_mean, smoothness_mean) failed normality tests ($p < 0.05$), indicating non-normal distributions. Therefore, non-parametric methods were selected for further analysis.

Methodology 2

Kruskal-Wallis Test

Kruskal-Wallis Test	Chi-Square	Df	P > ChSq
305.0811	1	<0.001	

Non-parametric analysis reveals a statistically significant difference in radius_mean between malignant and benign cases (Wilcoxon $Z = 11.0098$, $p < 0.0001$; Kruskal-Wallis $\chi^2 = 121.2209$, $p < 0.0001$).

This suggests that radius_mean is a very strong discriminative feature in breast cancer diagnosis.

smoothness_mean shows a significant difference between malignant and benign cases (Wilcoxon $Z = 8.8629$, $p = 0.0001$; Kruskal-Wallis $\chi^2 = 78.5564$, $p < 0.0001$).

This suggests that smoothness_mean is a moderate-strong discriminative feature in breast cancer diagnosis.

Non-Parametric One-Way ANOVA

Kruskal-Wallis Test	Chi-Square	Df	P > ChSq
121.2209	1	<0.001	

Non-parametric analysis reveals a statistically significant difference in radius_mean between malignant and benign cases (Wilcoxon $Z = 11.0098$, $p < 0.0001$).

This suggests that radius_mean is a very strong discriminative feature in breast cancer diagnosis.

Methodology 3

T-Test

Equality of Variances			
Welch F	20.000000	Df	1
Welch P	0.00000000	Pr > F	1.00000000

An independent samples T-test revealed a significant difference in radius_mean between malignant and benign cases ($t = -22.21$, $p < 0.0001$).

The mean radius was substantially higher in malignant tumors (12.845538) compared to benign (12.151), confirming its strong diagnostic value.

An independent samples T-test revealed a significant difference in texture_mean between malignant and benign cases ($t = -11.02$, $p < 0.0001$).

Malignant tumors had a higher mean texture (17.91) compared to benign (17.31), supporting its role as a discriminative feature in breast cancer diagnosis.

Interpretation & Conclusion

All three features—radius, texture, and smoothness—show statistically significant differences between malignant and benign tumors. Among them, radius_mean is the strongest discriminator, followed by texture_mean, with smoothness_mean showing moderate but meaningful separation. These findings support the diagnostic relevance of these features and their potential use in breast cancer classification.