

# A quantum-inspired classifier for clonogenic assay evaluations - Supplementary material

Giuseppe Sergioli<sup>1,\*</sup>, Carmelo Militello<sup>2</sup>, Leonardo Rundo<sup>3,4</sup>, Luigi Minafra<sup>2</sup>, Filippo Torrisi<sup>5</sup>, Giorgio Russo<sup>2</sup>, Keng Loon Chow<sup>1</sup>, and Roberto Giuntini<sup>1,6</sup>

<sup>1</sup>University of Cagliari, Cagliari, Italy.

<sup>2</sup>Institute of Molecular Bioimaging and Physiology, Italian National Research Council, Cefalú, Palermo, Italy.

<sup>3</sup>Department of Radiology, University of Cambridge, Cambridge, United Kingdom.

<sup>4</sup>Cancer Research UK Cambridge Centre, University of Cambridge, Cambridge, United Kingdom.

<sup>5</sup>Department of Biomedical and Biotechnological Sciences, University of Catania, Catania, Italy.

<sup>6</sup>Centro Linceo Interdisciplinare "Beniamino Segre", Accademia dei Lincei, Rome, Italy.

\*Corresponding author: giuseppe.sergioli@gmail.com

## ABSTRACT

This file contains all the supplementary material regarding the experimental trials described and carried out in the manuscript "A quantum-inspired classifier for clonogenic assay evaluations". In the first section, detailed descriptions of the extracted texture features are provided. In the second section, we show the experimental results obtained for each of the four cell lines MDA-MD-231, U87-MG, MCF7, and U251, by considering the best performing image feature. The third section is devoted to show the performance of the Helstrom Quantum Classifier (HQC) on a new unseen test set. Finally, in the last section we summarize the full results of the whole experiment, for all the investigated cell lines and image features.

## S1. The extracted Haralick's features

We start from the assumption that biomedical images contain information phenotype of the underlying physiopathology, which is not always easily identifiable by simple 'visual' inspection. These information can be revealed through quantitative analysis, by extracting the so called 'descriptors' in order to make it possible to acquire further knowledge on the dominion. The Gray-Level Co-occurrence Matrix (GLCM) computation is the first step to obtain the features.

Formally, let a GLCM with size  $L \times L$ , where  $L$  represents the maximum number of gray-levels according to a given quantization scheme, denote the second-order joint probability function  $p(i, j)$  of an image region (where  $i, j \in [0, 1, \dots, L-1]$  represent a gray-level pair) after the normalization by the total number of pixels. These descriptors are generally called Haralick's features<sup>1,2</sup>.

Given a squared window of size  $\omega \times \omega$  pixels sliding over the whole image<sup>3</sup>, we computed the following GLCM-based features (with  $i, j \in [0, 1, \dots, L-1]$ ):

- contrast  $\in [0, (L-1) \times (L-1)]$  yields a measure of the intensity contrast between neighboring pixels:

$$\text{contrast}(i, j) = \sum_{i=0}^{L-1} \sum_{j=0}^{L-1} |i-j|^2 \cdot p(i, j), \quad (1)$$

contrast = 0 for a constant image;

- correlation  $\in [-1, 1]$  indicates the degree of correlation between a pixel and its neighbor:

$$\text{correlation}(i, j) = \frac{\sum_{i=0}^{L-1} \sum_{j=0}^{L-1} (i - \mu_x)(j - \mu_y) \cdot p(i, j)}{\sigma_x \sigma_y}, \quad (2)$$

where:  $\mu_x = \sum_i \sum_j i \cdot p(i, j)$ ,  $\mu_y = \sum_i \sum_j j \cdot p(i, j)$ ,  $\sigma_x = \sum_i \sum_j (i - \mu_x) \cdot p(i, j)$ , and  $\sigma_y = \sum_i \sum_j (j - \mu_y) \cdot p(i, j)$  (with  $\sum_i$  and  $\sum_j$  denoting  $\sum_{i=0}^{L-1}$  and  $\sum_{j=0}^{L-1}$ , respectively). This feature is 1 or  $-1$  for a perfectly positively or negatively correlated image, respectively;

- energy  $\in [0, 1]$  calculates the sum of squared elements in the GLCM:

$$\text{energy}(i, j) = \sum_{i=0}^{L-1} \sum_{j=0}^{L-1} p(i, j)^2, \quad (3)$$

energy = 1 for a constant image;

- homogeneity  $\in [0, 1]$  Returns a value that measures the closeness of the distribution of elements in the GLCM to the GLCM diagonal:

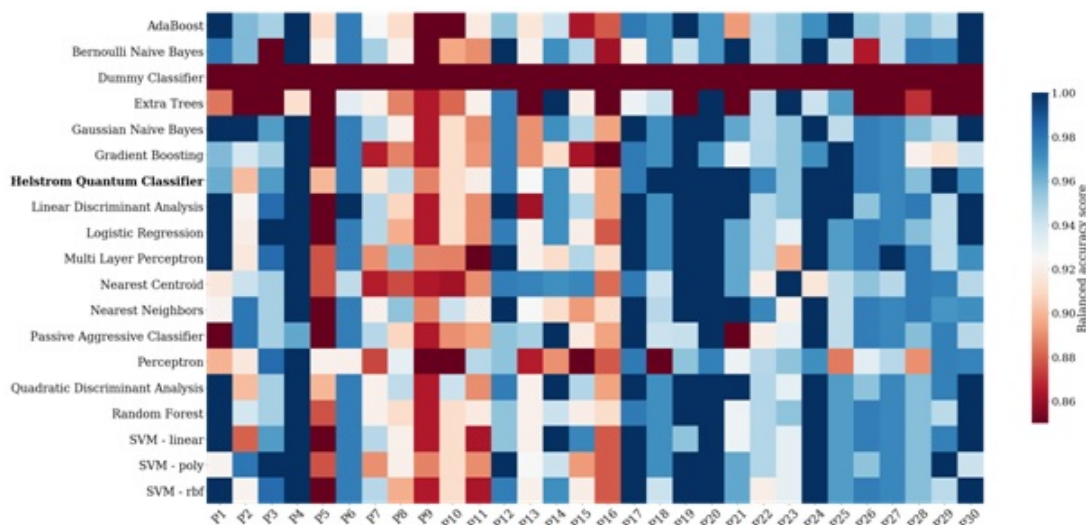
$$\text{homogeneity}(i, j) = \frac{\sum_{i=0}^{L-1} \sum_{j=0}^{L-1} p(i, j)}{(1 + |i - j|)}, \quad (4)$$

homogeneity = 1 for a diagonal GLCM.

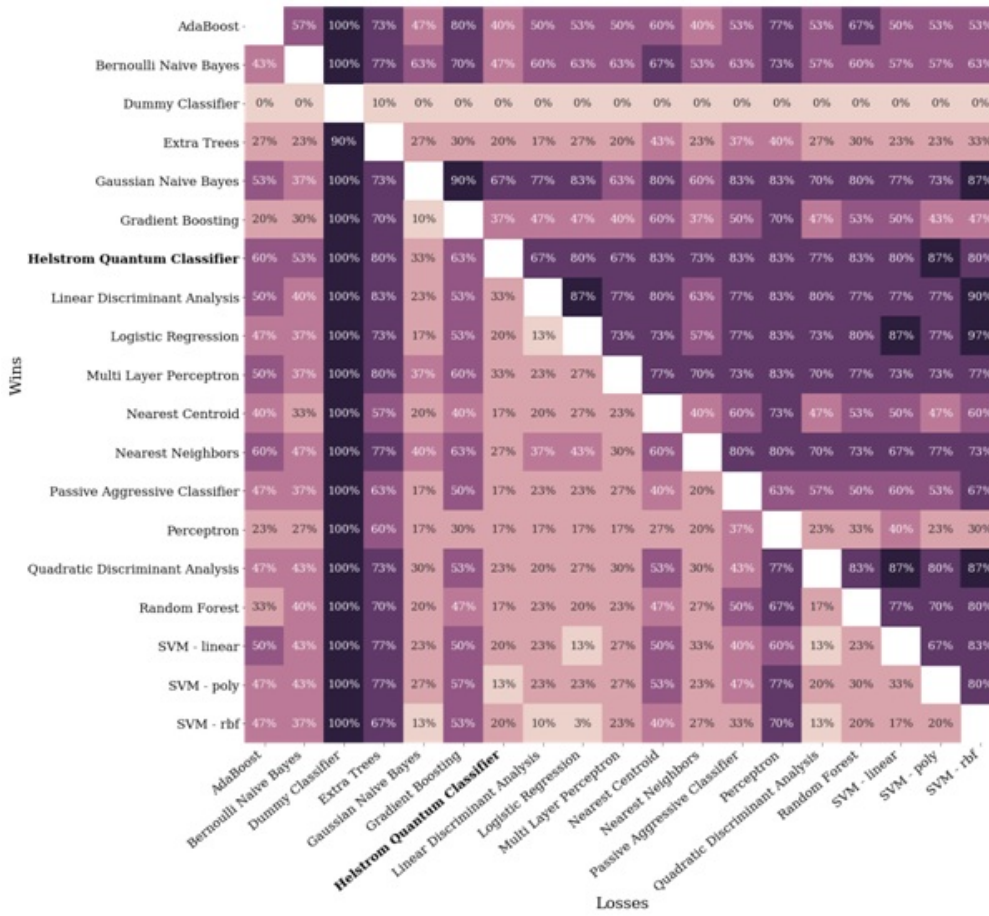
## S2. Experimental results

This section is divided into four groups, one for each cell line MDA-MD-231, U87-MG, MCF7 and U251, respectively. Each group contains two reports, the first shows: (1) the balanced accuracy score over 30 datasets (for the best performing image feature) for the HQC and the 18 competing classifiers, obtained by hypertuning the hyperparameters of each classifier in order to optimize the balanced accuracy score; (2) heatmaps of a classifier outperforming (“wins”) over another classifier (“losses”) out of the 30 datasets (for the best performing image feature); and (3) a table showing the averaged balanced accuracy score over the 30 datasets for each of the six image features, *RGB*,  $L^*u^*v^*$ , *contrast*, *correlation*, *energy* and *homogeneity*. The second report is the analogous of the first, where the role of the balanced accuracy is replaced by the AUROC score. All performance evaluation is performed using the test set. The aim of the experimental procedure is to find the most informative image feature in discriminating a pixel between a colony or a background, i.e., the image feature which maximizes the value of the balanced accuracy and the AUROC scores, respectively.

### S2.1. Cell line MDA-MD-231



**Fig. 2.1.1 | Balance accuracy score of 19 classifiers across 30 homogeneity image feature datasets for cell line MDA-MD-231.**



**Fig. 2.1.2 | Percentage of datasets where model A (“Wins”) outperformed model B (“Losses”) out of 30 homogeneity image feature datasets for cell line MDA-MD-231 (balanced accuracy score).**

Classifiers	Image features					
	RGB	L*u*v*	Contrast	Correlation	Energy	Homogeneity
AdaBoost	0.861 ± 0.088	0.878 ± 0.065	0.898 ± 0.065	0.738 ± 0.110	0.936 ± 0.052	0.942 ± 0.046
Bernoulli Naive Bayes	0.841 ± 0.060	0.855 ± 0.079	0.840 ± 0.097	0.630 ± 0.126	0.854 ± 0.087	0.940 ± 0.057
Dummy Classifier	0.507 ± 0.047	0.513 ± 0.056	0.506 ± 0.069	0.518 ± 0.042	0.497 ± 0.069	0.485 ± 0.060
Extra Trees	0.822 ± 0.120	0.839 ± 0.141	0.709 ± 0.163	0.616 ± 0.121	0.767 ± 0.169	0.851 ± 0.139
Gaussian Naive Bayes	0.855 ± 0.063	0.874 ± 0.084	0.886 ± 0.063	0.711 ± 0.102	0.923 ± 0.067	0.954 ± 0.045
Gradient Boosting	0.876 ± 0.076	0.874 ± 0.069	0.899 ± 0.063	0.835 ± 0.113	0.941 ± 0.047	0.933 ± 0.050
<b>Helstrom Quantum Classifier</b>	0.895 ± 0.058	0.883 ± 0.069	0.902 ± 0.070	0.775 ± 0.081	0.938 ± 0.047	<b>0.959 ± 0.036</b>
Linear Discriminant Analysis	0.850 ± 0.066	0.865 ± 0.080	0.837 ± 0.075	0.660 ± 0.122	0.860 ± 0.089	0.955 ± 0.049
Logistic Regression	0.882 ± 0.061	0.879 ± 0.086	0.883 ± 0.066	0.657 ± 0.131	0.932 ± 0.056	0.951 ± 0.045
Multi Layer Perceptron	0.890 ± 0.055	0.885 ± 0.069	0.901 ± 0.069	0.833 ± 0.086	0.942 ± 0.052	0.940 ± 0.091
Nearest Centroid	0.839 ± 0.058	0.868 ± 0.078	0.846 ± 0.070	0.686 ± 0.114	0.867 ± 0.063	0.941 ± 0.044
Nearest Neighbors	0.890 ± 0.069	0.875 ± 0.064	0.903 ± 0.073	0.808 ± 0.070	0.942 ± 0.049	0.953 ± 0.040
Passive Aggressive Classifier	0.831 ± 0.105	0.809 ± 0.113	0.828 ± 0.118	0.607 ± 0.137	0.907 ± 0.062	0.935 ± 0.052
Perceptron	0.790 ± 0.117	0.832 ± 0.106	0.831 ± 0.112	0.631 ± 0.103	0.914 ± 0.054	0.916 ± 0.060
Quadratic Discriminant Analysis	0.872 ± 0.065	0.876 ± 0.074	0.876 ± 0.076	0.731 ± 0.112	0.925 ± 0.066	0.957 ± 0.039
Random Forest	0.874 ± 0.078	0.883 ± 0.074	0.919 ± 0.058	0.827 ± 0.119	0.944 ± 0.043	0.951 ± 0.036
SVM - linear	0.880 ± 0.063	0.873 ± 0.078	0.880 ± 0.071	0.655 ± 0.141	0.929 ± 0.052	0.949 ± 0.046
SVM - poly	0.883 ± 0.055	0.892 ± 0.069	0.899 ± 0.065	0.811 ± 0.084	0.926 ± 0.050	0.950 ± 0.042
SVM - rbf	0.876 ± 0.062	0.874 ± 0.084	0.890 ± 0.069	0.662 ± 0.134	0.931 ± 0.052	0.948 ± 0.046

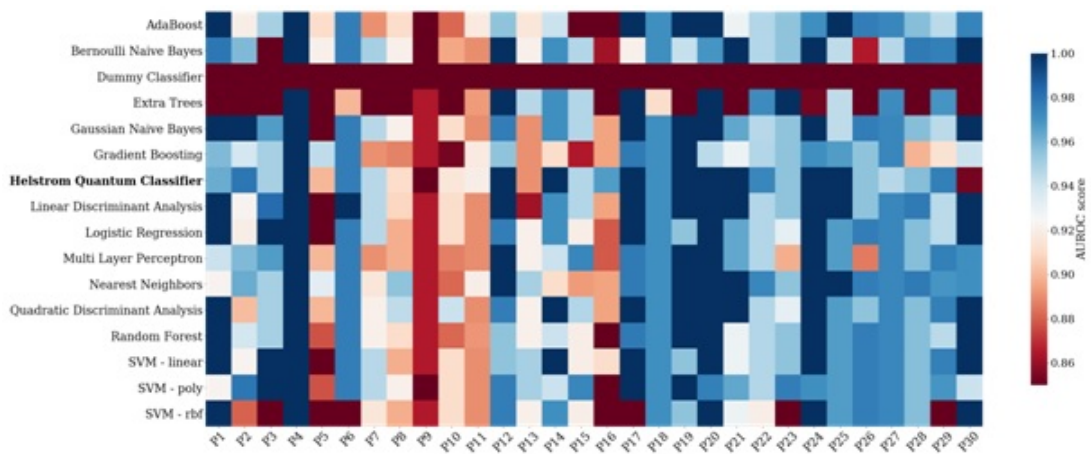


Fig. 2.1.3 | AUROC score of 19 classifiers across 30 homogeneity image feature datasets for cell line MDA-MD-231.

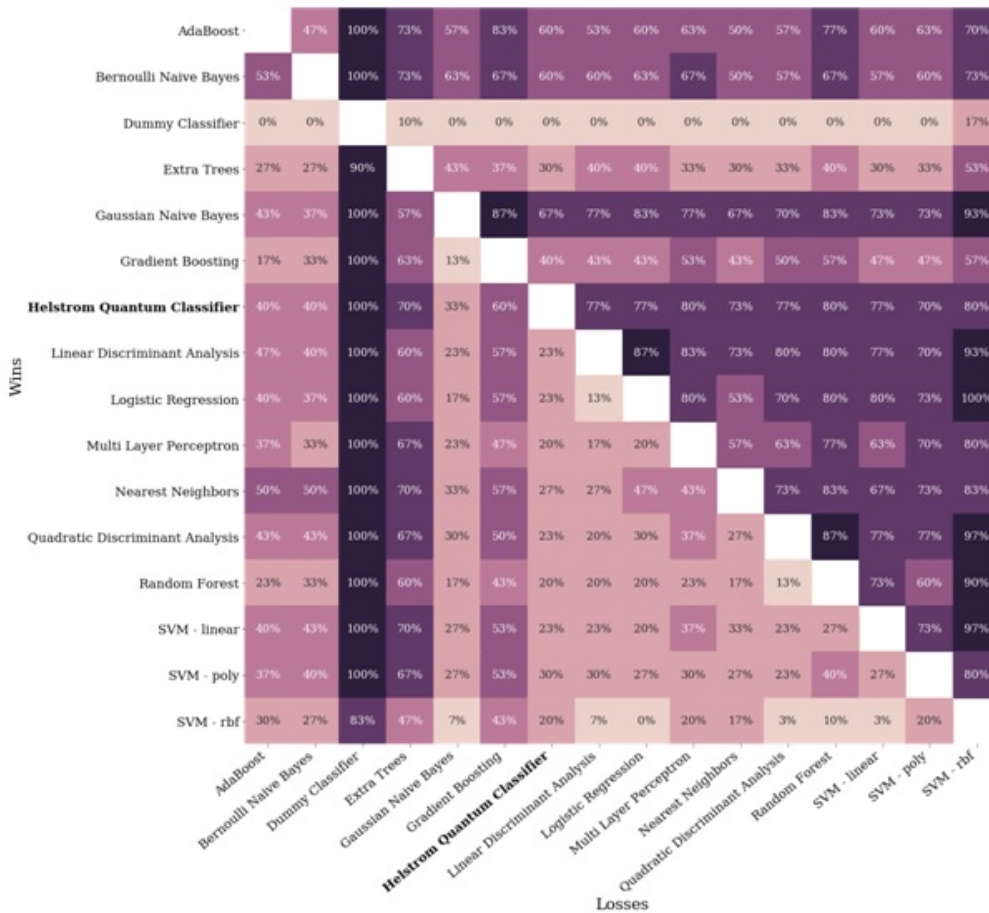


Fig. 2.1.4 | Percentage of datasets where model A (“Wins”) outperformed model B (“Losses”) out of 30 homogeneity image feature datasets for cell line MDA-MD-231 (AUROC score).

Classifiers	Image features					
	RGB	L*u*v*	Contrast	Correlation	Energy	Homogeneity
AdaBoost	0.874 ± 0.076	0.881 ± 0.067	0.898 ± 0.063	0.739 ± 0.130	0.932 ± 0.051	0.943 ± 0.048
Bernoulli Naive Bayes	0.841 ± 0.060	0.855 ± 0.079	0.840 ± 0.097	0.630 ± 0.126	0.854 ± 0.087	0.940 ± 0.057
Dummy Classifier	0.505 ± 0.027	0.498 ± 0.072	0.484 ± 0.046	0.497 ± 0.070	0.503 ± 0.077	0.497 ± 0.049
Extra Trees	0.813 ± 0.132	0.812 ± 0.159	0.678 ± 0.162	0.615 ± 0.127	0.740 ± 0.178	0.837 ± 0.159
Gaussian Naive Bayes	0.855 ± 0.063	0.874 ± 0.084	0.886 ± 0.063	0.711 ± 0.102	0.923 ± 0.067	0.954 ± 0.045
Gradient Boosting	0.880 ± 0.074	0.868 ± 0.076	0.902 ± 0.066	0.834 ± 0.114	0.930 ± 0.056	0.935 ± 0.039
<b>Helstrom Quantum Classifier</b>	0.879 ± 0.081	0.884 ± 0.066	0.905 ± 0.062	0.787 ± 0.082	0.927 ± 0.058	<b>0.954 ± 0.050</b>
Linear Discriminant Analysis	0.850 ± 0.066	0.865 ± 0.080	0.837 ± 0.075	0.660 ± 0.122	0.860 ± 0.089	0.955 ± 0.049
Logistic Regression	0.872 ± 0.072	0.874 ± 0.088	0.886 ± 0.068	0.660 ± 0.138	0.927 ± 0.051	0.950 ± 0.044
Multi Layer Perceptron	0.889 ± 0.056	0.889 ± 0.057	0.894 ± 0.065	0.804 ± 0.104	0.932 ± 0.041	0.946 ± 0.042
Nearest Neighbors	0.883 ± 0.070	0.877 ± 0.066	0.902 ± 0.064	0.800 ± 0.075	0.937 ± 0.049	0.956 ± 0.039
Quadratic Discriminant Analysis	0.872 ± 0.065	0.876 ± 0.074	0.876 ± 0.076	0.731 ± 0.112	0.925 ± 0.066	0.957 ± 0.039
Random Forest	0.881 ± 0.068	0.878 ± 0.077	0.912 ± 0.062	0.817 ± 0.117	0.943 ± 0.040	0.946 ± 0.043
SVM - linear	0.879 ± 0.064	0.876 ± 0.076	0.881 ± 0.073	0.648 ± 0.135	0.914 ± 0.049	0.953 ± 0.043
SVM - poly	0.880 ± 0.058	0.864 ± 0.082	0.897 ± 0.064	0.805 ± 0.087	0.920 ± 0.052	0.950 ± 0.047
SVM - rbf	0.845 ± 0.115	0.860 ± 0.104	0.831 ± 0.147	0.652 ± 0.132	0.879 ± 0.138	0.843 ± 0.193

Results exclude Nearest Centroid, Passive Aggressive Classifier and Perceptron.

## S2.2. Cell line U87-MG

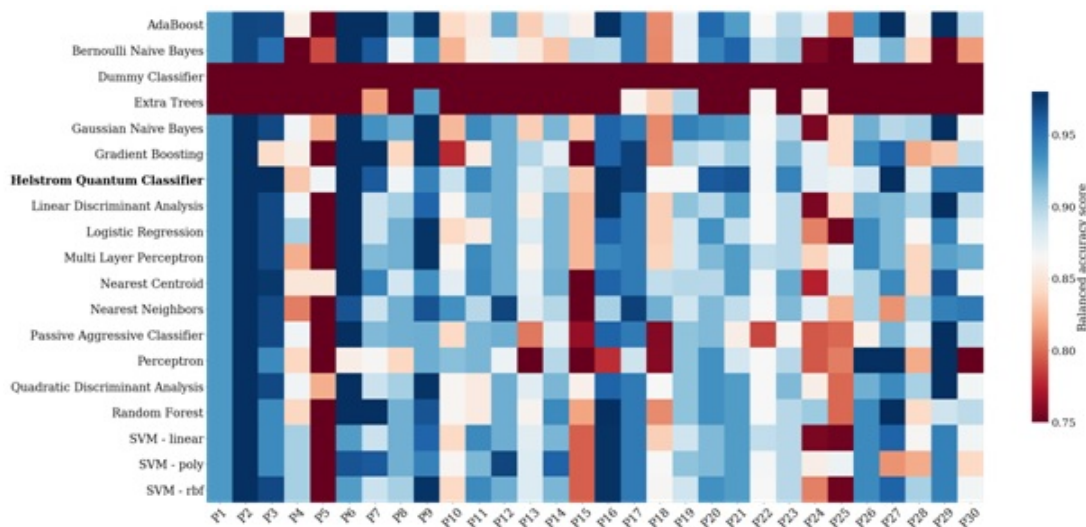


Fig. 2.2.1 | Balance accuracy score of 19 classifiers across 30 homogeneity image feature datasets for cell line U87-MG.



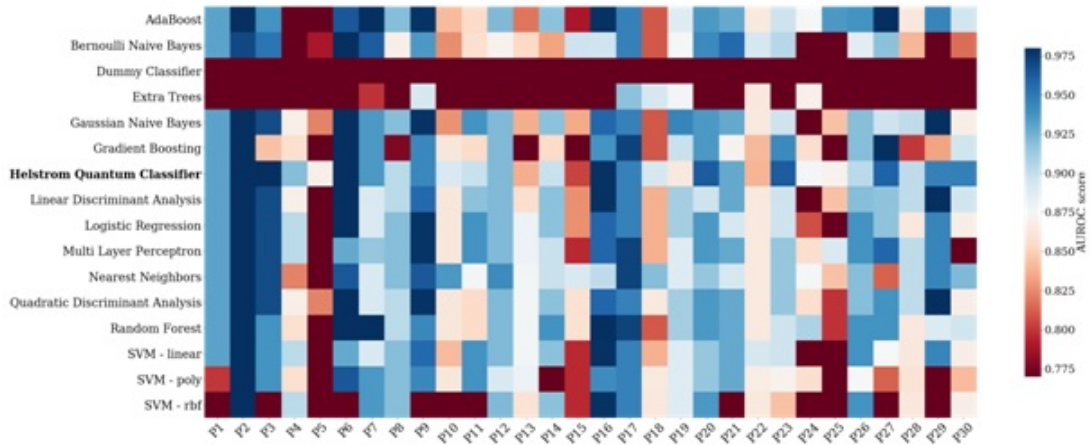


Fig. 2.2.3 | AUROC score of 19 classifiers across 30 homogeneity image feature datasets for cell line U87-MG.

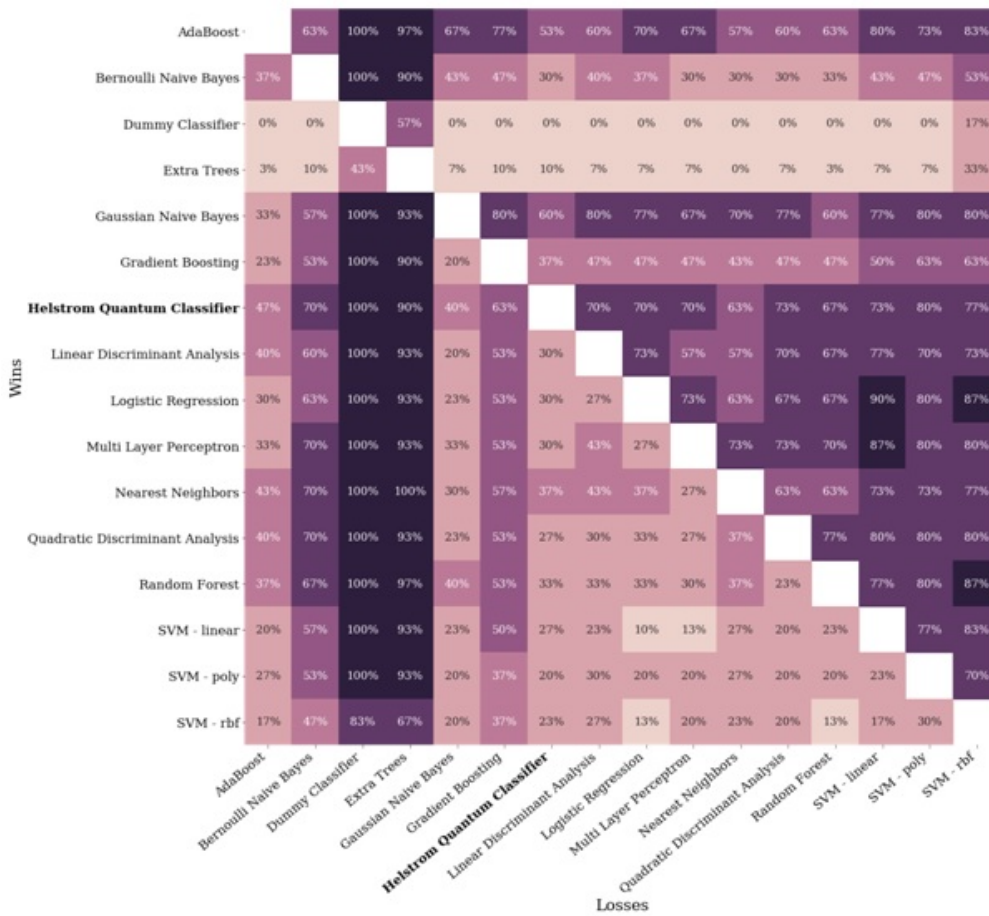


Fig. 2.2.4 | Percentage of datasets where model A (“Wins”) outperformed model B (“Losses”) out of 30 homogeneity image feature datasets for cell line U87-MG (AUROC score).

Classifiers	Image features					
	RGB	L*u*v*	Contrast	Correlation	Energy	Homogeneity
AdaBoost	0.776 ± 0.117	0.779 ± 0.110	0.864 ± 0.060	0.704 ± 0.105	0.873 ± 0.082	0.898 ± 0.077
Bernoulli Naive Bayes	0.765 ± 0.095	0.788 ± 0.084	0.841 ± 0.070	0.629 ± 0.088	0.771 ± 0.126	0.869 ± 0.082
Dummy Classifier	0.504 ± 0.050	0.504 ± 0.056	0.512 ± 0.055	0.480 ± 0.076	0.505 ± 0.055	0.516 ± 0.058
Extra Trees	0.577 ± 0.098	0.583 ± 0.120	0.529 ± 0.078	0.515 ± 0.043	0.525 ± 0.068	0.613 ± 0.156
Gaussian Naive Bayes	0.734 ± 0.089	0.752 ± 0.096	0.780 ± 0.086	0.674 ± 0.074	0.890 ± 0.065	0.905 ± 0.059
Gradient Boosting	0.815 ± 0.091	0.827 ± 0.085	0.877 ± 0.064	0.758 ± 0.095	0.879 ± 0.075	0.871 ± 0.085
<b>Helstrom Quantum Classifier</b>	0.791 ± 0.099	0.792 ± 0.090	0.838 ± 0.066	0.768 ± 0.075	0.902 ± 0.057	<b>0.917 ± 0.048</b>
Linear Discriminant Analysis	0.752 ± 0.095	0.736 ± 0.085	0.740 ± 0.076	0.639 ± 0.086	0.842 ± 0.107	0.900 ± 0.065
Logistic Regression	0.770 ± 0.100	0.764 ± 0.102	0.805 ± 0.085	0.636 ± 0.086	0.877 ± 0.071	0.897 ± 0.070
Multi Layer Perceptron	0.792 ± 0.109	0.784 ± 0.115	0.840 ± 0.067	0.768 ± 0.079	0.877 ± 0.099	0.901 ± 0.070
Nearest Neighbors	0.793 ± 0.105	0.793 ± 0.090	0.847 ± 0.044	0.739 ± 0.086	0.884 ± 0.076	0.902 ± 0.054
Quadratic Discriminant Analysis	0.773 ± 0.100	0.750 ± 0.102	0.782 ± 0.100	0.667 ± 0.074	0.879 ± 0.081	0.908 ± 0.050
Random Forest	0.808 ± 0.106	0.803 ± 0.100	0.861 ± 0.059	0.763 ± 0.083	0.900 ± 0.070	0.902 ± 0.068
SVM - linear	0.764 ± 0.100	0.751 ± 0.113	0.798 ± 0.088	0.591 ± 0.103	0.891 ± 0.065	0.888 ± 0.076
SVM - poly	0.745 ± 0.119	0.746 ± 0.111	0.780 ± 0.084	0.720 ± 0.091	0.865 ± 0.095	0.864 ± 0.090
SVM - rbf	0.756 ± 0.123	0.741 ± 0.114	0.682 ± 0.157	0.575 ± 0.096	0.859 ± 0.130	0.770 ± 0.179

Results exclude Nearest Centroid, Passive Aggressive Classifier and Perceptron.

### S2.3. Cell line MCF7

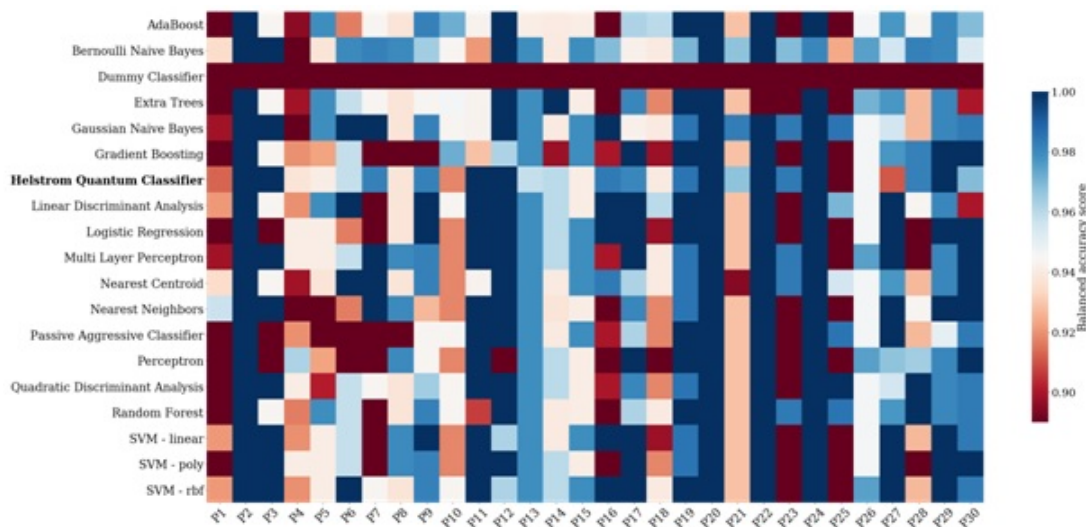


Fig. 2.3.1 | Balance accuracy score of 19 classifiers across 30 L\*u\*v\* image feature datasets for cell line MCF7.



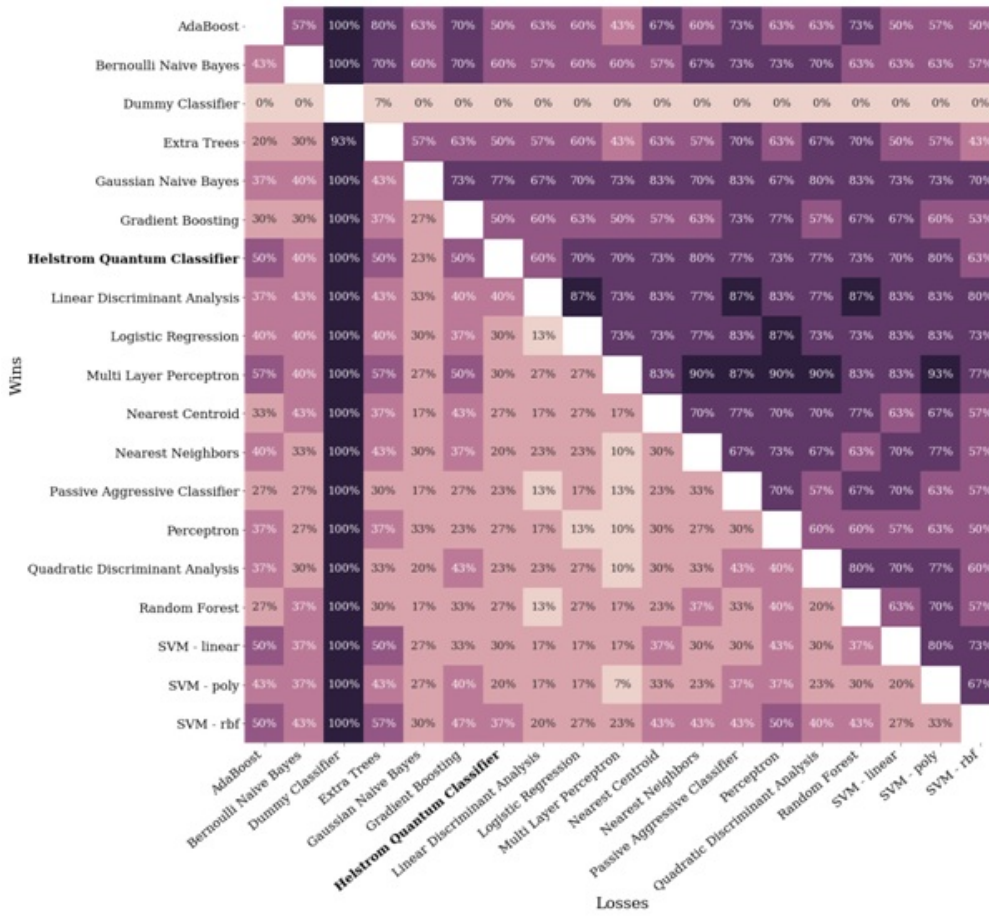


Fig. 2.3.2 | Percentage of datasets where model A (“Wins”) outperformed model B (“Losses”) out of 30 L\*u\*v\* image feature datasets for cell line MCF7 (balanced accuracy score).

Classifiers	Image features					
	RGB	L*u*v*	Contrast	Correlation	Energy	Homogeneity
AdaBoost	0.959 ± 0.046	0.940 ± 0.070	0.827 ± 0.106	0.699 ± 0.099	0.851 ± 0.111	0.875 ± 0.092
Bernoulli Naive Bayes	0.951 ± 0.048	0.964 ± 0.030	0.717 ± 0.124	0.661 ± 0.107	0.718 ± 0.133	0.837 ± 0.118
Dummy Classifier	0.505 ± 0.064	0.509 ± 0.041	0.494 ± 0.048	0.484 ± 0.044	0.500 ± 0.057	0.491 ± 0.044
Extra Trees	0.858 ± 0.150	0.914 ± 0.125	0.507 ± 0.037	0.526 ± 0.061	0.523 ± 0.078	0.585 ± 0.124
Gaussian Naive Bayes	0.943 ± 0.043	0.969 ± 0.034	0.641 ± 0.095	0.715 ± 0.091	0.886 ± 0.069	0.882 ± 0.078
Gradient Boosting	0.934 ± 0.080	0.939 ± 0.062	0.814 ± 0.103	0.706 ± 0.088	0.844 ± 0.093	0.866 ± 0.110
<b>Helstrom Quantum Classifier</b>	0.949 ± 0.052	<b>0.965 ± 0.033</b>	0.798 ± 0.109	0.775 ± 0.081	0.875 ± 0.059	0.892 ± 0.065
Linear Discriminant Analysis	0.947 ± 0.047	0.961 ± 0.047	0.626 ± 0.097	0.685 ± 0.098	0.781 ± 0.139	0.874 ± 0.095
Logistic Regression	0.946 ± 0.055	0.948 ± 0.060	0.638 ± 0.107	0.674 ± 0.095	0.866 ± 0.097	0.856 ± 0.096
Multi Layer Perceptron	0.946 ± 0.052	0.965 ± 0.042	0.760 ± 0.106	0.718 ± 0.089	0.867 ± 0.089	0.856 ± 0.111
Nearest Centroid	0.923 ± 0.056	0.964 ± 0.032	0.770 ± 0.117	0.739 ± 0.095	0.851 ± 0.068	0.890 ± 0.063
Nearest Neighbors	0.951 ± 0.056	0.947 ± 0.063	0.769 ± 0.109	0.759 ± 0.075	0.861 ± 0.075	0.854 ± 0.101
Passive Aggressive Classifier	0.945 ± 0.048	0.943 ± 0.051	0.690 ± 0.120	0.616 ± 0.145	0.787 ± 0.145	0.770 ± 0.154
Perceptron	0.938 ± 0.056	0.923 ± 0.094	0.652 ± 0.167	0.630 ± 0.145	0.774 ± 0.122	0.795 ± 0.149
Quadratic Discriminant Analysis	0.959 ± 0.050	0.961 ± 0.036	0.638 ± 0.096	0.710 ± 0.111	0.883 ± 0.072	0.872 ± 0.089
Random Forest	0.948 ± 0.064	0.956 ± 0.044	0.816 ± 0.112	0.740 ± 0.097	0.851 ± 0.113	0.860 ± 0.110
SVM - linear	0.955 ± 0.050	0.956 ± 0.052	0.654 ± 0.138	0.696 ± 0.107	0.868 ± 0.110	0.873 ± 0.108
SVM - poly	0.940 ± 0.054	0.949 ± 0.060	0.762 ± 0.108	0.730 ± 0.080	0.862 ± 0.078	0.874 ± 0.105
SVM - rbf	0.950 ± 0.053	0.962 ± 0.048	0.681 ± 0.134	0.697 ± 0.108	0.868 ± 0.112	0.877 ± 0.102

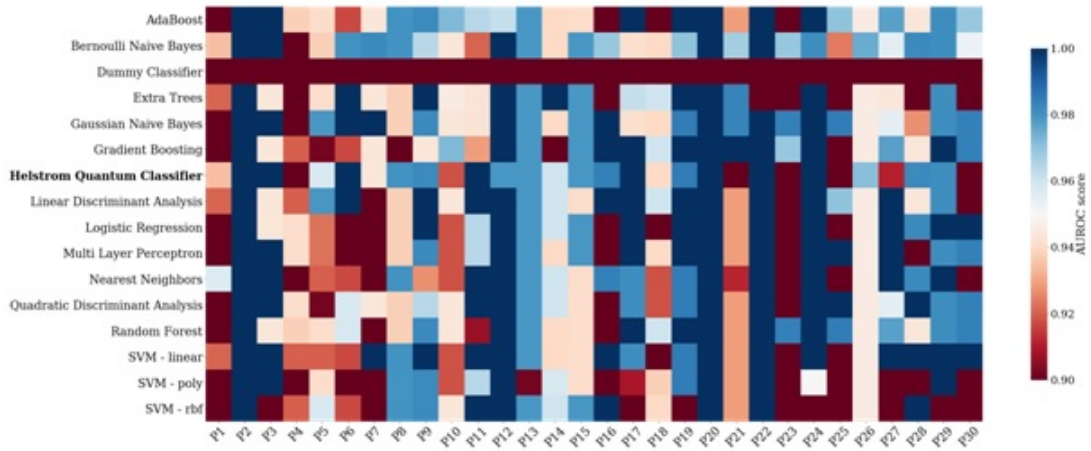


Fig. 2.3.3 | AUROC score of 19 classifiers across 30 L\*u\*v\* image feature datasets for cell line MCF7.

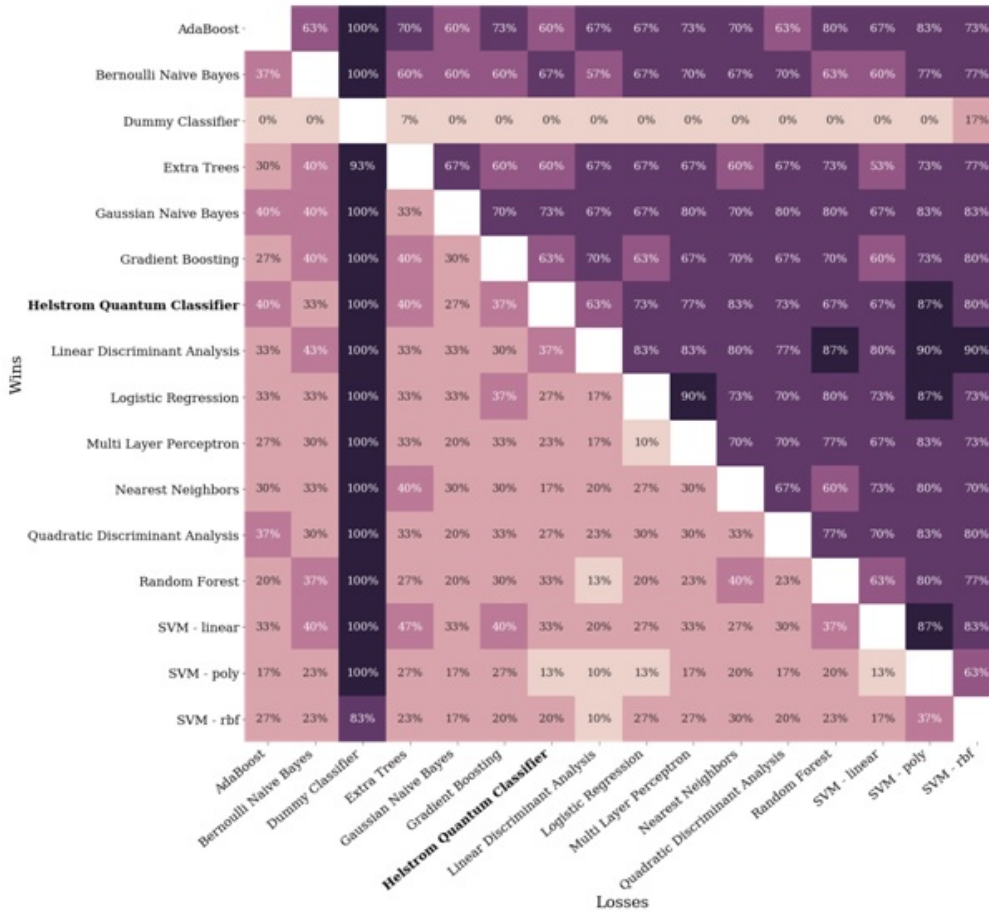


Fig. 2.3.4 | Percentage of datasets where model A (“Wins”) outperformed model B (“Losses”) out of 30 L\*u\*v\* image feature datasets for cell line MCF7 (AUROC score).

Classifiers	Image features					
	RGB	L*u*v*	Contrast	Correlation	Energy	Homogeneity
AdaBoost	0.944 ± 0.076	0.951 ± 0.058	0.821 ± 0.110	0.704 ± 0.107	0.845 ± 0.103	0.857 ± 0.102
Bernoulli Naive Bayes	0.951 ± 0.048	0.964 ± 0.030	0.717 ± 0.124	0.661 ± 0.107	0.718 ± 0.133	0.837 ± 0.118
Dummy Classifier	0.475 ± 0.066	0.506 ± 0.073	0.504 ± 0.041	0.506 ± 0.051	0.496 ± 0.069	0.493 ± 0.051
Extra Trees	0.856 ± 0.148	0.918 ± 0.129	0.505 ± 0.025	0.511 ± 0.038	0.509 ± 0.043	0.527 ± 0.068
Gaussian Naive Bayes	0.943 ± 0.043	0.969 ± 0.034	0.641 ± 0.095	0.715 ± 0.091	0.886 ± 0.069	0.882 ± 0.078
Gradient Boosting	0.911 ± 0.112	0.952 ± 0.060	0.773 ± 0.121	0.719 ± 0.091	0.820 ± 0.092	0.837 ± 0.117
<b>Helstrom Quantum Classifier</b>	0.955 ± 0.045	<b>0.960 ± 0.041</b>	0.753 ± 0.120	0.766 ± 0.090	0.866 ± 0.078	0.883 ± 0.071
Linear Discriminant Analysis	0.947 ± 0.047	0.961 ± 0.047	0.626 ± 0.096	0.685 ± 0.098	0.780 ± 0.141	0.874 ± 0.095
Logistic Regression	0.943 ± 0.062	0.950 ± 0.054	0.637 ± 0.108	0.672 ± 0.095	0.853 ± 0.115	0.848 ± 0.107
Multi Layer Perceptron	0.953 ± 0.056	0.950 ± 0.054	0.770 ± 0.120	0.700 ± 0.103	0.850 ± 0.082	0.869 ± 0.104
Nearest Neighbors	0.948 ± 0.045	0.947 ± 0.061	0.767 ± 0.115	0.751 ± 0.087	0.862 ± 0.080	0.850 ± 0.110
Quadratic Discriminant Analysis	0.959 ± 0.050	0.961 ± 0.036	0.638 ± 0.096	0.710 ± 0.111	0.883 ± 0.072	0.872 ± 0.089
Random Forest	0.949 ± 0.066	0.953 ± 0.057	0.811 ± 0.116	0.733 ± 0.100	0.838 ± 0.133	0.852 ± 0.104
SVM - linear	0.946 ± 0.054	0.960 ± 0.052	0.635 ± 0.135	0.687 ± 0.113	0.849 ± 0.126	0.858 ± 0.130
SVM - poly	0.887 ± 0.093	0.927 ± 0.063	0.713 ± 0.125	0.682 ± 0.113	0.841 ± 0.107	0.866 ± 0.121
SVM - rbf	0.872 ± 0.144	0.859 ± 0.176	0.586 ± 0.111	0.601 ± 0.121	0.839 ± 0.142	0.717 ± 0.200

Results exclude Nearest Centroid, Passive Aggressive Classifier and Perceptron.

#### 45 S2.4. Cell line U251

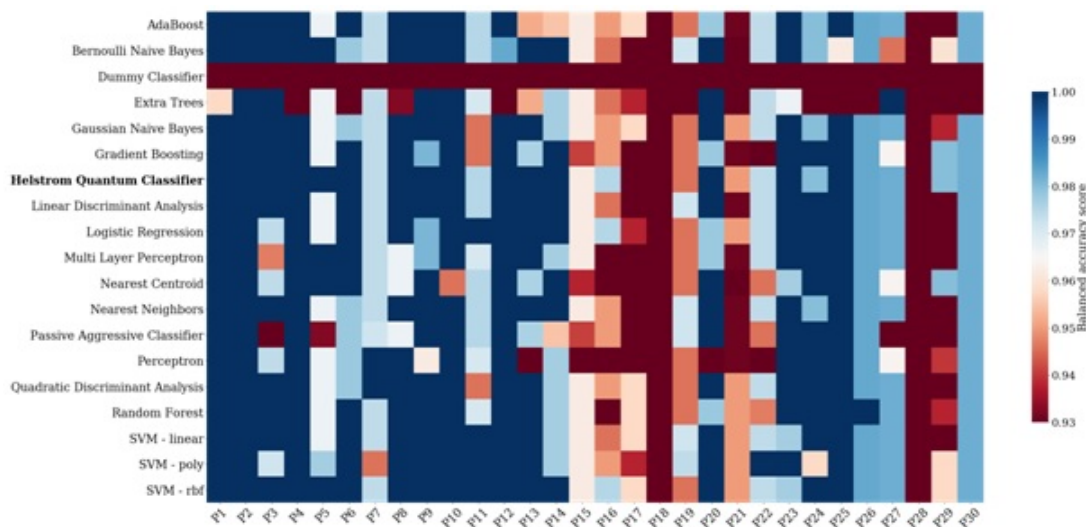


Fig. 2.4.1 | Balance accuracy score of 19 classifiers across 30 homogeneity image feature datasets for cell line U251.

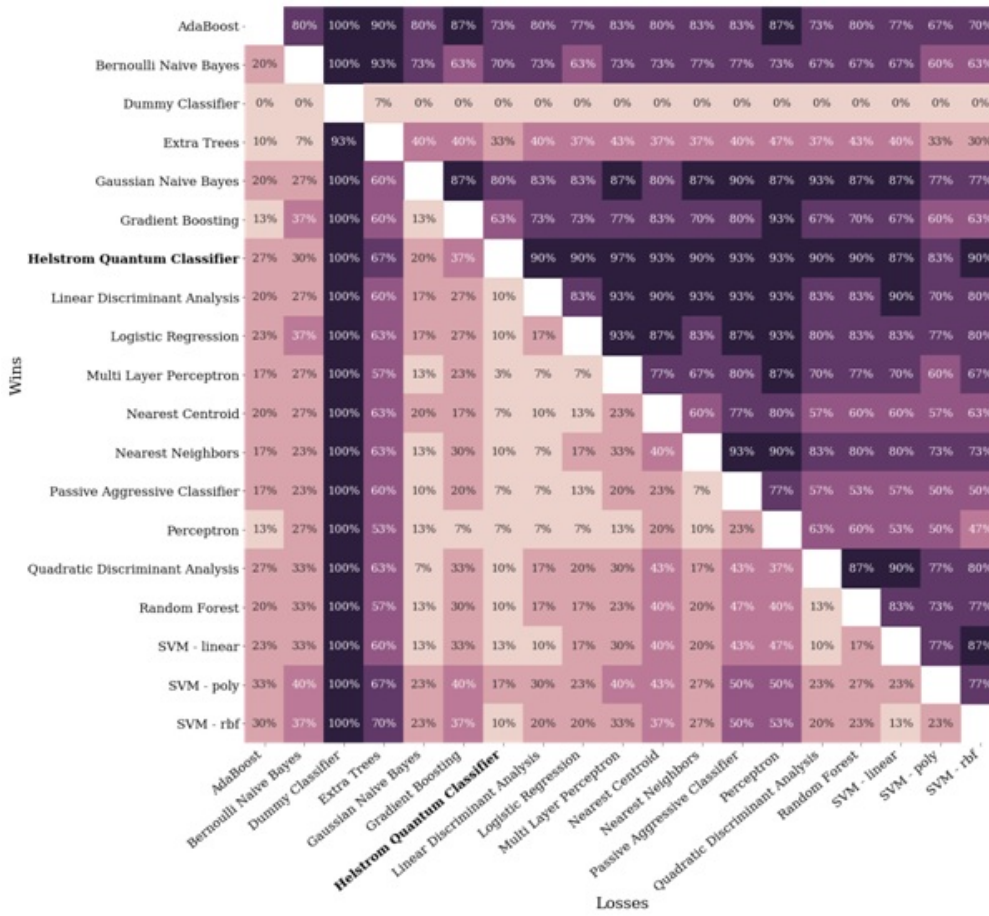


Fig. 2.4.2 | Percentage of datasets where model A (“Wins”) outperformed model B (“Losses”) out of 30 homogeneity image feature datasets for cell line U251 (balanced accuracy score).

Classifiers	Image features					
	RGB	L*u*v*	Contrast	Correlation	Energy	Homogeneity
AdaBoost	0.911 ± 0.049	0.909 ± 0.046	0.969 ± 0.027	0.771 ± 0.081	0.968 ± 0.045	0.972 ± 0.032
Bernoulli Naive Bayes	0.890 ± 0.047	0.903 ± 0.043	0.934 ± 0.055	0.649 ± 0.108	0.859 ± 0.118	0.967 ± 0.059
Dummy Classifier	0.504 ± 0.068	0.494 ± 0.058	0.495 ± 0.049	0.485 ± 0.057	0.495 ± 0.051	0.509 ± 0.069
Extra Trees	0.832 ± 0.126	0.865 ± 0.098	0.714 ± 0.188	0.571 ± 0.110	0.692 ± 0.206	0.867 ± 0.164
Gaussian Naive Bayes	0.908 ± 0.052	0.912 ± 0.051	0.969 ± 0.030	0.698 ± 0.084	0.964 ± 0.033	0.976 ± 0.027
Gradient Boosting	0.917 ± 0.049	0.907 ± 0.054	0.958 ± 0.034	0.845 ± 0.064	0.965 ± 0.039	0.970 ± 0.035
<b>Helstrom Quantum Classifier</b>	0.917 ± 0.045	0.907 ± 0.039	0.966 ± 0.032	0.792 ± 0.070	0.973 ± 0.024	<b>0.979 ± 0.029</b>
Linear Discriminant Analysis	0.892 ± 0.062	0.904 ± 0.047	0.916 ± 0.065	0.674 ± 0.087	0.935 ± 0.058	0.975 ± 0.036
Logistic Regression	0.911 ± 0.051	0.918 ± 0.044	0.965 ± 0.030	0.677 ± 0.087	0.964 ± 0.031	0.976 ± 0.031
Multi Layer Perceptron	0.915 ± 0.045	0.924 ± 0.046	0.965 ± 0.035	0.837 ± 0.087	0.968 ± 0.034	0.969 ± 0.035
Nearest Centroid	0.892 ± 0.051	0.892 ± 0.048	0.927 ± 0.033	0.694 ± 0.091	0.923 ± 0.048	0.967 ± 0.038
Nearest Neighbors	0.914 ± 0.046	0.907 ± 0.050	0.969 ± 0.032	0.836 ± 0.078	0.974 ± 0.036	0.974 ± 0.033
Passive Aggressive Classifier	0.875 ± 0.070	0.881 ± 0.070	0.951 ± 0.050	0.575 ± 0.111	0.958 ± 0.047	0.959 ± 0.046
Perceptron	0.902 ± 0.054	0.889 ± 0.060	0.945 ± 0.060	0.586 ± 0.117	0.951 ± 0.039	0.952 ± 0.056
Quadratic Discriminant Analysis	0.923 ± 0.053	0.924 ± 0.051	0.972 ± 0.028	0.712 ± 0.085	0.969 ± 0.033	0.976 ± 0.028
Random Forest	0.926 ± 0.048	0.918 ± 0.047	0.973 ± 0.033	0.815 ± 0.086	0.970 ± 0.044	0.974 ± 0.034
SVM - linear	0.916 ± 0.047	0.914 ± 0.051	0.970 ± 0.027	0.656 ± 0.103	0.975 ± 0.028	0.976 ± 0.035
SVM - poly	0.925 ± 0.052	0.911 ± 0.064	0.972 ± 0.027	0.821 ± 0.072	0.970 ± 0.031	0.978 ± 0.025
SVM - rbf	0.913 ± 0.051	0.917 ± 0.053	0.969 ± 0.032	0.667 ± 0.100	0.976 ± 0.027	0.980 ± 0.033

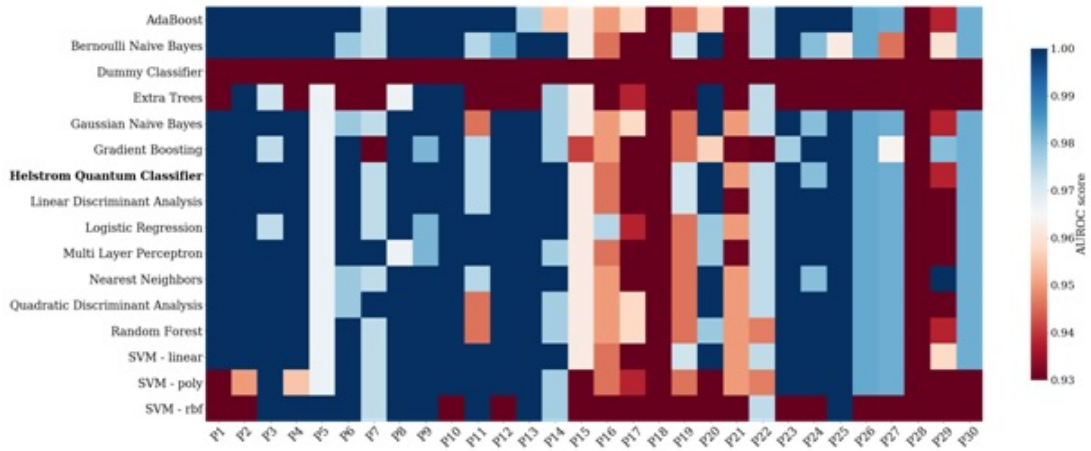


Fig. 2.4.3 | AUROC score of 19 classifiers across 30 homogeneity image feature datasets for cell line U251.

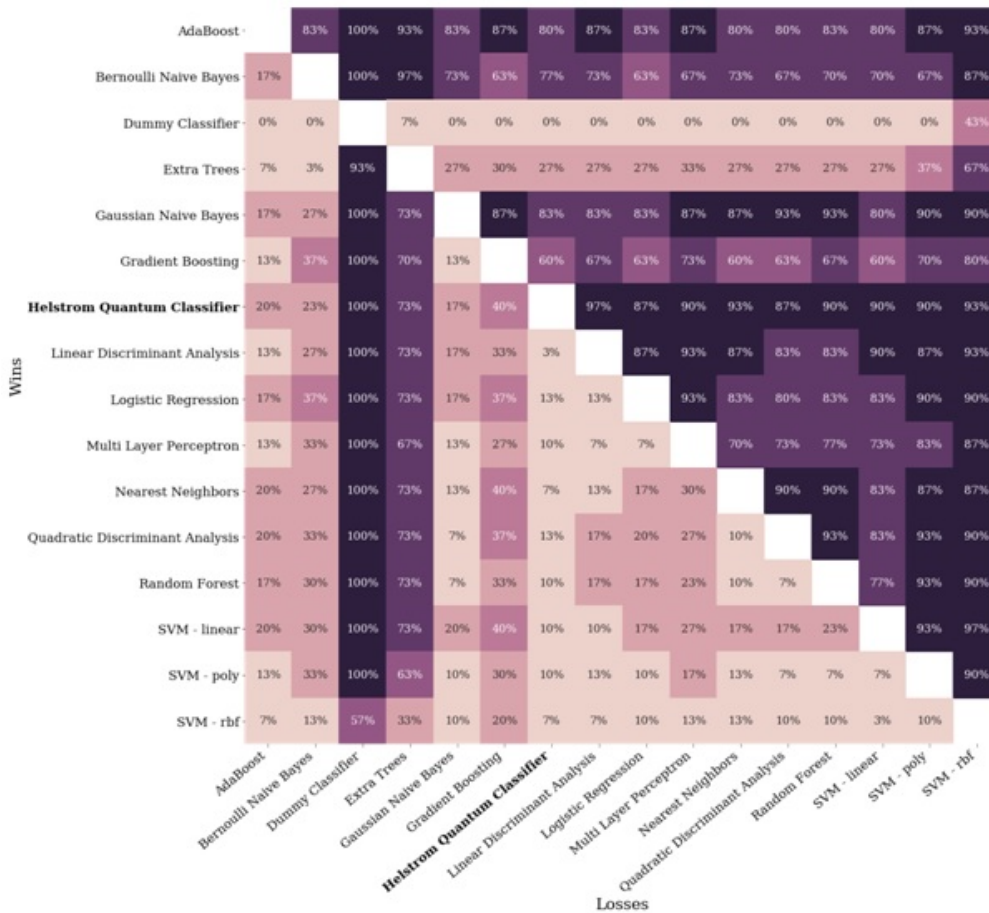


Fig. 2.4.4 | Percentage of datasets where model A (“Wins”) outperformed model B (“Losses”) out of 30 homogeneity image feature datasets for cell line U251 (AUROC score).

**Table 2.4.2 | The mean and standard deviation AUROC score (with respect to 30 datasets) for cell line U251**

Classifiers	Image features					
	RGB	L*u*v*	Contrast	Correlation	Energy	Homogeneity
AdaBoost	0.920 ± 0.046	0.911 ± 0.048	0.971 ± 0.027	0.741 ± 0.099	0.967 ± 0.045	0.975 ± 0.034
Bernoulli Naive Bayes	0.890 ± 0.047	0.903 ± 0.043	0.934 ± 0.055	0.649 ± 0.108	0.859 ± 0.118	0.967 ± 0.059
Dummy Classifier	0.500 ± 0.050	0.492 ± 0.049	0.499 ± 0.059	0.502 ± 0.054	0.503 ± 0.055	0.511 ± 0.055
Extra Trees	0.808 ± 0.152	0.843 ± 0.110	0.653 ± 0.177	0.558 ± 0.098	0.673 ± 0.186	0.838 ± 0.155
Gaussian Naive Bayes	0.908 ± 0.052	0.912 ± 0.051	0.969 ± 0.030	0.698 ± 0.084	0.964 ± 0.033	0.976 ± 0.027
Gradient Boosting	0.924 ± 0.052	0.906 ± 0.046	0.964 ± 0.031	0.830 ± 0.060	0.966 ± 0.040	0.968 ± 0.034
<b>Helstrom Quantum Classifier</b>	0.899 ± 0.057	0.907 ± 0.053	0.948 ± 0.063	0.797 ± 0.093	0.966 ± 0.031	<b>0.978 ± 0.027</b>
Linear Discriminant Analysis	0.892 ± 0.062	0.904 ± 0.047	0.916 ± 0.065	0.674 ± 0.087	0.935 ± 0.058	0.975 ± 0.036
Logistic Regression	0.911 ± 0.050	0.917 ± 0.046	0.962 ± 0.030	0.676 ± 0.088	0.962 ± 0.036	0.975 ± 0.034
Multi Layer Perceptron	0.921 ± 0.046	0.918 ± 0.051	0.965 ± 0.034	0.830 ± 0.093	0.962 ± 0.034	0.971 ± 0.040
Nearest Neighbors	0.914 ± 0.051	0.907 ± 0.050	0.964 ± 0.040	0.823 ± 0.062	0.972 ± 0.036	0.978 ± 0.028
Quadratic Discriminant Analysis	0.923 ± 0.053	0.924 ± 0.051	0.972 ± 0.028	0.712 ± 0.085	0.969 ± 0.033	0.976 ± 0.028
Random Forest	0.921 ± 0.050	0.925 ± 0.049	0.975 ± 0.025	0.836 ± 0.076	0.973 ± 0.038	0.972 ± 0.035
SVM - linear	0.916 ± 0.049	0.909 ± 0.047	0.962 ± 0.030	0.651 ± 0.103	0.971 ± 0.030	0.978 ± 0.035
SVM - poly	0.897 ± 0.063	0.901 ± 0.062	0.952 ± 0.061	0.801 ± 0.089	0.969 ± 0.030	0.955 ± 0.051
SVM - rbf	0.849 ± 0.150	0.833 ± 0.156	0.776 ± 0.221	0.634 ± 0.114	0.871 ± 0.189	0.723 ± 0.240

Results exclude Nearest Centroid, Passive Aggressive Classifier and Perceptron.

### S3. Performance of HQC on a new unseen test set (for the best image feature for each cell line)

In this sub-experiment the trained HQC model was tested on a new unseen test set extracted from the remaining 99.8% of the datasets. This experiment was done by randomly selecting 10 datasets (out of the 30 datasets) from the best performing image feature for each of the four cell lines. We show a comparison of the performance on this new unseen test set against the performance on the test set from the 0.2% random sample used in the main experiment.

**Table 3.1 | Balance accuracy and AUROC score for HQC on the test set used in the experiment and a new unseen test set for 10 randomly selected homogeneity image feature datasets for cell line MDA-MD-231**

Datasets	Balanced accuracy	Balanced accuracy	AUROC	AUROC
	Test set used in the experiment	New unseen test set	Test set used in the experiment	New unseen test set
P5	0.900	0.943	0.900	0.943
P6	0.977	0.921	0.977	0.921
P7	0.916	0.897	0.946	0.896
P8	0.944	0.937	0.911	0.943
P14	0.971	0.935	1.000	0.933
P15	0.921	0.869	0.947	0.888
P23	0.955	0.968	0.955	0.967
P26	0.977	0.947	0.955	0.944
P27	0.974	0.953	0.946	0.951
P30	0.971	0.951	0.853	0.811
Mean	0.951	0.932	0.939	0.920

**Table 3.2 | Balance accuracy and AUROC score for HQC on the test set used in the experiment and a new unseen test set for 10 randomly selected homogeneity image feature datasets for cell line U87-MG**

Datasets	Balanced accuracy	Balanced accuracy	AUROC	AUROC
	Test set used in the experiment	New unseen test set	Test set used in the experiment	New unseen test set
P1	0.931	0.947	0.931	0.947
P2	0.985	0.953	0.985	0.954
P5	0.869	0.908	0.869	0.908
P9	0.943	0.928	0.943	0.940
P10	0.891	0.894	0.891	0.897
P14	0.898	0.868	0.898	0.870
P21	0.964	0.910	0.929	0.901
P22	0.867	0.868	0.839	0.870
P25	0.871	0.893	0.871	0.893
P27	0.980	0.869	0.960	0.871
Mean	0.920	0.904	0.912	0.905

**Table 3.3 | Balance accuracy and AUROC score for HQC on the test set used in the experiment and a new unseen test set for 10 randomly selected L\*u\*v\* image feature datasets for cell line MCF7**

Datasets	Balanced accuracy	Balanced accuracy	AUROC	AUROC
	Test set used in the experiment	New unseen test set	Test set used in the experiment	New unseen test set
P6	0.958	0.955	1.000	0.958
P8	0.938	0.949	0.980	0.951
P9	0.982	0.958	0.982	0.955
P13	0.958	0.966	0.979	0.970
P14	0.960	0.978	0.960	0.978
P16	0.984	0.959	0.984	0.951
P19	0.985	0.977	0.985	0.976
P23	0.984	0.967	0.884	0.947
P25	0.875	0.867	0.879	0.886
P26	0.946	0.980	0.971	0.946
Mean	0.957	0.956	0.960	0.952

**Table 3.4 | Balance accuracy and AUROC score for HQC on the test set used in the experiment and a new unseen test set for 10 randomly selected homogeneity image feature datasets for cell line U251**

Datasets	Balanced accuracy	Balanced accuracy	AUROC	AUROC
	Test set used in the experiment	New unseen test set	Test set used in the experiment	New unseen test set
P2	1.000	0.985	1.000	0.977
P7	0.974	0.962	0.974	0.962
P15	0.962	0.956	0.962	0.898
P18	0.927	0.971	0.927	0.971
P20	1.000	0.960	1.000	0.959
P21	0.950	0.964	0.950	0.964
P22	0.974	0.941	0.974	0.948
P27	0.982	0.965	0.982	0.973
P29	0.980	0.926	0.938	0.932
P30	0.982	0.976	0.982	0.968
Mean	0.973	0.961	0.969	0.955

# S4. Experimental results - extended version

This section contains tables showing the experimental results for the balanced accuracy and AUROC scores for the six image features *RGB*, *L\*u\*v\**, *contrast*, *correlation*, *energy* and *homogeneity*, for each of the four cell lines MDA-MD-231, U87-MG, MCF7 and U251 respectively.

**Table 4.1 | Balanced accuracy score for 30 RGB image feature datasets for cell line MDA-MD-231**

Classifiers	P1	P2	P3	P4	P5	P6	P7	P8	P9	P10	P11	P12	P13	P14	P15	P16	P17	P18	P19	P20	P21	P22	P23	P24	P25	P26	P27	P28	P29	P30
AdaBoost	0.734	0.789	0.875	0.943	0.890	0.804	0.895	0.796	0.866	0.898	0.846	0.967	0.795	0.906	0.757	0.769	0.898	0.941	0.849	0.921	0.864	0.921	0.978	0.835	0.952	0.871	0.947	0.913	0.918	0.895
Bernoulli Naïve Bayes	0.904	0.866	0.844	0.907	0.906	0.804	0.784	0.750	0.701	0.712	0.918	0.944	0.692	0.859	0.741	0.824	0.932	0.878	0.842	0.941	0.727	0.899	0.938	0.881	0.871	0.946	0.863	0.918	0.946	
Decision Tree	0.732	0.756	0.755	0.897	0.889	0.850	0.850	0.850	0.850	0.850	0.850	0.850	0.850	0.850	0.850	0.850	0.850	0.850	0.850	0.850	0.850	0.850	0.850	0.850	0.850	0.850	0.850	0.850	0.850	
Extra Trees	0.855	0.577	0.500	0.835	0.917	0.742	0.938	0.938	0.938	0.938	0.938	0.938	0.938	0.938	0.938	0.938	0.938	0.938	0.938	0.938	0.938	0.938	0.938	0.938	0.938	0.938	0.938	0.938	0.938	
Gaussian Naïve Bayes	0.795	0.816	0.804	0.897	0.917	0.742	0.850	0.938	0.821	0.789	0.871	0.955	0.817	0.742	0.938	0.946	0.904	0.946	0.904	0.890	0.947	0.807	0.913	0.851	0.899	0.960	0.871	0.846	0.846	
Gradient Boosting	0.917	0.763	0.685	0.851	0.890	0.753	0.818	0.969	0.885	0.882	0.705	0.925	0.896	0.792	0.938	0.971	0.897	0.858	0.866	0.943	0.100	0.890	0.947	0.863	0.931	0.851	0.899	0.960	0.871	
Holdout Quantum Classifier	0.918	0.816	0.820	0.866	0.899	0.841	0.914	0.938	0.864	0.851	0.893	0.885	0.897	0.925	0.734	0.880	0.921	0.825	0.950	0.921	0.779	0.935	0.889	0.835	0.963	0.812	0.943	0.980	0.841	
Linear Discriminant Analysis	0.837	0.857	0.734	0.842	0.917	0.764	0.866	0.938	0.871	0.819	0.789	0.889	0.891	0.878	0.887	0.971	0.885	0.688	0.850	0.831	0.825	0.835	0.825	0.835	0.756	0.881	0.812	0.899	0.940	0.866
Logistic Regression	0.857	0.920	0.817	0.890	0.917	0.798	0.882	0.938	0.871	0.819	0.789	0.889	0.891	0.878	0.887	0.971	0.885	0.688	0.850	0.831	0.825	0.835	0.825	0.835	0.756	0.881	0.812	0.899	0.940	0.866
Multi Layer Perceptron	0.857	0.820	0.817	0.890	0.917	0.798	0.882	0.938	0.871	0.819	0.789	0.889	0.891	0.878	0.887	0.971	0.885	0.688	0.850	0.831	0.825	0.835	0.825	0.835	0.756	0.881	0.812	0.899	0.940	0.866
Nearest Centroid	0.797	0.812	0.804	0.784	0.889	0.741	0.897	0.906	0.827	0.844	0.867	0.849	0.916	0.791	0.742	0.897	0.946	0.827	0.716	0.850	0.764	0.825	0.890	0.811	0.808	0.913	0.812	0.856	0.918	0.846
Nearest Neighbors	0.857	0.688	0.702	0.870	0.917	0.855	0.839	0.929	0.844	0.867	0.849	0.909	0.921	0.814	0.834	0.946	0.827	0.716	0.850	0.764	0.825	0.890	0.811	0.808	0.913	0.812	0.856	0.918	0.846	
Passive Aggressive Classifier	0.795	0.734	0.888	0.914	0.863	0.508	0.890	0.890	0.762	0.411	0.732	0.685	0.878	0.744	0.778	0.918	0.946	0.885	0.750	0.770	0.979	0.916	0.833	0.890	0.963	0.827	0.834	1.000	0.959	
Perceptron	0.687	0.647	0.890	0.882	0.911	0.655	0.750	0.897	0.821	0.621	0.825	0.684	0.684	0.562	0.562	0.734	0.778	0.694	0.814	0.941	0.753	0.861	0.756	0.881	0.975	0.871	0.980	0.891	0.871	
Quadratic Discriminant Analysis	0.845	0.855	0.855	0.855	0.855	0.855	0.855	0.855	0.855	0.855	0.855	0.855	0.855	0.855	0.855	0.855	0.855	0.855	0.855	0.855	0.855	0.855	0.855	0.855	0.855	0.855	0.855	0.855	0.855	
Random Forest	0.875	0.798	0.614	0.914	0.917	0.798	0.669	0.838	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	
SVM - linear	0.857	0.788	0.884	0.914	0.917	0.798	0.669	0.838	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	
SVM - poly	0.835	0.784	0.868	0.859	0.917	0.798	0.669	0.838	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	
SVM - rbf	0.857	0.857	0.884	0.914	0.889	0.753	0.859	0.938	0.857	0.757	0.788	0.842	0.935	0.925	0.778	0.938	0.971	0.885	0.795	0.871	0.835	0.962	0.890	0.944	0.947	0.890	0.935	0.960	0.896	

**Table 4.2 | Balanced accuracy score for 30 L\*u\*v\* image feature datasets for cell line MDA-MD-231**

Classifiers	P1	P2	P3	P4	P5	P6	P7	P8	P9	P10	P11	P12	P13	P14	P15	P16	P17	P18	P19	P20	P21	P22	P23	P24	P25	P26	P27	P28	P29	P30
AdaBoost	0.734	0.789	0.875	0.943	0.890	0.804	0.895	0.796	0.866	0.898	0.846	0.967	0.795	0.906	0.757	0.769	0.898	0.941	0.849	0.921	0.864	0.921	0.978	0.835	0.952	0.871	0.947	0.913	0.918	0.895
Bernoulli Naïve Bayes	0.904	0.866	0.844	0.907	0.906	0.804	0.784	0.750	0.701	0.712	0.918	0.944	0.692	0.859	0.741	0.824	0.932	0.878	0.842	0.941	0.727	0.899	0.938	0.881	0.871	0.946	0.863	0.918	0.946	
Decision Tree	0.732	0.756	0.755	0.897	0.889	0.850	0.850	0.850	0.850	0.850	0.850	0.850	0.850	0.850	0.850	0.850	0.850	0.850	0.850	0.850	0.850	0.850	0.850	0.850	0.850	0.850	0.850	0.850	0.850	
Extra Trees	0.855	0.577	0.500	0.835	0.917	0.742	0.938	0.938	0.938	0.938	0.938	0.938	0.938	0.938	0.938	0.938	0.938	0.938	0.938	0.938	0.938	0.938	0.938	0.938	0.938	0.938	0.938	0.938	0.938	
Gaussian Naïve Bayes	0.795	0.816	0.804	0.897	0.917	0.742	0.850	0.938	0.821	0.789	0.871	0.955	0.817	0.742	0.938	0.946	0.904	0.946	0.904	0.890	0.947	0.807	0.913	0.851	0.899	0.960	0.871	0.846	0.846	
Gradient Boosting	0.917	0.763	0.685	0.851	0.890	0.753	0.818	0.969	0.885	0.882	0.705	0.925	0.896	0.792	0.938	0.971	0.897	0.858	0.866	0.943	0.100	0.890	0.947	0.863	0.931	0.851	0.899	0.960	0.871	
Holdout Quantum Classifier	0.918	0.816	0.820	0.866	0.899	0.841	0.914	0.938	0.864	0.851	0.893	0.885	0.897	0.925	0.734	0.880	0.921	0.825	0.950	0.921	0.779	0.935	0.889	0.835	0.963	0.812	0.943	0.980	0.841	
Linear Discriminant Analysis	0.837	0.857	0.734	0.842	0.917	0.764	0.866	0.938	0.871	0.819	0.789	0.889	0.891	0.878	0.887	0.971	0.885	0.688	0.850	0.831	0.825	0.835	0.825	0.835	0.756	0.881	0.812	0.899	0.940	0.866
Logistic Regression	0.857	0.920	0.817	0.890	0.917	0.798	0.882	0.938	0.871	0.819	0.789	0.889	0.891	0.878	0.887	0.971	0.885	0.688	0.850	0.831	0.825	0.835	0.825	0.835	0.756	0.881	0.812	0.899	0.940	0.866
Multi Layer Perceptron	0.857	0.820	0.817	0.890	0.917	0.798	0.882	0.938	0.871	0.819	0.789	0.889	0.891	0.878	0.887	0.971	0.885	0.688	0.850	0.831	0.825	0.835	0.825	0.835	0.756	0.881	0.812	0.899	0.940	0.866
Nearest Centroid	0.797	0.812	0.804	0.784	0.889	0.741	0.897	0.906	0.827	0.844	0.867	0.849	0.916	0.791	0.742	0.897	0.946	0.827	0.716	0.850	0.764	0.825	0.890	0.811	0.808	0.913	0.812	0.856	0.918	0.846
Nearest Neighbors	0.857	0.688	0.702	0.870	0.917	0.855	0.839	0.929	0.844	0.867	0.849	0.909	0.921	0.814	0.834	0.946	0.827	0.716	0.850	0.764	0.825	0.890	0.811	0.808	0.913	0.812	0.856	0.918	0.846	
Passive Aggressive Classifier	0.795	0.734	0.888	0.914	0.863	0.508	0.890	0.890	0.762	0.411	0.732	0.685	0.878	0.744	0.778	0.918	0.946	0.885	0.750	0.770	0.979	0.916	0.833	0.890	0.963	0.827	0.834	1.000	0.959	
Perceptron	0.687	0.647	0.890	0.882	0.911	0.655	0.750	0.897	0.821	0.621	0.825	0.684	0.684	0.562	0.562	0.734	0.778	0.694	0.814	0.941	0.753	0.861	0.756	0.881	0.975	0.871	0.980	0.891	0.871	
Quadratic Discriminant Analysis	0.845	0.855	0.855	0.855	0.855	0.855	0.855	0.855	0.855	0.855	0.855	0.855	0.855	0.855	0.855	0.855	0.855	0.855	0.855	0.855	0.855	0.855	0.855	0.855	0.855	0.855	0.855	0.855	0.855	
Random Forest	0.875	0.798	0.614	0.914	0.917	0.798	0.669	0.838	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	
SVM - linear	0.857	0.788	0.884	0.914	0.917	0.798	0.669	0.838	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	
SVM - poly	0.835	0.784	0.868	0.859	0.917	0.798	0.669	0.838	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	0.851	
SVM - rbf	0.857	0.857	0.884	0.914	0.889	0.753	0.859	0.938	0.857																					





**Table 4.10 | AUROC score for 30 contrast image feature datasets for cell line MDA-MB-231**

Classifiers	P1	P2	P3	P4	P5	P6	P7	P8	P9	P10	P11	P12	P13	P14	P15	P16	P17	P18	P19	P20	P21	P22	P23	P24	P25	P26	P27	P28	P29	P30	
AdaBoost	0.835	0.823	0.879	0.918	0.866	0.744	0.829	0.820	0.853	0.943	0.881	0.946	0.883	0.890	0.820	0.863	0.864	0.897	0.691	0.833	1.000	0.835	0.770	0.750	0.598	0.745	0.863	0.826	0.731	0.717	0.730
Bernoulli Naive Bayes	0.817	0.857	0.500	0.438	0.431	0.500	0.500	0.490	0.500	0.463	0.788	0.882	0.875	0.890	0.839	0.739	0.800	0.933	0.929	0.909	0.918	0.890	0.850	0.784	0.771	0.944	0.824	0.925	0.921	0.699	0.886
Dummy Classifier	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	
Extra Trees	0.817	0.886	0.862	0.918	0.789	0.782	0.881	0.888	0.875	0.821	0.914	0.975	0.890	0.839	0.818	0.800	0.933	0.945	0.969	0.909	0.974	1.000	0.921	0.899	0.866	0.977	0.857	0.941	0.850	1.000	0.955
Gaussian Naive Bayes	0.891	0.823	0.879	0.918	0.938	0.719	0.929	0.865	0.897	0.878	0.905	0.946	0.832	0.892	0.819	0.784	0.967	0.890	0.909	0.974	1.000	0.921	0.899	0.866	0.977	0.857	0.941	0.850	1.000	0.977	1.000
Gradient Boosting	0.911	0.765	0.912	0.944	0.820	0.816	0.881	0.911	0.906	0.929	0.952	0.950	0.836	0.839	0.861	0.784	0.833	0.952	0.909	0.972	0.972	0.896	0.920	0.821	0.967	0.857	0.941	0.850	0.945	1.000	0.977
Helium Quantum Classifier	0.800	0.795	0.643	0.917	0.695	0.807	0.802	0.855	0.847	0.857	0.882	0.900	0.865	0.861	0.800	0.867	0.842	0.772	0.972	0.762	0.875	0.724	0.771	0.967	0.794	0.950	0.859	0.965	0.909	0.909	
Linear Discriminant Analysis	0.873	0.838	0.895	0.890	0.789	0.715	0.887	0.888	0.878	0.907	0.888	0.925	0.921	0.890	0.839	0.800	0.945	0.818	0.972	0.895	0.950	0.843	0.821	0.977	0.824	0.975	0.929	1.000	0.955	1.000	0.955
Multi Layer Perceptron	0.891	0.886	0.845	0.890	0.839	0.715	0.897	0.888	0.858	0.964	0.929	0.921	0.890	0.839	0.800	0.945	0.818	0.972	0.895	0.950	0.920	0.843	0.821	0.977	0.824	0.975	0.929	1.000	0.955	1.000	0.955
Nearest-Neighbors	0.840	0.800	0.830	0.861	0.830	0.722	0.857	0.860	0.860	0.860	0.860	0.860	0.860	0.860	0.860	0.860	0.860	0.860	0.860	0.860	0.860	0.860	0.860	0.860	0.860	0.860	0.860	0.860	0.860	0.860	0.860
Nearest-Discriminant Analysis	0.817	0.868	0.839	0.890	0.839	0.715	0.897	0.888	0.858	0.964	0.929	0.921	0.890	0.839	0.800	0.945	0.818	0.972	0.895	0.950	0.920	0.843	0.821	0.977	0.824	0.975	0.929	1.000	0.955	1.000	0.955
Random Forest	0.855	0.823	0.895	0.918	0.883	0.759	0.927	0.857	0.911	0.869	0.910	0.869	0.885	0.865	0.800	0.833	0.945	0.909	0.974	0.972	0.950	0.941	0.896	0.973	0.853	0.944	0.881	0.900	0.881	1.000	0.977
SVM - linear	0.835	0.823	0.879	0.890	0.765	0.740	0.787	0.888	0.844	0.864	0.929	0.925	0.836	0.800	0.800	0.969	0.818	0.972	0.894	0.900	0.944	0.974	0.900	0.920	0.821	0.977	0.794	1.000	0.929	0.976	0.962
SVM - poly	0.835	0.823	0.845	0.917	0.906	0.807	0.914	0.888	0.838	0.943	0.952	0.950	0.836	0.800	0.800	0.969	0.818	0.972	0.894	0.900	0.944	0.974	0.900	0.920	0.821	0.977	0.794	1.000	0.929	0.976	0.962
SVM - rbf	0.817	0.823	0.879	0.917	0.796	0.749	0.787	0.888	0.800	0.943	0.952	0.975	0.890	0.839	0.784	0.500	0.500	0.945	0.773	0.500	0.918	0.900	0.920	0.864	0.846	0.977	0.794	1.000	0.929	1.000	0.955

Results include: Nearest Centroid, Passive Aggressive Classifier and Perceptron.

**Table 4.11 | AUROC score for 30 correlation image feature datasets for cell line MDA-MB-231**

Classifiers	P1	P2	P3	P4	P5	P6	P7	P8	P9	P10	P11	P12	P13	P14	P15	P16	P17	P18	P19	P20	P21	P22	P23	P24	P25	P26	P27	P28	P29	P30
AdaBoost	0.757	0.583	0.535	0.728	0.801	0.466	0.839	0.589	0.853	0.757	0.565	0.890	0.820	0.863	0.466	0.664	0.897	0.691	0.833	1.000	0.835	0.770	0.750	0.598	0.745	0.863	0.826	0.731	0.717	0.730
Bernoulli Naive Bayes	0.683	0.583	0.500	0.623	0.500	0.467	0.781	0.455	0.502	0.708	0.719	0.386	0.734	0.756	0.421	0.500	0.740	0.784	0.595	0.803	0.425	0.503	0.713	0.674	0.721	0.776	0.818	0.458	0.656	0.750
Dummy Classifier	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500
Extra Trees	0.840	0.800	0.830	0.861	0.830	0.722	0.857	0.860	0.860	0.860	0.860	0.860	0.860	0.860	0.860	0.860	0.860	0.860	0.860	0.860	0.860	0.860	0.860	0.860	0.860	0.860	0.860	0.860	0.860	0.860
Gaussian Naive Bayes	0.855	0.628	0.519	0.646	0.804	0.756	0.867	0.945	0.909	0.892	0.831	0.851	0.789	0.830	0.659	0.719	0.976	0.877	0.840	0.975	0.918	0.756	0.730	0.917	0.848	0.881	0.874	0.958	0.852	0.946
Gradient Boosting	0.833	0.483	0.500	0.652	0.566	0.615	0.754	0.538	0.500	0.681	0.669	0.650	0.882	0.677	0.620	0.867	0.974	0.894	0.816	0.946	0.753	0.741	0.820	0.765	0.795	0.844	0.881	0.819	0.914	0.784
Helium Quantum Classifier	0.833	0.483	0.500	0.652	0.566	0.615	0.754	0.538	0.500	0.681	0.669	0.650	0.882	0.677	0.620	0.867	0.974	0.894	0.816	0.946	0.753	0.741	0.820	0.765	0.795	0.844	0.881	0.819	0.914	0.784
Linear Discriminant Analysis	0.833	0.483	0.500	0.652	0.566	0.615	0.754	0.538	0.500	0.681	0.669	0.650	0.882	0.677	0.620	0.867	0.974	0.894	0.816	0.946	0.753	0.741	0.820	0.765	0.795	0.844	0.881	0.819	0.914	0.784
Logistic Regression	0.763	0.633	0.785	0.838	0.745	0.747	0.784	0.881	0.500	0.709	0.855	0.871	0.820	0.838	0.739	0.790	0.741	0.913	0.861	1.000	0.940	0.733	0.806	0.677	0.839	0.844	0.850	0.917	0.906	0.807
Multi Layer Perceptron	0.781	0.620	0.683	0.787	0.784	0.747	0.728	0.850	0.835	0.785	0.757	0.844	0.882	0.792	0.620	0.845	0.914	0.913	0.875	0.971	0.568	0.669	0.710	0.788	0.816	0.763	0.850	0.784	0.906	0.732
Nearest Neighbors	0.681	0.502	0.583	0.816	0.673	0.512	0.822	0.692	0.742	0.811	0.690	0.787	0.882	0.713	0.540	0.736	0.882	0.848	0.787	0.971	0.568	0.669	0.710	0.788	0.816	0.763	0.850	0.784	0.906	0.732
Quadratic Discriminant Analysis	0.850	0.589	0.535	0.920	0.843	0.708	0.867	0.795	0.853	0.838	0.635	0.851	0.789	0.885	0.679	0.610	0.969	0.935	0.920	1.000	0.815	0.699	0.742	0.844	0.881	0.881	0.850	0.958	0.969	0.865
Random Forest	0.750	0.502	0.500	0.652	0.500	0.591	0.782	0.538	0.500	0.690	0.605	0.748	0.442	0.500	0.710	0.828	0.500	0.853	0.547	0.689	0.800	0.653	0.817	0.744	0.850	0.850	0.500	0.500	0.730	0.730
SVM - linear	0.676	0.489	0.852	0.810	0.826	0.500	0.727	0.818	0.817	0.785	0.594	0.871	0.851	0.877	0.718	0.681	0.976	0.834	0.896	0.891	0.795	0.776	0.792	0.855	0.899	0.763	0.850	0.917	0.851	0.782
SVM - poly	0.750	0.489	0.852	0.810	0.826	0.500	0.727	0.818	0.817	0.785	0.594	0.871	0.851	0.877	0.718	0.681	0.976	0.834	0.896	0.891	0.795	0.776	0.792	0.855	0.899	0.763	0.850	0.917	0.851	0.782
SVM - rbf	0.750	0.489	0.852	0.810	0.826	0.500	0.727	0.818	0.817	0.785	0.594	0.871	0.851	0.877	0.718	0.681	0.976	0.834	0.896	0.891	0.795	0.776	0.792	0.855	0.899	0.763	0.850	0.917	0.851	0.782

Results include: Nearest Centroid, Passive Aggressive Classifier and Perceptron.

**Table 4.12 | AUROC score for 30 homogeneity image feature datasets for cell line MDA-MB-231**

Classifiers	P1	P2	P3	P4	P5	P6	P7	P8	P9	P10	P11	P12	P13	P14	P15	P16	P17	P18	P19	P20	P21	P22	P23	P24	P25	P26	P27	P28	P29	P30	
AdaBoost	0.871	0.885	0.877	0.929	1.000	0.836	0.943	1.000	0.788	0.866	0.778	0.971	0.969	0.944	0.890	0.920	0.935	0.929	1.000	0.972	0.976	0.946	0.980	0.943	0.975	0.909	0.947	0.888	0.864	0.858	
Bernoulli Naive Bayes	0.944	0.846	0.967	0.813	0.833	0.861	0.848	0.781	0.725	0.500	0.857	0.950	0.930	0.500	0.544	0.532	0.250	0.363	0.500	0.500	0.535	0.500	0.547	0.500	0.638	0.500	0.440	0.627	0.500	0.500	
Dummy Classifier	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	
Extra Trees	0.833	0.500	0.500	0.543	0.799	0.806	0.856	0.740	0.500	0.388	0.907	0.971	0.897	0.877	0.887	0.900	0.909	0.944	0.500	0.864	0.500	0.896	0.875	0.679	0.604	1.000	0.890	0.974	0.932	1.000	0.938
Gaussian Naive Bayes	0.944	0.885	0.840	0.893	0.944	0.889	0.899	0.899	0.839	0.816	0.929	0.971	0.969	0.944	0.81																



**Table 4.17 | Balanced accuracy score for 30 energy image feature datasets for cell line U87-MG**

Classifiers	P1	P2	P3	P4	P5	P6	P7	P8	P9	P10	P11	P12	P13	P14	P15	P16	P17	P18	P19	P20	P21	P22	P23	P24	P25	P26	P27	P28	P29	P30	
AdaBoost	0.927	0.940	0.984	0.852	0.752	0.920	0.877	0.760	0.920	0.778	0.882	0.933	0.778	0.920	0.938	0.849	0.856	0.835	0.825	0.933	1.000	0.855	0.877	0.870	0.878	0.833	0.971	0.818	0.878	0.900	0.921
Bernoulli Naïve Bayes	0.762	0.898	0.953	0.820	0.500	0.828	0.886	0.651	0.846	0.842	0.863	0.820	0.748	0.886	0.910	0.863	0.930	0.825	0.790	0.852	0.798	0.870	0.830	0.900	0.950	0.985	0.710	0.833	0.684	0.841	
Decision Tree	0.500	0.542	0.500	0.500	0.442	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	
Extra Trees	0.944	0.940	0.984	0.852	0.752	0.920	0.877	0.760	0.920	0.778	0.882	0.933	0.778	0.920	0.938	0.849	0.856	0.835	0.825	0.933	1.000	0.855	0.877	0.870	0.878	0.833	0.971	0.818	0.878	0.900	0.921
Gradient Boosting	0.871	0.940	0.984	0.852	0.752	0.920	0.877	0.760	0.920	0.778	0.882	0.933	0.778	0.920	0.938	0.849	0.856	0.835	0.825	0.933	1.000	0.855	0.877	0.870	0.878	0.833	0.971	0.818	0.878	0.900	0.921
Heuristic Quantum Classifier	0.889	0.898	0.984	0.852	0.752	0.920	0.877	0.760	0.920	0.778	0.882	0.933	0.778	0.920	0.938	0.849	0.856	0.835	0.825	0.933	1.000	0.855	0.877	0.870	0.878	0.833	0.971	0.818	0.878	0.900	0.921
Linear Discriminant Analysis	0.927	0.948	0.984	0.852	0.752	0.920	0.877	0.760	0.920	0.778	0.882	0.933	0.778	0.920	0.938	0.849	0.856	0.835	0.825	0.933	1.000	0.855	0.877	0.870	0.878	0.833	0.971	0.818	0.878	0.900	0.921
Logistic Regression	0.944	0.948	0.984	0.852	0.752	0.920	0.877	0.760	0.920	0.778	0.882	0.933	0.778	0.920	0.938	0.849	0.856	0.835	0.825	0.933	1.000	0.855	0.877	0.870	0.878	0.833	0.971	0.818	0.878	0.900	0.921
Multi Layer Perceptron	0.946	0.940	0.922	0.871	0.818	0.851	0.935	0.819	0.920	0.842	0.863	0.855	0.835	0.817	0.881	0.880	0.800	0.790	0.925	0.808	0.988	0.897	0.813	0.944	0.822	0.971	0.844	0.822	0.984	0.941	
Nearest Neighbors	0.944	0.938	1.000	0.875	0.710	0.920	0.911	0.891	1.000	0.827	0.844	0.955	0.814	0.922	0.838	0.871	0.922	0.838	0.871	0.925	0.808	0.988	0.897	0.813	0.944	0.822	0.971	0.844	0.822	0.984	0.941
Passive Aggressive Classifier	0.944	0.940	0.922	0.871	0.818	0.851	0.935	0.819	0.920	0.842	0.863	0.855	0.835	0.817	0.881	0.880	0.800	0.790	0.925	0.808	0.988	0.897	0.813	0.944	0.822	0.971	0.844	0.822	0.984	0.941	
Random Forest	0.944	0.940	0.922	0.871	0.818	0.851	0.935	0.819	0.920	0.842	0.863	0.855	0.835	0.817	0.881	0.880	0.800	0.790	0.925	0.808	0.988	0.897	0.813	0.944	0.822	0.971	0.844	0.822	0.984	0.941	
Support Vector Machine	0.944	0.940	0.922	0.871	0.818	0.851	0.935	0.819	0.920	0.842	0.863	0.855	0.835	0.817	0.881	0.880	0.800	0.790	0.925	0.808	0.988	0.897	0.813	0.944	0.822	0.971	0.844	0.822	0.984	0.941	
Weighted Majority Classifier	0.864	0.940	0.922	0.871	0.818	0.851	0.935	0.819	0.920	0.842	0.863	0.855	0.835	0.817	0.881	0.880	0.800	0.790	0.925	0.808	0.988	0.897	0.813	0.944	0.822	0.971	0.844	0.822	0.984	0.941	
Quadratic Discriminant Analysis	0.889	0.898	0.984	0.852	0.752	0.920	0.877	0.760	0.920	0.778	0.882	0.933	0.778	0.920	0.938	0.849	0.856	0.835	0.825	0.933	1.000	0.855	0.877	0.870	0.878	0.833	0.971	0.818	0.878	0.900	0.921
Random Forest	0.944	0.940	0.922	0.871	0.818	0.851	0.935	0.819	0.920	0.842	0.863	0.855	0.835	0.817	0.881	0.880	0.800	0.790	0.925	0.808	0.988	0.897	0.813	0.944	0.822	0.971	0.844	0.822	0.984	0.941	
SVM - linear	0.927	0.960	0.984	0.852	0.752	0.920	0.877	0.760	0.920	0.778	0.882	0.933	0.778	0.920	0.938	0.849	0.856	0.835	0.825	0.933	1.000	0.855	0.877	0.870	0.878	0.833	0.971	0.818	0.878	0.900	0.921
SVM - poly	0.927	0.938	0.984	0.852	0.752	0.920	0.877	0.760	0.920	0.778	0.882	0.933	0.778	0.920	0.938	0.849	0.856	0.835	0.825	0.933	1.000	0.855	0.877	0.870	0.878	0.833	0.971	0.818	0.878	0.900	0.921
SVM - rbf	0.927	0.960	0.984	0.852	0.752	0.920	0.877	0.760	0.920	0.778	0.882	0.933	0.778	0.920	0.938	0.849	0.856	0.835	0.825	0.933	1.000	0.855	0.877	0.870	0.878	0.833	0.971	0.818	0.878	0.900	0.921

**Table 4.18 | Balanced accuracy score for 30 homogeneity image feature datasets for cell line U87-MG**

Classifiers	P1	P2	P3	P4	P5	P6	P7	P8	P9	P10	P11	P12	P13	P14	P15	P16	P17	P18	P19	P20	P21	P22	P23	P24	P25	P26	P27	P28	P29	P30
AdaBoost	0.931	0.970	0.969	0.858	0.653	1.000	0.881	0.920	0.778	0.842	0.855	0.921	0.837	0.878	0.859	0.979	0.946	0.811	0.876	0.955	0.929	0.865	0.897	0.876	0.799	0.938	1.000	0.864	1.000	0.895
Bernoulli Naïve Bayes	0.931	0.970	0.952	0.744	0.789	1.000	0.962	0.869	0.935	0.826	0.838	0.871	0.854	0.833	0.897	0.896	0.946	0.810	0.876	0.942	0.927	0.893	0.904	0.757	0.715	0.886	0.918	0.840	0.611	0.816
Decision Tree	0.500	0.533	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500
Extra Trees	0.931	0.970	0.952	0.744	0.789	1.000	0.962	0.869	0.935	0.826	0.838	0.871	0.854	0.833	0.897	0.896	0.946	0.810	0.876	0.942	0.927	0.893	0.904	0.757	0.715	0.886	0.918	0.840	0.611	0.816
Gradient Boosting	0.931	0.970	0.952	0.744	0.789	1.000	0.962	0.869	0.935	0.826	0.838	0.871	0.854	0.833	0.897	0.896	0.946	0.810	0.876	0.942	0.927	0.893	0.904	0.757	0.715	0.886	0.918	0.840	0.611	0.816
Heuristic Quantum Classifier	0.931	0.970	0.952	0.744	0.789	1.000	0.962	0.869	0.935	0.826	0.838	0.871	0.854	0.833	0.897	0.896	0.946	0.810	0.876	0.942	0.927	0.893	0.904	0.757	0.715	0.886	0.918	0.840	0.611	0.816
Linear Discriminant Analysis	0.931	0.970	0.952	0.744	0.789	1.000	0.962	0.869	0.935	0.826	0.838	0.871	0.854	0.833	0.897	0.896	0.946	0.810	0.876	0.942	0.927	0.893	0.904	0.757	0.715	0.886	0.918	0.840	0.611	0.816
Logistic Regression	0.931	0.970	0.952	0.744	0.789	1.000	0.962	0.869	0.935	0.826	0.838	0.871	0.854	0.833	0.897	0.896	0.946	0.810	0.876	0.942	0.927	0.893	0.904	0.757	0.715	0.886	0.918	0.840	0.611	0.816
Multi Layer Perceptron	0.931	0.970	0.952	0.744	0.789	1.000	0.962	0.869	0.935	0.826	0.838	0.871	0.854	0.833	0.897	0.896	0.946	0.810	0.876	0.942	0.927	0.893	0.904	0.757	0.715	0.886	0.918	0.840	0.611	0.816
Nearest Neighbors	0.931	0.970	0.952	0.744	0.789	1.000	0.962	0.869	0.935	0.826	0.838	0.871	0.854	0.833	0.897	0.896	0.946	0.810	0.876	0.942	0.927	0.893	0.904	0.757	0.715	0.886	0.918	0.840	0.611	0.816
Passive Aggressive Classifier	0.931	0.970	0.952	0.744	0.789	1.000	0.962	0.869	0.935	0.826	0.838	0.871	0.854	0.833	0.897	0.896	0.946	0.810	0.876	0.942	0.927	0.893	0.904	0.757	0.715	0.886	0.918	0.840	0.611	0.816
Random Forest	0.931	0.970	0.952	0.744	0.789	1.000	0.962	0.869	0.935	0.826	0.838	0.871	0.854	0.833	0.897	0.896	0.946	0.810	0.876	0.942	0.927	0.893	0.904	0.757	0.715	0.886	0.918	0.840	0.611	0.816
SVM - linear	0.931	0.970	0.952	0.744	0.789	1.000	0.962	0.869	0.935	0.826	0.838	0.871	0.854	0.833	0.897	0.896	0.946	0.810	0.876	0.942	0.927	0.893	0.904	0.757	0.715	0.886	0.918	0.840	0.611	0.816
SVM - poly	0.931	0.970	0.952	0.744	0.789	1.000	0.962	0.869	0.935	0.826	0.838	0.871	0.854	0.833	0.897	0.896	0.946	0.810	0.876	0.942	0.927	0.893	0.904	0.757	0.715	0.886	0.918	0.840	0.611	0.816
SVM - rbf	0.931	0.970	0.952	0.744	0.789	1.000	0.962	0.869	0.935	0.826	0.838	0.871	0.854	0.833	0.897	0.896	0.946	0.810	0.876	0.942	0.927	0.893	0.904	0.757	0.715	0.886	0.918	0.840	0.611	0.816

**Table 4.19 | AUROC score for 30 RGB image feature datasets for cell line U87-MG**

Classifiers	P1	P2	P3	P4	P5	P6	P7	P8	P9	P10	P11	P12	P13	P14	P15	P16	P17	P18	P19	P20	P21	P22	P23	P24	P25	P26	P27	P28	P29	P30
AdaBoost	0.863	0.920	0.833	0.731	0.752	0.806	0.773	0.844	0.890	0.693	0.916	0.777	0.721	0.854	0.733	0.979	0.522	0.633	0.875	0.799	0.708	0.868	0.900	0.784	0.700	0.484	0.807	0.813	0.649	0.897
Bernoulli Naïve Bayes	0.907	0.772	0.773	0.833	0.636	0.780	0.693	0.916	0.827	0.590	0.935	0.763	0.788	0.709	0.793	0.885	0.407	0.713	0.855	0.755	0.733	0.762	0.855							







Table 4.33 | AUROC score for 30 contrast image feature datasets for cell line MCF7

Classifiers	P1	P2	P3	P4	P5	P6	P7	P8	P9	P10	P11	P12	P13	P14	P15	P16	P17	P18	P19	P20	P21	P22	P23	P24	P25	P26	P27	P28	P29	P30			
AdaBoost	0.834	0.881	0.800	0.864	0.677	0.718	0.897	0.826	0.857	0.724	0.811	0.806	0.671	0.795	0.733	0.704	0.469	0.832	0.690	0.705	0.864	0.724	0.831	0.469	0.844	0.500	0.626	0.686	0.722	0.820	0.891	0.881	0.906
Bernoulli Naive Bayes	0.762	0.890	0.864	0.696	0.719	0.835	0.853	0.855	0.722	0.706	0.699	0.745	0.665	0.650	0.521	0.569	0.741	0.423	0.730	0.614	0.500	0.844	0.500	0.813	0.500	0.715	0.786	0.744	0.807	0.874	0.807	0.822	0.862
Dummy Classifier	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	
Extra Trees	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	
Gaussian Naive Bayes	0.778	0.799	0.760	0.688	0.627	0.760	0.750	0.611	0.632	0.542	0.612	0.591	0.565	0.686	0.500	0.598	0.500	0.598	0.500	0.532	0.417	0.750	0.384	0.610	0.538	0.776	0.722	0.621	0.684	0.881	0.684	0.838	0.838
Gradient Boosting	0.792	1.000	0.891	0.620	0.700	0.897	0.871	0.898	0.500	0.500	0.691	0.861	0.724	0.775	0.784	0.833	0.627	0.675	0.890	0.525	0.417	0.750	0.384	0.610	0.538	0.776	0.722	0.621	0.684	0.881	0.684	0.838	0.838
Helstrom Quantum Classifier	0.820	0.981	0.871	0.694	0.699	0.857	0.891	0.917	0.722	0.759	0.848	0.861	0.627	0.757	0.686	0.584	0.897	0.445	0.595	0.725	0.750	0.722	0.650	0.609	0.807	0.610	0.715	0.786	0.698	0.798	0.731	0.744	0.744
Linear Discriminant Analysis	0.756	0.708	0.651	0.707	0.588	0.627	0.706	0.792	0.611	0.550	0.642	0.612	0.591	0.565	0.500	0.442	0.735	0.531	0.526	0.750	0.484	0.631	0.483	0.588	0.776	0.760	0.731	0.759	0.711	0.569	0.731	0.569	0.731
Logistic Regression	0.820	0.981	0.871	0.694	0.699	0.857	0.891	0.917	0.722	0.759	0.848	0.861	0.627	0.757	0.686	0.584	0.897	0.445	0.595	0.725	0.750	0.722	0.650	0.609	0.807	0.610	0.715	0.786	0.698	0.798	0.731	0.744	0.744
Multi Layer Perceptron	0.756	0.708	0.651	0.707	0.588	0.627	0.706	0.792	0.611	0.550	0.642	0.612	0.591	0.565	0.500	0.442	0.735	0.531	0.526	0.750	0.484	0.631	0.483	0.588	0.776	0.760	0.731	0.759	0.711	0.569	0.731	0.569	0.731
Nearest-Neighbors	0.820	0.981	0.871	0.694	0.699	0.857	0.891	0.917	0.722	0.759	0.848	0.861	0.627	0.757	0.686	0.584	0.897	0.445	0.595	0.725	0.750	0.722	0.650	0.609	0.807	0.610	0.715	0.786	0.698	0.798	0.731	0.744	0.744
Nearest-Neighbors	0.820	0.981	0.871	0.694	0.699	0.857	0.891	0.917	0.722	0.759	0.848	0.861	0.627	0.757	0.686	0.584	0.897	0.445	0.595	0.725	0.750	0.722	0.650	0.609	0.807	0.610	0.715	0.786	0.698	0.798	0.731	0.744	0.744
Nearest-Neighbors	0.820	0.981	0.871	0.694	0.699	0.857	0.891	0.917	0.722	0.759	0.848	0.861	0.627	0.757	0.686	0.584	0.897	0.445	0.595	0.725	0.750	0.722	0.650	0.609	0.807	0.610	0.715	0.786	0.698	0.798	0.731	0.744	0.744
Nearest-Neighbors	0.820	0.981	0.871	0.694	0.699	0.857	0.891	0.917	0.722	0.759	0.848	0.861	0.627	0.757	0.686	0.584	0.897	0.445	0.595	0.725	0.750	0.722	0.650	0.609	0.807	0.610	0.715	0.786	0.698	0.798	0.731	0.744	0.744
Nearest-Neighbors	0.820	0.981	0.871	0.694	0.699	0.857	0.891	0.917	0.722	0.759	0.848	0.861	0.627	0.757	0.686	0.584	0.897	0.445	0.595	0.725	0.750	0.722	0.650	0.609	0.807	0.610	0.715	0.786	0.698	0.798	0.731	0.744	0.744
Nearest-Neighbors	0.820	0.981	0.871	0.694	0.699	0.857	0.891	0.917	0.722	0.759	0.848	0.861	0.627	0.757	0.686	0.584	0.897	0.445	0.595	0.725	0.750	0.722	0.650	0.609	0.807	0.610	0.715	0.786	0.698	0.798	0.731	0.744	0.744
Nearest-Neighbors	0.820	0.981	0.871	0.694	0.699	0.857	0.891	0.917	0.722	0.759	0.848	0.861	0.627	0.757	0.686	0.584	0.897	0.445	0.595	0.725	0.750	0.722	0.650	0.609	0.807	0.610	0.715	0.786	0.698	0.798	0.731	0.744	0.744
Nearest-Neighbors	0.820	0.981	0.871	0.694	0.699	0.857	0.891	0.917	0.722	0.759	0.848	0.861	0.627	0.757	0.686	0.584	0.897	0.445	0.595	0.725	0.750	0.722	0.650	0.609	0.807	0.610	0.715	0.786	0.698	0.798	0.731	0.744	0.744
Nearest-Neighbors	0.820	0.981	0.871	0.694	0.699	0.857	0.891	0.917	0.722	0.759	0.848	0.861	0.627	0.757	0.686	0.584	0.897	0.445	0.595	0.725	0.750	0.722	0.650	0.609	0.807	0.610	0.715	0.786	0.698	0.798	0.731	0.744	0.744
Nearest-Neighbors	0.820	0.981	0.871	0.694	0.699	0.857	0.891	0.917	0.722	0.759	0.848	0.861	0.627	0.757	0.686	0.584	0.897	0.445	0.595	0.725	0.750	0.722	0.650	0.609	0.807	0.610	0.715	0.786	0.698	0.798	0.731	0.744	0.744
Nearest-Neighbors	0.820	0.981	0.871	0.694	0.699	0.857	0.891	0.917	0.722	0.759	0.848	0.861	0.627	0.757	0.686	0.584	0.897	0.445	0.595	0.725	0.750	0.722	0.650	0.609	0.807	0.610	0.715	0.786	0.698	0.798	0.731	0.744	0.744
Nearest-Neighbors	0.820	0.981	0.871	0.694	0.699	0.857	0.891	0.917	0.722	0.759	0.848	0.861	0.627	0.757	0.686	0.584	0.897	0.445	0.595	0.725	0.750	0.722	0.650	0.609	0.807	0.610	0.715	0.786	0.698	0.798	0.731	0.744	0.744
Nearest-Neighbors	0.820	0.981	0.871	0.694	0.699	0.857	0.891	0.917	0.722	0.759	0.848	0.861	0.627	0.757	0.686	0.584	0.897	0.445	0.595	0.725	0.750	0.722	0.650	0.609	0.807	0.610	0.715	0.786	0.698	0.798	0.731	0.744	0.744
Nearest-Neighbors	0.820	0.981	0.871	0.694	0.699	0.857	0.891	0.917	0.722	0.759	0.848	0.861	0.627	0.757	0.686	0.584	0.897	0.445	0.595	0.725	0.750	0.722	0.650	0.609	0.807	0.610	0.715	0.786	0.698	0.798	0.731	0.744	0.744
Nearest-Neighbors	0.820	0.981	0.871	0.694	0.699	0.857	0.891	0.917	0.722	0.759	0.848	0.861	0.627	0.757	0.686	0.584	0.897	0.445	0.595	0.725	0.750	0.722	0.650	0.609	0.807	0.610	0.715	0.786	0.698	0.798	0.731	0.744	0.744
Nearest-Neighbors	0.820	0.981	0.871	0.694	0.699	0.857	0.891	0.917	0.722	0.759	0.848	0.861	0.627	0.757	0.686	0.584	0.897	0.445	0.595	0.725	0.750	0.722	0.650	0.609	0.807	0.610	0.715	0.786	0.698	0.798	0.731	0.744	0.744
Nearest-Neighbors	0.820	0.981	0.871	0.694	0.699	0.857	0.891	0.917	0.722	0.759	0.848	0.861	0.627	0.757	0.686	0.584	0.897	0.445	0.595	0.725	0.750	0.722	0.650	0.609	0.807	0.610	0.715	0.786	0.698	0.798	0.731	0.744	0.744
Nearest-Neighbors	0.820	0.981	0.871	0.694	0.699	0.857	0.891	0.917	0.722	0.759	0.848	0.861	0.627	0.757	0.686	0.584	0.897	0.445	0.595	0.725	0.750	0.722	0.650	0.609	0.807	0.610	0.715	0.786	0.698	0.798	0.731	0.744	0.744
Nearest-Neighbors	0.820	0.981	0.871	0.694	0.699	0.857	0.891	0.917	0.722	0.759	0.848	0.861	0.627	0.757	0.686	0.584	0.897	0.445	0.595	0.725	0.750	0.722	0.650	0.609	0.807	0.610	0.715	0.786	0.698	0.798	0.731	0.744	0.744
Nearest-Neighbors	0.820	0.981	0.871	0.694	0.699	0.857	0.891	0.917	0.722	0.759	0.848	0.861	0.627	0.757	0.686	0.584	0.897	0.445	0.595	0.725	0.750	0.722	0.650	0.609	0.807	0.610	0.715	0.786	0.698	0.798	0.731	0.744	0.744
Nearest-Neighbors	0.820	0.981	0.871	0.694	0.699	0.857	0.891	0.917	0.722	0.759	0.848	0.861	0.627	0.757	0.686	0.584	0.897	0.445	0.595	0.725	0.750	0.722	0.650	0.609	0.807	0.610	0.715	0.786	0.698	0.798	0.731	0.744	0.744
Nearest-Neighbors	0.820	0.981	0.871	0.694	0.699	0.857	0.891	0.917	0.722	0.759	0.848	0.861	0.627	0.757	0.686	0.584	0.897	0.445	0.595	0.725	0.750	0.722	0.650	0.609									





Table 4.41 | Balanced accuracy score for 30 energy image feature datasets for cell line U251

Classifiers	P1	P2	P3	P4	P5	P6	P7	P8	P9	P10	P11	P12	P13	P14	P15	P16	P17	P18	P19	P20	P21	P22	P23	P24	P25	P26	P27	P28	P29	P30
AdaBoost	1.000	1.000	1.000	0.981	0.987	1.000	0.974	1.000	0.975	1.000	0.952	0.955	0.982	0.922	0.958	0.927	0.946	0.978	0.931	0.974	1.000	0.931	0.947	1.000	0.983	1.000	0.983	1.000	0.984	0.982
Bernoulli Naive Bayes	0.940	0.912	0.838	0.827	0.955	0.984	0.922	0.933	0.885	0.971	0.941	0.807	0.500	0.500	0.500	0.887	0.920	0.887	0.920	0.887	0.920	0.887	0.920	0.887	0.920	0.887	0.920	0.887	0.920	0.887
Decision Tree	0.988	0.988	0.988	0.988	0.988	0.988	0.988	0.988	0.988	0.988	0.988	0.988	0.988	0.988	0.988	0.988	0.988	0.988	0.988	0.988	0.988	0.988	0.988	0.988	0.988	0.988	0.988	0.988	0.988	0.988
Extra Trees	0.933	0.500	0.859	0.500	0.494	0.500	0.484	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	
Gaussian Naive Bayes	0.917	1.000	1.000	1.000	0.944	1.000	0.944	1.000	1.000	1.000	0.944	1.000	0.944	1.000	0.944	1.000	0.944	1.000	0.944	1.000	0.944	1.000	0.944	1.000	0.944	1.000	0.944	1.000	0.944	
Gradient Boosting	1.000	1.000	1.000	1.000	0.933	1.000	0.974	1.000	1.000	1.000	0.933	1.000	0.974	1.000	1.000	0.933	1.000	0.974	1.000	1.000	0.933	1.000	0.974	1.000	1.000	0.933	1.000	0.974	1.000	
Helstrom Quantum Classifier	0.960	1.000	1.000	1.000	1.000	0.974	1.000	0.967	0.981	0.971	0.941	0.983	0.976	0.967	0.941	0.960	0.958	0.927	0.946	0.978	0.931	0.918	1.000	0.958	0.981	0.983	0.964	0.979	0.920	0.982
Linear Discriminant Analysis	1.000	1.000	1.000	1.000	1.000	0.929	0.972	0.933	0.921	0.941	0.981	0.976	0.967	0.941	0.960	0.958	0.927	0.946	0.978	0.931	0.918	1.000	0.958	0.981	0.983	0.964	0.979	0.920	0.982	
Logistic Regression	1.000	1.000	1.000	1.000	1.000	0.929	0.972	0.933	0.921	0.941	0.981	0.976	0.967	0.941	0.960	0.958	0.927	0.946	0.978	0.931	0.918	1.000	0.958	0.981	0.983	0.964	0.979	0.920	0.982	
Multi Layer Perceptron	1.000	1.000	1.000	1.000	1.000	0.929	0.972	0.933	0.921	0.941	0.981	0.976	0.967	0.941	0.960	0.958	0.927	0.946	0.978	0.931	0.918	1.000	0.958	0.981	0.983	0.964	0.979	0.920	0.982	
Nearest Centroid	0.960	0.941	0.865	0.923	0.955	1.000	0.917	0.933	0.942	0.916	0.941	0.950	0.952	0.933	0.938	0.782	0.938	0.927	0.946	0.978	0.931	0.893	1.000	0.958	0.981	0.983	0.964	0.979	0.920	0.982
Nearest Neighbors	1.000	1.000	1.000	1.000	1.000	0.964	0.972	0.987	0.971	0.986	1.000	1.000	1.000	1.000	1.000	0.975	0.975	0.975	0.975	0.975	0.975	0.975	0.975	0.975	0.975	0.975	0.975	0.975	0.975	0.975
Passive Aggressive Classifier	1.000	1.000	1.000	1.000	1.000	0.981	0.987	0.981	0.987	0.981	0.987	0.981	0.987	0.981	0.987	0.981	0.987	0.981	0.987	0.981	0.987	0.981	0.987	0.981	0.987	0.981	0.987	0.981	0.987	0.981
Passive Aggressive Classifier	1.000	1.000	1.000	1.000	1.000	0.981	0.987	0.981	0.987	0.981	0.987	0.981	0.987	0.981	0.987	0.981	0.987	0.981	0.987	0.981	0.987	0.981	0.987	0.981	0.987	0.981	0.987	0.981	0.987	0.981
Quadratic Discriminant Analysis	0.917	1.000	1.000	1.000	1.000	0.944	0.918	0.967	0.969	1.000	0.944	0.929	0.940	0.944	0.929	0.940	0.929	0.940	0.944	0.929	0.940	0.944	0.929	0.940	0.944	0.929	0.940	0.944	0.929	0.982
Random Forest	1.000	1.000	1.000	1.000	1.000	0.974	1.000	0.974	1.000	1.000	0.974	1.000	0.974	1.000	0.974	1.000	0.974	1.000	0.974	1.000	0.974	1.000	0.974	1.000	0.974	1.000	0.974	1.000	0.974	1.000
SVM - linear	1.000	1.000	1.000	1.000	1.000	0.964	1.000	0.967	0.981	1.000	0.971	1.000	1.000	1.000	0.962	0.946	0.958	0.927	0.946	1.000	0.931	0.893	1.000	0.981	0.983	1.000	0.983	1.000	0.984	0.982
SVM - poly	1.000	1.000	1.000	1.000	1.000	0.964	1.000	0.967	0.981	1.000	0.971	1.000	1.000	1.000	0.962	0.946	0.958	0.927	0.946	1.000	0.931	0.893	1.000	0.981	0.983	1.000	0.983	1.000	0.984	0.982
SVM - rbf	1.000	1.000	1.000	1.000	1.000	0.974	1.000	0.967	0.981	1.000	0.971	1.000	1.000	1.000	0.962	0.946	0.958	0.927	0.946	1.000	0.931	0.893	1.000	0.981	0.983	1.000	0.983	1.000	0.984	0.982

Table 4.42 | Balanced accuracy score for 30 homogeneity image feature datasets for cell line U251

Classifiers	P1	P2	P3	P4	P5	P6	P7	P8	P9	P10	P11	P12	P13	P14	P15	P16	P17	P18	P19	P20	P21	P22	P23	P24	P25	P26	P27	P28	P29	P30
AdaBoost	1.000	1.000	1.000	1.000	1.000	0.974	1.000	0.974	1.000	1.000	0.975	1.000	0.952	0.955	0.982	0.922	0.946	0.978	0.931	0.974	1.000	0.931	0.947	1.000	0.983	1.000	0.983	1.000	0.984	0.982
Bernoulli Naive Bayes	1.000	1.000	1.000	1.000	1.000	0.978	0.974	1.000	0.975	0.983	1.000	0.975	0.983	1.000	0.962	0.946	0.917	0.969	0.972	1.000	0.926	0.974	1.000	0.980	0.962	0.983	0.946	0.681	0.960	0.982
Decision Tree	1.000	1.000	1.000	1.000	1.000	1.000	1.000	1.000	1.000	1.000	1.000	1.000	1.000	1.000	1.000	1.000	1.000	1.000	1.000	1.000	1.000	1.000	1.000	1.000	1.000	1.000	1.000	1.000	1.000	1.000
Extra Trees	1.000	1.000	1.000	1.000	1.000	0.974	1.000	0.974	1.000	1.000	0.974	1.000	0.974	1.000	0.974	1.000	0.974	1.000	0.974	1.000	0.974	1.000	0.974	1.000	0.974	1.000	0.974	1.000	0.974	1.000
Gaussian Naive Bayes	1.000	1.000	1.000	1.000	1.000	0.987	1.000	0.987	1.000	0.987	1.000	0.987	1.000	0.987	1.000	0.987	1.000	0.987	1.000	0.987	1.000	0.987	1.000	0.987	1.000	0.987	1.000	0.987	1.000	0.987
Gradient Boosting	1.000	1.000	1.000	1.000	1.000	0.987	1.000	0.987	1.000	0.987	1.000	0.987	1.000	0.987	1.000	0.987	1.000	0.987	1.000	0.987	1.000	0.987	1.000	0.987	1.000	0.987	1.000	0.987	1.000	0.987
Helstrom Quantum Classifier	1.000	1.000	1.000	1.000	1.000	0.974	1.000	0.974	1.000	1.000	0.974	1.000	0.974	1.000	0.974	1.000	0.974	1.000	0.974	1.000	0.974	1.000	0.974	1.000	0.974	1.000	0.974	1.000	0.974	1.000
Linear Discriminant Analysis	1.000	1.000	1.000	1.000	1.000	0.967	1.000	0.974	1.000	1.000	0.967	1.000	1.000	1.000	0.962	0.946	0.917	0.927	0.972	1.000	0.931	0.893	1.000	0.980	0.983	1.000	0.983	1.000	0.984	0.982
Logistic Regression	1.000	1.000	1.000	1.000	1.000	0.967	1.000	0.974	1.000	1.000	0.967	1.000	1.000	1.000	0.962	0.946	0.917	0.927	0.972	1.000	0.931	0.893	1.000	0.980	0.983	1.000	0.983	1.000	0.984	0.982
Multi Layer Perceptron	1.000	1.000	1.000	1.000	1.000	0.967	1.000	0.974	1.000	1.000	0.967	1.000	1.000	1.000	0.962	0.946	0.917	0.927	0.972	1.000	0.931	0.893	1.000	0.980	0.983	1.000	0.983	1.000	0.984	0.982
Nearest Centroid	1.000	1.000	1.000	1.000	1.000	0.974	1.000	0.967	0.981	1.000	0.971	1.000	1.000	1.000	0.962	0.946	0.917	0.927	0.972	1.000	0.931	0.893	1.000	0.981	0.983	1.000	0.983	1.000	0.984	0.982
Nearest Neighbors	1.000	1.000	1.000	1.000	1.000	0.974	1.000	0.967	0.981	1.000	0.971	1.000	1.000	1.000	0.962	0.946	0.917	0.927	0.972	1.000	0.931	0.893	1.000	0.981	0.983	1.000	0.983	1.000	0.984	0.982
Passive Aggressive Classifier	1.000	1.000	1.000	1.000	1.000	0.967	1.000	0.974	1.000	1.000	0.967	1.000	1.000	1.000	0.962	0.946	0.917	0.927	0.972	1.000	0.931	0.893	1.000	0.981	0.983	1.000	0.983	1.000	0.984	0.982
Passive Aggressive Classifier	1.000	1.000	1.000	1.000	1.000	0.967	1.000	0.974	1.000	1.000	0.967	1.000	1.000	1.000	0.962	0.946	0.917	0.927	0.972	1.000	0.931	0.893	1.000	0.981	0.983	1.000	0.983	1.000	0.984	0.982
Quadratic Discriminant Analysis	0.917	1.000	1.000	1.000	1.000	0.944	0.918	0.967	0.969	1.000	0.944	0.929	0.940	0.944	0.929	0.940	0.929	0.940	0.944	0.929	0.940	0.944	0.929	0.940	0.944	0.929	0.940	0.944	0.929	0.982
Random Forest	1.000	1.000	1.000	1.000	1.000	0.974	1.000	0.974	1.000	1.000	0.974	1.000	0.974	1.000	0.974	1.000	0.974	1.000	0.974	1.000	0.974	1.000	0.974	1.000	0.974	1.000	0.974	1.000	0.974	1.000
SVM - linear	1.000	1.000	1.000	1.000	1.000	0.964	1.000	0.967	0.981	1.000	0.971	1.000	1.000	1.000	0.962	0.946	0.917	0.927	0.972	1.000	0.931	0.893	1.000	0.981	0.983	1.000	0.983	1.000	0.984	0.982
SVM - poly	1.000	1.000	1.000	1.000	1.000	0.964	1.000	0.967	0.981	1.000	0.971	1.000	1.000	1.000	0.962	0.946	0.917	0.927	0.972	1.000	0.931	0.893	1.000	0.981	0.983	1.000	0.983	1.000	0.984	0.982
SVM - rbf	1.000	1.																												



## References

1. Haralick, R. M., Shanmugam, K. *et al.* Textural features for image classification. *IEEE Trans. Syst. Man Cybern.* **SMC-3**, 610–621, DOI: [10.1109/TSMC.1973.4309314](https://doi.org/10.1109/TSMC.1973.4309314) (1973).
2. Haralick, R. M. Statistical and structural approaches to texture. *Proc. IEEE* **67**, 786–804, DOI: [10.1109/PROC.1979.11328](https://doi.org/10.1109/PROC.1979.11328) (1979).
3. Rundo, L. *et al.* HaraliCU: GPU-powered Haralick feature extraction on medical images exploiting the full dynamics of gray-scale levels. In Malyshkin, V. (ed.) *Parallel Computing Technologies (PaCT)*, vol. 11657 of *LNCS*, 304–318, DOI: [978-3-030-25636-4\\_24](https://doi.org/10.1007/978-3-030-25636-4_24) (Springer International Publishing, Cham, Switzerland, 2019).