

# INVESTIGATE CANCER

## A STEM INTEGRATED AP BIOLOGY UNIT

DAY 1 - 3

GENETIC-BASIS OF CANCER

### Topics and Related Student Activities:

- Cell cycle function
- Digital Activity
- Oncogenes and tumor suppressor genes
- Card sort and classification
- Identification on chromosomes
- Mutations and effects of mutations
- Reading and response on factors that cause mutations

DAY 4

PHYSICAL DESCRIPTION OF TUMORS

### Topics and Related Student Activities:

- How cell cycle function causes tumors
- Tumors and nodules (specifically lung)
- Reading and response on physical characteristics of nodules
- Characteristics of benign and malignant nodules
- Card sort of CT Images- "Benign or Malignant?"
- Data analysis of sort results
- Imaging tools for identification of nodules

DAY 5

STEM CAREER CONNECTION

### Topics and Related Student Activities:

- Using computer programming to evaluate CT Scans
- Computer Vision and CT Scans
- Machine Learning for nodule classification
- Data output and the meaning of those data
- "How Well is the Model Working?" statistical analysis sheet
- Other connections to careers in computer vision and machine learning

### Final Student Reflection

- Potential Field Trip to UCF



# Integrating Computer Vision in AP Biology

@ Lake Nona High School

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**Integration Theme:** How Low-Dosage Computed Tomography (LDCT) Scans and Computer-Assisted Detection (CAD) systems aid in accurate identification of potentially life threatening lung cancer nodules.

## Mission

As part of Research Experience for Educators (RET) the primary mission of the project is to expose Advanced Placement Biology high school students to the field of computer vision by providing them with a learning experience that both advances their knowledge in a specific biological topic while simultaneously integrating concepts and tools typically encountered in the field of computer vision.

## ACCURACY OF LUNG CANCER DIAGNOSIS- HUMAN VISION VS. "COMPUTER VISION"

### "THE HUMAN MODEL"

Students classify nodules using the table of characteristics and run statistical analyses on the results.

Table 2. Evaluating Probability of Malignancy using Characterization of Nodules

Characteristic	Likely Benign ≤ 1.5 cm	Possibly Malignant 1.5-2.5 cm	Highly Likely Malignant ≥ 2.5 cm
Diameter	Smooth	Spiculated	Corona radiate or spiculated
Nodule Margins (border)	This Wall	Thick Wall	
Calcification (Hole/dark in the center)	Concentric, Central, lamellar, popcorn or homogeneous patterns	Non-calcified or eccentric calcification	
Calcification (Calcium Deposits)	Subsolid	Non-solid = "ground glass"	
Density	Round, Oval or Polygonal	Irregular	
Shape			

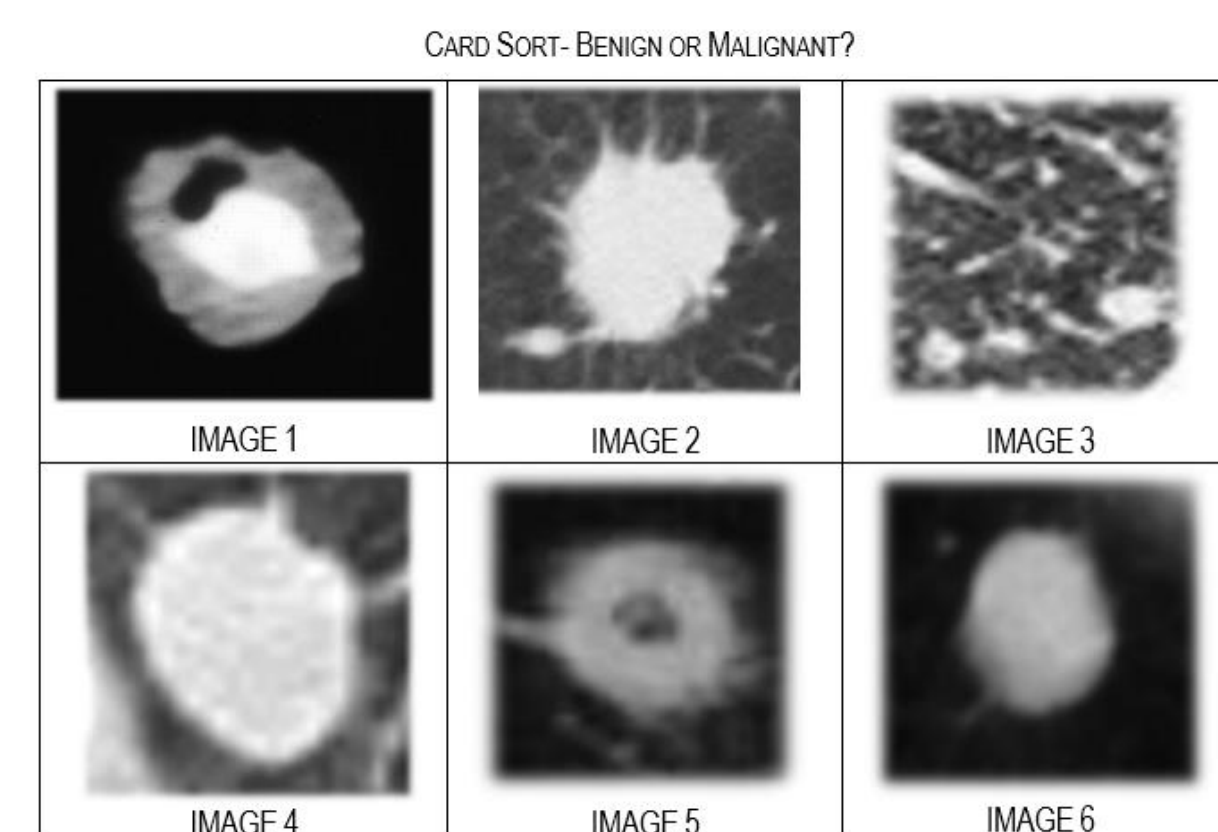


Table 1: Confusion Matrix

$\text{Accuracy} = \frac{\sum TP + TN}{\sum P + N (\sum TP + FN + FP + FN)}$	TP = TRUE POSITIVE (Known = Predicted) TN = TRUE NEGATIVE (Known = Predicted)
$\text{Sensitivity} = \frac{\sum TP}{\sum TP + FN}$	FP = FALSE POSITIVE (Known = Negative; Predicted = Positive)
$\text{Specificity} = \frac{\sum TN}{\sum TN + FP}$	FN = FALSE NEGATIVE (Known = Positive; Predicted = Negative)
$\text{Precision} = \frac{\sum TP}{\sum TP + FP}$	

True Positives Total predicted as positive (malignant) and labeled as positive (malignant)	False Positives Total predicted as positive (malignant) and labeled as negative (benign)
False Negatives Total predicted as negative (benign) and labeled as positive (malignant)	True Negatives Total predicted as negative (benign) and labeled as negative (benign)

## VERSUS

### "THE COMPUTER MODEL"

Students run statistical analyses on the results of the program and compare them with their results from the "human model".

**Lesson Objective:** Students will explain how computer vision and machine learning tools can improve accuracy and speed of lung cancer diagnoses.

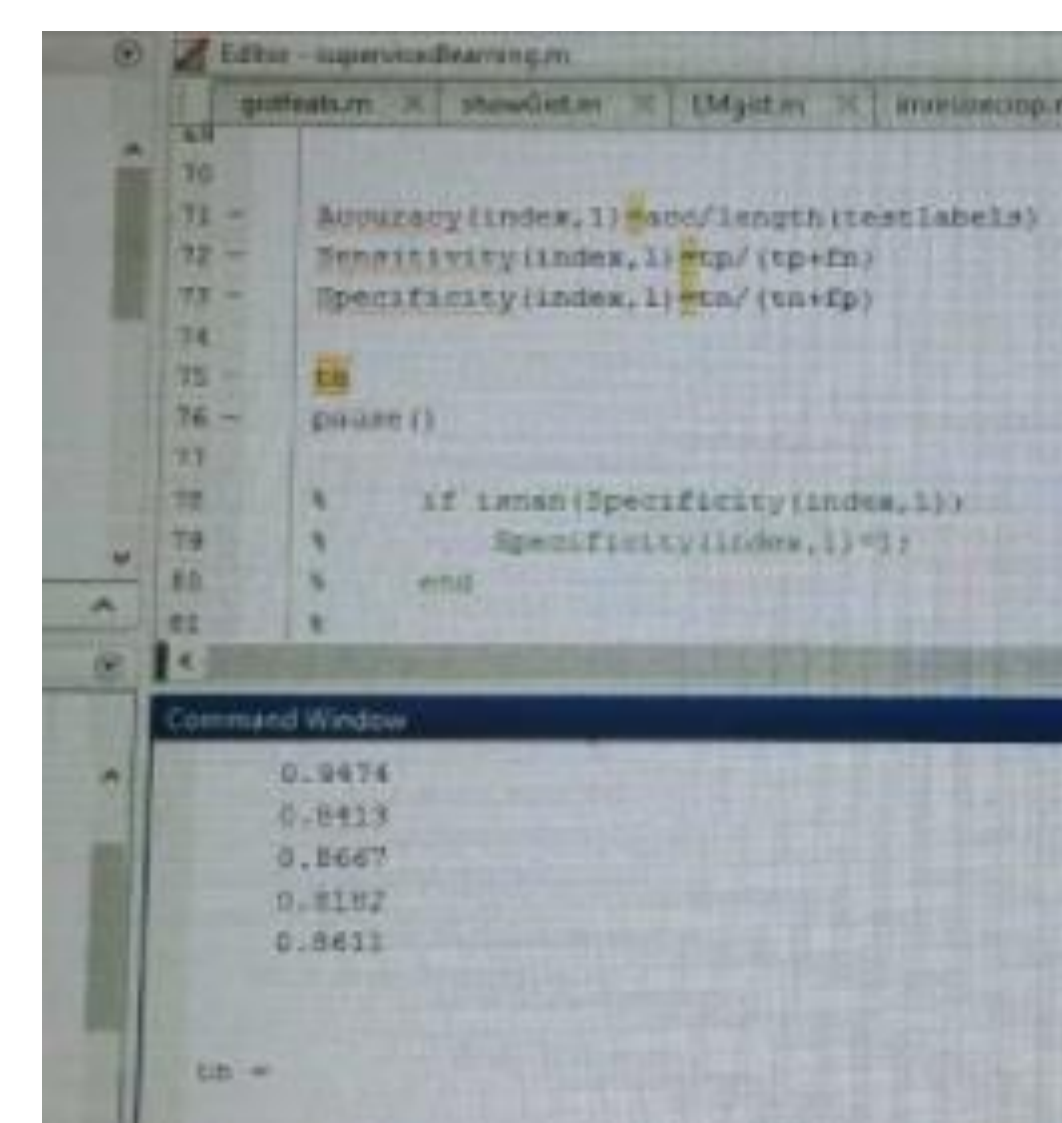
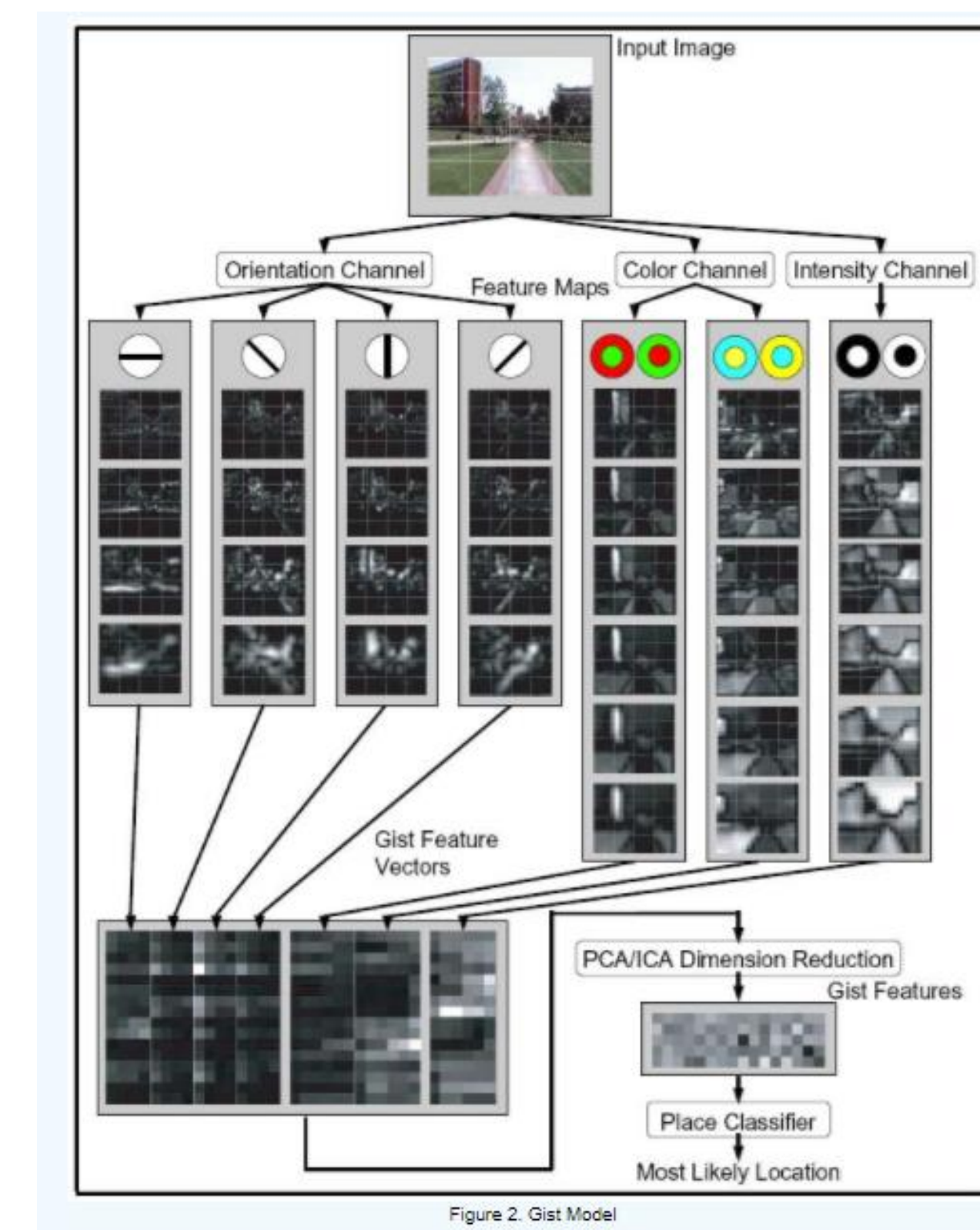


Figure 1: Schematic representation of 5-fold cross-validation.

Table 2: Confusion Matrix from Program Output

True Positives Total predicted as positive (malignant) and labeled as positive (malignant)	False Positives Total predicted as positive (malignant) and labeled as negative (benign)
False Negatives Total predicted as negative (benign) and labeled as positive (malignant)	True Negatives Total predicted as negative (benign) and labeled as negative (benign)

Table 3: Confusion Matrix from Program Output

True Positives Total predicted as positive (malignant) and labeled as positive (malignant)	False Positives Total predicted as positive (malignant) and labeled as negative (benign)
False Negatives Total predicted as negative (benign) and labeled as positive (malignant)	True Negatives Total predicted as negative (benign) and labeled as negative (benign)

Table 4: Confusion Matrix from Program Output

True Positives Total predicted as positive (malignant) and labeled as positive (malignant)	False Positives Total predicted as positive (malignant) and labeled as negative (benign)
False Negatives Total predicted as negative (benign) and labeled as positive (malignant)	True Negatives Total predicted as negative (benign) and labeled as negative (benign)

Table 5: Confusion Matrix from Program Output

True Positives Total predicted as positive (malignant) and labeled as positive (malignant)	False Positives Total predicted as positive (malignant) and labeled as negative (benign)
False Negatives Total predicted as negative (benign) and labeled as positive (malignant)	True Negatives Total predicted as negative (benign) and labeled as negative (benign)

### EXTENSION GOALS

- Help students understand that "science" is not done in a vacuum- there are connections to mathematics, technology, etc. AND there is a huge collaboration piece
- Have interested students meet on a Saturday at the College of Computer Vision for tour and information
- Provide a resource for students interested in computer vision focused science fair project
- Make connections to computer vision technology in other areas of the curriculum (bacteria counting, population analysis, etc.)

## Predicting Cancer Mutation Rate

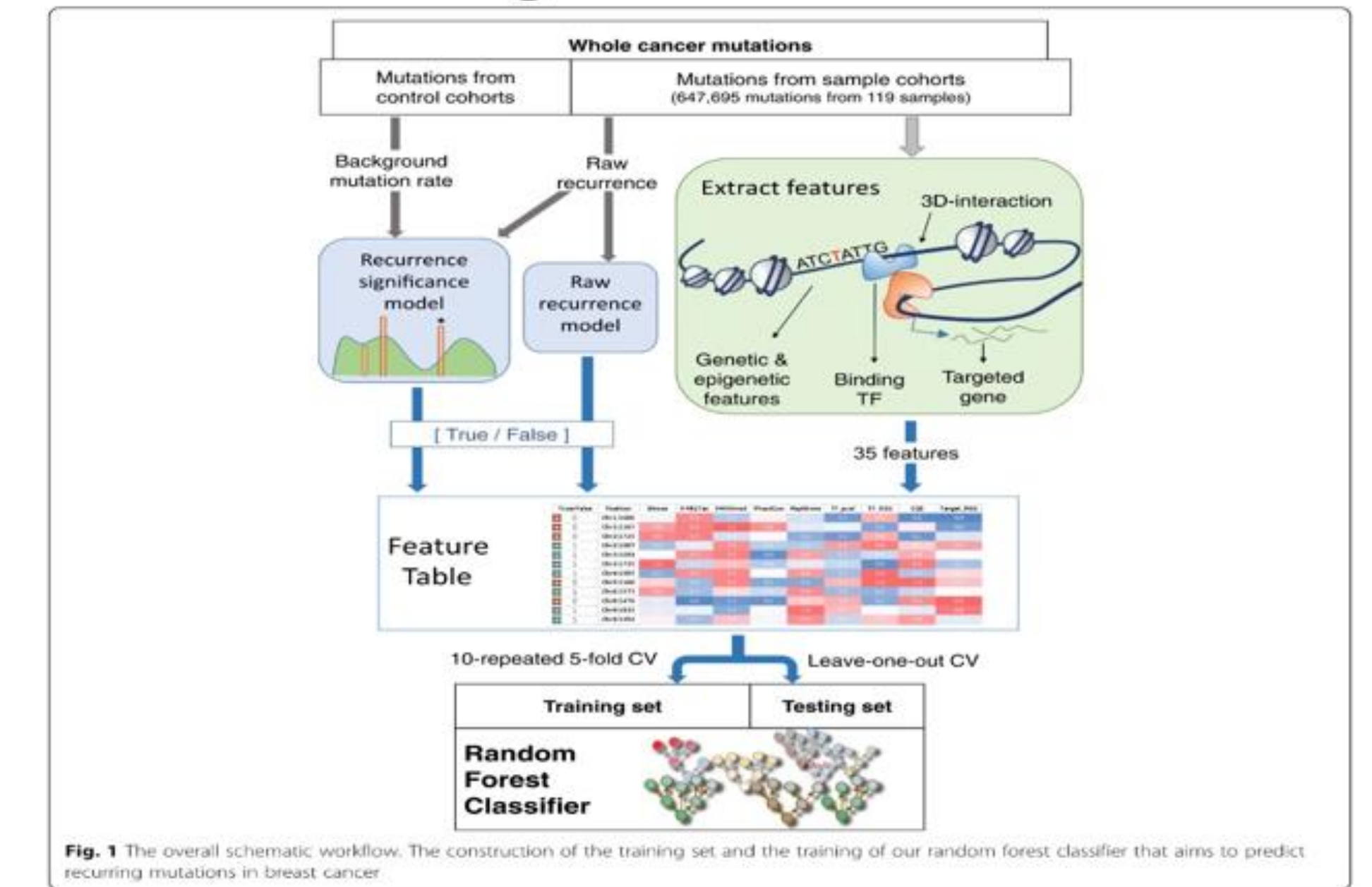


Fig. 1 The overall schematic workflow. The construction of the training set and the training of our random forest classifier that aims to predict recurring mutations in breast cancer.

### AP BIOLOGY LEARNING OBJECTIVES:

- LO 3.6 . . .predict how a change in a specific DNA or RNA sequence can result in changes in gene expression
- LO 3.7 . . .make predictions about natural phenomena occurring during the cell cycle.
- LO 3.8 . . .describe the events that occur in the cell cycle.
- LO 3.21 . . .use representations to describe how gene regulation influences cell products and function.
- LO 3.22 . . .explain how signal pathways mediate gene expression, including how this process can affect protein production.
- LO 3.23 . . .use representations to describe mechanisms of the regulation of gene expression.

### AP BIOLOGY REQUIRED SCIENCE PRACTICES:

- SP 2.1 . . .justify the selection of a mathematical routine to solve problems.
- SP 2.2 . . .apply mathematical routines to quantities that describe natural phenomena.
- SP 5.1 . . .analyze data to identify patterns or relationships.
- SP 5.2 . . .refine observations and measurements based on data analysis.
- SP 5.3 . . .evaluate the evidence provided by data sets in relation to a particular scientific question.

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