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

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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 $i, j \in [0, 1, ..., L-1]$ represent a gray-level pair) after the normalization by the total number of pixels. These descriptors are generally called Haralick's features^{1,2}.

Given a squared window of size $\omega \times \omega$ pixels sliding over the whole image³, we computed the following GLCM-based features (with $i, j \in [0, 1, ..., L-1]$):

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

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

contrast = 0 for a constant image;

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

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

where: $\mu_x = \sum_i \sum_j i \cdot p(i, j), \ \mu_y = \sum_i \sum_j j \cdot p(i, j), \ \sigma_x = \sum_i \sum_j (i - \mu_x) \cdot p(i, j), \ \text{and} \ \sigma_y = \sum_i \sum_j (j - \mu_y) \cdot p(i, j)$ (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:

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

energy = 1 for a constant image;

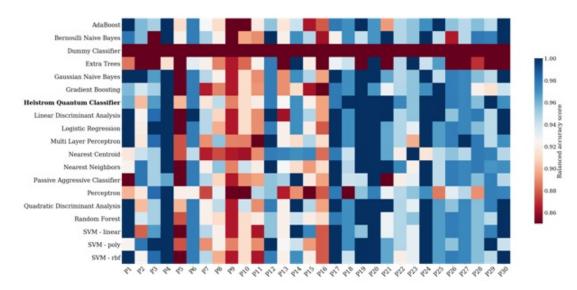
homogeneity ∈ [0,1] Returns a value that measures the closeness of the distribution of elements in the GLCM to the GLCM diagonal:

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

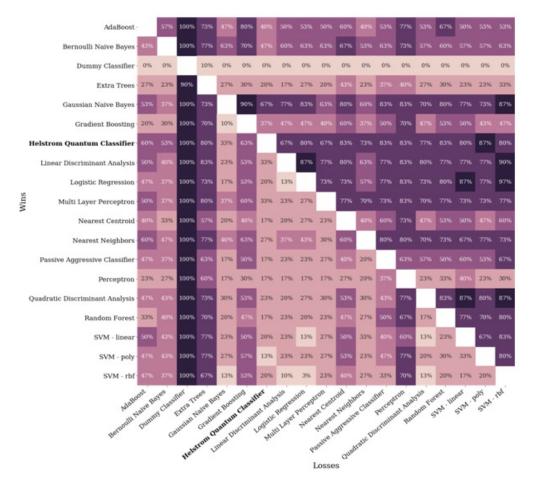


Fig. 2.1.2 | Percentage of datasets where model A ("Wins") outperformed model B ("Losses") out of 30 homogeneity image feature datasets for cell line MDA-MD-231 (balanced accuracy score).

	Image features					
Classifiers	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
Helstrom Quantum Classifier	0.895 ± 0.058	0.883 ± 0.069	0.902 ± 0.070	0.775 ± 0.081	0.938 ± 0.047	0.959 ± 0.036
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

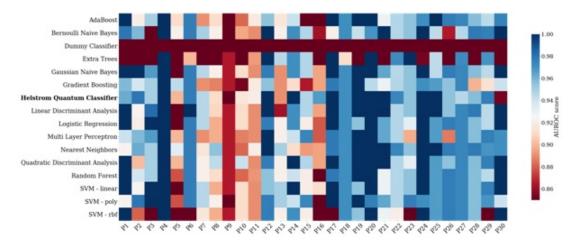


Fig. 2.1.3 | AUROC score of 19 classifiers across 30 homogeneity image feature datasets for cell line MDA-MD-231.

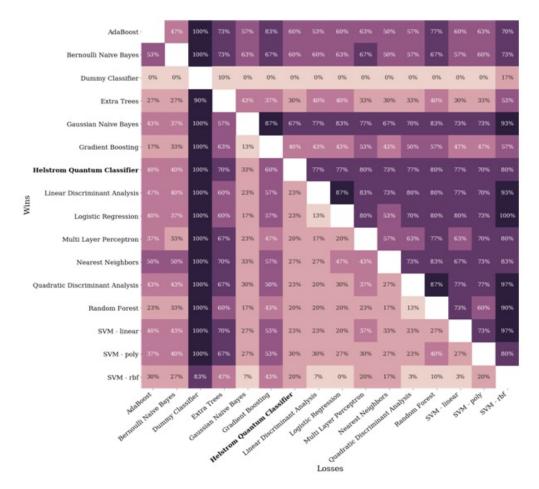


Fig. 2.1.4 | Percentage of datasets where model A ("Wins") outperformed model B ("Losses") out of 30 homogeneity image feature datasets for cell line MDA-MD-231 (AUROC score).

	Image features	5				
Classifiers	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
Helstrom Quantum Classifier	0.879 ± 0.081	0.884 ± 0.066	0.905 ± 0.062	0.787 ± 0.082	0.927 ± 0.058	0.954 ± 0.050
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

Table 2.1.2 | The mean and standard deviation AUROC score (with respect to 30 datasets) for cell line MDA-MD-231

S2.2. Cell line U87-MG

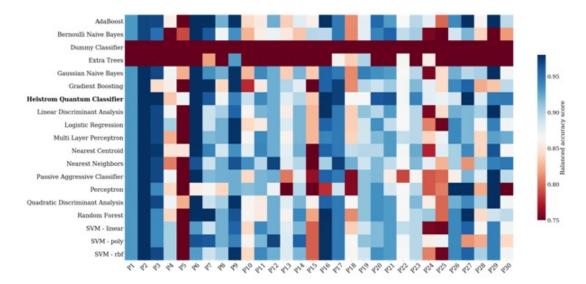


Fig. 2.2.1 | Balance accuracy score of 19 classifiers across 30 homogeneity image feature datasets for cell line U87-MG.

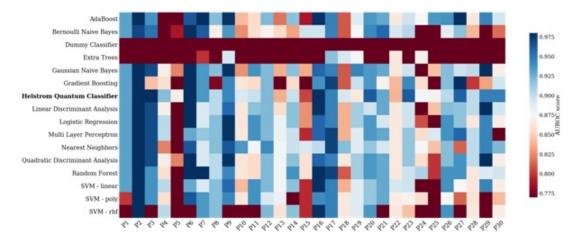


Fig. 2.2.3 | AUROC score of 19 classifiers across 30 homogeneity image feature datasets for cell line U87-MG.

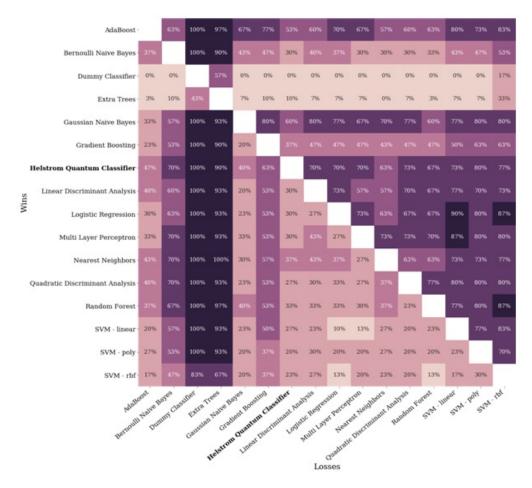


Fig. 2.2.4 | Percentage of datasets where model A ("Wins") outperformed model B ("Losses") out of 30 homogeneity image feature datasets for cell line U87-MG (AUROC score).

	Image features					
Classifiers	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
Helstrom Quantum Classifier	0.791 ± 0.099	0.792 ± 0.090	0.838 ± 0.066	0.768 ± 0.075	0.902 ± 0.057	0.917 ± 0.048
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

S2.3. Cell line MCF7

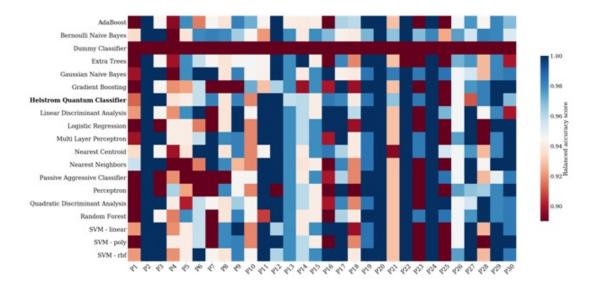


Fig. 2.3.1 | Balance accuracy score of 19 classifiers across 30 L*u*v* image feature datasets for cell line MCF7.

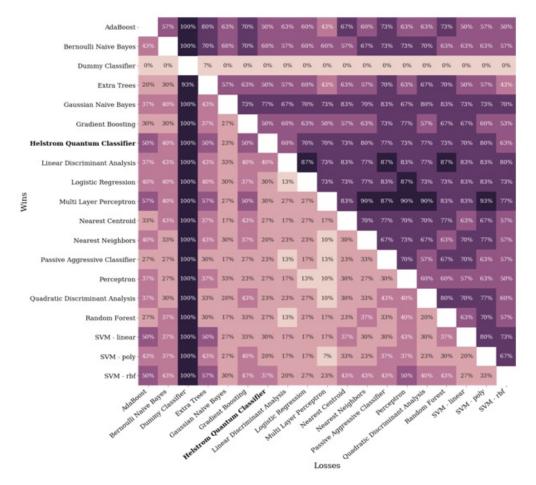


Fig. 2.3.2 | Percentage of datasets where model A ("Wins") outperformed model B ("Losses") out of 30 L*u*v* image feature datasets for cell line MCF7 (balanced accuracy score).

	Image features					
Classifiers	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
Helstrom Quantum Classifier	0.949 ± 0.052	0.965 ± 0.033	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

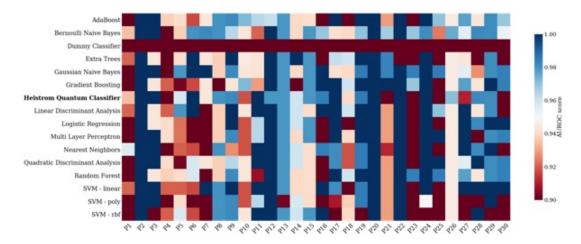


Fig. 2.3.3 | AUROC score of 19 classifiers across 30 L*u*v* image feature datasets for cell line MCF7.

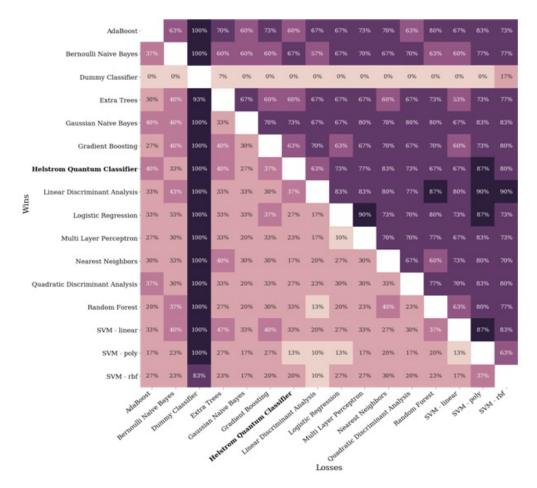


Fig. 2.3.4 | Percentage of datasets where model A ("Wins") outperformed model B ("Losses") out of 30 L*u*v* image feature datasets for cell line MCF7 (AUROC score).

Table 2.3.2 The mean and standard deviation AUROC score (with respect to 30 datasets) for cell line MCF7
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	Image features					
Classifiers	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
Helstrom Quantum Classifier	0.955 ± 0.045	0.960 ± 0.041	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

45 S2.4. Cell line U251

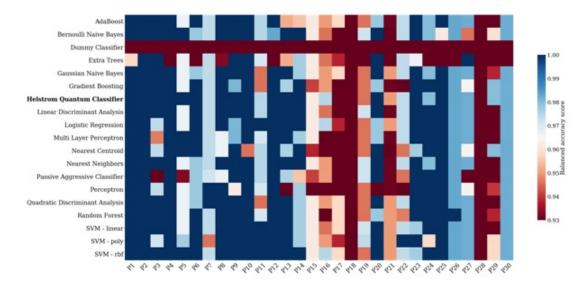


Fig. 2.4.1 | Balance accuracy score of 19 classifiers across 30 homogeneity image feature datasets for cell line U251.

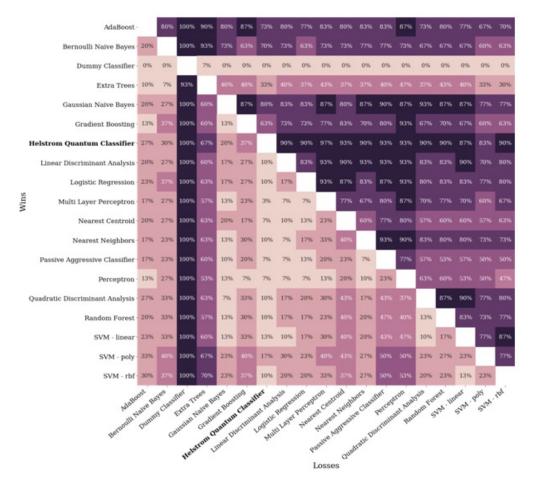


Fig. 2.4.2 | Percentage of datasets where model A ("Wins") outperformed model B ("Losses") out of 30 homogeneity image feature datasets for cell line U251 (balanced accuracy score).

	Image features					
Classifiers	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
Helstrom Quantum Classifier	0.917 ± 0.045	0.907 ± 0.039	0.966 ± 0.032	0.792 ± 0.070	0.973 ± 0.024	0.979 ± 0.029
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

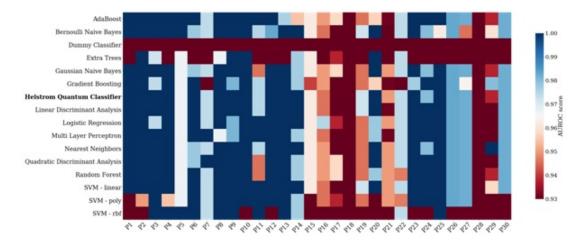


Fig. 2.4.3 | AUROC score of 19 classifiers across 30 homogeneity image feature datasets for cell line U251.

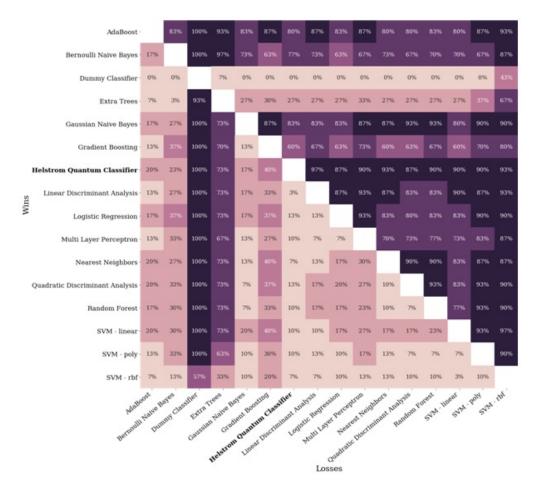


Fig. 2.4.4 | Percentage of datasets where model A ("Wins") outperformed model B ("Losses") out of 30 homogeneity image feature datasets for cell line U251 (AUROC score).

	Image features					
Classifiers	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
Helstrom Quantum Classifier	0.899 ± 0.057	0.907 ± 0.053	0.948 ± 0.063	0.797 ± 0.093	0.966 ± 0.031	0.978 ± 0.027
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

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

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.

experime	Balance accuracy a ent and a new unsee atasets for cell line	en test set for 10 rai			experime	Balance accuracy ent and a new unsee atasets for cell line	en test set for 10 rar		
Datasets	Balanced accuracy Test set used in the experiment	Balanced accuracy New unseen test set		AUROC New unseen test set	Datasets	Balanced accuracy Test set used in the experiment	Balanced accuracy New unseen test set		AUROC New unseen test set
P5	0.900	0.943	0.900	0.943	P1	0.931	0.947	0.931	0.947
P6	0.977	0.921	0.977	0.921	P2	0.985	0.953	0.985	0.954
P7	0.916	0.897	0.946	0.896	P5	0.869	0.908	0.869	0.908
P8	0.944	0.937	0.911	0.943	P9	0.943	0.928	0.943	0.940
P14	0.971	0.935	1.000	0.933	P10	0.891	0.894	0.891	0.897
P15	0.921	0.869	0.947	0.888	P14	0.898	0.868	0.898	0.870
P23	0.955	0.968	0.955	0.967	P21	0.964	0.910	0.929	0.901
P26	0.977	0.947	0.955	0.944	P22	0.867	0.868	0.839	0.870
P27	0.974	0.953	0.946	0.951	P25	0.871	0.893	0.871	0.893
P30	0.971	0.951	0.853	0.811	P27	0.980	0.869	0.960	0.871
Mean	0.951	0.932	0.939	0.920	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	New unseen test	Test set used in	New unseen test
	the experiment	set	the experiment	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	New unseen test	Test set used in	New unseen test
	the experiment	set	the experiment	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 Balancel accuracy score for 30 KCB image feature datasets for cell databaset P To P To P To P A AddBoort 23 047 058 027 050 031 041 058 051 041 058 051 05	Jobie 4.1 Balanced accuracy score for 50. J were image features datasets for control datasets. Jobie 4.1 Balanced accuracy score for 50. J were image features datasets for control datasets. Addition: P1 P1 P1 P1 P1 P1 Addition: Data Nume Datasets for 50 OPEN TO	Disk 4.1 Bialanced accuracy scores for the selective activation of the selective activation
y score for P1 0.855 0.855 0.855 0.855 0.912 0.912 0.912 0.912 0.915 0.917 0.917 0.917 0.917 0.918 0.917 0.918 0.918 0.91	91 0004 000 0004 000 0004 0000 0004 0000 0004 00000 0004 00000 00000 00000 00000 00000 00000 0000	y score for P1 P1 0.757 0.757 0.757 0.757 0.757 0.757 0.757 0.8413 0.801 0.813 0.802 0.8013 0.813 0.813 0.813 0.813 0.813 0.813 0.813 0.813 0.813 0.813 0.813 0.813 0.813 0.813 0.773 0.773 0.775 0.773 0.775 0.773 0.775 0.773 0.775 0.775 0.775 0.775 0.775 0.775 0.775 0.775 0.775 0.775 0.750 0.775
P2 P2 0.647 0.556 0.577 0.577 0.577 0.577 0.816 0.816 0.816 0.815 0.815 0.816 0.815 0.826 0.826 0.826 0.826 0.667 0.564 0.667 0.564 0.677 0.667 0.667 0.677 0.566 0.577 0.576 0.577 0.576 0.577 0.576 0.577 0.575 0.5775 0.5756 0.5775 0.5756 0.5775 0.5775 0.5775 0.5775 0.5775 0.5775 0.5775 0.5775 0.5775 0.5775 0.5775 0.5775 0.5775 0.5775 0.5775 0.5775 0.5775 0.5777 0.5775 0.5775 0.5775 0.5775 0.5775 0.5775 0.5777 0.5775 0.5777 0.5775 0.5775 0.5775 0.5775 0.5775 0.5775 0.5777 0.57750 0.57750 0.57750 0.57750 0.57750 0.57750 0.57750 0.57750 0.57750 0.57750 0.57750 0.57750 0.57750 0.57750 0.57750 0.577500 0.57750000000000	(10) (10) (10) (10) (10) (10) (10) (10)	P2 P2 P2 0.583 0.583 0.583 0.583 0.500 0.483 0.483 0.483 0.415 0.453 0.453 0.453 0.453 0.453 0.559 0.559 0.553 0.555 0.555 0.555 0.555 0.555 0.555 0.555 0.555 0.555 0.555 0.555
P3 P3 P3 0.586 0.586 0.586 0.556 0.560 0.500 0.500 0.817 0.817 0.817 0.817 0.817 0.817 0.817 0.817 0.817 0.817 0.817 0.817 0.817 0.817 0.817 0.817 0.817 0.817 0.817 0.817 0.817 0.817 0.817 0.817 0.817 0.817 0.817 0.817 0.817 0.817 0.817 0.817 0.817 0.814 0.817 0.814 0.817 0.818 0.818 0.818 0.818 0.818 0.818 0.818	Para Image 0517 0517 0517 </td <td>Antion lima P3 P3 P3 0.519 0.500 0.500 0.570 0.570 0.570 0.570 0.570 0.570 0.570 0.570 0.570 0.570 0.570 0.574 0.574 0.574 0.574 0.574 0.560 0.560 0.560 0.560 0.560 0.560 0.560 0.560 0.560 0.560 0.560 0.560 0.560 0.560</td>	Antion lima P3 P3 P3 0.519 0.500 0.500 0.570 0.570 0.570 0.570 0.570 0.570 0.570 0.570 0.570 0.570 0.570 0.574 0.574 0.574 0.574 0.574 0.560 0.560 0.560 0.560 0.560 0.560 0.560 0.560 0.560 0.560 0.560 0.560 0.560 0.560
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P18 0.858 0.858 0.868 0.484 0.903 0.897 0.903 0.827 0.827 0.923 0.828 0.942 0.942 0.904 0.9023 0.9033 0.9033 0.9042 0.9042 0.9042 0.9042 0.9043	814 1000 1	P18 0.784 0.784 0.789 0.719 0.719 0.784 0.784 0.784 0.784 0.784 0.784 0.784 0.784 0.784 0.784 0.783 0.806 0.733 0.806 0.533 0.805 0.834 0.834
P19 P19 0.7875 0.761 0.795 0.795 0.875 0.875 0.875 0.750 0.770 0.770 0.770 0.770 0.770 0.770 0.770 0.770 0.770 0.770 0.775 0.875 0.875 0.775 0.875 0.875 0.875 0.775 0.875 0.875 0.775 0.775 0.875 0.755	P10 0.849 0.843 0.843 0.843 0.843 0.843 0.843 0.843 0.843 0.843 0.843 0.843 0.843 0.843 0.844 0.844 0.844 0.844 0.844 0.845 0.845 0.845 0.845 0.845 0.845 0.845 0.845 0.845 0.845 0.845 0.845 0.845 0.845 0.845 0.845 0.946 0.946 0.947 0.948 0.949 0.949 0.949 0.949 0.949 0.949 0.949 0.949 0.949 <td>P19 0.833 0.5833 0.5833 0.5835 0.5855 0.55530 0.5550 0.5550 0.55500 0.55500 0.55500000000</td>	P19 0.833 0.5833 0.5833 0.5835 0.5855 0.55530 0.5550 0.5550 0.55500 0.55500 0.55500000000
P20 0.865 0.865 0.875 0.900 0.900 0.950 0.9000 0.900 0.90000 0.90000 0.90000 0.90000 0.90000 0.90000 0.90000 0.900000 0.90000000000	027 027 027 027 027 027 027 027	P20 0.900 0.900 0.803 0.925 0.975 0.975 0.975 0.975 0.975 0.971 0.882 0.971 1.000 0.837 0.946 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 0.9757 0.07577 0.07577 0.07577 0.07577 0.07577 0.07577 0.07577 0.07577 0.07577 0.07577 0.07577 0.075770 0.075770 0.075770 0.075770 0.075770 0.0757700 0.075700 0.075700 0.07570000000000
P21 0.978 0.8257 0.8187 0.8371 0.764 0.8371 0.8353 0.764 0.8353 0.764 0.8353 0.764 0.8353 0.764 0.8353 0.9453 0.8353 0.8354 0.8353 0.83550 0.83550 0.83550 0.83550 0.83550 0.835500 0.835500 0.83550000000000000000000000000000000000	124 124 125 125 125 125 125 125 125 125	P21 0.877 0.877 0.407 0.467 0.467 0.467 0.773 0.467 0.505 0.550 0.550 0.550 0.550 0.550 0.554 0.554 0.554 0.554 0.554 0.554 0.554 0.554 0.554 0.554 0.554 0.556 0.556 0.556 0.556 0.556 0.556 0.556 0.556 0.556 0.556 0.556 0.556 0.556 0.556 0.556 0.556 0.556 0.557 0.556 0.
P22 1,000 0,845 0,500 0,845 0,845 0,979 0,979 0,979 0,979 0,979 0,979 0,979 0,979 0,979 0,979 0,979 0,979 0,979	P23 0.971 0.971	P22 0.770 0.500 0.500 0.575 0.7756 0.7756 0.7756 0.7476 0.877 0.877 0.446 0.446 0.446 0.776 0.776 0.776 0.776 0.776
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P24 0.917 0.863 0.560 0.560 0.560 0.889 0.817 0.9218 0.833 0.861 0.917 0.9290 0.92100 0.92100 0.92100 0.92100 0.92100 0.92100000000000000000000000000000000000	P.24 (0.000) (0.0	P24 0.598 0.5798 0.6714 0.578 0.788 0.788 0.788 0.7550 0.7550 0.7550 0.7550 0.7550 0.7550 0.7550 0.75500 0.7550000000000
P25 0.865 0.865 0.864 0.807 0.807 0.865 0.755 0.865 0.885 0.885 0.885 0.885 0.885 0.885 0.885 0.885 0.865 0.	P25 P25 0.0581 0.0581 0.0972 0.0972 0.0972 0.0972 0.0972 0.0971 0.09	P25 0.745 0.745 0.7245 0.721 0.816 0.816 0.816 0.816 0.816 0.837 0.839 0.837 0.837 0.837 0.837 0.837 0.837 0.837 0.837 0.837 0.837 0.837 0.837
P26 0.881 0.884 0.589 0.428 0.913 0.	0.120 0.	P26 0.363 0.776 0.500 0.770 0.744 0.744 0.744 0.770 0.770 0.770 0.779 0.778 0.779 0.779 0.779 0.779 0.779 0.779 0.779 0.776 0.777 0.776 0.7776 0.7776 0.77776 0.77776 0.7777777777
P27 0.796 0.789 0.789 0.820 0.812 0.812 0.812 0.812 0.812 0.812 0.8210 0.82100000000000000000000000000000000000	P27 0947 0946 0946 0946 0972 0972 0972 0972 0972 0972 0972 0972	P27 0.850 0.8150 0.816 0.810 0.810 0.850 0.810 0.8500 0.8500 0.8500 0.8500 0.8500 0.8500 0.8500 0.8500 0.85000 0.850000000000
P28 0.855 0.855 0.8556 0.8566 0.8907 0.8907 0.8977 0.8914 0.877 0.943 0.943 0.943 0.943 0.943 0.943 0.943 0.943 0.943 0.943 0.943 0.835 0.935 0.835 0.9350 0.9350 0.9350 0.9350 0.9350 0.9350 0.9350 0.9350 0.9350 0.9350 0.9350 0.9350 0.9350 0.9350 0.9350 0.93500 0.93500 0.93500 0.93500000000000000000000000000000000000	P:28 0.915 0.900 0.900 0.900 0.915 0.915 0.915 0.915 0.915 0.915 0.915 0.915 0.929 0.920 0.9290 0.9290 0.9290 0.929000 0.9290000000000	P28 0.665 0.455 0.458 0.500 0.500 0.9795 0.9795 0.9795 0.9795 0.415 0.415 0.415 0.415 0.415 0.417 0.500 0.9798 0.9798 0.9798 0.9798 0.9798 0.9798 0.9778 0.9788 0.97980 0.9798 0.97980 0.97980 0.97980 0.97980 0.97980 0.97980 0.97980 0.97980 0.97980 0.97980 0.97980 0.97980 0.97980 0.97980 0.97980 0.97980000000000000000000000000000000000
P29 0.980 0.378 0.378 0.378 0.980 0.918 0.918 0.980 0.990 0.9800 0.980000000000	P2:9 0.918 0.911 0.929 0.929 0.929 0.929 0.929 0.929 0.929 0.929 0.929 0.929 0.929 0.929 0.929 0.929 <td>P29 0.710 0.500 0.500 0.500 0.6302 0.6302 0.577 0.577 0.570 0.5500 0.5500 0.5500 0.5500 0.5500000000</td>	P29 0.710 0.500 0.500 0.500 0.6302 0.6302 0.577 0.577 0.570 0.5500 0.5500 0.5500 0.5500 0.5500000000
P30 0.807 0.871 0.871 0.871 0.846 0.846 0.846 0.846 0.891 0.981 0.89100000000000000000000000000000000000	00.44 00.64 00	P30 0.730 0.730 0.730 0.730 0.730 0.730 0.730 0.738 0.738 0.738 0.738 0.738 0.738 0.738 0.738 0.738 0.738 0.738 0.738 0.738 0.738 0.733 0.733 0.730

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P28 0.8555 0.8555 0.5905 0.9522 0.9555 0.8644 0.9555 0.9555 0.9555 0.9555 0.9555 0.9555 0.9555 0.9555 0.9555 0.8868 0.8865 0.8855 0.8855 0.8855 0.8855 0.8855 0.8855 0.8855 0.8855 0.8855 0.8855 0.8855 0.8555 0.9555 0.8555 0.9555 0.8555 0.9555 0.8555 0.9555 0.8555 0.9555 0.9555 0.8555 0.9555 0.8555 0.9555 0.8555 0.8555 0.9555 0.8555 0.9555 0.8555 0.8555 0.9555 0.8555 0.9555 0.8555 0.9555 0.8555 0.9555 0.8555 0.9555 0.8555 0.8555 0.9555 0.85550 0.85550 0.85550 0.85550 0.85550 0.855500 0.855500 0.85550000000000	P28 0.957 0.500 0.500 0.957 0.957 0.978 0.978 0.978 0.978 0.957 0.957 0.957 0.957	P28 0.885 0.586 0.500 0.500 0.500 0.907 0.907 0.907 0.907 0.907 0.907 0.907 0.907 0.907	P28 0.885 0.863 0.604 0.500 0.921 0.921 0.921 0.829 0.863 0.863 0.863 0.863 0.863 0.863 0.863 0.869 0.869 0.869
P27 0.947 0.948 0.944 0.974 0.974 0.974 0.974 0.974 0.974 0.974 0.974 0.974 0.974 0.974 0.974 0.974	P27 0.946 0.946 0.500 0.974 0.974 0.974 0.974 0.974 0.974 0.974 0.974 0.974 0.974 0.974	P27 0.796 0.789 0.558 0.558 0.820 0.820 0.821 0.822 0.821 0.822 0.821 0.823 0.851 0.851 0.851 0.851 0.851 0.852 0.851 0.852 0.852 0.852	P27 0.947 0.946 0.946 0.974 0.946 0.946 0.946 0.946 0.947 0.947 0.947 0.972 0.972
P26 0.955 0.547 0.547 0.540 0.540 0.540 0.540 0.540 0.540 0.540 0.923 0.923 0.923 0.935 0.935 0.935	P26 0.955 0.518 0.518 0.977 0.977 0.977 0.977 0.977 0.977 0.977 0.977 0.977 0.977 0.977 0.977	P26 0.931 0.894 0.500 0.500 0.913 0.913 0.913 0.913 0.913 0.913 0.913 0.913 0.913 0.913 0.913 0.913 0.913 0.963	P26 0.929 0.871 0.685 0.500 0.500 0.821 0.823 0.964 0.823 0.929 0.857 0.857 0.857 0.857 0.857
P25 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 0.975 0.975	P25 1,000 0,944 0,967 0,944 1,000 1,000 1,000 0,944 0,944 0,947 0,947 0,967 0,967 0,967 0,967	P25 0.863 0.807 0.784 0.784 0.786 0.786 0.786 0.756 0.836 0.863 0.863 0.758 0.758 0.758 0.758 0.758 0.758 0.758 0.758 0.758 0.7580 0.863 0.863 0.863 0.863 0.863 0.863 0.863 0.863 0.863 0.863 0.863 0.863 0.863 0.863 0.863 0.863 0.863 0.863 0.8830 0.8830 0.8830 0.8830 0.8830 0.8830 0.8830 0.8830 0.8830 0.8830 0.8830 0.8830 0.8830 0.88330 0.88330 0.88330 0.88330 0.883300 0.88330000000000	P25 0.952 0.881 0.464 0.905 0.976 0.922 0.922 0.922 0.925 0.922 0.925 0.925 0.925 0.952
P24 0.943 0.943 0.943 0.964 0.964 0.964 0.964 0.964 0.964 0.964 0.964 0.964 0.964 0.964 0.964	P24 0.971 1.000 0.500 0.941 1.000 1.000 1.000 1.000 1.000 1.000 1.000 1.000 1.000 1.000 1.000 1.000	P24 0.892 0.863 0.500 0.510 0.917 0.917 0.917 0.917 0.917 0.917 0.917	P24 0.890 0.938 0.914 0.914 0.914 0.918 0.908 0.914 0.914 0.918 0.918 0.918 0.918
P23 0.938 0.9708 0.343 0.917 0.958 0.958 0.958 0.958 0.958 0.958 0.958 0.958 0.958	P23 0.955 0.955 0.500 0.955 0.955 0.955 0.955 0.955 0.932 0.932 0.932 0.932 0.932 0.932	P23 0.890 0.477 0.890 0.890 0.890 0.935 0.890 0.9350 0.9350 0.9350 0.9350 0.9350 0.9350 0.9350 0.9350 0.0350000000000000000000000000000000	P23 0.964 0.899 0.705 0.957 0.957 0.957 0.957 0.957 0.951 0.978 0.97750 0.97750 0.97750 0.97750 0.97750 0.97750 0.97750 0.97750 0.97750 0.97750 0.97750 0.97750 0.97750 0.97750 0.977500 0.97750000000000000000000000000000000000
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P20 0.972 0.972 0.965 0.966 0.982 0.918 0.972 0.972 0.918 0.918 0.918 0.920 0.972 0.972 0.972 0.972	P20 0.969 0.969 0.969 0.500 0.500 1.000	P20 0.971 0.971 0.5875 0.5801 0.900 0.900 0.950 0.950 0.950 0.950 0.950 0.921 0.856 0.921 0.866 0.921 0.866 0.920 0.920 0.92	P20 0.921 0.941 0.941 0.946
P19 1.000 1.000 0.500 0.500 1.000 0.935 0.935 0.935 0.935 0.935 0.957 0.9550 0.9550 0.9550 0.9550000000000	P19 1.000 0.942 0.353 0.318 1.000 1.000 1.000 1.000 1.000 0.942 0.0000 0.0000 0.0000 0.000000	P19 0.875 0.761 0.500 0.500 0.588 0.795 0.875 0.875 0.875 0.875 0.875 0.875 0.875 0.875 0.875 0.875 0.875	P19 0.871 0.871 0.842 0.845 0.814 0.815 0.849 0.385 0.845 0.385 0.863 0.863 0.863 0.863 0.863 0.863 0.863
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P17 0.916 0.916 0.516 0.516 0.981 0.981 0.981 0.982 0.942 0.942 0.942 0.981 0.982 0.981 0.981 0.981 0.981 0.981 0.981	P17 0.978 0.978 0.929 0.929 0.978 0.978 0.978 0.978 0.978 0.978 0.978 0.978 0.978 0.978 0.978 0.978 0.978 0.978	P17 0.946 0.946 0.510 0.946 0.946 0.946 0.946 0.946 0.946 0.946 0.946 0.946	P17 0.898 0.932 0.932 0.932 0.932 0.932 0.932 0.932 0.932 0.932 0.932 0.932 0.932 0.932
P16 0.528 0.528 0.528 0.500 0.588 0.923 0.928 0.928 0.858 0.858 0.858 0.726 0.726 0.726 0.583 0.726 0.583 0.583 0.583 0.583 0.583	P16 0.879 0.862 0.526 0.526 0.895 0.895 0.895 0.879 0.912 0.895 0.895 0.895 0.895 0.895 0.879 0.879 0.879	P16 0.938 0.4878 0.4878 0.4878 0.938 0.938 0.938 0.938 0.938 0.938 0.938 0.938	P16 0.643 0.824 0.824 0.500 0.500 0.506 0.526 0.656 0.656 0.656 0.656 0.656 0.656 0.656 0.658 0.658 0.658 0.668
P15 0.864 0.864 0.512 0.512 0.867 0.867 0.867 0.867 0.867 0.865 0.814 0.814 0.813 0.823 0.823 0.823 0.823 0.823 0.823 0.823 0.823 0.823 0.823 0.0233 0.823 0.833 0.833 0.833 0.9320 0.833 0.9320 0.833 0.0320 0.833 0.0320 0.833 0.0320 0.833 0.0320 0.833 0.0320 0.833 0.0320 0.8330 0.8330 0.8330 0.8330 0.83300 0.83300 0.83300 0.8330000000000	P15 0.864 0.947 0.512 0.920 0.947 0.921 0.921 0.921 0.920 0.918 0.918 0.918 0.918 0.918 0.918 0.918 0.918	P15 0.756 0.778 0.742 0.742 0.778 0.778 0.778 0.778 0.778 0.778 0.778 0.778	PIS 0.816 0.741 0.741 0.741 0.850 0.850 0.707 0.821 0.821 0.787 0.785 0.785 0.785 0.785 0.782 0.782 0.782
P14 0.944 0.943 0.517 0.944 0.944 0.944 0.944 0.944 0.944 0.944 0.944 0.944 0.944 0.944 0.944 0.944 0.944 0.944 0.944 0.944	P14 0.941 0.971 0.500 0.971 0.971 0.971 0.971 0.971 0.971 0.971 0.971 0.971 0.971 0.971 0.971 0.971 0.971 0.091 0.091 0.091 0.091 0.091 0.091 0.091 0.091 0.097 0.000 0.097 0.0000 0.0000 0.0000 0.0000 0.0000 0.0000 0.0000 0.0000 0.0000 0.00000 0.0000 0.00000 0.000000	P14 0.925 0.866 0.866 0.876 0.876 0.896 0.921 0.921 0.921 0.925 0.925 0.925	P14 0.906 0.859 0.882 0.929 0.926 0.906 0.906 0.906 0.906 0.906 0.906 0.906
P13 0.938 0.938 0.938 0.969 0.969 0.969 0.969 0.969 0.969 0.969 0.969 0.969 0.969 0.969 0.969	P13 0.916 0.926 0.550 0.550 0.881 0.925 0.925 0.925 0.925 0.925 0.921 0.921 0.921 0.921 0.921 0.921 0.921 0.921	P13 0.858 0.858 0.916 0.916 0.916 0.916 0.916 0.916 0.916 0.916 0.916 0.916	P13 0.795 0.692 0.692 0.765 0.775 0.775 0.775 0.775 0.775 0.775 0.775 0.775 0.775 0.775 0.775 0.775 0.775 0.772
P12 0.971 0.451 0.451 1.000 0.971 0.971 0.971 0.971 0.971 0.971 1.000 0.971 1.000 1.000	P12 0.955 0.427 0.977 0.977 0.977 0.977 0.977 0.977 0.977 0.977 0.955 0.955 0.955 0.955 0.955 0.955	P12 0.885 0.891 0.500 0.500 0.871 0.899 0.895 0.895 0.895 0.897 0.863 0.897 0.897 0.897 0.893	P12 0.967 0.944 0.947 0.967 0.967 0.967 0.967 0.967 0.967 0.967 0.967 0.963 0.933
P11 0.978 0.354 0.354 0.929 0.929 0.929 0.929 0.929 0.929 0.929 0.929 0.929 0.929 0.929 0.929 0.929 0.929 0.927 0.929 0.927 0.929 0.927 0.929 0.927 0.929 0.927 0.929 0.927 0.929 0.927 0.929 0.927 0.929 0.929 0.929 0.929 0.929 0.929 0.929 0.929 0.929 0.929 0.929 0.929 0.929 0.929 0.929 0.929 0.929 0.929 0.928 0.929 0.9280 0.92800 0.9280 0.92800 0.92800 0.92800 0.92800 0.928000 0.92800000000000000000000000000000000000	P11 0.918 0.890 0.380 0.380 0.892 0.920 0.890 0.890 0.890 0.890 0.890 0.890 0.890 0.891 0.864 0.864	P11 0.865 0.814 0.500 0.500 0.787 0.787 0.814 0.787 0.833 0.783 0.783 0.787 0.833 0.787 0.833 0.787 0.833 0.787 0.814 0.787 0.814 0.787	P11 0.920 0.918 0.500 0.918 0.918 0.918 0.918 0.918 0.918 0.946 0.946 0.946 0.944 0.944
P10 0.866 0.856 0.500 0.500 0.735 0.837 0.837 0.925 0.841 0.841 0.841 0.841 0.841 0.841 0.840 0.866 0.866 0.866 0.866	P10 0.826 0.566 0.566 0.582 0.912 0.912 0.912 0.912 0.912 0.912 0.912 0.912 0.912 0.912 0.912	P10 0.789 0.781 0.500 0.500 0.827 0.827 0.827 0.827 0.812 0.757 0.851 0.757 0.851 0.757 0.853 0.757 0.854 0.757 0.850 0.757 0.850 0.757 0.850 0.757 0.850 0.778 0.789 0.778 0.789 0.778 0.77777 0.7777 0.7777 0.7777 0.7777 0.77777 0.77777 0.7777 0.77777 0.77777 0.77777 0.77777777	P10 0.878 0.712 0.712 0.772 0.772 0.772 0.772 0.772 0.772 0.772 0.772 0.772 0.772 0.773 0.773 0.773 0.773
P9 0.725 0.500 0.500 0.839 0.839 0.839 0.833 0.726 0.867 0.756 0.867 0.867 0.867 0.868 0.867 0.868 0.867 0.868 0.867 0.867 0.867 0.839 0.839 0.839 0.839 0.839 0.839 0.839 0.839 0.839 0.839 0.867 0.756 0.867 0.756 0.867 0.756 0.867 0.756 0.867 0.756 0.839 0.833 0.835 0.833 0.835 0.833 0.835 0.835 0.835 0.835 0.835 0.835 0.835 0.835 0.835 0.833 0.835 0.833 0.835 0.833 0.833 0.835 0.833 0.8350 0.8350 0.8350 0.83500000000000000000000000000000000000	P9 0.832 0.832 0.832 0.832 0.835 0.865	P9 0.885 0.8857 0.893 0.893 0.823 0.823 0.835 0.835 0.835 0.835 0.835 0.835 0.835 0.835 0.835 0.835	P9 0.866 0.701 0.473 0.859 0.820 0.897 0.897 0.897 0.897 0.897 0.897 0.897 0.897 0.897 0.897 0.835 0.835 0.835 0.845 0.845 0.845 0.8550 0.8550 0.8550 0.8550 0.8550000000000
31 P8 1,000 1,000 1,000 1,000 1,000 0,812 0,812 1,000 0,969 0,812 1,000 0,969 0,968 0	MD-231 P8 P8 0.9211 0.9211 0.9288 0.9288 0.944 0.944 0.9550 0.9550 0.9550 0.9550 0.9550 0.9550 0.9550 0.9550 0.95500 0.955000 0.9550000000000	P8 0.906 0.500 0.530 0.938 0.938 0.938 0.938 0.938 0.938 0.938 0.938 0.938 0.938	P8 0.775 0.750 0.750 0.771 0.771 0.771 0.776 0.796 0.796 0.796 0.796 0.796 0.796 0.796 0.796 0.796 0.796 0.796 0.796 0.796
P7 P7 0.943 0.943 0.848 0.848 0.848 0.899 0.899 0.921 0.899 0.899 0.899 0.899 0.899 0.899 0.899 0.899 0.899 0.899 0.899 0.899 0.899 0.899 0.899 0.899 0.899 0.899 0.921 0.899 0.899 0.921 0.899 0.921 0.899 0.921 0.899 0.9210 0.9210 0.9210 0.92100000000000000000000000000000000000	Def MDA. P7 P7 0.925 0.925 0.921 0.946 0.946 0.946 0.946 0.946 0.946 0.946 0.946 0.946 0.946 0.921 0.946 0.921 0.946 0.921 0.946 0.921 0.946 0.921 0.946 0.921 0.921 0.921 0.921 0.946 0.921	P7 0.882 0.951 0.951 0.951 0.914 0.914	31 P7 0.895 0.784 0.784 0.865 0.805 0.805 0.805 0.833 0.833 0.833 0.833 0.833 0.833 0.833 0.864 0.833 0.865 0
Image Image <th< td=""><td>Der cell II 201 0.977 0.977 0.933 0.9777 0.97777 0.97777 0.97777 0.97777 0.977777 0.9777777777777777777777777777777777777</td><td> P.6 P.6 0.764 0.720 0.720 0.775 0.776 0.7</td><td>P6 P6 0.804 0.804 0.7844 0.7844 0.7844 0.78440000000000000000000000000000000000</td></th<>	Der cell II 201 0.977 0.977 0.933 0.9777 0.97777 0.97777 0.97777 0.97777 0.977777 0.9777777777777777777777777777777777777	 P.6 P.6 0.764 0.720 0.720 0.775 0.776 0.7	P6 P6 0.804 0.804 0.7844 0.7844 0.7844 0.78440000000000000000000000000000000000
P5 P5 1.000 1.000 0.555 0.555 0.944 1.000 0.918 0.944 0.946 0.946 0.946 0.946 0.946 0.946 0.946 0.946 0.946 0.946 0.946 1.000 0.972 0.972 0.972 0.972 0.972 0.972 0.972 0.972 0.972 0.972 0.972 0.972 0.972 0.972 0.972 0.972 0.975 0.97	Proj Proj <th< td=""><td>PS 0.917 0.917 0.917 0.889 0.917 0.917 0.917</td><td>ell line M PS 0.890 0.906 0.914 0.914 0.918 0.906 0.938 0.906 0.938 0.906 0.938 0.906 0.938 0.906</td></th<>	PS 0.917 0.917 0.917 0.889 0.917 0.917 0.917	ell line M PS 0.890 0.906 0.914 0.914 0.918 0.906 0.938 0.906 0.938 0.906 0.938 0.906 0.938 0.906
P4 P4 0.929 0.501 0.501 0.503 0.893 0.893 0.893 0.893 0.893 0.893 0.929 0.92	gg fatur P4 P4 1000 1.000 0.913 1.0000 1.0000 1.0000 1.0000 1.0000 1.0000 1.00000	Is for cell P4 0.887 0.887 0.500 0.501 0.502 0.897 0.500 0.897 0.896 0.695	P4 P4 0.943 0.907 0.943 0.9445 0.9445 0.9445 0.9445 0.9445 0.9455 0.9455 0.9455 0.9455 0.9455 0.9455 0.9455 0.9455 0.9455 0.9455 0.9455 0.9455 0.94555 0.94555 0.94555555555555555555555555555555555555
image fe P3 0.857 0.567 0.560 0.560 0.250 0.857 0.769 0.769 0.857 0.840 0.929 0.857 0.840 0.929 0.857 0.857 0.857 0.857 0.786 0.7786 0.78866 0.78866 0.78866 0.78866 0.78866 0.78866 0.7886		rre datase P3 0.618 0.755 0.765 0.769 0.769 0.769 0.769 0.769 0.769 0.769 0.769 0.769 0.769 0.769 0.769 0.769 0.768 0.768 0.768 0.765 0.768 0.765 0.765 0.804 0.775 0.755 0.804 0.775 0.768 0.775 0.804 0.775 0.768 0.776 0.804 0.775 0.775 0.804 0.775 0.785	Returne data P3 0.338 0.338 0.338 0.338 0.359 0.500 0.500 0.500 0.775 0.938 0.775 0.938 0.775 0.338 0.775 0.338 0.791 0.791 0.792 0.338 0.792 0.8225 0.8225 0.8223 0.8225 0.8223 0.8236 0.8223 0.8237 0.8223 0.8238 0.8223 0.8237 0.8223 0.8238 0.8223 0.8237 0.8223 0.8238 0.8223 0.8238 0.8233 0.8238 0.8233 0.8238 0.8233 0.8238 0.8233 0.8238 0.8238
20 energy 22 0 923 0 923 0 923 0 620 0 620 0 846 0 846	30 homog P2 P2 0.958 0.958 0.958 0.907 0.917 0.917 0.917 0.979 0.979 0.979 0.979 0.979 0.979 0.979 0.979 0.973 0.972 0.973 0.973 0.973 0.972 0.973 0.972	P2 0.804 0.804 0.756 0.500 0.500 0.500 0.585 0.857 0.857 0.920 0.784 0.784 0.784 0.784 0.784 0.784 0.784 0.7988 0.7988 0.7988 0.7988 0.7988 0.79888 0.79888 0.7988 0.7988 0.7988 0.79888 0.79888 0.7988	F image (e P2 P2 0.851 0.855 0.866 0.855 0.866 0.865 0.749 0.632 0.749 0.822 0.885 0.885 0.885 0.885 0.85
P1 P1 0.944 0.944 0.944 0.944 0.944 0.944 0.944 0.944 0.944 0.944 0.944 0.944 0.944 0.944 0.944 0.944 0.944	P1 P1 P1 100 0.979 0.345 0.0855 1.000 0.978 0.958 0.0953 0.958 0.0973 0.9973 0.0973 0.978 0.0973 0.973 0.0973 0.973 0.1000 11.000 1.1000 11.000 1.1000 11.000 0.923 0.923 0.923 0.923 0.923 0.923 0.923 0.923 0.923 0.923 0.917 0.923 0.917 0.923 0.917 0.923 0.917 0.923 0.917 0.923 0.917 0.923 0.923 0.923 0.923 0.923 0.923 0.923	30.RGB ir P1 0.855 0.500 0.505 0.857 0.855 0.738 0.738 0.738 0.738 0.738 0.738 0.738 0.738 0.738 0.738 0.738 0.738 0.738 0.738 0.738 0.755 0.857 0.755 0.857 0.755 0.857 0.855 0.755 0.857 0.755 0.8570 0.8570 0.85700000000000000000000000000000000000	30 L*u*v' P1 0.825 0.904 0.735 0.735 0.735 0.735 0.735 0.736 0.736 0.736 0.736 0.736 0.736 0.736 0.736 0.736 0.844 0.840 0.940 0.840 0.7350 0.73500 0.73500 0.75500 0.75500000000000000000000000
accuracy s Classifier nalysis assifier t Analysis	accuracy s Lassifier nalysis assifier Analysis	score for s cLassifier nalysis t Analysis t Analysis	score for s ralysis f Analysis f Analysis
Table 4.5 Balanceria scorate for 30 energy image feature datasets for each constraints. P	Table A.C. IBAlanced accuracy serves for 30 homores AddBoot P. AddBoot P. AddBoot 1000 953 AddBoot 1000 953 Dammed Native Bayes 1000 953 Dammy Chandifier 1000 953 Dammy Chandifier 0.315 0.303 East The Posterging 0.315 0.303 East Theorem 0.315 0.903 East Theorem 0.315 0.903 East Theorem 0.315 0.903 East National Assistion 0.903 0.917 Monti Lyser Presention 0.901 0.901 Paster Agressive Chandier 0.903 0.917 Paster Agressive Chandier 0.903 0.912 Paster Agressive Chandier 0.903 0.903 Paster Agressive Chandier 0.903 0.903	Table 4.7 ALTROC score for -00 RGB image feature datasets for cell line MDA. Table 5.7 ALTROC score for -00 RGB image feature datasets for cell line MDA. AddBiost P.2 P.3	Table 4.81, ALTROC score for 40.1 ^{an} v ⁺¹ image feature datasets for cell line AD1 Ad4Boost PA PS PS PA PS PA PS PA
Table 4.5.1 [] Table 4.5.1 [] AdaBoost AdaBoost AdaBoost AdaBoost Extra Trees Extra Trees Extra Trees Extra Trees Extra Trees Caratient Boo Gradient Boo Caratient Boo Caratient Boo Caratient Boo Caratient Boo Caratient Boo Caratient Boo Caratient Boo Linear Disciple Cara Multi Layer F. Multi Layer Cara Multi Cara Mul	Table 4.0 [Jah Lastflers: Bernull Narre Bernull Narre Estuar Danie (Carlo Carlo Carlo Carlo Extra Trees Extra Trees Extra Trees Gradiern Board Lager Regen Mult Layer For Arter Cherno Mult Layer For Perception Quadran Forts Perception Quadran Forts Perception Strike Ages Perception Strike Ages Perception Strike Strike Ages Perception Strike Ages Perception Strike Strike Ages Perception Strike Ages Perception Strike Strike Ages Perception Strike Ages Perception Strike Ages Perception Strike Ages Perception Strike Ages Perception Strike Ages Perception Strike Ages Perception Strike Ages Perception Strike Ages Perception Strike Ages Perception Strike Ages Perception Strike Ages Perception Strike	Table 4.71 AU Classifiers AdaBost AdaBost Extra Trees Extra Trees Extra Trees Gardan Trees Gardan Laye Pe North Laye Pe North Laye Pe North Laye Pe Chadhar Des Chadhar Des Ch	Table 481 M. Table 481 M. Adabari Bata and a second the

Table 4.9 AUROC score for 30 contrast image feature datasets for cell line MDA-	0 contrast im	age featur	e datasets	tor cell li	ne MDA-	MD-231																	l						i
Classifiers					5 P6	P7	P8	6d	P10	IId	P12	P13	P14	PIS	P16	P17	P18	P19 F	P20 P	P21 P2	P22 P2	P23 P2	P24 P2	P25 P26	5 P27	P28	P29	P30	
AdaBoost	0.835 0.8	0.823 0.879		0.918 0.8	0.866 0.7	44 0.9						0.863	0.890	0.784	0.800	0.933			-										
Bernoulli Narve Bayes					8.0 CO/.0	0.0						0.890	0.567	0./39	608.0	0.955													0.0
Extra Trees						.6 0 65						0 500	0.811	169.0	0 500	0 500													
Gaussian Naive Baves						82 0.8						0.890	0.839	0.878	0.800	0.933													
Gradient Boosting					0.938 0.7	19 0.9.						0.836	0.892	0.819	0.784	0.967													0
Helstrom Quantum Classifier		0.765 0.912			0.820 0.8	16 0.8						0.836	0.839	0.864	0.784	0.833			-										-
Linear Discriminant Analysis						107 0.8						0.863	0.865	0.801	0.800	0.867			-								-		6
Logistic Regression						15 0.7						0.890	0.839	0.843	0.800	006.0			-						-				~
Muu Layer Perception	149.0	200 0 000 0 000 0			1.0 606.0	0.0 CI						90.00	C0/.0	100.0	0.000	CC4.0													
Treatest tregulous						2.0 2.0						190.0	010.0	01010	0.000														
Quadrance Laborationality Antalysis	0.0 110.0	0.000 0.000			2.0 101.0	0.0 00						190.0	570 0	0.004	0.000	CCC.0													
CAPACITY OF CAL						0.0 0V						100.0	598.0	0.810	0.000	000 0													
CTTM and		2/0.0 2/0.0		1.0 0.00	0 0 0000	0.0 01						0.026	202.0	610.0	0.000	0.023													
VINI - POIV	5.0 C58.0					6.0 01						0.830	0.920	0.764	0.800	0.500													0 1
O VM = 101 0.017 0.017 0.017 0.017 0.017 0.017 0.017 Results exclude Nearest Centroid, Passive Azzressive Classifier and Perceptro	Azzressive Class	ifier and Perce	d		0	10 44						0.00.0	6000	10/.0	000.0	0000													
	3																												
Table 4.10 AUROC score for 30 correlation image feature datasets for cell line M	30 correlatio	n image fe	ature data	tsets for c	ell line M	DA-MD-231																							
Classifiers	PI P2	P3	P4	PS	5 P6							P13	P14	PIS	P16	P17													
AdaBoost	0.757 0.5			0.728 0.801	0							0.820	0.863	0.466	0.664	0.897													0
Bernoulli Naive Bayes					0							0.734	0.756	0.421	0.500	0.740											Ŭ		0
Dummy Classifier					0							0.500	0.425	0.500	0.598	0.500											Č		-
Extra Trees					0							0.500	0.607	0.603	0.500	0.664											-		0
Gaussian Naive Bayes					0							0.781	0.778	0.481	0.648	0.780				-							Ŭ		0
Gradient Boosting					0							0.789	0.820	0.659	0.719	0.976											Ŭ		2
Helstrom Quantum Classifier					0						-	0.851	0.727	0.620	0.867	0.914											Ŭ		
Linear Discriminant Analysis					0 0	.615 0.754	54 0.558	58 0.500	00 0.681	0.669	0.650	0.882	0.677	0.481	0.576	0.741	0.784	0.553 0	0.882 0	0.505 0.	0.640 0.0	0.756 0.6	0.686 0.8	0.816 0.744	44 0.850	0 0.500	0.577	0.703	
Logistic Regression					0 0							0.906	0.655	0.481	0.433	0.741													
Multi Layer Perceptron					0 0							0.820	0.828	0./39	06/.0	0./41													
Nearest Neighbors					0 0							0.882	0.792	0.020	0.845	0.914													2
Quadratic Discriminant Analysis					0 0							0.882	0./13	0.540	0./30	0.882													~
Kandom Forest	2.0 028.0	CEC.0 68C.0	076-0 000		0.845 0.7	0.0						68/.0	C8870	6/0.0	0.600	606.0													
SVIM - mica						10 01						198.0	0 877	0 718	189.0	0.076													
stud - tetf				0.705 0.5	0.5666 0.5	00 07						052.0	0.748	0.460	190.0	0120													
2 V.M 101 0.100 0.100 0.100 0.100 0.100 Presitive Agreective Classifier and Percentrum	Antrecity Clace	ifier and Percen			000	00			10.0 0	0000		00110	0.710	001-0	0000	7117													
	3																												
Table 4.11 AUROC score for 30 energy image feature datasets for cell line MDA-	30 energy im	lage featur	e datasets	s for cell li	ine MDA-	NID-23																							
Classifiers	PI P2	P3	P4	P5	P6							P13	P14	PIS	P16	P17													
AdaBoost					0.00 0.8	36 0.9						0.969	0.944	0.890	0.920	0.935		-	-								-		00
Bernoulli Naive Bayes					0.833 0.8	61 0.8	-	-				0.938	0.933	0.841	0.528	0.916		-	-								-		-
Dummy Classifier						00 0.568	68 0.496	96 0.567	57 0.500	0.500	0.565	0.500	0.500	0.544	0.532	0.250	0.363	0.500 0	0.500 0	0.500 0.	0.535 0.1	0.500 0.5	0.547 0.5	0.500 0.638	38 0.500	0 0.417	7 0.505	5 0.500	0
Extra Trees						0.8	-					0.897	0.877	0.892	0.500	606.0			-								-		
Gaussian Narve Bayes					0.944 0.8	89 0.8						696.0	0.944	0.80/	0.088	186.0													
Gradient Boosting	20 1760	676.0 059.0 0 59.0 0			8.0 2/6.0 8.0	C.0 00						606.0	++6.0	1120	076.0	706.0													
Terstrom Quantum Classmer						CO 000						200.0	176.0	110.0	005.0	106.0													
Linear Discriminant Analysis		0.00 0.180			1 000 0.8	A-0 00 00						0/6.0	116.0	0.007	0.000	016.0													0 0
Multi I aver Percentron						80 08						696.0	0 944	298.0	0.983	0 981													
Nearest Neighbors						44 0.9	-					0.938	0.921	0.788	0.983	0.981			-										
Onadratic Discriminant Analysis						80 08						696.0	0 944	0.841	0.688	186.0													
Random Forest					1.000 0.9	17 0.9						0.969	0.944	0.865	0.938	0.981		-	-										60
SVM - linear	0.944 0.8	885 0.786			000 0.8	89 0.8						0.969	0.944	0.867	0.858	0.981		-	-										
SVM - poly		0.825 0.786			0.944 0.9	46 0.9						696.0	0.888	0.841	0.983	0.897		-									-		•
SVM - rbf	0.889 0.5	0.500 0.500			.000 0.8	89 0.9.						0.969	0.944	0.867	0.858	0.981													
Results exclude Nearest Centroid, Passive Aggressive Classifier and Perceptron	- Aggressive Class	ifter and Perce	eptron.																										
Table 4.12.1 AUROC score for 30 homogoneity image feature datasets for cell line	30 homogone	offy imago	feature da	itseate for	roll line	ma-Am-	120																l	l	l	l	l	l	ì
Classifiers	PI P1	P3	P4	P5	P6	PT	1					P13	P14	PIS	P16	P17	PIS	17	Γ		Γ	1				1	1		
AdaBoost	1.000 0.5	0.920 0.950	[000 0.9	0.911 0.9	77 0.8	[0.916	0.941	0.838	0.840	1.000	0.971	[-					-	-			5
Bernoulli Naive Bayes				1.000 0.921	0		0					0.921	1/6.0	0.947	0.862	0.921	126.0	0								-			0
Dummy Classifier	0.500 0.5	0.500 0.526	26 0.467		0.500 0.4		0					0.500	0.490	0.570	0.500	0.461	0.500	0	-			Ŭ				-	Ŭ		5
Extra Trees					0		-					0.946	0.971	0.947	0.500	1.000	0.912	-				-			-	-	-		2
Gaussian Naive Bayes	1.000 1.0	1.000 0.967	1	.000 0.821	0		Č					0.891	0.971	0.947	0.895	1.000	0.971	-								-	Ĩ		0
Gradient Boosting	0.958 0.5	0.938 0.950	-	000 000	0.944 0.9		0					0.891	0.912	0.865	0.895	0.978	1/20	Ű				Ŭ				-	Ĩ		1
Helstrom Quantum Classifier			-		0		0					0.891	1.000	0.947	0.967	1.000	1/20	-			-					-	Ŭ		
Linear Discriminant Analysis					1.		0					0.862	0.971	0.947	0.895	1.000	1/6.0	-								Ĩ	Ŭ		0
Logistic Regression					0		Ű					0.921	0.971	0.920	0.879	1.000	1/6.0	-	-							-			0
Multi Layer Perceptron					0		0					0.921	0.941	0.974	0.879	0.978	1/6-0	-	-							-			-
Nearest Neighbors			-		0		<u> </u>					0.950	0.912	0.893	0.895	0.978	1/6.0	-								-	-		-
Quadratic Discriminant Analysis					0 0							0.921	1.000	0.947	0.895	1.000	116.0												
SAMA linear	1 000 0.5	00001 500.0										176.0	146.0	0200	0.840	1 000	1/6.0												
SVM - poly				0.0 0.8	0	977 0.946		0.795	5 0.912	0.890	0.977	0.950	0.941	0.974	0.840	1.000	179.0	1.000 0	0.976 0	0.964 0.	0.947 0.9	0.977 0.9	0.971 0.5	779.0 7967	77 0.974	4 0.957	0.969	0.941	
SVM - rbf		-			0							0.921	0.971	0.920	0.500	0.500	179.0			-	-				-	-			0
Results exclude Nearest Centroid, Passive Aggressive Classifier	Aggressive Class	ifier and Perceptro	eptron.																										

230 0.887 0.887 0.897 0.844 0.844 0.9210 0.9210 0.9210 0.9210 0.92100000000000000000000000000000000000	P30 P30 0.832 0.832 0.832 0.832 0.837 0.856 0.856	P50 2.5724 2.5250 2.5250 2.7110 2.774 2.7744 2.7744 2.7744 2.7744 2.7762 2.773 2.773 2.773 2.774
P29 F P29 F P20 P2 P20 P20 P2 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P20 P	9459 1 958 98 98	P29 F2 0.754 0 0.754 0 0.754 0 0.750 0 0.500 0 0.500 0 0.702 0 0.702 0 0.702 0 0.702 0 0.753 0 0.753 0 0.753 0 0.753 0 0.753 0 0.753 0 0.753 0 0.753 0 0.753 0 0.753 0 0.754 0 0.852 0 0.853 0 0.854 0 0.855 0 0.856 0 0.8602 0
P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P28 P	Res Sec 2000 2000 2000	P38 0.0278 0.0278 0.0279 0.0200 0.0250 0.0259 0.05520 0.05520 0.05520 0.05520 0.05520 0.05520 0.05520 0.05520 0.055200 0.05520000000000
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P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P26 P	Astronom	P36
1255 1000 100	1 1 <th1< th=""> <th1< th=""> <th1< th=""> <th1< th=""></th1<></th1<></th1<></th1<>	P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P35 P
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		P24 9 0.611 7 0.661 7 0.661 7 0.661 7 0.661 8 0.530 9 0.530 9 0.530 9 0.530 9 0.530 9 0.535 9 0.532 9 0.532 9 0.532 9 0.532 9 0.532 9 0.532 9 0.541 9 0.541 9 0.541 9 0.541 9 0.541 9 0.541 9 0.541 9 0.541 9 0.541 9 0.541 9 0.541 9 0.541 9 0.541 9 0.541 9 0.541 9 0.541
P23 8 0.911 8 0.911 8 0.911 8 0.911 9 0.911 9 0.911 9 0.933 9 0.933 9 0.933 9 0.933 9 0.933 9 0.933 9 0.933 9 0.933 9 0.933 9 0.933 9 0.933 9 0.933 9 0.933 9 0.933 9 0.936 9 0.936 9 0.937 9 0.937 9 0.937 9 0.938 9 0.938 9 0.938 9 0.938 9 0.938 9 0.938 9 0.938 9 0.938	P23 P23 200 0.001 200 0.001 200 0.001 200 0.001 200 0.001 200 0.001 200 0.001 201	P23 20 0.878 20 0.6478 20 0.6478 20 0.6478 20 0.6478 20 0.6478 20 0.6478 20 0.6478 20 0.6478 20 0.6478 21 0.6478 22 0.6479 23 0.878 24 0.878 25 0.878 26 0.4753 28 0.6753 29 0.878 29 0.878 29 0.878 20 0.753 20 0.753 20 0.753 20 0.753 20 0.753 20 0.753 20 0.753 20 0.753 20 0.753 20 0.753 20 0.753 20 0.753 <tr< td=""></tr<>
P22 2 0.868 3 0.506 6 0.200 5 0.500 6 0.750 5 0.750 5 0.750 6 0.750 6 0.750 7 0.750 6 0.750 7 0.750 7 0.750 7 0.750 7 0.750 7 0.750 7 0.751 7 0.841 7 0.842 7 0.842 7 0.753 7 0.754 7 0.754 7 0.842 7 0.842 7 0.842 7 0.842	P23 P23 0.759 0.779 0.719 0.717 0.719 0.719 0.717 0.717 0.711 0.717 0.717 0.713 0.717 0.717 0.713 0.717 0.017 0.017 0.717 0.017 0.017 0.717 0.017 0.017 0.717 0.017 0.017 0.717 0.017 0.017 0.717 0.017 0.017 0.717 0.017 0.017 0.717 0.017 0.017 0.717 0.017 0.017 0.717 0.017 0.017 0.717 0.017 0.017 0.717 0.017 0.017 0.717 0.017 0.017 0.717 0.017 0.017 0.717 0.017 0.017 0.717 0.017 0.017 0.717 0.017 0.017 0.717 0.0	P23 6 732 6 0.732 6 0.732 7 0.623 8 0.740 9 0.773 9 0.773 9 0.773 9 0.773 9 0.773 9 0.773 9 0.773 9 0.775 9 0.775 9 0.775 9 0.776 9 0.776 9 0.776 9 0.776 1 0.776 7 0.776 7 0.776 7 0.777 7 0.777 7 0.777 7 0.777
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P20 0382 0782 0782 0782 0782 0.857 0.857 0.857 0.857 0.857 0.857 0.857 0.854 0.854 0.854 0.854 0.854 0.854 0.854 0.854 0.854 0.854 0.854 0.854 0.854 0.8570 0.8570 0.8570 0.8570 0.8570000000000	P10 P10 P10 P10 P10 P10 P10 P10	P20 0.734 0.754 0.547 0.547 0.547 0.547 0.547 0.547 0.547 0.547 0.5770 0.57700 0.57700 0.57700 0.57700 0.57700 0.57700 0.577000 0.5770000000000
P19 0.877 0.8875 0.8855 0.8813 0.8813 0.8813 0.8833 0.8833 0.8833 0.8833 0.8835 0.8855 0.88350 0.88350 0.88350000000000000000000000000000000000	P10 0.723 0.723 0.723 0.723 0.723 0.725 0.72	P19 0.754 0.0754 0.0500 0.0500 0.0570 0.0677 0.0677 0.0700 0.7750 0.7750 0.7750 0.7700 0.7700 0.7700 0.7700 0.7700 0.7700 0.7700 0.7700 0.7700 0.7700 0.7700 0.7700 0.7700 0.7700 0.7700 0.7700 0.7700 0.7700 0.7710 0.77000 0.77000 0.77000 0.77000 0.7700000000
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P16 0.885 0.885 0.580 0.885 0.5808 0.5808 0.923 0.923 0.923 0.923 0.923 0.923 0.923 0.923 0.923 0.923 0.923 0.885 0.923 0.885 0.923 0.885 0.923 0.885 0.923 0.885 0.923 0.885 0.923 0.885 0.923 0.923 0.923 0.923 0.923 0.923 0.923 0.923 0.923 0.923 0.923 0.923 0.923 0.923 0.923 0.930 0.930 0.930 0.805 0.500 0.805 0.500 0.805 0.500 0.805 0.500 0.805 0.500 0.805 0.500 0.805 0.500 0.805 0.500 0.805 0.500 0.805 0.500 0.805 0.9300 0.930 0.93000 0.93000 0.93000 0.93000 0.93000 0.930000000000	P16 916 0 880 0 980 0 992 0 992 0 984 0 988 0 978 0 9788 0 9788 0 9788 0 9788 0 9788 0 9788 0 9788 0 9788 0 978	P16 0.716 0.644 0.424 0.424 0.644 0.644 0.510 0.914 0.914 0.914 0.780 0.916 0.780 0.780 0.780 0.865 0.673 0.653 0.653 0.663 0.663 0.663 0.663 0.664 0.770 0.770 0.770 0.770 0.770 0.770 0.770 0.770 0.770 0.770 0.770 0.770 0.770 0.770 0.770 0.771 0.770 0.771 0.772 0.7720 0.77700 0.77200 0.77200 0.77700 0.77700 0.77700 0.77700 0.77700 0.77700 0.77700 0.77700 0.77700 0.77700 0.77700 0.77700 0.777000 0.77700 0.77700000000
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P13 0.721 0.728 0.590 0.644 0.788 0.788 0.788 0.788 0.738 0.738 0.755 0.755 0.755 0.755 0.755 0.765 0.765 0.765 0.765 0.765 0.765	P13 P13 0.760 0.760 0.576 0.576 0.577 0.576 0.576 0.576 0.577 0.576 0.577 0.576 0.577 0.577 0.577 0.576 0.577 0.577 0.577	P13 0.721 0.585 0.585 0.585 0.585 0.586 0.586 0.586 0.573 0.617 0.633 0.506 0.617 0.667 0.560 0.560 0.500 0.500
P12 0.755 0.763 0.543 0.543 0.612 0.763 0.541 0.763 0.840 0.781 0.784 0.781 0.784 0.783 0.784 0.783 0.784 0.783 0.784 0.783 0.784 0.783 0.783 0.783 0.783 0.783 0.783 0.783 0.784 0.783 0.784 0.783 0.784 0.784 0.784 0.784 0.784 0.784 0.784 0.784 0.784 0.784 0.784 0.784 0.784 0.7755 0.784 0.7755 0.7750 0.77550 0.77550 0.77550 0.77550 0.77550 0.77550 0.77550 0.77550 0.77500 0.77500 0.77500 0.77500 0.7750000000000	P12 P12 0 8301 0.8301 0 8301 0.8301 0 8301 0.8301 0 8301 0.8301 0 8301 0.7364 0 8301 0.7364 0 8301 0.7364 0 8301 0.8312 0 8301 0.8312 0 8301 0.8312 0 8301 0.8312 0 9301 0.8312 0 0.841 0.8412 0 0.841 0.8412 0 0.841 0.8414 0 0.841 0.8414 0 0.841 0.8414 0 0.841 0.8414 0 0.841 0.8414 0 0.841 0.8414 0 0.841 0.8414 0 0.8414 0.8414 0 0.8414 0.8414	P12 0.744 0.500 0.500 0.500 0.674 0.674 0.674 0.674 0.674 0.677 0.688 0.677 0.671 0.871 0.811 0.742 0.677 0.677
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Ime U87. P6 0.806 0.806 0.411 0.411 0.500 0.733 0.411 0.734 0.500 0.741 0.506 0.741 0.506 0.741 0.6596 0.741 0.6596 0.741 0.741 0.741 0.741 0.741 0.741 0.741 0.767 0.741 0.767 0.741 0.763 0.741 0.767 0.741 0.767 0.768 0.774 0.774 0.774	All line 13 P. Self P. Self	r cell line P6 0.772 0.500 0.500 0.500 0.500 0.817 0.817 0.817 0.817 0.585 0.585 0.585 0.595 0.500 0.500
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Table 4.1.31 Balanced scenacy score for 30. NCB image feature directed for edit AdaBoost P1 P2 P3 P3 P3 AdaBoost 0.87 0.87 0.81 0.91 P2 P3 P3 P5 AdaBoost 0.87 0.72 0.73 0.73 0.73 0.75	Table 4.1 Balanced actures store for 30.1 ······ inage forture datasets for Allocation acturates store for 30.1 ······ inage forture datasets for Allocation acturates acturates for allocation acturates acturates for allocation acturates acturates for allocation acturates a	Table 4.16 [] Balanced accuracy score for 30 correlation image feature chravet for Additional accuracy score for 30 correlation image feature chravet for Additional Xare Party 2019 Party
Table 4.13. Balanced Classifiers Additors Damay Classifier Baronill Nucke Bayes Damay Classifier Eran Trees Canacian Nucke Bayes Gaussian Nucke Bayes Gaussian Nucke Bayes Gaussian Nucke Class Paster Angester Descrimtant Paster Angester Class Paster Angester Angester Angester Class Paster Angester Angester Class Paster Angester Angester Angester Class Paster Angester Angester Angester Angester Class Paster Angester Anges	Table 4.14 [Bahanced Lessifiers and the second seco	Table 4.16 [Balmeed Classifiers Addition: Barrond Nicke Bayes Darmon (Lassifier Exama Trees) Darmon (Lassifier Exama Trees) Candent Boyening Candent Boyening Darger (Candent Boyening Candent Boyening Candent Boyening Candent Boyening Candent Boyening Candent Boyening Candent Boyening Candent Development Candent Boyening Candent Development Candent Boyening Candent Development Candent Boyening Candent Development Candent Boyening Candent Development Candent Boyening Candent Development Candent Develop
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P30 P31 P31 P32 P33 P34	P30 P31 1 <td>P30 9 0.837 7 0.507 7 0.500 7 0.500 7 0.500 7 0.500 7 0.500 7 0.500 7 0.500 7 0.500 7 0.500 7 0.500 7 0.500 7 0.500 7 0.500 7 0.500 8 0.914 9 0.914 9 0.914 9 0.914 9 0.914 9 0.914 9 0.914 9 0.914 9 0.914 9 0.914 9 0.935 9 0.935</td> <td> P30 P31 P32 P32</td>	P30 9 0.837 7 0.507 7 0.500 7 0.500 7 0.500 7 0.500 7 0.500 7 0.500 7 0.500 7 0.500 7 0.500 7 0.500 7 0.500 7 0.500 7 0.500 7 0.500 8 0.914 9 0.914 9 0.914 9 0.914 9 0.914 9 0.914 9 0.914 9 0.914 9 0.914 9 0.914 9 0.935 9 0.935	 P30 P31 P32 P32
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III line U8 P6 0.920 0.920 0.920 0.503 0.503 0.500 0.500 0.501 0.502 0.502 0.503 0.500 0.528 0.920 0.920 0.921 0.922 0.923 0.923 0.923 0.923 0.923 0.923 0.923 0.923 0.923 0.923 0.933 0.933 0.933 0.933 0.933 0.933 0.933 0.933 0.933 0.933	for cell II P6 1.1000 1.1000 0.500 0.500 0.500 0.500 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 0.959 0.929 0.929 0.929 0.929	MG P6 0.806 0.780 0.780 0.780 0.780 0.760 0.760 0.760 0.760 0.760 0.760 0.760 0.760 0.760 0.760 0.776	7.3.MG P.0 0.804 0.804 0.682 0.577 0.577 0.538 0.557 0.558 0.558 0.557 0.557 0.557 0.558 0.557 0.558 0.557 0.558 0.557 0.558 0.558 0.557 0.558 0.557 0.558 0.557 0.5588 0.5588 0.5588 0.5588 0.5588 0.5588 0.5588 0.5
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thure data P4 0.852 0.852 0.500 0.500 0.952 0.915 0.915 0.915 0.935 0.919 0.988 0.988 0.958 0.919 0.958 0.919	ge featury P4 0.588 0.588 0.588 0.500 0.500 0.500 0.858 0.858 0.858 0.828 0.828 0.828 0.828 0.828 0.828 0.828 0.828 0.828 0.828 0.828 0.828 0.828 0.828 0.828 0.828 0.828 0.829 0.829 0.829 0.841 0.841 0.828 0.829 0.829 0.829 0.829 0.829 0.829 0.829 0.829 0.828 0.828 0.828 0.828 0.828 0.828 0.828 0.828 0.828 0.828 0.828 0.828 0.856 0.858 0.856 0.858 0.856 0.866 0.9000 0.866 0.90000 0.866 0.90000000000000000000000000000000000	Is for cell P4 P4 0.731 0.731 0.731 0.731 0.731 0.750 0.750 0.750 0.750 0.750 0.750 0.771 0.771 0.771 0.771 0.771 0.771 0.770 0.750 0.750 0.750 0.750 0.750 0.750 0.750 0.750 0.750 0.750 0.750 0.750 0.750 0.750 0.750 0.771 0.771 0.750 0.750 0.750 0.750 0.750 0.750 0.750 0.751 0.750 0.770 0.750 0.770 0.750 0.750 0.750 0.750 0.750 0.750 0.750 0.750 0.750 0.770 0.750 0.770 0.770 0.770 0.770 0.770 0.770 0.770 0.770 0.770	sets for c P4 0.781 0.781 0.594 0.594 0.594 0.594 0.773 0.773 0.773 0.773 0.773 0.775 0.863 0.763 0.763 0.764 0.763 0.764 0.764 0.7754 0.7764 0.7763 0.7764 0.7763 0.7764 0.7763 0.7764 0.7763 0.7774 0.7773 0.7774 0.7773 0.7774 0.7773 0.7774 0.7773 0.77747 0.77747 0.77747 0.77747 0.7774777777777777777777777777777777777
Image fea P3	emetry ima P3 0.955 0.955 0.956 0.956 0.625 0.625 0.625 0.625 0.969 0.969 0.966 0.966 0.966 0.966 0.966 0.966 0.966 0.966 0.966 0.966 0.966 0.966 0.966 0.966 0.965 0.966 0.966 0.864 0.844 0.966 0.966 0.844 0.966 0.966 0.864 0.864 0.866 0.966 0.9388 0.9388 0.9388 0.9388 0.9388 0.9388 0.9388 0.9388	rre datase P3 0.833 0.0773 0.0768 0.0568 0.7500 0.710 0.710 0.710 0.712 0.813 0.813 0.813 0.813 0.813 0.813 0.813 0.813 0.875 0.835 0.750 0.770 0.750 0.833 0.750 0.833 0.833 0.833 0.835 0.833 0.835 0.833 0.835 0.833 0.835 0.833 0.8355 0.83550 0.83550 0.83550 0.83550000000000000000000000000000000000	dure data P3 0.900 0.500 0.500 0.500 0.500 0.500 0.931 0.931 0.931 0.931 0.931 0.950000000000
30 energy P2 0.940 0.940 0.502 0.542 0.542 0.542 0.958 0.918 0.958 0.918 0.918 0.9988 0.99888 0.99888 0.99888 0.99888 0.99888 0.99888 0.99888 0.99888 0.998888 0.998888 0.9988888 0.99888888 0.9988888888888888888888888888888888888	30 homog P2 0.970 0.970 0.985 0.	mage featurage f	P2 P2 P2 0.875 0.875 0.830 0.610 0.610 0.610 0.610 0.610 0.955 0.955 0.955 0.955 0.750 0.750 0.750 0.750 0.750 0.7550 0.7550 0.7550 0.7550 0.7550000000000
score for 3 P1 0.927 0.927 0.927 0.9264 0.944 0.944 0.944 0.946 0.944 0.946 0.944 0.946 0.944 0.946 0.944 0.946 0.944 0.946 0.944 0.944 0.946 0.944 0.946 0.944 0.944 0.946 0.944 0.944 0.944 0.946 0.944 0.946 0.944 0.944 0.946 0.944 0.946 0.944 0.946 0.944 0.946 0.944 0.944 0.944 0.944 0.944 0.944 0.944 0.944 0.944 0.944 0.946 0.944 0.944 0.946 0.944 0.946 0.944 0.946 0.944 0.946 0.944 0.944 0.946 0.944 0.946 0.944 0.946 0.944 0.944 0.946 0.944 0.946 0.944 0.946 0.944 0.946 0.944 0.946 0.946 0.944 0.946 0.944 0.946 0.944 0.944 0.946 0.944 0.927 0.944 0.927 0.944 0.927 0.9	xcore for 3 xcore for 3 yrl 0031 0031 0031 0031 0031 0031 0031 003	30 RGB in P1 0.865 0.865 0.512 0.512 0.512 0.512 0.512 0.512 0.512 0.512 0.512 0.513 0.535 0.53	30.L.u.*** P1 0.7793 0.7710 0.7710 0.750 0.772 0.7750 0.772 0.772 0.772 0.7750 0.7550 0.7750 0.7750 0.7750 0.7750 0.7750 0.7750 0.7550 0.7550 0.7550 0.7550 0.7550 0.7550 0.7550 0.7550 0.7550 0.7550 0.7550 0.7550 0.7550 0.7550 0.7550 0.7550 0.7550 0.75500 0.75500 0.75500 0.75500 0.7550000000000
accuracy bysis sifier haalysis Analysis	accuracy lysis sifier Analysis	score for . lysis . Analysis .	score for : lysis Analysis bid, Passive A
Balanced re Bayes sifter e Bayes string nantum Cl minant Ana minant Ana thors thors thors thors thors thors thore tho thore thore tho thore thore tho thore thore tho thore thore tho thore tho thore tho thore tho thore tho thore tho thore tho tho tho tho thore tho tho tho tho tho tho tho tho tho tho	Balanced sifer ve Bayes sifer Bayes ve Bayes antum Cl minant Ana ession bioors essive Clas essive clas	AUROC ve Bayes sifier sting sting minant Ana ession terceptron terceptron terceptron terceptron terceptron terceptron	AUROC ve Bayes sifier stree Bayes stree Bayes stree ession renception terception terception terception terception terception terception terception
Table 4.1 Tl Balanced accursts stores for 30 energy image feature datasets for ex- tractifications P.2 P.3	Table A.S. Balanced accuracy secsores for 20 monusplations P 2 Adations 91 92 Adations 031 070 Bernund ker: Bayes 030 035 Kara Franker 030 035 Kara Trees 030 035 Kara Trees 031 036 Guddan Davier Bayes 031 036 Kara Tecnola 031 036 Mult Jaye Pereption 031 035 Mult Jaye Pereption 031 035 Vector Adjaction Gammer 031 035 Vector Adjaction Gammer 031 035 Paster Adjaction Gammer 031 035	Table 4.10 AUKOC score for 30 RCB image feature datasets for cell line 1453. Table 4.10 AUKOC score for 30 RCB image feature datasets for cell line 1453. AddBooxt P2 P3 P4 P5 AddBooxt 0930 0730 0731 0733 0731 0733 Dammy Classifies 0930 0730 0733 0731 0733 0731 0733 0731 0733 0731 0733 0731 0733 0731 0733 0731 0733 0731 0733 0731 0733 0731 0733 0731 0733 0731 0733 0730 0730 0730 0730 0730 0730 0730 0730 0730 0731<	Table 4.201 AUROC score for 30 L * 1** image feature datasets for cell that 13 Additions P.2 P.3 P.3 P.3 P.3 Ord
N S S S R B B R N N N C E H J S N N R R B R S S S S	S S S S S S S S N N N N N N N N N N N N	W S S S S S S S S S S S S S S S S S S S	

Table 4.21 AUROC score for 30 contrast image feature datasets for cell line US7	30 contrast i	mage feat	ure datase	ets for cell	l line U87-	MG							14	1	N.	- 10	010	010		1					24	-			040
Classifiers	Id Los o	A 000	A Det	P4 P3	A LIG	P/ 0010		Na 100	014 D000	LIT CONT	714 J	P13	P14	CIA	0 TOD	/I/	814	VIA	074	171	77.4	6.740	P.24	C7.4	074	17.4	87.4	67.4	P30
Partoviti Noire Princ			0 200 0		140.0	0.0 0.0							110.0	101.0	0 737	0.720	0.02	1/0.0	192.0	1000	0.644	0.042	0.000	1.945	100.0	751.0			0.10.0
Dummer Charifter													9250	0050	101.0	105.0	0.570	0.500	10/00	0.534	0.500	1950	CL 10	1190	100.0	0 500			0.500
Futer Trans													0000	002.0	0000	0 631	0000	207.0	00000	009.0	0.643	0000	712.0	0.600	005.0	0 500			002.0
Gaussian Nativa Rauss					0 2020								00800	0.710	102.0	10C-0	102.0	0.00.0	0.750	0.885	158.0	0.731	0.780	0.735	P0.8.0	0.688			0.000
Gradiant Rooting					o e								0.800	0 866	171.0	0.850	0.235	0.847	0.944	0.885	0.800	0.843	1 877	0.055	0.870	0.757			0.787
Haletrom Quantum Classifier		0 800			o e								0 806	0.816	107.65	0.881	0.780	12200	0.950	128.0	0.874	198.0	0.875	079.0	70.80	0.688			181.0
I inear Discriminant Analysis													0 764	0.660	102.0	0 787	0 704	0.834	002.0	10.764	202.0	0.697	008.0	0.735	0.874	0.647			0.806
Logistic Regression					0								0 800	0 782	0 765	0 881	0 724	0 899	0 800	0.885	0 890	0 731	0.835	0 735	0 874	0.688			0 808
Multi Laver Perceptron	0.835 0.5	0.900 0.900	0.927 0.	0.800 0.6	0								0.800	0.787	0.765	0.881	0.835	0.877	0.900	0.907	0.826	0.843	0.811	0.955	0.879	0.688			0.808
Nearest Neighbors					0		-	-					0.907	0.871	0.798	0.826	0.835	0.863	0.850	0.835	0.826	0.881	0.882	0.860	0.895	0.730	-		0.782
Quadratic Discriminant Analysis	0.625 1.0	1.000 0.3	0.889 0.	0.700 0.6	0								0.764	0.719	0.721	0.786	0.760	0.921	0.750	0.885	0.780	0.692	0.851	0.610	0.824	0.688			0.781
Random Forest					0.795 0.		Ē						0.849	0.841	0.776	0.826	0.835	0.877	0.781	0.885	0.914	0.881	0.890	0.860	0.879	0.752			0.782
SVM - linear	0.772 0.5	0.900 0.900	0.833 0.		0								0.835	0.812	0.755	0.857	0.760	0.877	0.750	0.885	0.945	0.710	0.811	0.845	0.824	0.688			0.808
SVM - poly	0.730 0.8				0	833	-	-					0.800	0.799	0.721	0.857	0.760	0.899	0.800	0.871	0.866	0.899	0.851	0.625	0.824	0.647			0.806
SVM - rbf	0.813 0.5	0.500 0.	0.500 0.		0.500 0.	667 0.8							0.835	0.812	0.732	0.857	0.649	0.877	0.500	0.500	0.976	0.615	0.811	0.500	0.500	0.688			0.808
Results exclude Nearest Centroid, Passive Aggressive Classifier and Perceptro	: Aggressive Class	ifier and Perc	eptron.																										
1 able 4.22 AUKOC score for 30 correlation image feature datasets for cell line US	30 correlatio	n image l	eature dat	asets lor v	cell line L	5IV-/1	Ľ			I	ľ	Ľ		I													ľ	L	
Classifiers		La La	P3	P4 P5	P5 P6								P14	PIS	PIO	LIA	PIS	PIO	P20	121	121	P23	P24	P25	P20	124			P30
AdaBoost													0.753	0.00/	0./10	1000	0.814	0.754	0./34	0./19	0.752	0.85/	0.031	0.500	600.0	0.500			0.724
Bernoulli Narve Bayes						0.0 0.0							0.702	C2C.0	0.044	0.552	0/00	0.014	0.051	100.0	0.032	0.04/	0.001	00000	6000	0.520			0.805
Dummy Classifier					0.5/2 0.								005.0	875.0	955.0	0.500	075.0	005.0	875.0	0.403	965.0	005.0	00000	0.500	000.0	0.42/			0.400
EXUA Irees						C.U 000							005.0	00000	0.045	00000	0000	00000	00000	00000	262.0	0000	00000	00000	00000	005.0			180.0
Gaussian Naive Bayes	0./00 0./00		0./00 0.	0.0 140.0	0.045 0.0	700 0.0	0.0 202.0	C0070 21070	C60.U C0	/90.0 0	110.0 7	00000	201.0	/00:0	0.044	762.0	10/ 0	660.0	078.0	C60.0	10/-0	20.0	6712 0	00000	600.0	CC0.0	67/ 0	20/.0	0.650
Crament poosmig		10 1000			.0 10/.0	1.0 641							0+0.0	111.0	010.0	767.0	6400	70/.0	140.0	10/.0	101.0	CT0.0	11/.0	110.0	cc/.0	0+/-0			200.0
Heistrom Quantum Classiner			0.466 0.0	0.73 0.4		1.0 211							10/-0	0./1/	0.772	500 D	640.0	618.0	0.070	C0/ 0	500.0	816.0	10/.0	+C/ .0	1+0.0	878.0			10/10
Linear Discrimiant Analysis					0 2/20	1.0 280							70/ 0	180.0	0.123	100.0	170.0	160.0	0.070	6007 0	10/-0	0.000	870.0	002.0	CC0.0	202.0			+//-D
Logistic Regression						1.0 000							70/ 0	19000	440'D	1000	170'0	160.0	0.016	200.0	10/-0	0.40.0	00L 0	212.0	022.0	220 0			++/ · 0
Num Layer Perception					0.60.6	10 161							70/ 0	0.650	176.0	0.610	202.0	4C/ 0	016.0	071.0	0.627	0 909	0.765	01/10	400.0	0.000			P1/10
Contractive designed is			0 701.0	0 103 0		1.0 100							07/10	0000	10.0	610.0	01.0	00/-0	0.070	01410	750.0	0.757 0	C01.0	0.000	0000	0.000			701.0
Quamanc Discriminani Analysis						10 01/							70/.0	070.0	10.0	700.0	640°0	171.0	0.000	010.0	70/ 0	0000	700.0	700.0	600.0	C00.0			61/.0
Nandom Forest	0.500 0.500	0.700 0.700	100.0	0.600 0.20	0.500	1.0 CC8							171.0	0.697	007.0	CC/ 0	+6/ 0	0/0.0	0.012	07/10	0 707	0.695	44/ 0	C70'0	600.0	00/10			724
CATA Carden						1.0 DOC							0 700	209.0	0.000	2020	0.712	00200	0.016	TOD TO	0 702	0.762	01470	192.0	0.652	0 600			0.656
VIDI - INT OF				10 002 0	0 200 0 20	2.0 002							0.500	0.500	0.500	100.0	0.500	0.500	01210	1010	201.0	009.0	0.600	10/-0	005.0	0.500			0C0.0
SVIM = 101 0.200 0.125 0.000 Partite and Darante Cantrold Daratic American Classifier and Daranteon	America Clear	ifiar and Dare			000	10 000							0000	0000	000.0	670.0	0000	0000	0000	110.0	0.101	0.020	0000	000.0	000.0	000.0			
Table 4.23 AUROC score for 30 energy image feature datasets for cell line U87-1	30 energy im	iage featu	re dataset	ts for cell h	line U87-N	IC																							
Classifiers	PI P2	P	5 P.	4 P:	5 P.	5 P7							P14	PIS	P16	P17	P18	P19	P20	P21	P22	P23	P24	P25	P26	P27		29	P30
AdaBoost					752 0.	920 0.9							0.920	0.938	0.871	0.834	0.875	0.888	1.000	0.730	0.898	0.916	0.837	0.833	0.985	0.818		.800	0.925
Bernoulli Naive Bayes					0.500 0.	828 0.8	0						0.886	0.591	0.863	0.703	0.825	0.708	0.522	0.692	0.798	0.671	0.900	0.500	0.985	0.710	-	.684	0.841
Dumny Classifier					0.500 0.	500 0.5	Ű	-					0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.590	0.689	0.490	0.500	0.500	0.334	Ŭ	544	0.612
Extra Trees						500 0.5		-					0.500	0.500	0.693	0.500	0.500	0.577	0.500	0.500	0.500	0.500	0.797	0.500	0.500	0.500		200	0.500
Gaussian Naive Bayes		-				920 0.5	<u> </u>						0.869	0.906	0.871	0.682	0.850	0.876	1.000	0.877	0.838	0.839	0.920	0.785	0.985	0.916		.900	0.866
Gradient Boosting					0.752 0.0	938 0.5							0.920	606.0	0.828	0.812	0.832	176.0	0.980	0.772	0.850	10.00	0.898	0.833	C86.0	100.0		000	176.0
Heistrom Quantum Classmer		0 000 0	0.984 0.	2.0 202.0		200 00X							0.850	162.0	106.0	0.605	C76.0	0.8/0	0.500	07610	0.057	808.0	0.880	CUY.U	200.0	0.200		606	0.940
Lucar Discriminant Analysis I onietic Damazzion	0.077 0.0				0 757 0	255.0 0.0		0.00 0.000	408 0 6/ 6/	116.0 1	0.033	0.800	0.24.0	776.0	0.835	C4C-0	0000	0/8/0	0.058	0.813	108.0	0.858	0.830	00000	286.0	002.0	0 610 0	008.0	018/1
Multi Laver Perceptron						920 0.9							0.841	906 0	0.814	0.747	0.946	0.944	0.980	0.813	0.878	0.897	0.857	0.500	0.985	0.844		006	0.921
Nearest Neighbors					0.793 0.	920 0.9							0.841	0.938	0.828	0.747	0.946	0.888	0.960	0.877	0.795	0.832	0.900	0.718	1.000	0.844		984	0.971
Quadratic Discriminant Analysis						938 0.9.	0						0.869	0.891	0.871	0.682	0.850	0.853	1.000	0.918	197.0	0.858	0.920	0.618	1.000	0.871		006.0	0.866
Random Forest	0.871 0.5				0.752 0.	938 0.9	Ű						0.920	0.969	0.849	0.790	0.950	0.944	1.000	0.833	0.835	0.916	0.898	0.901	0.985	0.818	Ŭ	.800	0.950
SVM - linear		-			835 0.	920 0.9	Ű						0.903	906.0	0.835	0.703	0.900	0.898	0.958	0.813	0.858	0.858	0.900	0.833	0.985	0.935	Ŭ	006.	0.841
SVM - poly					0.752 0.	920	0				-		0.903	0.922	0.871	0.682	0.875	0.944	0.833	0.835	0.837	0.878	0.900	0.500	0.971	0.825		006'	0.841
SVM - rbt 0.92/ 0.900 0.984	0.92/ 0.5	0.900 0.0		0.903 0.3	835 0.	670 076							0.903	006:0	0.835	0.500	\$76.0	0.898	866.0	0.835	0.858	0.820	0.8/8	0.833	C86.0	0.500		006	0.841
THE REPORT OF THE PARTY AND THE PARTY AND THE PARTY	and Anternation of		monde																										
Table 4.24 AUROC score for 30 homog	ē	eity image	feature d	sets	r cell line	U87-MG																							
Classifiers	PI P2	P2	P3	P4	P5 P6	PT PT						P13	P14	PIS	PI6	P17	PIS	6Id	P20	P21	P22		P24	P25	P26	P27			P30
AdaBoost					0	904 0.5						0.819	0.918	0.789	6/6/0	0.940	0.810	0.888	0.935	676.0	0.802		0.8/0	C56.0	0.938	1.000			C68.0
Dernoull Naive Bayes	2.0 126.0	.0 0/610	0 005 0	0.144 0.1	0.789 1.	2.0 001						+025.0	0.835	1620	0.460	0+6.0	0.500	0.8/0	246.0	1003 0	0.695		101.0	CI/.0	0.500	0.500	-		0.510
Form Treas						2.0 005						005.0	1990	9590	204-0	0.010	0.803	C04-0	0 500	0.500	0.000		019 0	005.0	0.500	005.0			775-0
Gaussian Naive Baves					5 -							0.837	0.018	0.835	856.0	0.946	0.810	0.944	0.035	0.00	0.865		0.757	0.844	0.00	0.897			8980
Gradient Boosting												669 0	0.855	629.0	0.938	626.0	0.810	808.0	550 0	0.871	0.841		0.857	0.760	0.920	1 000			0.895
Helstrom Quantum Classifier							-					0.837	0.898	0.804	0.979	0.946	0.893	0.865	0.962	0.929	0.839		0.876	0.871	0.903	0.960			0.947
Linear Discriminant Analysis					1.	_	-		-			0.857	0.918	0.827	0.979	0.946	0.838	0.911	0.897	0.929	0.865		0.757	0.844	0.920	0.917			0.895
Logistic Regression	0.931 0.5	0.985 0.	0 696.0	0.903 0.6	0.653 1.	_	-		-			0.881	0.918	0.827	0.958	0.946	0.838	0.911	0.935	0.893	0.865		0.807	0.753	0.938	0.917			0.868
Multi Layer Perceptron				0.903 0.6	0		-		-		-	0.881	0.898	0.796	0.958	0.972	0.838	0.888	0.935	0.929	0.865		0.857	0.890	0.938	0.958	Ŭ		0.763
Nearest Neighbors					0.		-		-		-	0.878	0.898	0.890	0.902	0.972	0.920	0.888	0.916	0.893	0.865		0.876	0.844	0.920	0.813	Ŭ .		0.921
Quadratic Discriminant Analysis	2.0 155.0				-i -							0.8/8	0.918	668.0	866.0	0.940	C08.0	116.0	250.0	676.0	C08.0		10000	66/ 0	0.920	0.938			0.808
SVM - linear		0 285	0.938 0.	0.903 0.6	0.608 0.0							0.878	0.918	0.796	0.979	0.946	0.838	0.888	0.916	0.929	0.893		0.757	0.753	0.938	0.875			0.868
SVM - poly					0	964 0.935	-	0.920 0.943	43 0.863	3 0.938	8 0.891	0.881	0.708	0.796	0.941	0.946	0.865	0.888	0.916	0.929	0.867	0.871	0.857	0.753	0.875	0.813	0.861 0	0.722	0.842
SVM - rbf	0.500 1.0	0.000			0							0.861	0.918	0.796	0.979	0.946	0.864	0.888	0.935	0.500	0.865		0.757	0.753	0.938	0.500			0.868
Results exclude Nearest Centroid, Passive Aggressive Classifier and Perceptro	e Aggressive Class	ifier and Perc	eptron.																										

50 1000 10	20 20 20 20 20 20 20 20 20 20	230 0.591 0.591 0.544 0.421 0.421 0.422 0.422 0.422 0.522 0.522 0.522 0.522 0.522 0.522 0.522 0.522 0.522 0.522 0.522 0.522 0.522 0.522 0.522 0.522 0.522 0.421 0.421 0.522 0.526 0.526 0.526 0.526 0.526 0.526 0.526 0.526 0.526 0.56
Part 1 Part 	1 1	229 21 21 22 21 22 21 21 21 21 22 20 23 20 24 22 25 20 25 20 25 20 25 20 25 20 25 20 25 20 25 20 26 20 27 20 26 20 27 20 26 20 27 20 27 20 27 20 27 20 27 20 27 20 27 20 27 20 27 20 27 20 27 20 27 20 27 20 27
P28 P2 P28 10 0382 11 0382 12 0382 13 0382 13 0382 0.53 0353 0.53 0353 0.53 0354 13 0355 0.3 0354 10 0355 10 0358 10 0358 10 0358 10 0358 10 0358 10 0358 10 0358 10 0358 10 0358 10 0358 10 0358 10 0358 10 0359 10 0358 10 0358 10 0359 10 0359 10 0359 10 0359 10 0359 10 0359<	P25 P25 P25 P26 0.902 0.902 0.902 0.908 0.902 0.902 0.902 0.908 0.902 0.902 0.902 0.908 0.902 0.902 0.902 0.908 0.902 0.902 0.902 0.902 0.902 0.902 0.902 0.91 0.027 0.902 0.902 0.927 0.902 0.902 0.902 0.927 0.902 0.902 0.902 0.927 0.902 0.902 0.902 0.927 1.000 0.927 1.00 0.927 1.000 0.927 1.00 0.927 1.000 0.927 1.00 0.927 1.000 0.927 1.00 0.928 0.928 0.92 0.92 0.929 0.92 0.92 0.92 0.929 0.92 0.92 0.92 0.929 0.92 0.92 0.	P28 P2 7.72 0.172 0.172 7.72 0.550 0.550 0.550 7.533 0.172 0.172 0.172 7.530 0.550 0.550 0.550 0.550 7.533 0.17 0.15 0.16 0.16 0.16 7.538 0.17 0.16 0.17 0.16 0.16 0.17 0.16 0.17 0.16 0.16 0.17 0.16 0.17 0.16 0.17 0.16 0.17 0.16 0.17 0.16 0.17 0.16 0.17 0.16 0.17 0.17 0.16 0.17 0.16 0.17 0.16 0.17 0.16 0.17 0.16 0.16 0.17 0.16 0.16 0.16 0.16 0.16 0.16 0.16 0.16 0.17 0.16 0.16 0.16 0.16 0.16 0.16 0.16 0.16 0.16 0.16 0.16 0.16 0.16 0.16 0.16 0.16
P27 P27 P27 100000 10000 10000 10000 10000 10000 10000 10000 10000 10000 10000 10000 10000	P27 977 0.977 0.977 0.977 0.977 0.977 0.977 0.977 0.977 0.977 0.977 0.977 0.977 0.977 0.977 0.977 0.977 0.977 0.977 0.977 0.977 0.975 0.977 0.975 0.976 0.9776 0.9776 0.7766 0.9776 0.7776 0.9776 0.7776 0.9776 0.7776 0.9776 0.7776 0.9776 0.7776 0.9776 0.7776 0.9776 0.7776 0.9776 0.7776 0.9776 0.7776 0.9776 0.7776 0.9776 0.7786 0.9776 0.7786 0.9776 0.7786 0.9776 0.7786 0.9776 0.7786 0.9776 0.7786 0.9786	P27 5 0.306 6 0.500 1 0.500 2 0.500 2 0.500 3 0.500 3 0.550 3 0.553 3 0.553 3 0.553 3 0.553 3 0.553 3 0.553 3 0.553 3 0.553 4 0.556 5 0.563 5 0.563 6 0.563 7 0.563 7 0.563 6 0.563 7 0.563 7 0.563 7 0.563 7 0.563 7 0.563 7 0.563 7 0.563 7 0.563 7 0.563 7 0.563 7 0.563
P26 0.9471 0.9471 0.9471 0.9416 0.9416 0.9416 0.9416 0.9416 0.9416 0.9416 0.9416 0.9416 0.9416 0.9416 0.9416 0.975 0.975	P36 0.949 0.946 0.946 0.946	P26 0.765 0.765 0.563 0.563 0.569 0.543 0.543 0.543 0.551 0.555 0.551 0.5550 0.5550 0.5550 0.5550 0.5550 0.5
P25 0.970 0.5950 0.5050 0.5050 0.5050 0.5050 0.5050 0.5455 0.5455 0.5455 0.5455 0.5455 0.5455 0.9750 0.97000 0.9700 0.97000 0.97000 0.97000 0.97000 0.97000 0.97000 0.97000 0.970000000000	 233 234 235 235 236 236	P25 0.500 0.570 0.570 0.5000 0.500 0.50000 0.50000 0.50000 0.50000 0.50000 0.500000 0.50000 0.500000000 0.50000000000
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P23 0.884 0.884 0.884 0.4259 0.4259 0.4259 0.4259 0.4259 0.4259 0.784 0.785 0.786 0.787 0.77844 0.77844 0.77844 0.77844 0.77844 0.77844 0.77844 0.77844 0.77844 0.77844	P23 0.889 0.889 0.889 0.889 0.884 0.884 0.784 0.984 0.784 0.984 0.784 0.984 0.784 0.984 0.784 0.984 0.784 0.984 0.784 0.984 0.784 0.984 0.784 0.984 0.784 0.984 0.784 0.984 0.784 0.984 0.784 0.984 0.784 0.984 0.784 0.984 0.784 0.984 0.784 0.784 0.784 0.784 0.784 0.784 0.784 0.784 0.784 0.784 0.784 0.784 0.784 0.784 0.784 0.784 0.784 0.784 0.784 0.784 0.784 0.784 0.794 0.794	P23 0.469 0.469 0.500 0.584 0.584 0.584 0.584 0.584 0.584 0.584 0.584 0.784 0.784 0.784 0.784 0.784 0.784 0.5000 0.5000 0.5000 0.5000 0.5000 0.5000 0.5000 0.5000 0.5000 0.50000 0.500000000
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P21 0.929 0.929 0.929 0.531 0.929 0.912 0.923 0.929 0.9200 0.9200 0.9200 0.9200 0.9200 0.9200 0.9200 0.9200 0.9200 0.92000 0.92000 0.920000000000	P21 0.929 0.929 0.920 0.929 0.921 0.926 0.921 0.926 0.921 0.926 0.921 0.926 0.921 0.926 0.926 0.926 0.926 0.926 0.926 0.926	P21 0.762 0.762 0.762 0.5740 0.512 0.812 0.812 0.812 0.812 0.869 0.740 0.740 0.740 0.740 0.740 0.740 0.740 0.740 0.740 0.740 0.762 0.762 0.762 0.762 0.762 0.762 0.762 0.762 0.762 0.762 0.772 0.772 0.772 0.772 0.772 0.772 0.772 0.772 0.7720 0.7740 0.7770 0.7740 0.7770 0.7740 0.7770 0.7740 0.7770 0.7740 0.7770 0.7740 0.7770 0.77000 0.77000 0.77000 0.7700000000
P20 1.000 1.000 0.500 0.500 1.000 1.000 1.000 1.000 1.000 1.000 1.000 1.000 1.000 1.000 1.000 1.000	P20 1000 1000 1000 11000	P20 0.699 0.590 0.500 0.500 0.697 0.697 0.697 0.697 0.697 0.697 0.697 0.697 0.697 0.697 0.697 0.697 0.697 0.725 0.697 0.500 0.697 0.500 0.697 0.500 0.6970
P19 P100 1,000	P19 0 970 0 970 0 970 0 970 0 970 0 970 0 970 0 970 0 970 0 970 0 985 0 973 0 974 0 974 <td>P19 0.784 0.784 0.394 0.549 0.549 0.549 0.784 0.784 0.581 0.583 0.583 0.580 0.583 0.580 0.57800 0.57800 0.57800 0.57800 0.57800 0.57800 0.57800 0.57800000000000000000000000000000000000</td>	P19 0.784 0.784 0.394 0.549 0.549 0.549 0.784 0.784 0.581 0.583 0.583 0.580 0.583 0.580 0.57800 0.57800 0.57800 0.57800 0.57800 0.57800 0.57800 0.57800000000000000000000000000000000000
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P17 1.000 1.000 0.500 0.942 0.942 0.942 0.942 0.942 0.942 0.945 0.945 0.9550 0.9550 0.9550 0.9550000000000	P17 0.982 0.982 0.982 0.982 0.982 0.982 0.982 0.982 0.982 0.982 0.982 0.982 0.982 0.982 0.982 0.982 0.982 0.982 0.982 0.983 0.982 0.983 0.983 0.983 0.984 0.985 <td>P17 0.734 0.567 0.560 0.5667 0.5667 0.5682 0.832 0.832 0.832 0.832 0.767 0.767 0.767 0.767 0.767 0.767 0.768 0.560 0.5</td>	P17 0.734 0.567 0.560 0.5667 0.5667 0.5682 0.832 0.832 0.832 0.832 0.767 0.767 0.767 0.767 0.767 0.767 0.768 0.560 0.5
P16 0.800 0.509 0.500 0.500 0.9000 0.9000 0.9000 0.9000 0.9000 0.9000 0.9000 0.9000 0.9000 0.900000000	P16 P16 0.700 0.999 0.6969 0.6969 0.9900 0.9900 0.9900 0.9900 0.9900 0.9900 0.9900 0.559 0.550 0.5590000000000	P16 0.469 0.500 0.500 0.500 0.569 0.569 0.569 0.569 0.569 0.758 0.560 0.758 0.560 0.758 0.560 0.738 0.560 0.500 0.5500 0.5500 0.5500 0.5500 0.5500 0.5500 0.5500 0.5500 0.5500 0.5500000000
P15 0.9410.941 0.94100000000000000000000000000000000000	PI3 0.973 0.973 0.973 0.973 0.973 0.973 0.974 0.973 0.974 0.974 0.975 <td>P15 0.676 0.683 0.500 0.550 0.550 0.557 0.557 0.557 0.557 0.557 0.557 0.557 0.557 0.557 0.557 0.557 0.556 0.550 0.568 0.568 0.568 0.568 0.568 0.568 0.550 0.550 0.550 0.550 0.550 0.550 0.555 0.550 0.555 0.550 0.5550 0.5550 0.5550 0.5550 0.5550 0.55500000000</td>	P15 0.676 0.683 0.500 0.550 0.550 0.557 0.557 0.557 0.557 0.557 0.557 0.557 0.557 0.557 0.557 0.557 0.556 0.550 0.568 0.568 0.568 0.568 0.568 0.568 0.550 0.550 0.550 0.550 0.550 0.550 0.555 0.550 0.555 0.550 0.5550 0.5550 0.5550 0.5550 0.5550 0.55500000000
P14 0.590 0.552 0.552 0.958 0.958 0.958 0.940 1.000 1.000 0.960 0.952 0.952 0.955 0.956 0.960 0.9560 0.95600 0.95600 0.95600 0.95600000000000000000000000000000000000	Plat 0.940 0.94	P14 0.733 0.757 0.500 0.500 0.775 0.777 0.777 0.777 0.777 0.777 0.775 0.775 0.775 0.775 0.775 0.775 0.775
P13 0.941 0.677 0.6675 0.6979 0.979 0.979 0.979 0.979 0.979 0.979 0.979 0.979 0.979 0.979	Pla 0.979 0.970 0.970 0.970 0.970 0.970 0.970 0.970 0.970 0.970 0.970 0.970 0.970 0.970 0.970 0.970 <td>P13 0.795 0.709 0.707 0.819 0.819 0.819 0.819 0.742 0.742 0.742 0.742 0.742 0.742 0.742 0.742 0.742 0.742</td>	P13 0.795 0.709 0.707 0.819 0.819 0.819 0.819 0.742 0.742 0.742 0.742 0.742 0.742 0.742 0.742 0.742 0.742
P12 1,000 0,500 0,500 0,973 0,973 0,973 1,000 1,000 1,000 1,000 1,000 1,000 1,000 1,000 1,000	P13 1.000 <td>P12 0.780 0.780 0.490 0.651 0.651 0.742 0.742 0.742 0.742 0.742 0.742 0.742 0.742 0.742 0.742 0.742 0.653 0.653 0.653 0.653 0.653 0.653 0.653 0.653 0.653 0.653 0.653 0.653 0.653 0.653 0.653 0.653 0.653 0.653 0.653 0.760 0.653 0.760 0.770 0.770 0.770 0.770 0.770 0.770 0.770 0.770 0.770 0.770 0.770 0.77000 0.77000 0.77000 0.77000 0.77000 0.7700000000</td>	P12 0.780 0.780 0.490 0.651 0.651 0.742 0.742 0.742 0.742 0.742 0.742 0.742 0.742 0.742 0.742 0.742 0.653 0.653 0.653 0.653 0.653 0.653 0.653 0.653 0.653 0.653 0.653 0.653 0.653 0.653 0.653 0.653 0.653 0.653 0.653 0.760 0.653 0.760 0.770 0.770 0.770 0.770 0.770 0.770 0.770 0.770 0.770 0.770 0.770 0.77000 0.77000 0.77000 0.77000 0.77000 0.7700000000
P11 0.957 0.907 0.500 0.500 0.907 0.907 0.978 0.978 0.978 0.978 0.978 0.978 0.978 0.978 0.978	III 0000 0000 0000 0000 0000 0000 0000	P11 0.761 0.756 0.756 0.756 0.763 0.719 0.719 0.719 0.719 0.719 0.719 0.711 0.711 0.718 0.771 0.778 0.778 0.778
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P15 0.837 0.816 0.500 0.500 0.795 0.833 0.833 0.833 0.833 0.854 0.833 0.853 0.853 0.853 0.853 0.853 0.853 0.833 0.833 0.833 0.833	P15 P15 0.854 0.854 0.854 0.875 0.875 0.875 0.780 0.780 0.780 0.783 0.553 0.603 0.854 0.553 0.854 0.854 0.854 0.854 0.854 0.854 0.854	P15 0.941 0.941 0.941 0.941 0.941 0.941 0.979 0.979 0.979 0.979 0.979 0.979 0.979 0.979	P15 0.979 0.979 0.979 0.979 0.979 0.979 0.979 0.979 0.979 0.941 0.941 0.941 0.941 0.941 0.941 0.941 0.941 0.979
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sets for c P5 0.756 0.756 0.756 0.756 0.795 0.812 0.812 0.812 0.812 0.756 0.756 0.813 0.756 0.756 0.756 0.756 0.756 0.776 0.776 0.776 0.776 0.776 0.775	e dataset: P5 0.812 0.812 0.812 0.812 0.756 0.854 0.854 0.875 0.875 0.812 0.875 0.812 0.875 0.812 0.875 0.875 0.854 0.875 0.854 0.854	Ine AIC P5 0.941 0.958 0.923 0.923 0.923 0.923 0.923 0.923 0.923 0.923 0.923 0.923 0.923 0.920	ell line M P5 0.941 0.958 0.979 0.979 0.979 0.979 0.979 0.979 0.979 0.979 0.979 0.979 0.979 0.979 0.973 0.973 0.973 0.973 0.973 0.973 0.973 0.973 0.973 0.973 0.973 0.973 0.973 0.973 0.973 0.974 0.975 0.974 0.975 0.974 0.975 0.974 0.975 0.974 0.975 0.97700 0.9770 0.9770 0.97700 0.97700 0.97700 0.97700 0.97700 0.97700 0.97700 0.97700 0.97700 0.97700 0.97700 0.97700 0.97700 0.977000 0.9770000000000
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Table 4.30 Balanced accuracy score for 20 suergy image feature datasets for a databout P2 P3 P4 P5 AddBoot 071 012 070 054 074 075 AddBoot 071 012 070 054 074 075 AddBoot 071 0120 070 054 074 076 Dummy Charlier 0736 077 070 050 050 050 053 053 Dummy Charlier 0736 073 079 036	Table 4.20 Balaneed accurst score for 30 homose these flows: P P AdaBowr P P P P AdaBowr P P P P AdaBowr P P P P Baroud Mixer Esysts P P P P Baroud Mixer Esysts P P P P P Baroud Mixer Esysts P	Table 4.31 (AUROC score for 30 RCB image features datasets for cell line AUC Table 4.31 (AUROC score for 30 RCB image features datasets for cell line AUC Adablost P2 P3 P4 P5 Adablost 0.211 (200 0.944) 0.944 0.821 0.941 0.941 Adablost 0.321 (200 0.944) 0.941 0.820 0.941 0.941 0.941 Dimmy diativity line participant 0.371 (100 0.944) 0.941 0.921 0.941<	Table 4.31 AUROC score for 30 L w ⁺⁺ image feature datasets for cell line M Additors P
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Matrix I <th>Table 4.33 AUROC score for 30 contrast image feature datasets for cell line MCI</th> <th>30 contrast i</th> <th>mage feat</th> <th>ure datase</th> <th>ets for cell</th> <th>line MCI</th> <th></th>	Table 4.33 AUROC score for 30 contrast image feature datasets for cell line MCI	30 contrast i	mage feat	ure datase	ets for cell	line MCI																							
Math Math <th< th=""><th>Classifiers</th><th></th><th></th><th></th><th></th><th></th><th></th><th></th><th></th><th></th><th></th><th></th><th></th><th></th><th>P16</th><th>714</th><th>PIS</th><th>P19</th><th>P20</th><th>P21</th><th>P22</th><th>P23</th><th>P24</th><th>P25</th><th>P26</th><th>P27</th><th>P28</th><th>P29</th><th>P30</th></th<>	Classifiers														P16	714	PIS	P19	P20	P21	P22	P23	P24	P25	P26	P27	P28	P29	P30
Matrix Matrix<	AdaBoost														0.769	0.885	0.713	0.955	0.893	0.812	0.944	0.669	0.826	0.500	0.715	0.820	0.891	0.881	0.906
Mathematic Mathema	Dummy Classifier														0 406	0 500	00500	0.455	0 500	0 500	0.520	0 500	0.520	0 500	0 500	0 500	0 577	0 396	0.500
0 0	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 0	Gaussian Naive Bayes							0	-		-				0.500	0.598	0.503	0.735	0.532	0.417	0.750	0.584	0.631	0.610	0.538	0.776	0.722	0.631	0.684
0 0	Gradient Boosting	0.792						-					-		0.853	0.832	0.627	0.955	0.890	0.812	0.663	0.669	0.807	0.610	0.715	0.786	0.835	0.844	0.838
0 0	Helstrom Quantum Classifier	0.820							-						0.584	0.897	0.445	0.595	0.725	0.779	0.700	0.722	0.650	0.955	0.681	0.686	0.798	0.731	0.744
0 0	Linear Discriminant Analysis							-							005.0	002.0	744-0	202.0	100.0	075-0	00/-0	+80.0	0.00.0	010.0	0000	0///0	00/-0	16/-0	6000
900	Multi Torrer Decomposition														002.0	200 0	201-0	0.610	100.0	201.0	16/-0	10000	100.0	204.0	0007-0	01/0	0.017	10721	0000
0 0	Newset Neithhore														0 460	0 960	2020	0.770	0 784	0 740	0.813	0.584	0 663	509.0	0.837	0 786	0 767	0 944	0 877
10 <th>Ouadratic Discriminant Analysis</th> <th></th> <th>0.500</th> <th>0.624</th> <th>0.462</th> <th>0.735</th> <th>0.560</th> <th>0.488</th> <th>0.731</th> <th>0.584</th> <th>0.650</th> <th>0.610</th> <th>0.538</th> <th>0.776</th> <th>0.704</th> <th>0.681</th> <th>0.584</th>	Ouadratic Discriminant Analysis														0.500	0.624	0.462	0.735	0.560	0.488	0.731	0.584	0.650	0.610	0.538	0.776	0.704	0.681	0.584
1 1	Random Forest							0							0.569	0.878	0.650	0.970	0.918	0.812	0.794	0.600	0.813	0.500	0.857	0.786	0.762	0.931	0.838
10 10<	SVM - linear							0							0.500	0.500	0.462	0.735	0.529	0.433	0.650	0.500	0.631	0.500	0.538	0.776	0.780	0.731	0.600
N Col	SVM - poly														0.500	0.722	0.420	0.610	0.838	0.669	0.550	0.553	0.650	0.500	0.710	0.776	0.780	0.731	0.722
Image: black I	SVM - rbf	0.770 0	500 0.	~	-										0.500	0.533	0.462	0.860	0.529	0.500	0.550	0.500	0.631	0.500	0.504	0.500	0.762	0.531	0.500
M N	Results exclude Nearest Centroid, Passiv	e Aggressive Class	ifter and Perc	ceptron.																									
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0 0	Table 4.34 AUNUC SCORE IO	DU COLTEIAII	n image I	eature dat	asets lor	Cell IIIe N																							
000 <th0< th=""><th>Classifiers</th><th></th><th></th><th></th><th>4</th><th></th><th></th><th></th><th></th><th></th><th></th><th></th><th></th><th></th><th>PIO</th><th>11</th><th>814</th><th>PIV -</th><th>074</th><th>124</th><th></th><th>67.A</th><th>P.24</th><th>CL I</th><th>074</th><th>17.4</th><th>874</th><th>P.29</th><th>P30</th></th0<>	Classifiers				4										PIO	11	814	PIV -	074	124		67.A	P.24	CL I	074	17.4	874	P.29	P30
No <th>Adaboost</th> <th></th> <th></th> <th></th> <th></th> <th>o' '</th> <th>670</th> <th></th> <th></th> <th></th> <th></th> <th></th> <th></th> <th></th> <th>60+.0</th> <th>0.852</th> <th>0.00</th> <th>cn/ .0</th> <th>0.804</th> <th>671.0</th> <th>0.831</th> <th>60+.0</th> <th>0.844</th> <th>0000</th> <th>070'0</th> <th>0.080</th> <th>771.0</th> <th>610.0</th> <th>160.0</th>	Adaboost					o' '	670								60+.0	0.852	0.00	cn/ .0	0.804	671.0	0.831	60+.0	0.844	0000	070'0	0.080	771.0	610.0	160.0
0 0	Bernoulli Narve Bayes					5 ¢	10								00000	10.00	262.0	0.249	0.754	0.740	10/10	00000	0.752	0.200	0.031	0.830	0.055	0.402	0.022
0 0	Dummy Classiner					- · ·	9 8								0.000	040.0	00000	000.0	CIC.0	00000	660.0	000.0	000.0	000.0	000.0	080.0	c.cc.0	000.0	00000
300	EXTra Irees					5 ¢									005.0	00010	0000	005.0	00000	00000	00000	00000	0000	00000	0000	00000	00000	00000	00000
000	Gaussian Naive Bayes					5 0									0000	0.652	750.0	207.0	C7/-0	719.0	0.750	0.660	0.844	579.0	90000	0.676	0./80	0/50	770.0
000	Gradient Boosting	171.0				5 0									0.405	060.0	0.000	c0/.0	0.890	06/ 0	00/0	600.0	+6/.0	C70'0	0/0/0	0/0/0	1000	0.744	0000
000 <th0< th=""><th>Helstrom Quantum Classifier</th><th>0.790</th><th></th><th></th><th></th><th>o' '</th><th></th><th></th><th></th><th></th><th></th><th></th><th></th><th></th><th>0.859</th><th>0.677</th><th>0.590</th><th>0.814</th><th>0.808</th><th>0.812</th><th>0.894</th><th>0.738</th><th>0.813</th><th>0.955</th><th>0.676</th><th>0.653</th><th>0.873</th><th>0.733</th><th>0.628</th></th0<>	Helstrom Quantum Classifier	0.790				o' '									0.859	0.677	0.590	0.814	0.808	0.812	0.894	0.738	0.813	0.955	0.676	0.653	0.873	0.733	0.628
000	Linear Discriminant Analysis					o ,									0.600	0.832	0.547	0.595	0.697	0.812	0.781	0.569	0.794	0.500	0.543	0.653	0.817	0.676	0.622
No <th>Logistic Regression</th> <th></th> <th></th> <th></th> <th></th> <th>0, 0</th> <th></th> <th></th> <th></th> <th></th> <th></th> <th></th> <th></th> <th></th> <th>0.600</th> <th>0.832</th> <th>0.567</th> <th>0.595</th> <th>0.670</th> <th>0.669</th> <th>0.731</th> <th>0.584</th> <th>0.794</th> <th>0.500</th> <th>0.543</th> <th>0.620</th> <th>0.853</th> <th>0.594</th> <th>0.622</th>	Logistic Regression					0, 0									0.600	0.832	0.567	0.595	0.670	0.669	0.731	0.584	0.794	0.500	0.543	0.620	0.853	0.594	0.622
1000000000000100100100100	Multi Layer Perceptron					5 0									0000	01/10	0.485	080.0	0.805	1900 0	16/-0	0.700	0.844	C70'0	61/-0	0.050	0.080	000.0	770.0
10 <th>Nearest Neignbors</th> <th></th> <th></th> <th></th> <th></th> <th>5 0</th> <th></th> <th></th> <th></th> <th></th> <th></th> <th></th> <th></th> <th></th> <th>0.784</th> <th>77/0</th> <th>0.0/3</th> <th>0.545</th> <th>10.00</th> <th>006.0</th> <th>+6/.0</th> <th>00/.0</th> <th>0.844</th> <th>0000</th> <th>C0/.0</th> <th>6000</th> <th>C28.0</th> <th>61.0</th> <th>0./00</th>	Nearest Neignbors					5 0									0.784	77/0	0.0/3	0.545	10.00	006.0	+6/.0	00/.0	0.844	0000	C0/.0	6000	C28.0	61.0	0./00
000	Quadratic Discriminant Analysis					5 ¢									0.200	0.024	0.54/	0.550	160.0	0./40	0.831	0./84	0.820	C70.0	C80.0	500.0	1760	0.720	0000
m m	Kandom Forest						108 0.1								0.584	18/.0	0.008	0.845	0.835	0.1/4	0.800	0.584	0.794	0.000	CE/ 0	0.142	0./00	1881	0.000
000	SVIN - mean						750 0.5								002.0	202.0	010.0	202.0	01010	678.0	0 721	00270	0 076	202.0	100.0	0.620	1190	0029.0	770.0
3 1	Viot - INV S						-0 0C/								002.0	0.76/	744-0	C0/-0	01010	0.600	16/-0	005.0	07010	005.0	109.0	0.500	110.0	0/0.0	770.0
111 <th1< th=""><th>Davids acclude Navaet Cantroid Passio</th><th>- American Clear</th><th>ifiar and Darr</th><th></th><th></th><th>0</th><th>000</th><th></th><th></th><th></th><th></th><th></th><th></th><th></th><th>0000</th><th>0000</th><th>0000</th><th>1000</th><th>1110</th><th>00000</th><th>0000</th><th>0000</th><th>101</th><th>0000</th><th>100:0</th><th>000.0</th><th>202.0</th><th>0.1.0</th><th>0000</th></th1<>	Davids acclude Navaet Cantroid Passio	- American Clear	ifiar and Darr			0	000								0000	0000	0000	1000	1110	00000	0000	0000	101	0000	100:0	000.0	202.0	0.1.0	0000
77899																													
6FFBDDD <thd< th="">DDDDDD</thd<>	Table 4.35 AUROC score for	30 energy in	lage featur	re dataset	ts for cell h	ine MCF																							
900	Classifiers	PI P1	P	3 P4	4 P:	S P									P16	P17	P18	P19	P20	P21	P22	P23	P24	P25	P26	P27	P28	P29	P30
000	AdaBoost			964 0.	774 0.	756 0.		-					-		0.700	0.760	0.778	0.845	0.944	0.829	1.000	0.769	0.881	0.500	0.862	0.898	0.891	0.900	0.891
000 <th0< th=""><th>Bernoulli Naive Bayes</th><th></th><th></th><th></th><th></th><th></th><th>069</th><th>-</th><th></th><th></th><th></th><th></th><th></th><th></th><th>0.500</th><th>0.865</th><th>0.695</th><th>0.500</th><th>0.868</th><th>0.779</th><th>0.500</th><th>0.500</th><th>0.652</th><th>0.500</th><th>0.626</th><th>0.864</th><th>0.575</th><th>0.565</th><th>0.753</th></th0<>	Bernoulli Naive Bayes						069	-							0.500	0.865	0.695	0.500	0.868	0.779	0.500	0.500	0.652	0.500	0.626	0.864	0.575	0.565	0.753
30300	Dummy Classifier						Ĩ	-					-		0.500	0.502	0.500	0.500	0.402	0.707	0.500	0.422	0.415	0.500	0.500	0.395	0.500	0.500	0.500
11103033033036033033033033034034034033033033034	Extra Trees						500 0.5	-							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
1110011013<	Gaussian Naive Bayes						917 0.8	-							0.853	0.878	0.778	0.955	0.946	0.900	1.000	0.953	0.913	0.750	0.866	0.876	0.929	0.876	0.844
77.70.010.300.	Gradient Boosting	0.742					813 0.8	-							0.669	0.806	0.757	0.955	0.944	0.757	1.000	0.884	0.844	0.610	0.832	0.898	0.891	0.844	0.922
7770.9400.9400.9610.0710.0300.3610.3600.	Helstrom Quantum Classifier	0.826					792 0.5	-							0.953	0.923	0.800	606.0	0.920	0.867	0.781	0.884	0.926	0.924	0.950	0.876	0.873	0.876	0.797
100.460.5600.5600.5670.5840.5740.5730.5840.5730.5840.5730.5840.5840.5730.5840.5840.5740.5840.5940.5860.5940.5860.5940.5860.5940.5860.5940.5860.5940.5860.5940.5860.5940.5950.5940111 <td< th=""><th>Linear Discriminant Analysis</th><th>0.870</th><th></th><th></th><th></th><th></th><th>772 0.5</th><th>-</th><th></th><th></th><th></th><th></th><th></th><th></th><th>0.600</th><th>0.904</th><th>0.673</th><th>0.485</th><th>0.920</th><th>0.795</th><th>0.800</th><th>0.500</th><th>0.550</th><th>0.500</th><th>0.744</th><th>0.865</th><th>0.780</th><th>0.881</th><th>0.900</th></td<>	Linear Discriminant Analysis	0.870					772 0.5	-							0.600	0.904	0.673	0.485	0.920	0.795	0.800	0.500	0.550	0.500	0.744	0.865	0.780	0.881	0.900
110.490.50	Logistic Regression						897 0.5	-							0.884	0.878	0.778	0.830	0.946	0.795	0.950	0.500	0.863	0.500	0.891	0.898	0.946	0.863	0.838
30030030030030033	Multi Layer Perceptron						CI 2 012								50.0	169.0	CI/.0	0.830	0.803	718.0	000T	60/0	505.0	00/-0	168.0	0.8/0	0540	156.0	0.658
5 6 0	Nearest Neignbors						833 0.1								0.809	7470	CT/-0	CC6.0	0+4.0	0.885	0.881	006.0	0/9/0	C70'0	146.0	C09.0	70/.0	506.0	006.0
9.0 0.00	Quadratic Discriminant Analysis						0.0 856								0.833	8/8/0	86/.0	0.830	0.940	006.0	1.000	1.000	0.913	00/.0	168.0	0.898	6760	0.894	0.844
More More <th< th=""><th>Kandom Forest</th><th></th><th></th><th></th><th></th><th></th><th>1.0 000</th><th></th><th></th><th></th><th></th><th></th><th></th><th></th><th>600.0</th><th>0.010</th><th>7000</th><th>0/6.0</th><th>716.0</th><th>006.0</th><th>0060</th><th>00000</th><th>505.0</th><th>002.0</th><th>148.0</th><th>0.074</th><th>046.0</th><th>004.0</th><th>856.0</th></th<>	Kandom Forest						1.0 000								600.0	0.010	7000	0/6.0	716.0	006.0	0060	00000	505.0	002.0	148.0	0.074	046.0	004.0	856.0
Mo Mo<	The motion						0.0 0.0								00200	C720	0 757 0	2200	04400	100.0	0.000	10000	0.012	0000	1000	0.076	042.0	0.040	0 001
MCF No	SVM - rbf						980 0.8								0.500	0.500	0.778	0.955	0.946	0.867	1.000	0.584	0.944	0.500	0.925	0.876	0.929	0.826	0.844
NUT Nut <th>Results exclude Nearest Centroid, Passiv</th> <th>e Aggressive Class</th> <th>ifier and Perc</th> <th>reptron.</th> <th></th>	Results exclude Nearest Centroid, Passiv	e Aggressive Class	ifier and Perc	reptron.																									
M. M																													
1 1	Table 4.30 AUKUC score lo	730 homogen	eity image	e leature d	latasets lo	r cell line	MCF/	1	Ľ	ľ	Ľ	1	Ľ	Ľ	100	-	010	014	000	144			144		244		040	000	000
0.00 0.01 <th< th=""><th>AdaBoost</th><th></th><th></th><th></th><th></th><th>254 D</th><th>170 0.5</th><th></th><th></th><th></th><th></th><th></th><th></th><th></th><th>0144</th><th>0.837</th><th>0.758</th><th>0 845</th><th>0.944</th><th>0.870</th><th>0.081</th><th>0.960</th><th>0.063</th><th>0.500</th><th>0.971</th><th>0.900</th><th>0.835</th><th>0.900</th><th>0.038</th></th<>	AdaBoost					254 D	170 0.5								0144	0.837	0.758	0 845	0.944	0.870	0.081	0.960	0.063	0.500	0.971	0.900	0.835	0.900	0.038
0.00 0.01 <th< th=""><th>Demonth Morine Device</th><th></th><th></th><th></th><th></th><th></th><th>0 212</th><th></th><th></th><th></th><th></th><th></th><th></th><th></th><th>0.620</th><th>200.0</th><th>0.750</th><th>0 760</th><th>910.0</th><th>0.000</th><th>10.04</th><th>0 500</th><th>P D AAA</th><th>0 500</th><th>0.050</th><th>200.0</th><th>1100</th><th>02.0</th><th>0 020</th></th<>	Demonth Morine Device						0 212								0.620	200.0	0.750	0 760	910.0	0.000	10.04	0 500	P D AAA	0 500	0.050	200.0	1100	02.0	0 020
000 000 001 001 001 000 <th>Dummy Classifier</th> <th></th> <th></th> <th></th> <th></th> <th></th> <th>482 0.3</th> <th></th> <th></th> <th></th> <th></th> <th></th> <th></th> <th></th> <th>0.453</th> <th>0.500</th> <th>0.500</th> <th>0.500</th> <th>0.539</th> <th>0.500</th> <th>0.500</th> <th>0 500</th> <th>0 500</th> <th>0.470</th> <th>0.531</th> <th>0.474</th> <th>0 562</th> <th>0.500</th> <th>0 344</th>	Dummy Classifier						482 0.3								0.453	0.500	0.500	0.500	0.539	0.500	0.500	0 500	0 500	0.470	0.531	0.474	0 562	0.500	0 344
0 0	Extra Trees						200 005								0.500	0 500	0.500	0 500	0 500	0.500	0 500	0.500	0 500	0 500	0.500	0 508	0.500	0 500	0 500
070 070 054 073 054 073 050 074 051 073 070 051 070 051 070 051 070 051 071 070 051 071 071 070 051 071 070 051 071 070 051 071 071 070 051 071 <th>Gaussian Naive Baves</th> <th></th> <th></th> <th></th> <th></th> <th>0</th> <th></th> <th></th> <th></th> <th></th> <th></th> <th></th> <th></th> <th></th> <th>0.869</th> <th>0 897</th> <th>0 695</th> <th>0 970</th> <th>0.918</th> <th>678 0</th> <th>0 981</th> <th>0.869</th> <th>0 963</th> <th>0 750</th> <th>0.916</th> <th>0 898</th> <th>0.873</th> <th>0 894</th> <th>0.891</th>	Gaussian Naive Baves					0									0.869	0 897	0 695	0 970	0.918	678 0	0 981	0.869	0 963	0 750	0.916	0 898	0.873	0 894	0.891
0581 0584 077 078 0587 0	Gradient Boosting														0 769	0 741	1757 0	0.970	0.835	0.500	0 981	1 000	0.913	0.610	0 774	0 909	0 946	0.881	0 969
011 101 101 101 101 101 101 101 101 101	Helstrom Quantum Classifier	0.870				0	-			-					0.853	0.839	0.758	0.955	0.946	0.900	0.944	0.953	0.913	0.955	0.950	0.932	0.873	0.900	0.828
0.820 1000 1000 074 075 0.87 089 099 079 099 029 069 022 088 027 073 047 073 076 073 070 079 045 100 079 071 1000 079 091 050 071 078 083 035 056 070 056 070 059 070 079 071 078 086 055 050 074 058 035 056 050 074 058 035 056 050 059 050 059 056 050 074 058 055 050 059 050 059 050 059 056 050 059 050 059 056 050 059 050 059 056 050 059 050 059 050 059 050 059 050 059 050 059 050 059 050 059 050 059 050 059 050 059 050 050	Linear Discriminant Analysis	0.812				0		0		-					0.784	0.897	0.695	0.970	0.890	0.900	0.981	0.869	0.963	0.610	0.921	0.932	0.817	0.913	0.891
0.820 0.00 0.964 0.74 0.75 0.857 0.960 0.950 0.566 0.75 0.820 0.744 0.53 0.75 0.75 0.851 0.75 0.861 0.750 0.550 0.550 0.550 0.550 0.550 0.750 0.550	Logistic Regression					0	Ĩ	-	-		-		-		0.684	0.897	0.673	0.970	0.890	0.845	1.000	0.769	0.913	0.500	0.774	0.898	0.835	0.863	0.906
0 856 0 081 0 999 0 73 0 812 0 853 0 949 0 730 0 836 0 776 0 200 0 784 0 838 0 873 0 737 0 797 0 944 0 500 1 000 0 700 0 509 0 500 0 259 0 888 0 817 0 505 0 500 0	Multi Layer Perceptron					0	Ĩ		-	-			-		0.869	0.916	0.695	0.970	0.920	0.917	1.000	0.784	0.963	0.500	0.891	0.898	0.835	0.863	0.953
0.850 1000 0564 0.76 0.75 0.57 0.93 0.930 0.931 0.932 0.840 0.73 0.940 0.73 0.73 0.760 0.73 0.797 0.930 0.590 0.591 0.590 0.850 0.137 0.901 0.500 0.501 0.501 0.500 0.500 0.501 0.500 0.50	Nearest Neighbors		_			0		Ű		-					0.669	0.858	0.717	0.970	0.946	0.900	1.000	0.700	0.963	0.500	0.950	0.898	0.817	0.963	0.806
	Quadratic Discriminant Analysis					0	-	0							0.869	0.878	0.735	0.970	0.890	0.900	0.981	0.800	0.963	0.625	0.891	0.898	0.817	0.913	0.791
· Usak Usak Usak Usak Uzak Usak Usak Usak Usak Usak Uzak Uzak Uzak Uzak Uzak Uzak Uzak Usak Usak Uzak Uzak Uzak Uzak Uzak Uzak Uzak Uz	Random Forest					0 0		<u> </u>		-	-				0.769	0.878	0.673	0.970	0.944	0.883	0.981	0.700	0.963	0.500	0.862	0.898	0.817	0.900	0.938
000 0-01 0-01 0-01 0-01 0-01 0-01 0-01	SVM - mear						106								690 U	0.016	560.0	0/6/0	0.000	0.000	1 000	0.584	C04.0	005.0	208.0	0.027	0.873	199.0	0.940
	SVM - the					o e	2005								0.500	0.897	0.500	0.500	0.918	0000	0.500	0.684	005.0	0 500	0.832	220.0	005.0	0.913	906.0

P30 0.87100000000000000000000000000000000000	0¢4 0¢4 065 065 065 065 065 065 065 065	P30 0.871 0.6871 0.507 0.507 0.750 0.770 0.750 0.7700 0.7700 0.7700 0.7700 0.7700 0.7700 0.7700 0.7700 0.7700 0.7700 0.7700 0.7700 0.7700 0.7700 0.7700 0.77000 0.77000 0.77000 0.7700000000
P29 0.795 0.500 0.500 0.750 0.792 0.835 0.835 0.837 0.837 0.837 0.818 0.833 0.833 0.833 0.833 0.833 0.833 0.833 0.833 0.833 0.833 0.833	P2.0 0.739 0.739 0.813 0.813 0.813 0.815 0.815 0.815 0.815 0.815 0.815 0.815 0.815 0.815 0.815 0.815 0.815 0.815 0.815 0.815 0.815 0.815 0.815 0.815 0.815 0.815 0.815 0.815 0.815 0.815 0.815 0.815 0.815 0.815 0.817 0.815 0.817 0.815 0.817 0.815 0.817 0.815 0.817 0.815 0.918 0.916 0.918 0.916 0.918 0.916 0.918 0.916 0.918 0.916 0.918 0.916 0.918 0.916 0.918 0.916 0.918 0.916 <td>P29 0.813 0.813 0.753 0.753 0.773 0.773 0.7710 0.7710 0.773 0.775 0.775 0.775 0.775 0.775 0.7730 0.730 0.730 0.730 0.730 0.730 0.730 0.730 0.730</td>	P29 0.813 0.813 0.753 0.753 0.773 0.773 0.7710 0.7710 0.773 0.775 0.775 0.775 0.775 0.775 0.7730 0.730 0.730 0.730 0.730 0.730 0.730 0.730 0.730
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Tokk 4.3.1 Balanced accuracy score for 30 KCB Image feature datasets for est and allowing the score for 30 KCB Image feature datasets for est AdaBost P	Total 4.58. Balanced accuracy score for M_1 - M_2 - M_2 and	Table 4.61 Balanced accuracy score for 30 correlation image feature Classifiers P 27 P3
Table 4.37 Table 4.37 Ectables 2.35 Benold Xa Benold Xa Benold Xa Content Boo Helstrom Q Helstrom Q Helstrom Q Helstrom Q Helstrom Q Market S Market S Marke	Table 4.38 Classifiers is AdaBoort Trees Classifier Boo Burnmy Glass Trees Eran Trees Eran Trees Eran Trees Eran Discr Helteron G. Paster Agg Praster Agg Classifiers Trees Cable 100 (1997) 100 (1997) 10	Table 4.40) Table 4.40 Cassificrs Bernoull Xa Bernoull Xa Bernoull Xa Bernoull Xa Extra Trees Extra T

P1 P2 P1 P1 P2 1 1.000 1.000 1000 0.940 0.912 0 0.388 0.524 0 0.318 0.500 0 0.917 1.000 1 1.000 1.000 1	m Classifier 1.000 1.000 rt Analysis 0.960 1.000 a 1.000 1.000 1.000 ptron 1.000 1.000 9,000 1.000	1.000 1.000 0 1.000 1.000 0	1.000	1.000 1.000	Table 4.42 Balanced accuracy score for 30 homogene [lassifiers p] p3 p3	AdaBoost 1.000 1.000 1.000	0.500 0.500	1.000	um Classifier 1.000 1.000 um Classifier 1.000 1.000	dysis 1.000 1.000 1.000 1.000	Multi Layer Perceptron 1.000 1.000 0.947 Nearest Centroid 1.000 1.000 0.974	1.000 1.000 1.000	assive Aggressive Classifier 1.000 1.000 0.314 Perceptron 1.000 1.000 0.974	Quadratic Discriminant Analysis 1.000 1.000 1.000 Random Forest 1.000 1.000 1.000 1.000	1.000 1.000 1.000	00T 000T 000T	B image featu P2	0.917 0.950 ayes 0.917 0.875	0.500 0.440 0.917 0.875	Gaussian Naive Bayes 0.897 0.925 0.944 Gradient Boschief 0.917 0.950 1.000	um Classifier 0.917 0.950 (# Analysis 0.917 0.875 (0.917 0.950	0.917 0.925	Quadratic Discriminant Analysis 0.897 0.975 0.972 Random Forest 0.917 0.925 0.972	0.917 0.950 0.944 0.917 0.921 0.889	SVM - rbf 0.917 0.950 0.583 Results exclude Nearest Centroid, Passive Aggressive Classifier and Perceptron	Table 4.44 AUROC score for 30 L ⁺ u ⁺ v ⁺ image feature datasets for cell line U3 Classifiets P3 P3 P3 P3 P4 P5	0.917 0.950	0.500	Extra Trees 0.875 0.875 0.861 Gaussian Naive Baves 0.897 0.950 0.944	0.877 0.925 1	n Classifier 0.917 0.900 Analysis 0.917 0.900	0.897 0.950 0.897 0.950	0.917 0.900 0.975 0.975	0.917 0.925	719.0 029.0 /28.0 719.0 718.0 719.0 719.0 700.0 700.0	sive Classifier and F
Patentine Patentine Patentine Patentine Patentine Patentine Patentine 0.881 Patentine 0.827 Patentine 0.827 Patentine 0.827 Patentine 0.827 Patentine 0.500 Patentine 0.500 Patentine 0.500 Patentine 0.500 Patentine 0.500 Patentine 0.500				1.000 1.000	eneity image feature dataset D3 D4 D5	1.000 1.000				00 1.000 74 1.000	47 1.000 74 1.000			00 1.000 00 1.000		1.000	datasets for ce P4					44 0.955			44 0.955 89 0.909		re datasets for P4			61 0.818 44 0.955						0.955	ti di
P5 P5 0.967 0.955 0.494 0.967 1.000 1.000 0.933	1.000 1.000 1.000 0.955 0.955	1.000	1.000	1.000	re datase	0.967	0.500	796.0	1.000	0.967	1.000	0.967	CC670	0.967	796.0	0001	I line U2 P5	0.933	0.500	776.0	0.833	0.933	0.967	0.944	0.900	0.933	cell line l	0.933	0.967	0.833	196.0	0.867	0.967	006.0	1.000	0.867	
P6 P6 1.000 0.964 0.500 0.500 1.000 1.000	1.000 0.929 0.929 1.000 1.000	1.000	1.000	0.964	-	1.000	0.500	0.978	1.000	1.000	1.000	0.978	0.978	0.978	1.000		_	0.943	0.635	0.957	0.935	0.978	0.943	0.978	0.957 0.943	0.978		0.943	0.913	0.978	0.943	0.935	0.978 0.921	0.943	0.943	0.943	
251 P7 0.974 0.972 0.946 0.946 0.944	0.974 0.972 1.000 1.000 0.917	0.912	0.974	0.974	line U251	0.974	0.500	0.974	0.974	0.974	0.974	0.974	1.000	1.000	0.974	0.9/4	P7	0.895	0.377	0.895	0.895	0.947	0.921	0.947	0.921	0.947	14	0.920	0.895	0.895	0.920	0.947	0.947 0.947	0.921	0.920	0.947	
PS 1.000 0.933 0.933 0.967 1.000	0.967 0.933 0.967 0.967 0.933	796.0 796.0	1.000	1.000	ž	1.000	0.500	1.000	1.000	1.000	0.967	1.000	1.000	1.000	1.000	1.000	P8	0.886	0.494	0.955	606-0	0.932	0.932	0.955	0.932	0.909	84	0.888	0.500	0.921	0.888	0.932	0.921 0.932	0.898	0.921	0.932	
P9 1.000 0.885 0.500 0.981 1.000 1.000	0.981 0.923 0.962 0.962 0.942	1.000	1.000	1.000	g	1.000	0.329	1.000	1.000	0.981	0.981	1.000	0.962	1.000	1.000	1.000	P9	0.955	0.500	0.935	1.000	0.962	1.000	0.962	1.000	1.000	8	0.981	0.500	0.935	0.962	cc4:0	186.0	1.000	186.0	556.0 2550	
P10 1.000 0.971 0.500 0.941 1.000 1.000	0.971 0.971 0.971 0.916 0.916	0.900	1.000	1.000	OId	1.000	0.510	1.000	1.000	1.000	1.000	1.000	1.000	1.000	1.000	1.000	P10	0.866	0.857	1.000	0.950	0.975	0.975	1.000	0.975	0.975	P10	1/6.0	0.606	0.916 0.946	1/6.0	056.0	0.950	0.950	1.000	0.950	
P11 0.946 0.941 0.500 0.853 0.941 0.975	0.971 0.941 0.941 0.941 0.941	0.912 0.941	0.941	0.971	IId	0.975	0.500	0.946	0.975	0.975	0.975	0.975	176.0	0.946 0.971	1.000	1:000	111	0.900	0.500	0.900	0.925	0.866	0.900	0.925	0.871	0.896	IId	168.0	0.500	006.0	168.0	0.841	0.896	0.925	168.0	006-0	
P12 1.000 0.807 0.500 1.000 1.000	1.000 0.983 1.000 0.950	1.000	1.000	1.000	214	1.000	0.707	1.000	1.000	1.000	1.000	1.000	1.000	1.000	1.000	1.000	P12	0.929	0.929	0.929	0.929	0.929	0.929	0.929	0.929	0.929	P12	0.929	0.500	0.929	0.929	0.929	0.929	0.929	0.929	0.929	
P13 1.000 0.500 0.929 1.000	1.000 0.976 1.000 1.000 0.952	1.000	1.000	1.000	PI3	0.952	0.500	1.000	1.000	1.000	1.000	1.000	0.929	1.000	1.000	1.000	P13	0.906	0.500	0.945	906.0	0.897	0.882	0.921	0.945	0.945	P13	0.921	0.500	0.747 0.890	0.897	0.906 0.906	0.874	0.906	0.945	068.0	
P14 0.977 0.967 0.548 0.977 0.977 0.977	1.000 0.967 0.977 0.933	0.977	0.977 0.977	0.977	P14	0.955	0.500	776.0	1.000	1.000	0.977	1.000	116.0	779.0	772.0	1.000	P14	0.932	0.921	0.921	0.955	0.955	0.955	776.0	0.955 0.855	0.955	p14	0.888	0.955	0.955	116.0	0.955	0.977	0.888	0.944	cc60	
P15 0.962 0.920 0.500 0.923 0.962 0.962 0.941	0.920 0.917 0.941 0.941 0.938	0.941 0.885	0.962	0.920	PIS	0.962	0.659	0.962	0.962	0.962	0.938	0.962	0.864	0.962	0.962	706.0	PIS	0.923	0.540	0.902	0.923	0.923	0.923	0.923	0.923	0.923	PIS	0.923	0.500	0.885	0.902	0.923	0.923	0.920	0.923	0.902	
P16 0.925 0.887 0.500 0.528 0.928 0.950	0.946 0.887 0.900 0.946 0.782	0.946	0.950 0.925	0.946	91d	0.950	0.565	0.950	0.975	0.946	0.921	0.950	006-0	0.950	0.946 0.950	C/670	P16	0.896 0.846	0.500	0.896	006.0	0.871	0.900	0.871	0.871	0.871	P16	0.841	0.846	0.871	0.866	0.900	0.871 0.871	0.871	0.896	0.871	
P17 0.938 0.958 0.500 0.671 0.938 0.938	0.920 0.928 0.958 0.958 0.958	0.766	0.958	0.958	P17	0.958	0.623	0.958	07670	0.938	0.878	0.899	0.804	0.958	0.958	806.0	P17	0.917	0.500	0.896	0.819	0.837	0.798	0.875	0.854 0.854	0.500	P17	0.896	0.854	0.896	0.857	0.810	0.896	0.816	968.0	c/8.0 862.0	
P18 10.927 0.927 0.615 0.615 0.500 0.500 0.500 0.927 0.500 0.927 0	0.927 0.927 0.927 0.927	0.927	0.927	0.927	PIS	0.927	0.542	0.927	0.927	0.927	0.927	0.927	0.927	0.927	0.927	176.0	P18	0.927	0.500	0.927	0.889	0.891	0.889	0.909	0.927	0.927	PIS	0.927	0.500	0.833	606.0	0.889	0.891	0.889	0.927	0.889	
P19 1 0.946 0.946 0.732 0.732 0.500 0.500 0.500 0.946 0.920 0.946	0.974 0.762 0.946 0.946 0.948	0.946	0.974	0.946	plq	0.946	0.462	0.946	0.946	0.946	0.946	0.972	0.946	0.946	0.972	0.940	P19	0.867	0.500	0.841	0.789	0.893	0.921	0.893	0.895 0.842	0.842	919	0.893	0.865	0.737	0.892	0.868	0.867	0.868	0.893	0.842	
P20 I 1.000 (0.848 (0.0.848 (0.0.786 (0.0.786 (0.0.786 (0.0.786 (0.0.786 (0.0.786 (0.0.786 (0.0.786 (0.0.978 (0															1.000						0.929								0.893 0.467 0								
P21 F 0.931 0 0.652 0 0.500 0 0.500 0 0.531 0 0.531 0 0.531 0 0.531 0 0.531 0 0.531 0 0.931 0													-		0.950 0						0.850 0								0.409 0								
P2.2 P 0.947 1 0.917 0 0.917 0 0.974 0 0.974 0 0.974 0 0.972 1 0.972 1 0.920 0											- 0				0.974 0						0.842 0								0.895 0	-	-	-					
P23 P23 P2 1.000 1. 1.886 0. 1.886 0. 1.2886 0. 1.2886 0. 1.000 0. 1.000 0. 0.000 0.000 0. 0.0000 0.0000 0. 0.0000 0.0000 0.0000 0.00000000			-						-			-			1.000 0.						0 7961								0.406 0.					-			
P24 P2 1.000 1 0.980 0 0.612 0 0.500 0 0.580 1 0.980 1 0.980 0								-							1.000 1						0.920 0								0.550 0								
P25 P 1.000 0 0.729 0 0.500 0 0.500 0 0.500 0 0.500 0 0.981 0														-	0 000.1						0 606.0								0.955 0								
P2.6 P 2.6 P 2.0.983 1.0.983 1.0.0.879 0.0.0.1455 0.0.0.0.1455 0.0.0.0.0.1455 0.0.0.0.0.0.1283 0.0.0.0.0.0.0.0.0.0.0.0.0.0.0.0.0.0.0.	000000		0 - 0					- 0 0		00	00			00	0.983 0.						0.840 0.								0.636 0.								
27 Pr 2000 0. 282 0. 2500 0. 264 0															982 0.						944 0.					-			0.413 0.					-			
28 2.505 2.500 2.500 2.500 2.500 2.895 2.824															0.840 0.897 0.912 0.958						0.752 0.813								0.786 0.813								
P29 0.835 0.817 0.527 0.500 0.500 0.938 0.938																																					

Matrix I I I I <th>Table 4.45 AUROC score for 30 contrast image fea</th> <th>r 30 contrast</th> <th>image fea</th> <th>nture datas</th> <th>ature datasets for cell line U251</th> <th>ill line U2:</th> <th>_</th> <th></th>	Table 4.45 AUROC score for 30 contrast image fea	r 30 contrast	image fea	nture datas	ature datasets for cell line U251	ill line U2:	_																							
Image: black Image: black <th< th=""><th>Classifiers</th><th>PI I</th><th>2</th><th></th><th></th><th></th><th></th><th></th><th></th><th></th><th></th><th></th><th></th><th></th><th></th><th></th><th></th><th></th><th></th><th>P20</th><th>P21</th><th>P22</th><th>P23</th><th>P24</th><th>P25</th><th>P26</th><th>P27</th><th>P28</th><th>P29</th><th>P30</th></th<>	Classifiers	PI I	2																	P20	P21	P22	P23	P24	P25	P26	P27	P28	P29	P30
model model <th>AdaBoost Remonifi Natire Rause</th> <th></th> <th></th> <th></th> <th></th> <th>0.933</th> <th></th> <th></th> <th></th> <th></th> <th></th> <th></th> <th></th> <th>-</th> <th></th> <th></th> <th></th> <th></th> <th></th> <th>0.957</th> <th>0.950</th> <th>0.918</th> <th>0.944</th> <th>0.958</th> <th>1.000</th> <th>0.983</th> <th>0.889</th> <th>0.983</th> <th>0.938</th> <th>0.982</th>	AdaBoost Remonifi Natire Rause					0.933								-						0.957	0.950	0.918	0.944	0.958	1.000	0.983	0.889	0.983	0.938	0.982
Math Math <th< th=""><th>Dummy Classifier</th><th></th><th></th><th></th><th></th><th>0.552</th><th>-</th><th>-</th><th></th><th></th><th></th><th>-</th><th></th><th>-</th><th></th><th>-</th><th></th><th></th><th></th><th>0.500</th><th>0.500</th><th>0.513</th><th>0.500</th><th>0.548</th><th>0.500</th><th>0.460</th><th>0.500</th><th>0.500</th><th>0.500</th><th>0.500</th></th<>	Dummy Classifier					0.552	-	-				-		-		-				0.500	0.500	0.513	0.500	0.548	0.500	0.460	0.500	0.500	0.500	0.500
0 0	Extra Trees						500	-				-		-		-				0.821	0.500	0.808	0.677	0.500	0.500	0.500	0.500	0.500	0.500	0.500
1 0	Gaussian Naive Bayes						000	-						-		-				0.929	0.981	0.974	0.977	0.960	1.000	0.983	0.964	0.950	0.875	0.982
0 0	Gradient Boosting	1.000	116				978	-								-				0.957	0.950	0.946	116.0	0.958	1.000	0.983	0.889	0.967	0.938	0.982
100 100 <th>Linear Discriminant Analysis</th> <th>0.958</th> <th></th> <th></th> <th></th> <th></th> <th>978</th> <th></th> <th>576-0</th> <th>0.900</th> <th>146.0</th> <th>116.0</th> <th>0.980</th> <th>0.000</th> <th>C06.0</th> <th>0 889</th> <th>0.643</th> <th>2/0.0</th> <th>0.815</th>	Linear Discriminant Analysis	0.958					978													576-0	0.900	146.0	116.0	0.980	0.000	C06.0	0 889	0.643	2/0.0	0.815
100 100 <th>Logistic Regression</th> <th></th> <th></th> <th></th> <th></th> <th></th> <th>978</th> <th>-</th> <th></th> <th></th> <th></th> <th>-</th> <th></th> <th>-</th> <th></th> <th>-</th> <th></th> <th></th> <th></th> <th>0.929</th> <th>0.950</th> <th>0.947</th> <th>0.944</th> <th>0.980</th> <th>1.000</th> <th>0.983</th> <th>0.944</th> <th>0.967</th> <th>0.875</th> <th>0.982</th>	Logistic Regression						978	-				-		-		-				0.929	0.950	0.947	0.944	0.980	1.000	0.983	0.944	0.967	0.875	0.982
0 0	Multi Layer Perceptron						978													0.957	1.000	0.947	176.0	0.980	0.981	0.983	0.944	0.967	0.855	0.982
0 100 011 010 011	Nearest Neighbors						507									-				0.964	0.950	0.974	116.0	0.938	1.000	0.983	1.000	0.857	0.875	0.982
1 100 000	Quadratic Discriminant Analysis					7.967		-												0.964	0.981	0.974	116.0	0.980	1.000	0.983	0.964	0.967	0.875	0.982
1 100	Random Forest						978	-						-		-				1.000	0.950	0.947	0.977	0.958	1.000	0.983	0.944	0.967	0.938	0.982
M M	SVM - linear						978	-												0.964	0.950	0.947	0.944	0.960	1.000	0.983	0.944	0.967	0.897	0.982
N N	SVM - poly	1.000 0					000	-						-		-				0.929	0.950	0.947	0.944	0.938	1.000	0.983	0.944	0.950	0.875	0.667
N N	SVM - rbf	0.500 0	500	0.500 1			978							-						0.964	0.500	0.974	0.500	0.980	0.818	0.500	0.500	0.643	0.855	0.500
M M	tesults exclude Nearest Centroid, Passi	e Aggressive Cla	ssifier and Pe	rceptron.																										
MMM																														
N F N	Table 4.46 AUROC score for	r 30 correlati	on image	feature da	atasets for	r cell line	_																							
0 0	Classifiers	PI F	12	3													P17	PIS	PI9	P20	P21	P22	P23	P24	P25	P26	P27	P28	P29	P30
000	daBoost						607	-		-						-	0.801	0.744	0.759	0.820	0.700	0.700	0.653	0.750	0.572	0.840	606.0	0.521	0.710	0.871
0 0	semoulli Naive Bayes						200	-		-						-	0.763	0.556	0.548	0.592	0.681	0.620	0.655	0.672	0.689	0.736	0.500	0.450	0.753	0.669
0 0	hummy Classifier						500	-									0.500	0.377	0.500	0.500	0.500	0.405	0.500	0.500	0.500	0.493	0.500	0.500	0.488	0.617
10 <th>xtra Trees</th> <th></th> <th></th> <th></th> <th></th> <th></th> <th>500</th> <th>-</th> <th></th> <th>-</th> <th></th> <th></th> <th></th> <th>-</th> <th></th> <th>-</th> <th>0.538</th> <th>0.500</th> <th>0.500</th> <th>0.500</th> <th>0.500</th> <th>0.753</th> <th>0.533</th> <th>0.500</th> <th>0.500</th> <th>0.500</th> <th>0.500</th> <th>0.500</th> <th>0.500</th> <th>0.500</th>	xtra Trees						500	-		-				-		-	0.538	0.500	0.500	0.500	0.500	0.753	0.533	0.500	0.500	0.500	0.500	0.500	0.500	0.500
11 11<	aussian Naive Bayes					0.742		-								-	0.739	0.502	0.577	0.635	0.813	0.730	0.755	0.732	0.591	0.769	0.760	0.643	0.730	0.760
11 <th>radient Boosting</th> <th>0.733</th> <th></th> <th></th> <th></th> <th></th> <th>921</th> <th>-</th> <th></th> <th>-</th> <th></th> <th></th> <th></th> <th></th> <th></th> <th>-</th> <th>0.798</th> <th>0.891</th> <th>0.788</th> <th>0.856</th> <th>0.794</th> <th>0.835</th> <th>0.730</th> <th>0.897</th> <th>0.864</th> <th>0.786</th> <th>0.871</th> <th>0.769</th> <th>0.710</th> <th>0.815</th>	radient Boosting	0.733					921	-		-						-	0.798	0.891	0.788	0.856	0.794	0.835	0.730	0.897	0.864	0.786	0.871	0.769	0.710	0.815
111	felstrom Quantum Classifier	0.525					728	-		-						-	0.753	0.891	0.784	0.899	0.863	0.756	0.844	0.710	0.825	0.912	0.871	0.643	0.733	606.0
0 0	near Discriminant Analysis						585	-	-	-				-		-	0.662	0.484	0.550	0.542	0.794	0.649	0.709	0.648	0.617	0.840	0.722	0.643	0.710	0.760
100.10	ogistic Regression						585	-		-						-	0.662	0.484	0.550	0.542	0.794	0.649	0.709	0.648	0.617	0.840	0.722	0.643	0.710	0.760
0 0	lulti Layer Perceptron						849	-	-	-				-		-	0.780	0.982	0.813	0.842	0.744	0.807	0.855	0.877	0.591	0.840	0.964	0.648	0.730	0.964
10 0	earest Neighbors						849										C6/.0	0./80	0.80/	0.803	0.803	65/.0	C08.0	158.0	0.//3	0.840	0.944	75/.0	0.730	0.835
0 0	uadratic Discriminant Analysis						728	-	-			-		-		-	0.756	0.593	0.604	0.679	0.826	0.703	0.755	0.752	0.636	0.752	0.742	0.610	0.730	0.760
0 0.11 0.	andom Forest						893	-								-	0.840	606.0	0.709	0.877	0.863	0.728	0.798	0.835	0.734	0.857	0.982	0.714	0.813	0.815
No No<	VM - Imear						200	-								-	0.623	0.484	0.550	0.542	0.763	0.648	0.721	0.708	0.617	0.840	0.500	0.500	0.730	09/.0
0 011 020 011 020 011 020 021	VIM - Poly						104										0.724	606.0	0.788	196.0	0.703	0.810	0.821	0.938	0.545 002 0	0.709	176.0	0./19	0.752	0.853
6 Pr 96 Pr 96 Pr 97 </th <th>V.M FDI</th> <th>0 0 000</th> <th>0 7 7 0 T</th> <th></th> <th></th> <th>077.0</th> <th></th> <th>0000</th> <th>070'0</th> <th>000.0</th> <th>C8C.0</th> <th>6/.0</th> <th>0.0/3</th> <th>0.152</th> <th>/00/0</th> <th>7/0.0</th> <th>701.0</th> <th>0000</th> <th>1/0.0</th> <th>0./30</th> <th>0./00</th>	V.M FDI	0 0 000	0 7 7 0 T			077.0											0000	070'0	000.0	C8C.0	6/.0	0.0/3	0.152	/00/0	7/0.0	701.0	0000	1/0.0	0./30	0./00
6 Fr 8 Pi Pi <th></th> <th>the supervise of</th> <th>T PUT PUT PUT PUT</th> <th>mandan</th> <th></th>		the supervise of	T PUT PUT PUT PUT	mandan																										
6FF	able 4.47 AUROC score fo	r 30 energy i	mage feat	ure datase	ets for cell	l line U25	_																							
0001 <th>lassifiers</th> <th>PI F</th> <th>2</th> <th>3</th> <th>P4 1</th> <th>54</th> <th>90</th> <th>[</th> <th></th> <th>P20</th> <th>P21</th> <th>P22</th> <th>P23</th> <th>P24</th> <th>P25</th> <th>P26</th> <th>P27</th> <th>P28</th> <th>P29</th> <th>P30</th>	lassifiers	PI F	2	3	P4 1	54	90	[P20	P21	P22	P23	P24	P25	P26	P27	P28	P29	P30
000	daBoost					1.967								-		-			-	1.000	0.931	0.947	1.000	1.000	1.000	0.983	1.000	0.824	0.835	0.982
0 0	ernoulli Naive Bayes					1.955	Ĩ							-		-			-	0.848	0.652	0.917	0.886	0.980	0.729	0.879	0.782	0.505	0.817	0.742
000	ummy Classifier					0.518	Ĩ					-		-		-			-	0.500	0.533	0.500	0.473	0.447	0.500	0.500	0.613	0.400	0.500	0.500
00 014 100 100 100 100 100 100 100 000 031 034 030 031 034 030	xtra Trees					0.800		Ŭ						-		-			-	0.929	0.500	0.974	0.500	0.500	0.500	0.500	0.500	0.500	0.500	0.500
00 031 030	aussian Naive Bayes	-			-	1.000								-					-	0.978	0.931	0.972	1.000	0.980	1.000	0.912	0.964	0.895	0.918	0.927
99907037037037037034044036	iradient Boosting	1.000 1	000				000							-		-			-	1.000	0.931	0.920	1.000	0.980	1.000	0.983	0.964	0.824	0.918	0.982
979 073 034 <th>Ielstrom Quantum Classifier</th> <th>0.917</th> <th></th> <th></th> <th></th> <th></th> <th>964</th> <th>Ű</th> <th></th> <th></th> <th></th> <th></th> <th></th> <th></th> <th></th> <th>-</th> <th></th> <th></th> <th></th> <th>0.978</th> <th>0.944</th> <th>0.974</th> <th>1.000</th> <th>0.980</th> <th>0.962</th> <th>0.983</th> <th>0.964</th> <th>0.862</th> <th>0.960</th> <th>0.927</th>	Ielstrom Quantum Classifier	0.917					964	Ű								-				0.978	0.944	0.974	1.000	0.980	0.962	0.983	0.964	0.862	0.960	0.927
39 101 031	inear Discriminant Analysis						929	<u> </u>						-		-			-	0.891	0.844	0.944	606.0	0.958	1.000	0.967	0.964	0.769	0.920	0.982
30.010.070.09.00.0010.000.010.00 <t< td=""><th>ogistic Regression</th><td></td><td></td><td></td><td></td><td></td><td>926</td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td>-</td><td></td><td></td><td>-</td><td>156.0</td><td>0.931</td><td>0.918</td><td>1.000</td><td>0.958</td><td>186.0</td><td>0.983</td><td>0.964</td><td>0.824</td><td>0.940</td><td>0.982</td></t<>	ogistic Regression						926									-			-	156.0	0.931	0.918	1.000	0.958	186.0	0.983	0.964	0.824	0.940	0.982
0 0	nuu Layet Fercepuon						06.1													0001	10.000	172.0	0.077	0.050	1 000	C00.0	1000	100.0	0 000	702.0
000 011 0100 0	tradication Disordinational Analysis	0.017				000														0.079	120.0	1 000	10001	00000	0.025	C00.0	100.0	10.005	00000	702.0
96 100 060 091 100 091 000 033 034 0391 0341 0391 0341	andom Forest	1.000				000								-		-			-	1.000	0.931	0.920	1.000	1.000	0.981	0.983	1.000	0.824	0.938	0.982
901 001 <th>VM - linear</th> <td></td> <td></td> <td></td> <td></td> <td>1.000</td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td>-</td> <td></td> <td></td> <td>-</td> <td>0.957</td> <td>0.931</td> <td>0.974</td> <td>1.000</td> <td>0.958</td> <td>1.000</td> <td>0.983</td> <td>0.964</td> <td>0.895</td> <td>0.898</td> <td>0.982</td>	VM - linear					1.000										-			-	0.957	0.931	0.974	1.000	0.958	1.000	0.983	0.964	0.895	0.898	0.982
30 100 087 083 071 0.50 100 100 030 100 <th>VM - poly</th> <td>-</td> <td>000</td> <td></td> <td>-</td> <td>1.000</td> <td></td> <td>Ŭ</td> <td>-</td> <td></td> <td></td> <td></td> <td></td> <td>-</td> <td></td> <td>-</td> <td></td> <td></td> <td>-</td> <td>1.000</td> <td>0.931</td> <td>0.974</td> <td>1.000</td> <td>0.958</td> <td>1.000</td> <td>0.983</td> <td>0.964</td> <td>0.879</td> <td>0.938</td> <td>0.982</td>	VM - poly	-	000		-	1.000		Ŭ	-					-		-			-	1.000	0.931	0.974	1.000	0.958	1.000	0.983	0.964	0.879	0.938	0.982
Name Name Par Par </td <th>VM - rbf</th> <td>1.000</td> <td>000</td> <td></td> <td>1.000</td> <td>1.000</td> <td></td> <td>-</td> <td>0.935</td> <td>0.931</td> <td>0.974</td> <td>1.000</td> <td>0.500</td> <td>1.000</td> <td>0.983</td> <td>0.500</td> <td>0.895</td> <td>0.940</td> <td>0.500</td>	VM - rbf	1.000	000		1.000	1.000													-	0.935	0.931	0.974	1.000	0.500	1.000	0.983	0.500	0.895	0.940	0.500
Item Item <th< td=""><th>esuits exclude Nearest Centroid, Pass.</th><td>e Aggressive Ula</td><td>ssifter and Pe</td><td>srceptron.</td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td></th<>	esuits exclude Nearest Centroid, Pass.	e Aggressive Ula	ssifter and Pe	srceptron.																										
No Pr Ns Po Po<	able 4.48 AUROC score fo	r 30 homogei	neity imag	e feature	datasets fi	or cell line	a U251																							
100 100 <th>Classifiers</th> <td>P1 F</td> <td>2</td> <td>P3 1</td> <td>P4 1</td> <td>54</td> <td>90</td> <td></td> <td>P19</td> <td>P20</td> <td>P21</td> <td>P22</td> <td>P23</td> <td>P24</td> <td>P25</td> <td>P26</td> <td>P27</td> <td>P28</td> <td>P29</td> <td>P30</td>	Classifiers	P1 F	2	P3 1	P4 1	54	90												P19	P20	P21	P22	P23	P24	P25	P26	P27	P28	P29	P30
1.00 1.00 <th< td=""><th>daBoost</th><td></td><td></td><td></td><td></td><td></td><td>000</td><td>-</td><td></td><td></td><td></td><td></td><td>-</td><td>Ŭ</td><td></td><td></td><td></td><td>Ű</td><td>0.946</td><td>0.957</td><td>0.931</td><td>0.974</td><td>1.000</td><td>1.000</td><td>1.000</td><td>0.983</td><td>0.982</td><td>0.845</td><td>0.938</td><td>0.982</td></th<>	daBoost						000	-					-	Ŭ				Ű	0.946	0.957	0.931	0.974	1.000	1.000	1.000	0.983	0.982	0.845	0.938	0.982
0431 100 0200 0500 0500 0500 0530 0531 0540 0530 0436 0536 0530 0530 0530 0530 0530 0530 05	semoulli Naive Bayes				-		978	-			-				-	-		0	0.972	1.000	0.926	0.974	1.000	0.980	0.962	0.983	0.946	0.681	0.960	0.982
0.11 01 097 077 056 057 056 057 100 100 100 100 100 100 051 078 077 052 051 051 051 051 051 071 052 051 051 051 052 053 051 051 051 051 051 051 051 051 051 051	Jummy Classifier						533	0		-	-	-	-		-	-	-	0	0.572	0.500	0.470	0.500	0.500	0.610	0.500	0.500	0.522	0.471	0.530	0.500
1100 1000 <th< td=""><th>xtra Trees</th><td></td><td></td><td>0.</td><td></td><td>0.967</td><td></td><td>0</td><td></td><td></td><td>-</td><td></td><td>-</td><td></td><td></td><td>-</td><td>-</td><td><u> </u></td><td>0.500</td><td>1.000</td><td>0.750</td><td>0.974</td><td>0.900</td><td>0.917</td><td>0.682</td><td>0.500</td><td>0.778</td><td>0.500</td><td>0.583</td><td>0.833</td></th<>	xtra Trees			0.		0.967		0			-		-			-	-	<u> </u>	0.500	1.000	0.750	0.974	0.900	0.917	0.682	0.500	0.778	0.500	0.583	0.833
100 100 100 001 100 007 100 007 100 007 100 007 100 007 000 007 000 007 000 007 000 007	Jaussian Naive Bayes					/06/0													0.940	1.000	056.0	0.9/4	1.000	0.980	1.000	0.983	786.0	C68.0	0.938	286.0
100 100 100 100 100 001 100 001 100 001 100 100 001 100 100 001 001 001 001 001 001 001 001 001 001 001 001 001 000 001 001 000 001 001 000 001 001 000 000 001 000 000 001 000 000 001 000	I oletrom Quantum Classifice	1 000		+/6.0		106.0													046.0	10001	156.0	076.0	116.0	0001	1 000	0.962	+06'0	208.0	0.980	786.0
100 100 031 100 031 100 031 100 031 100 <th>inear Discriminant Analysis</th> <th>1 000</th> <th></th> <th>000</th> <th></th> <th>1967</th> <th></th> <th>226.0</th> <th>1 000</th> <th>0.931</th> <th>0 974</th> <th>1 000</th> <th>1 000</th> <th>1 000</th> <th>0 983</th> <th>0 982</th> <th>0.840</th> <th>0.918</th> <th>0 982</th>	inear Discriminant Analysis	1 000		000		1967													226.0	1 000	0.931	0 974	1 000	1 000	1 000	0 983	0 982	0.840	0.918	0 982
100 100 100 100 100 097 100 100 097 100 100 100 100 100 077 096 097 096 097 096 097 100 100 100 100 100 100 100 100 100 10	ogistic Regression					1.967									-	-	-	0	0.946	0.978	0.950	0.974	1.000	1.000	1.000	0.983	0.982	0.840	0.918	0.982
100 100 100 100 007 077 077 100 100 100	Aulti Layer Perceptron			-		1.967		0	Ĩ					-	Ĩ	-	-	0	0.946	0.978	0.931	0.974	1.000	1.000	1.000	0.983	0.982	0.840	0.897	0.982
1000 1000 1000 1000 0597 1097 1097 1097 1000 1000 1000 1094 1000 1000 0577 0582 0593 0537 0544 1000 1000 1000 1000 0530 0532 0532 0535 0535 0537 0544 1000 1000 1000 1000 1000 0530 0532 0532 0535 0535 0535 1000 1000 1000 1000 1000	Vearest Neighbors	-					978	-			-				-	-	-	0	0.946	1.000	0.950	0.974	1.000	0.980	1.000	0.983	0.982	0.912	1.000	0.982
100 1000	Quadratic Discriminant Analysis						978	-			-			-	-	-	-	0	0.946	1.000	0.950	0.974	1.000	1.000	1.000	0.983	0.982	0.895	0.918	0.982
. 100 100 100 100 100 0.974 100 100 100 100 100 100 100 100 100 10	Kandom Forest					0.967					-					-			0.946	0.978	0.950	0.947	1.000	1.000	1.000	0.983	0.982	0.862	0.938	0.982
	s V.M Imear					/06'0													276.0	1.000	0.050 0	6/6/0	1 000	1 000	1 000	0.083	286.0	0.840	866.0	0.833
	VJM - rhf					000													0.500	0.500	0.500	0.974	0.500	0.500	1.000	0.500	0.500	0.840	0.500	0.500
		20																												

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