Additional file 1 — Supplementary methods

The eight NCI-60 cell-lines were used as the training data for marker set sizes 8, 16, 32, 64, and 96. The nine primary tumor cohorts from the TCGA data set were normalized to the NCI-60 data set using quantile normalization using NCI-60 data as the target distribution.

As discussed in the Validation Strategy section, a nested 5-fold cross validation with repeating the outer cross validation 100 times was utilized to train the NCI-60 data set, using the same feature selection and classifier methods described in the main text. Before running the nested k-fold cross validation, the training data was filtered to the top 1,000 features ranked using the F-statistic. The resulting top models used for feature selection and classification is shown in Supplementary Table ??. For each marker set size, the NCI-60 training data is used to extract the features and used as the reference data set for classification. The resulting accuracy of the validation experiment is defined as the number of correct predictions divided by the number of total predictions.

${\bf Additional\ file\ 1-Supplementary\ tables}$

Table 1: Datatypes Downloaded from CellMiner Data Type CellMiner Data Type Normalization Option CellMiner File Name

mRNA RNA: Affy HuEx 1.0 GCRMA nci60_RNA_Affy_HuEx_1.0_GCRMA.txt.zip CNV DNA: aCGH Agilent 44K AgelentFE nci60_DNA_aCGH_Agilent_44K_AgilentFE.txt		mRNA CNV microRNA	DNA: aCGH Agilent 44K RNA: microRNA OSU V3 chip	AgelentFE log2	nci60_DNA_aCGH_Agilent_44K_AgilentFE.txt. nci60_RNA_microRNA_OSU_V3_chip_log2.txt.z	
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Data files and normalization strategy used for each data set

Table 2: Cell Lines used in Analysis

Data Type	Breast	CNS	Colon	Non-Small Cell Lung	Leukemia	Melanoma	Ovarian	Renal
mRNA	BR.MCF7 BR. MDA_MB_231 BR. HS578T BR.BT_549 BR.T47D	CNS.SF_268 CNS.SF_295 CNS.SNB_19 CNS.SNB_75 CNS.U251	CO.COLO205 CO.HCC.2998 CO.HCT.116 CO.HCT.15 CO.HT29 CO.KM12 CO.SW_620	LC.A549	LE.CCRF_CEM LE.HL_60 LE.K_562 LE.MOLT_4 LE.RPMI_8226 LE.SR	ME.LOXIMVI ME.MALME.3M ME.MI4 ME.SK.MEL.2 ME.SK.MEL.28 ME.SK.MEL.28 ME.SK.MEL.5 ME.UACC.257 ME.UACC.62 ME.MDA.MB.435 ME.MDA.MB.435	OV.IGROV1 OV.OVCAR.3 OV.OVCAR.4 OV.OVCAR.5 OV.OVCAR.8 OV.SK.OV.3 OV.NCI_ADR_RES	RE.786_0 RE.A498 RE.ACHN RE.CAKI_1 RE.RXF_393 RE.SNI_2C RE.TK_10 RE.UO_31
Protein	BR.MCF7 BR.MDA_MB_231 BR.HS578T BR.BT_549 BR.T47D	CNS.SF_268 CNS.SF_295 CNS.SF_539 CNS.SNB_19 CNS.SNB_75 CNS.U251	CO.COLO205 CO.HCC.2998 CO.HCT.116 CO.HCT.15 CO.HT29 CO.KM12 CO.SW_620	LC.A549 LC.EKVX LC.HOP_62 LC.HOP_92 LC.NCI_H226 LC.NCI_H23 LC.NCI_H322M LC.NCI_H460 LC.NCI_H522	LE.CCRF_CEM LE.HL_60 LE.K_562 LE.MOLT_4 LE.RPMI_8226 LE.SR	ME.LOXIMVI ME.MALME_3M ME.M14 ME.SK_MEL_2 ME.SK_MEL_28 ME.SK_MEL_28 ME.UACC_257 ME.UACC_62 ME.MDA_MB_435 ME.MDA_MB_435	OV.IGROV1 OV.OVCAR_3 OV.OVCAR_4 OV.OVCAR_5 OV.OVCAR_8 OV.SK_OV_3 OV.NCI_ADR_RES	RE.786_0 RE.A498 RE.ACHN RE.CAKI_1 RE.RXF_393 RE.SNI_2C RE.TK_10 RE.UO_31
microRNA	BR.MCF7 BR.MDA_MB_231 BR.HS578T BR.BT_549 BR.T47D	CNS.SF_268 CNS.SF_295 CNS.SF_539 CNS.SNB_19 CNS.SNB_75 CNS.U251	CO.COLO205 CO.HCC.2998 CO.HCT.116 CO.HCT.15 CO.HT29 CO.KM12 CO.SW_620	LC.A549 LC.EKVX LC.HOP_62 LC.HOP_92 LC.NCI_H226 LC.NCI_H23 LC.NCI_H322M LC.NCI_H460 LC.NCI_H522	LE.CCRF_CEM LE.HL.60 LE.K.562 LE.MOLT_4 LE.RPMI_8226 LE.SR	ME.LOXIMVI ME.MALME.3M ME.M14 ME.SK.MEL.2 ME.SK.MEL.28 ME.SK.MEL.28 ME.UACC.257 ME.UACC.62 ME.MDA_MB.435 ME.MDA_MB.435	OV.IGROV1 OV.OVCAR.3 OV.OVCAR.4 OV.OVCAR.5 OV.OVCAR.8 OV.SK.OV.3 OV.NCI_ADR_RES	RE.786_0 RE.A498 RE.ACHN RE.CAKI_1 RE.RXF_393 RE.SN12C RE.TK_10 RE.UO_31
SNP	BR.MCF7 BR.MDA_MB_231 BR.HS578T BR.BT_549 BR.T47D	CNS.SF_268 CNS.SF_295 CNS.SF_539 CNS.SNB_19 CNS.SNB_75 CNS.U251	CO.COLO205 CO.HCC.2998 CO.HCT.116 CO.HCT.15 CO.HT29 CO.KM12 CO.SW_620	LC.A549 LC.BKVX LC.HOP_62 LC.HOP_92 LC.NCI_H226 LC.NCI_H23 LC.NCI_H322M LC.NCI_H460 LC.NCI_H522	LE.CCRF_CEM LE.HL_60 LE.K_562 LE.MOLT_4 LE.RPMI_8226 LE.SR	ME.LOXIMVI ME.MALME.3M ME.M14 ME.SK.MEL.2 ME.SK.MEL.28 ME.SK.MEL.28 ME.UACC.257 ME.UACC.62 ME.MDA.MB.435 ME.MDA.MB.435	OV.IGROV1 OV.OVCAR.3 OV.OVCAR.4 OV.OVCAR.5 OV.OVCAR.8 OV.SK.OV.3 OV.NCI_ADR_RES	RE.786_0 RE.A498 RE.ACHN RE.CAKI_1 RE.RXF_393 RE.SN12C RE.TK_10 RE.UO_31
CNV	BR.MCF7 BR.MDA_MB_231 BR.HS5581 BR.BT_549 BR.T47D	CNS.SF_268 CNS.SF_295 CNS.SF_539 CNS.SNB_19 CNS.SNB_75 CNS.U251	CO.COLO205 CO.HCC_2998 CO.HCT_116 CO.HCT_15 CO.HT29 CO.KM12 CO.SW_620		LE.CCRF_CEM LE.HL_60 LE.K_562 LE.MOLT_4 LE.RPMI_8226 LE.SR	ME.LOXIMVI ME.MALME_3M ME.ML ME.SK_MEL_2 ME.SK_MEL_2 ME.SK_MEL_5 ME.UACC_257 ME.UACC_62 ME.MDA_MB_435 ME.MDA_N	OV.IGROV1 OV.OVCAR.3 OV.OVCAR.4 OV.OVCAR.5 OV.OVCAR.8 OV.SK_OV.3 OV.NCI_ADR_RES	RE.786_0 RE.A498 RE.ACHN RE.SN12C RE.TK_10 RE.UO_31

The cell lines used in the cross validation experiment.

Table 3: Feature Selection Parameters

FS Metho	od Parameter Descript	ion Values	
Anova	NA	No parameters	
RFE	Estimator	The supervised learning estimator	Support Vector Classification (SVC) with a linear kernal for the decision functions
	Estimator Parameter - C	The penalty parameter of the error term	0.25, 1, 4, 16, 64, 256
Random Forest	Max Features	Function to determine the number of features to consider when looking for best split	sqrt and log2
	N Estimators	The number of trees to be used in the forest	10, 50, 100, 250
Extra Trees	Max Features	Function to determine the number of features to consider when looking for best split	sqrt and log2
	N Estimators	The number of trees to be used in the forest	10, 50, 100, 250

The tested parameters for each feature selection algorithm.

Table 4: Classification Parameters

CL Method	Parameter	Description	Values
Support Vector Machine	Kernel RBF Gamma RBF C Linear C	Function to use as a decision function Kernel coefficient for RBF Penalty Parameter of the error term Penalty Parameter of the error term	RBF and Linear 0.03125, 0.125, 0.5, 2, 8, 32, 128, and 512 0.03125, 0.125, 0.5, 2, 8, 32, 128, and 512 0.03125, 0.125, 0.5, 2, 8, 32, 128, and 512
Logistic Regression	Penalty C	Norm used in the penalization Inverse of regularization strength; smaller values specify stronger regularization	12 0.25, 1, 4, 16, 64, and 256
Decision Trees	Max Features	Function to determine the number of features to consider when looking for best split	sqrt and log2
Random Forest	Max Features	Function to determine the number of features to consider	sqrt and log2
	N Estimators	when looking for best split Number of trees to be used in the forest	10, 50, 100, 250
Extra Trees	Max Features	Function to determine the number of features to consider when looking for best split	sqrt and log2
	N Estimators	Number of trees to be used in the forest	10, 50, 100, 250
Gradient Boosting	N Boosting Stages Max Depth	Number of boosting stages to perform Maximum depth of the individual regression estimators	100, 250, and 500 3, 5, 7
K-nearest Neighbors	Compute Nearest Neighbor	Algorithm used to compute the nearest neighbors	BallTree and KDtree
	Distance Function Weight Function	Function used to calculate the distance A function to apply weights to the points	Euclidean and Manhattan Uniform and Inverse weighting based on the distance to their neighbors; the closer the distance the better the score.
Cosine	NA	No parameters	
Correlation	NA	No parameters	

The tested parameters for each classification algorithm.

Table 5: Mean AUC per Model Grouped by Platform and Marker Set Size

		•		nean ne	•		•								
			Mark	er Set S	Size n	ıRNA	Protein	micro	RNA S	NP CI	1V				
8	RFE	ET	0.9585	RFE	ET	0.9426	RFE	RF	0.8352	RFE	ET	0.8598	RFE	LR	0.7198
Ü	RFE	RF	0.9554	ET	ET	0.9394	RFE	SVM	0.8352	RFE	RF	0.8591	ET	LR	0.7115
	RFE	SVM	0.9521	RFE	RF	0.9382	RFE	KNN	0.8295	RFE	SVM	0.8321	RF	$_{ m LR}$	0.71
	RFE	LR	0.951	RF	ET	0.9376	RFE	ET	0.8275	ET	ET	0.8295	RFE	ET	0.691
	$_{ m RFE}$	KNN Cos	0.9467 0.9397	RF	RF RF	0.9312 0.9272	Anova Anova	SVM LR	0.8089 0.8051	RF	RF ET	0.8247 0.8177	RFE RFE	RF SVM	0.6802 0.6702
	RFE	Corr	0.9397	RFE	SVM	0.9272 0.922	RF	ET	0.8028	RF	RF	0.8177	ET	ET	0.6698
	ET	LR	0.9114	RF	SVM	0.9187	RF	RF	0.8027	RFE	KNN	0.8161	RFE	KNN	0.6681
	RF	$_{ m LR}$	0.9113	ET	$_{\rm SVM}$	0.9177	RFE	$_{ m LR}$	0.8021	RF	$_{ m LR}$	0.816	ET	RF	0.6648
	RF	ET	0.9093	RF	LR	0.9168	RF	LR	0.802	RFE	LR	0.8138	RF	ET	0.6631
	RFE	GB	0.9076	ET ET	LR	0.9165	RFE	GB	0.7962	ET	LR	0.8114	RF ET	RF	0.663
	$_{ m ET}$	$_{ m RF}$	0.9061 0.9029	RF	KNN KNN	$0.9161 \\ 0.9156$	RF ET	$_{ m RF}$	0.7951 0.7947	RF	SVM SVM	0.8006 0.7955	RFE	KNN GB	0.6543 0.6528
	RF	RF	0.9006	Anova	ET	0.9076	ET	ET	0.7883	ET	KNN	0.7826	RF	KNN	0.6463
	RF	SVM	0.8969	RFE	KNN	0.9039	RF	KNN	0.7878	RFE	Cos	0.7719	ET	SVM	0.6304
	$_{\rm ET}$	SVM	0.8875	RFE	$_{\rm LR}$	0.8986	RFE	Cos	0.7867	RF	KNN	0.7712	RF	$_{\mathrm{GB}}$	0.6292
	ET	KNN	0.8806	Anova	RF	0.8925	Anova	ET	0.7867	RFE	Corr	0.7698	RF	SVM	0.6217
	RF	KNN	0.8788	Anova	LR	0.8808	Anova	RF	0.7848	Anova	RF	0.7659	ET	GB	0.6188
	$_{ m RF}$	$_{\text{Cos}}$	0.8738 0.8647	Anova RFE	KNN Cos	$0.8701 \\ 0.8693$	RFE Anova	Corr KNN	0.7812 0.7806	Anova Anova	ET SVM	0.7604 0.753	Anova RFE	LR Corr	0.5883 0.5756
	RF	GB	0.8619	Anova	SVM	0.8598	RF	Corr	0.7762	RF	GB	0.7289	Anova	KNN	0.5722
	ET	Cos	0.8603	RF	Cos	0.852	ET	KNN	0.7738	ET	Corr	0.7239	RF	Corr	0.5721
	RF	Corr	0.8518	ET	Cos	0.8502	ET	SVM	0.7724	ET	Cos	0.722	RFE	DT	0.5663
	ET	Corr	0.8475	RFE	Corr	0.8381	Anova	Corr	0.7722	ET	GB	0.7197	ET	Corr	0.5646
	RFE	DT	0.7807	ET	Corr	0.8202	Anova	Cos	0.7662	Anova	LR	0.7179	Anova	ET	0.5637
	ET RF	$_{ m DT}$	0.7325 0.7319	RF ET	Corr GB	0.8132 0.8132	RF ET	$_{ m LR}^{ m Cos}$	$0.7624 \\ 0.7602$	RF Anova	$_{ m KNN}$	0.7172 0.7166	Anova ET	$_{ m DT}$	$0.5636 \\ 0.5628$
	Anova	SVM	0.6732	RF	GB	0.8101	ET	GB	0.7537	RFE	GB	0.7146	RF	DT	0.5587
	Anova	ET	0.6588	RFE	GB	0.8062	RF	GB	0.7487	RF	Corr	0.701	Anova	SVM	0.5549
	Anova	RF	0.651	Anova	Cos	0.7996	Anova	$_{\mathrm{GB}}$	0.7484	Anova	Corr	0.6686	Anova	$_{\mathrm{GB}}$	0.554
	Anova	LR	0.6368	Anova	GB	0.7656	ET	Corr	0.7441	Anova	Cos	0.6673	RFE	Cos	0.5457
	Anova	Corr	0.6191	Anova	Corr	0.7562	ET	Cos	0.7422	RFE	DT	0.6321	RF	Cos	0.5424
	Anova Anova	KNN GB	0.6184 0.6146	RFE RF	$_{ m DT}$	0.7197 0.7152	RFE Anova	$_{ m DT}$	0.6584 0.6473	Anova RF	$_{ m DT}$	0.6317 0.6195	ET Anova	$_{ m DT}^{ m Cos}$	0.5392 0.5343
	Anova	Cos	0.6134	ET	DT	0.7115	ET	DT	0.6389	ET	DT	0.6173	Anova	Corr	0.5213
	Anova	DT	0.5647	Anova	DT	0.6799	RF	DT	0.6385	Anova	DT	0.5769	Anova	Cos	0.5164
16	RFE	ET	0.972	RFE	ET	0.9666	RFE	SVM	0.8758	RFE	ET	0.922	ET RFE	LR	0.7616 0.7607
	RFE	$_{ m RF}$	0.9709 0.9681	ET RFE	$_{ m RF}$	$0.9582 \\ 0.9565$	RFE RFE	KNN RF	0.8704 0.8671	RFE RFE	RF SVM	0.9162 0.9111	RF	$_{ m LR}$	0.7468
	RFE	SVM	0.968	RF	ET	0.9539	RFE	ET	0.8597	RFE	KNN	0.9033	RFE	ET	0.7159
	RFE	Cos	0.9663	ET	RF	0.9482	RFE	$_{ m LR}$	0.8535	ET	ET	0.8997	RFE	RF	0.7086
	RFE	KNN	0.9604	ET	KNN	0.9435	Anova	SVM	0.8496	RFE	$_{ m LR}$	0.897	RFE	SVM	0.7067
	RF	LR	0.9588	RF	RF	0.9414	Anova	LR	0.8439	RF	ET	0.896	ET	ET	0.7009
	$_{ m RFE}$	$_{ m ET}^{ m Corr}$	$0.9576 \\ 0.9514$	RF ET	KNN SVM	0.9413 0.9407	RFE Anova	$_{ m RF}$	0.8407 0.8361	RF	RF	0.8914 0.8861	ET RF	$_{ m ET}$	$0.7007 \\ 0.6995$
	ET	LR	0.9508	RF	SVM	0.9381	Anova	KNN	0.8349	RF	LR	0.878	RF	RF	0.699
	ET	ET	0.9482	RFE	SVM	0.938	RFE	Corr	0.8325	ET	LR	0.8767	RFE	GB	0.6899
	ET	RF	0.9468	ET	LR	0.9305	Anova	$_{\rm ET}$	0.8323	ET	SVM	0.8766	ET	SVM	0.6896
	RF	RF	0.9453	Anova	$_{\rm ET}$	0.9302	RF	$_{\rm LR}$	0.8295	RF	SVM	0.8675	ET	KNN	0.6827
	RF	SVM	0.9393	RFE	KNN	0.925	ET	RF	0.8273	ET	KNN	0.8593	RF	SVM	0.6799
	RFE RF	$_{ m GB}$	$0.9306 \\ 0.9242$	Anova RF	$_{ m LR}$	0.9193 0.9192	RF RF	ET KNN	0.8265 0.8256	RFE RF	Corr KNN	0.8591 0.8545	RF RFE	KNN KNN	0.6743 0.6739
	ET	KNN	0.9242	RFE	LR	0.9086	Anova	Corr	0.8252	RFE	Cos	0.8508	ET	GB	0.6585
	ET	SVM	0.9233	Anova	LR	0.9076	RF	RF	0.8243	Anova	RF	0.8142	RF	GB	0.6566
	RF	Cos	0.923	Anova	KNN	0.902	RF	SVM	0.8234	Anova	$_{\rm ET}$	0.7982	RFE	Corr	0.621
	RF	Corr	0.9224	Anova	SVM	0.8914	Anova	Cos	0.8226	ET	Corr	0.7898	Anova	LR	0.6089
	ET	GB	0.915	RFE	Cos	0.8871	RF	Corr	0.8212	RF	Cos	0.7885	RFE	Cos	0.6075
	RF ET	$_{\text{Cos}}$	0.9145 0.9087	ET RF	Cos Cos	0.8836 0.8744	ET ET	ET KNN	0.8156 0.8137	Anova	Cos SVM	0.7812 0.7779	RF ET	Corr Corr	$0.6066 \\ 0.6061$
	ET	Corr	0.906	RFE	Corr	0.8641	RFE	GB	0.8058	RF	Corr	0.7717	Anova	KNN	0.5878
	RFE	DT	0.8129	ET	Corr	0.8618	ET	SVM	0.803	RF	$_{\mathrm{GB}}$	0.7596	Anova	RF	0.5829
	ET	DT	0.794	RF	Corr	0.8552	ET	LR	0.7982	Anova		0.7511	RFE	DT	0.5768
	RF	DT	0.7857	Anova	Cos	0.8424	RF	Cos	0.7968	ET	GB KNN	0.7426	Anova	ET SVM	0.5764
	Anova Anova	SVM ET	0.7232 0.7192	Anova ET	$_{\rm GB}$	0.8266 0.8189	ET ET	Corr Cos	0.7945 0.7897	Anova RFE	GB	0.733 0.7078	Anova RF	DT	$0.575 \\ 0.5726$
	Anova	LR	0.7157	RFE	GB	0.8169	Anova	GB	0.7858	Anova	Cos	0.6791	Anova	GB	0.57
	Anova	RF	0.705	RF	$_{\mathrm{GB}}$	0.8069	ET	$_{\mathrm{GB}}$	0.7715	Anova	Corr	0.6732	ET	DT	0.5688
	Anova	Corr	0.6893	Anova	GB	0.7795	RF	$^{\mathrm{GB}}$	0.7609	Anova	GB	0.6581	RF	Cos	0.5569
	Anova	KNN	0.6761	ET	DT	0.7277	RFE	DT	0.674	RF	DT	0.6481	ET	Cos	0.5552
	Anova Anova	Cos	0.6749 0.6646	RFE RF	DT DT	0.725 0.716	Anova ET	$_{ m DT}$	0.6684 0.6563	RFE ET	DT DT	$0.6396 \\ 0.638$	Anova Anova		0.5402 0.5335
	Anova		0.5911	Anova		0.6945	RF	DT	0.6529	Anova		0.6007	Anova		0.5163
32	RFE	$_{ m LR}$	0.9759	RFE	\mathbf{ET}	0.9792	RFE	KNN		RFE	$_{ m LR}$	0.9685	RFE	$_{ m LR}$	0.8194
	RFE	ET	0.9757	ET	ET	0.969	RFE	RF	0.8801	RFE	SVM		ET	LR	0.7904
	RF RFE	$_{\text{Cos}}$	0.9747 0.9736	RFE RFE	RF KNN	0.969 0.9689	RFE RFE	ET SVM	0.8717 0.8679	RFE RFE	KNN ET	$0.966 \\ 0.9646$	RF RFE	LR SVM	0.788 0.7607
	RFE	RF	0.9734	RF	ET	0.9645	RFE	LR	0.866	RFE	RF	0.9577	RFE	ET	0.743
	RFE	SVM	0.9734	Anova		0.9636		LR	0.8538	ET	ET	0.9518	ET	ET	0.7278
	ET	$_{\rm LR}$	0.9683	ET	KNN	0.9617	Anova	SVM	0.8528	RF	ET	0.9504	RFE	RF	0.7261
	RF	ET	0.9676	RFE	SVM	0.9602	RFE	Corr	0.8525	RF	LR	0.9477	RF	ET	0.7259
	RFE	KNN	0.9672	ET	RF	0.9594	Anova	KNN	0.851	RF	RF	0.9448	RFE	KNN	0.7257
	RF RFE	RF Corr	$0.9666 \\ 0.9662$	ET RF	SVM KNN	0.9587 0.9583	RFE RF	$_{ m Cos}$	$0.85 \\ 0.8479$	ET ET	$_{ m RF}$	0.9438 0.9412	RF ET	RF SVM	0.7209 0.7206
	ET	RF	0.9658	Anova	KNN	0.9573	RF	Corr	0.8457	RF	SVM		RF	SVM	0.7196
	ET	ET	0.9625	RF	SVM	0.9562	RF	$_{ m LR}$	0.8435	ET	SVM	0.9384	ET	RF	0.7159
	RF	SVM	0.9543	Anova	RF	0.9559	RF	RF	0.8408	RF	KNN	0.9352	RFE	$_{\mathrm{GB}}$	0.7097
	RF	Cos	0.9541	RF	RF	0.9547	Anova	RF	0.8406	ET	KNN	0.9312	RF	KNN	0.7088
	RF RF	Corr KNN	0.9524	ET RFE	$_{ m LR}$	0.9543	ET RF	RF ET	0.8402	RFE RFE	Corr	0.9225 0.9122	ET RFE	KNN	0.7034
	ET	SVM	0.9462 0.9439	Anova		0.9537 0.9532	ET	KNN	0.8395 0.8381	Anova	$_{ m RF}$	0.9122 0.8604	ET	$_{\mathrm{GB}}^{\mathrm{Cos}}$	0.6879 0.6712
	ET	KNN	0.9438	Anova	SVM	0.9505	Anova	Corr	0.8362	ET	Corr	0.8601	RFE	Corr	0.669
	ET	Cos	0.9424	RF	$_{ m LR}$	0.9479	RF	SVM	0.8303	ET	Cos	0.8589	RF	$_{\mathrm{GB}}$	0.6665
	RFE	GB	0.9414	ET	Cos	0.9015	Anova	ET	0.83	RF	Cos	0.8565	RF	Corr	0.6321
	ET ET	Corr	0.9405	Anova RFE	Cos	0.9007	Anova	Cos	0.8291	RF	Corr	0.8503	Anova ET	LR	0.6272
	L I	GB	0.9397 Conti	nued on	Cos	0.8979	ET	LR	0.8282	Anova	ET	0.8425	P.T.	Corr	0.6218

Continued on next page

				ntinued f		evious pa	^{ige} Protein	micro	RNA S	NP CN	1V				
	RF RFE Anova RF ET Anova Anova Anova Anova Anova Anova Anova	DT DT SVM ET RF Cos Corr KNN GB	0.9386 0.8215 0.8094 0.8039 0.8033 0.7995 0.7935 0.7844 0.7654 0.7455 0.7424	Anova RF ET RF RFE ET RF Anova RFE ET RFE RF Anova	Corr Cos Corr Corr GB GB GB GB DT DT	0.8922 0.8918 0.8891 0.8893 0.8797 0.8238 0.8109 0.8088 0.8087 0.723 0.723 0.7189	ET ET RF ET Anova RFE ET RF RFE Anova ET RF	Corr ET Cos SVM Cos GB GB GB GB GB DT DT	0.8266 0.8257 0.824 0.8229 0.8186 0.7926 0.7897 0.7728 0.7563 0.6703 0.6658 0.6546	Anova RF ET Anova RFE Anova Anova RFE Anova Anova RF ET RFE Anova	SVM LR GB GB KNN GB GB Cosr DT DT DT	0.8053 0.7989 0.7812 0.765 0.7539 0.7014 0.6999 0.6932 0.6861 0.6541 0.6445 0.6211	RF Anova Anova Anova RFE Anova ET RF ET Anova Anova	Cos RF KNN SVM ET DT GB Cos DT DT Corr DT Cos	0.592 0.5902 0.5898 0.5838 0.5825 0.5797 0.5766 0.5767 0.5741 0.5713 0.5474 0.5452
64	RF RFE RFE RF ET RFE ET RFE ET RFE ET RF ET ET Anova		0.9789 0.9777 0.977 0.977 0.977 0.9775 0.9755 0.9737 0.9733 0.9738 0.9798 0.9656 0.9647 0.9651 0.9555 0.9576 0.9548 0.9501 0.9648 0.9544 0.8743 0.8705 0.8352 0.8334 0.8705 0.8352 0.8334 0.8226 0.8273 0.8266 0.8173	RFE RFE RFF RFF ET RF ET RF ET Anova ET ET Anova ET Anova ET Anova RFE Anova RFE ET Anova RFE ET Anova RFE ET Anova	ET LR LR KNN ET ET ET RF SVM KNN LR SVM KNN KNN KNN Cos Coor Coor Cor Cor Cor Cor Cor Cor Cor C	0.979 0.9782 0.9781 0.9727 0.9721 0.9719 0.9709 0.9695 0.9693 0.9688 0.9662 0.9635 0.9693 0.9613 0.9613 0.9613 0.9608 0.9601 0.9581 0.9167 0.922 0.8928 0.8942 0.8935 0.8945 0.8055 0.71203 0.7126		KNN LR RF LR COTT ET KNN KNN LR LR COTT KNN COS RF SVM SVM RF COS SVM RF COS ET COS ET COS ET COT COS ET	0.8746 0.8688 0.8682 0.8595 0.8578 0.8578 0.8564 0.8564 0.8557 0.8539 0.8536 0.852 0.8492 0.84453 0.8426 0.8422 0.8407 0.8538 0.8426 0.8422 0.8407 0.8536 0.870 0.870 0.870 0.870 0.870 0.870 0.870 0.870 0.870 0.870 0.870 0.870 0.870 0.870 0.870 0.870 0.870	RFE RFE RFF RFE RFF RFF RFF RFF RFF RFF	KNN LR LR SVM LR ET KNN ET SVM ET KNN ET KNN ET Corr Cor Cor Cor Cor Cor Cor Cor Cor Co	0.9911 0.9892 0.9862 0.9843 0.9837 0.9833 0.9771 0.9778 0.9776 0.9762 0.9748 0.9763 0.9712 0.9233 0.9227 0.9215 0.919 0.8378 0.7938 0.7938 0.7938 0.7938 0.9227 0.9215 0.919 0.8479 0.8479 0.8479 0.8786 0.7794 0.7786 0.7794 0.7786 0.7794 0.7786 0.7253 0.7214 0.7208	RFE ET RF RFE ET RF ET RF ET RFE RFE RFE RFE RFE RFE Anova Anova Anova Anova Anova Anova	SVM	0.8376 0.8136 0.8135 0.7658 0.7475 0.7474 0.7471 0.7386 0.7331 0.7295 0.7157 0.7076 0.6891 0.67025 0.6711 0.6514 0.6502 0.646 0.5932 0.6015 0.5932 0.5932 0.5762 0.5762 0.5675 0.5466
96	RF RFE RFE RFE RFE RFE RFE RFE RFF RF RF ET RF ET RF ET RF ET RF Anova	ET SVM RF KNN GB Cos Corr DT DT DT	0.9808 0.9787 0.9774 0.9776 0.9761 0.9766 0.9744 0.9723 0.9699 0.9618 0.9616 0.9639 0.9618 0.9616 0.9639 0.9618 0.9616 0.9639 0.9618 0.9616 0.9639 0.9618 0.9616 0.9545 0.9487 0.9487 0.9487 0.9487 0.9487 0.9488 0.9548 0.8558 0.8568 0.8568 0.8329 0.8473 0.8474 0.8473 0.8474	RF RFE ET RFE ET Anova RFE ET Anova RFE Anova RFE Anova RFE RF Anova RFE RFE Anova RFE RFE RF Anova RFE RFE RF Anova RFE RF RF Anova RFE RF RF Anova RFE RF Anova	Cos Corr Cos Cos Corr Corr Corr GB GB GB GB DT DT	0.979 0.9779 0.9779 0.9778 0.9765 0.9734 0.973 0.971 0.9675 0.9662 0.9662 0.9624 0.9623 0.9589 0.9581 0.9574 0.9825 0.99182 0.90475 0.9182 0.90475 0.8845 0.88713 0.8175 0.8865 0.8056 0.8056 0.80669	RFE RF RF ET RFE ET Anova RFE ET RFE RF RFE RF RFE RF Anova RF Anova Anova Anova Anova Anova Anova	Cos Cos SVM ET Corr GB	0.8697 0.8657 0.8643 0.8634 0.8633 0.8629 0.8629 0.8613 0.8596 0.8483 0.8477 0.8447 0.8447 0.8447 0.8467 0.845 0.8493 0.8397 0.8493 0.8397 0.8493 0.8397 0.8493 0.8394 0.8494 0.8494 0.8410 0.8410	RFE RF RFE ET RF ET RFE ET ET ET RFE ET RFE ET RFE ET Anova Anova Anova Anova Anova Anova Anova RF ET Anova Anova RF ET Anova	ET LR SVM KNN GB GB Corr GB GB DT DT	0.9933 0.9918 0.9916 0.9909 0.9877 0.987 0.9887 0.9852 0.9852 0.9822 0.9821 0.9806 0.9703 0.9481 0.9481 0.9480 0.9503 0.9488 0.9476 0.9107 0.883 0.7483 0.7485 0.7496 0.7496 0.7384 0.7271 0.6667 0.6667 0.666467 0.66282	RFE ET RF RFE ET RFE ET RFE	Corr Cos Cos KNN SVM GB RF ET Corr DT DT DT	0.847 0.8292 0.8285 0.7715 0.7663 0.7602 0.7514 0.7441 0.7347 0.7336 0.7326 0.7292 0.7264 0.7023 0.6797 0.6797 0.6794 0.6487 0.6374 0.6148 0.6114 0.6097 0.6988 0.5758 0.5756 0.5745 0.5745 0.5745

Table 6: External Validation: Best performing models

Marker Set Size	Feature Selection	Classifier	AUC
8	RFE	Extra Trees	0.938 + / - 0.015
16	RFE	Logistic Regression	0.9676 + / - 0.112
32	RFE	Logistic Regression	0.9849 + / - 0.0055
64	RFE	Logistic Regression	0.9904 + / - 0.0036
96	RFE	Logistic Regression	0.9920 + / - 0.025

The top performing feature selection and classification algorithms selected by AUC in the external validation experiment.

Table 7: External Validation: NCI-60 to TCGA mapping

NCI-60 Cell Line	TCGA Cell lines
Central Nervous System	Glioblastoma multiforme (GBM)
	Brain Lower Grade Glioma (LGG)
Lung	Lung adenocarcinoma (LUAD)
	Lung squamous cell carcinoma (LUSC)
Colon	Colon adenocarcinoma (COAD)
	Rectum adenocarcinoma (READ)
Ovarian	Serous Cystadenocarcinoma (OV)
Renal	Kidney renal clear cell carcinoma (KIRC)
	Kidney renal papillary cell carcinoma (KIRP)

This table indicates which feature selection and classification algorithm was ranked highest by AUC during the model selection and training phase of the external validation experiment.

Table 8: Accuracy per Cancer Type Grouped by Marker Set Size

-				N	CI-60 C	Cell Lin	es		
TCGA Samples	Marker	Set Siz	e ME	LE C	O CNS	RE I	BR OV	LC	
CNS	8	0.0	0.0	0.0	1.0	0.0	0.0	0.0	0.0
CO	8	0.0144	0.0048	0.1779	0.0769	0.0048	0.0192	0.4135	0.2885
LC	8	0.0291	0.0058	0.0058	0.3081	0.1977	0.0174	0.0	0.436
OV	8	0.4074	0.1852	0.0	0.0	0.0	0.0	0.0	0.4074
RE	8	0.0	0.0	0.0649	0.7013	0.0	0.0	0.1039	0.1299
CNS	16	0.0	0.0	0.0	0.8246	0.1754	0.0	0.0	0.0
CO	16	0.0	0.0	0.9856	0.0	0.0144	0.0	0.0	0.0
LC	16	0.0	0.0058	0.064	0.0523	0.1512	0.0407	0.0	0.686
OV	16	0.0	0.2222	0.037	0.0	0.0	0.0	0.037	0.7037
RE	16	0.013	0.026	0.0649	0.0519	0.7403	0.013	0.0649	0.026
CNS	32	0.0	0.0877	0.0	0.9123	0.0	0.0	0.0	0.0
CO	32	0.0	0.0	1.0	0.0	0.0	0.0	0.0	0.0
LC	32	0.0	0.0581	0.2151	0.1279	0.2791	0.0814	0.0	0.2384
OV	32	0.0	0.7037	0.1481	0.0	0.0	0.037	0.1111	0.0
RE	32	0.0	0.0779	0.0	0.013	0.8571	0.0	0.039	0.013
CNS	64	0.0	0.386	0.0	0.5965	0.0175	0.0	0.0	0.0
CO	64	0.0	0.0	1.0	0.0	0.0	0.0	0.0	0.0
LC	64	0.0	0.0872	0.064	0.0058	0.7035	0.0233	0.0	0.1163
OV	64	0.0	0.4074	0.1481	0.0	0.1111	0.0741	0.0741	0.1852
RE	64	0.013	0.0909	0.0	0.0	0.8961	0.0	0.0	0.0
CNS	96	0.0	0.4386	0.0	0.5351	0.0263	0.0	0.0	0.0
CO	96	0.0	0.0	1.0	0.0	0.0	0.0	0.0	0.0
LC	96	0.0	0.1802	0.2209	0.0174	0.5233	0.0465	0.0	0.0116
OV	96	0.0	0.4074	0.2222	0.0741	0.037	0.0	0.2593	0.0
RE	96	0.0	0.0	0.0	0.0	1.0	0.0	0.0	0.0

This table indicates the percentage of each TCGA cancer type that was classified to each NCI-60 cell line.

The TCGA samples are grouped by marker set size. These predictions were made using the model specified in Supplemental Table 6.

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