PS C:\Users\Aniket\OneDrive\Desktop\marvellousinfo\Assignments\Assignment_34> & C:/Users/Aniket/AppData/Local/Programs/Python/Python313/python.exe c:/Users/Aniket/OneDrive/Desktop/marvellousinfo/Assignments/Assignment_34/BreastCanc e.py

mean radius mean texture mean perimeter ... worst concave points worst symmetry worst fractal dimension

0	17.99	10.38	122.80	0.2654	0.4601	0.11890
1	20.57	17.77	132.90	0.1860	0.2750	0.08902
2	19.69	21.25	130.00	0.2430	0.3613	0.08758
3	11.42	20.38	77.58	0.2575	0.6638	0.17300
4	20.29	14.34	135.10	0.1625	0.2364	0.07678
564	21.56	22.39	142.00	0.2216	0.2060	0.07115
565	20.13	28.25	131.20	0.1628	0.2572	0.06637
566	16.60	28.08	108.30	0.1418	0.2218	0.07820
567	20.60	29.33	140.10	0.2650	0.4087	0.12400
568	7.76	24.54	47.92	0.0000	0.2871	0.07039

[569 rows x 30 columns]

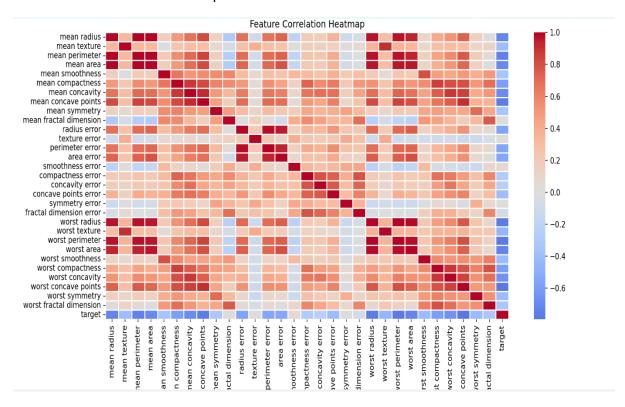
statestical summery of data:

mean radius mean texture mean perimeter ... worst symmetry worst fractal dimension target

dimension target									
count 569.000000 569.000000 569.000000 569.000000 569.000000									
mean 0.627		19.289649	91.969033	0.290076	0.083946				
std	3.524049	4.301036	24.298981	0.061867	0.018061 0.483918				
min	6.981000	9.710000	43.790000	0.156500	0.055040 0.000000				
25% 0.000	11.700000 000	16.170000	75.170000	0.250400	0.071460				
50% 1.000	13.370000 000	18.840000	86.240000	0.282200	0.080040				
75% 1.000	15.780000 000	21.800000	104.100000	0.317900	0.092080				
max 1.000	28.110000 000	39.280000	188.500000	0.663800	0.207500				

[8 rows x 31 columns]

Feature Correlation on heatmap:



now feature correlation with targets

we will show with only top 10 feature:

Top 10 features correlated with target:

smoothness error 0.067016

mean fractal dimension 0.012838

texture error 0.008303

symmetry error 0.006522

fractal dimension error -0.077972

concavity error -0.253730

compactness error -0.292999

worst fractal dimension -0.323872

mean symmetry -0.330499

mean smoothness -0.358560

Name: target, dtype: float64

After scaling: X_scaled shape = (569, 30) y shape = (569,)

model build on training set

model build on testing set

Accuracy: 98.24561403508771

Confusion Matrix:

[[41 1]

[171]]

Classification Report:

precision recall f1-score support

THE END OF PROGRAM (BREAST CANCER CASE STUDY)

Conclusion

- The classification model built on the Breast Cancer dataset performs **very well** with almost **perfect accuracy**.
- The results indicate that the model can be **reliably used for breast cancer tumor classification** (malignant vs benign).
- Only 2 out of 114 predictions were wrong, showing strong generalization ability.
- With such high performance, this model can be an excellent starting point for medical diagnostic support systems, though real-world deployment should include further validation on unseen clinical data.