

Features in the Dataset

The dataset consists of 30 *features*, categorized based on the measurements of the cell nuclei:

1. Radius:

radius_mean, radius_se, radius_worst:

- Represents the mean, standard error, and worst-case measurements of the distance from the center to the perimeter of the cell.
- Larger radii are often associated with malignant tumours due to irregular and larger cell growth.

2. Texture:

texture_mean, texture_se, texture_worst:

- Captures the standard deviation of pixel intensity values within the cell.
- Malignant tumours exhibit greater variability in texture due to irregular structures.

3. Perimeter:

perimeter_mean, perimeter_se, perimeter_worst:

- Measures the length of the outer boundary of the cell nucleus.
- Malignant cells tend to have larger and more irregular perimeters.

4. Area:

area_mean, area_se, area_worst:

- Represents the area of the cell nucleus.
- Larger cell areas are common in malignant tumours.

5. Smoothness:

smoothness_mean, smoothness_se, smoothness_worst:

- Measures the variation in the radius lengths, indicating how smooth the cell edges are.
- Malignant cells often have less smooth (more irregular) edges.

6. Compactness:

compactness_mean, compactness_se, compactness_worst:

- Calculated as $[\text{Perimeter}^2 / (\text{Area} - 1)]$, this measures how compact the cell is.
- Malignant tumours are typically more compact due to uncontrolled cell division.

7. Concavity:

concavity_mean, concavity_se, concavity_worst:

- Describes the severity of concave areas in the cell nucleus boundary.
- Malignant tumours often have deeper concavities.

8. Concave Points:

concave_points_mean, concave_points_se, concave_points_worst:

- The number of concave portions in the boundary of the nucleus.
- More concave points are a strong indicator of malignancy.

9. Symmetry:

symmetry_mean, symmetry_se, symmetry_worst:

- Measures the asymmetry in the shape of the nucleus.
- Malignant tumours are usually less symmetric.

10. Fractal Dimension:

***fractal_dimension_mean, fractal_dimension_se,
fractal_dimension_worst:***

- Quantifies the complexity of the boundary of the nucleus.
- A higher fractal dimension indicates irregular and complex boundaries, which are more likely in malignant cells.