

Supplemental Table 1. Cohort sizes and demographics of patients with embryonal tumors, high-grade glioma (HGG) and ependymoma (EP).

		Embryonal							HGG		EP
		ETMR	HGNET	NB	PB	NOS	ATRT	GBM	AA		
N		7	3	3	7	7	23	75	52	54	
% Total		3.03%	1.30%	1.30%	3.03%	3.03%	9.96%	32.47%	22.51%	23.38%	
Age	Mean	64.7	138.5	56.2	122.9	72.4	46.1	147.9	124.1	87.3	
	SD	55.9	97.4	19.6	88.5	68.2	61.7	79.1	65.2	65.3	
Sex	Female	2	3	1	5	1	9	33	30	26	
	%	28.6%	100.0%	33.3%	71.4%	14.3%	39.1%	44.0%	57.7%	48.1%	
	Male	5	0	2	2	6	14	42	22	28	
	%	71.4%	0.0%	66.7%	28.6%	85.7%	60.9%	56.0%	42.3%	51.9%	

Supplemental Table 2. Performance metrics of the best classifier for each binary evaluation on the holdout test set. Accuracy is statistically compared to No Information Rate.

	Classifier	Positive Class	Sens	Spec	PPV	NPV	AUC	Accuracy	95% CI	NIR	p
Embryonal - HGG	LR	Embryonal	0.8461	0.9062	0.7857	0.9354	0.9807	0.8888	0.8000 - 0.9778	0.7176	< 0.0001***
Embryonal - EP	XGB	Embryonal	0.9285	0.6923	0.7647	0.9	0.8241	0.8148	0.6296 - 0.963	0.5238	0.0012**
EP - HGG	NN	EP	0.8181	0.9428	0.8181	0.9428	0.9558	0.9130	0.8261 - 0.9783	0.7017	< 0.0001***

p < 0.01, * p < 0.001

Supplemental Table 3. Listing of contributing institutions by pathology.

	Embryonal							HGG		EP
	ATRT	ETMR	HGNET	NB	PB	NOS	GBM	AA		
Lurie Children's Hospital of Chicago	16	0	3	1	0	5	23	28	25	
Children's Hospital of Orange County	1	0	0	1	1	1	0	0	0	
Dayton Children's Hospital	1	1	0	0	0	0	5	3	2	

Indiana University Riley Hospital for Children	0	0	0	0	0	0	1	8	1	0
Seattle Children's Hospital	0	0	0	0	0	0	0	0	0	5
Stanford Lucile Packard Children's Hospital	4	6	0	1	6	0	0	29	17	15
Intermountain Primary Children's Hospital	1	0	0	0	0	0	0	10	3	7

Supplemental Table 4a. A list of the variables identified by feature reduction and submitted for model training in each binary pairing.

T1-MRI Features	T2-MRI Features	Demographics
Embryonal vs HGG		
t1_log-sigma-3-mm-3D_glcm_InverseVariance	t2_log-sigma-1-mm-3D_glcm_ClusterShade	Age
t1_original_shape_Flatness	t2_log-sigma-1-mm-3D_glcm_InverseVariance	
t1_wavelet-HHH_glszm_SizeZoneNonUniformityNormalized	t2_log-sigma-3-mm-3D_glszm_LargeAreaHighGrayLevelEmphasis	
t1_wavelet-HHH_glszm_SmallAreaEmphasis	t2_original_glcm_Imc2	
t1_wavelet-HHH_glszm_SmallAreaLowGrayLevelEmphasis	t2_original_shape_Flatness	

t1_wavelet-HLH_firstorder_Mean	t2_wavelet-HHH_glcM_Idn	
t1_wavelet-LHL_firstorder_Median	t2_wavelet-HHH_glszm_ZoneVariance	
t1_wavelet-LHL_glcM_Imc2	t2_wavelet-HHL_firstorder_Median	
t1_wavelet-LLH_glcM_Idn	t2_wavelet-HHL_glcM_MCC	
	t2_wavelet-HHL_glrIm_LongRunHighGrayLevelEmphasis	
	t2_wavelet-HLH_firstorder_Skewness	
	t2_wavelet-LLH_firstorder_Skewness	
	t2_wavelet-LLL_firstorder_Skewness	
Embryonal vs EP		
t1_log-sigma-1-mm-3D_glcM_Imc2	t2_log-sigma-5-mm-3D_firstorder_Median	Age
t1_wavelet-HLL_firstorder_Skewness	t2_wavelet-LLL_firstorder_Kurtosis	
EP vs HGG		
t1_log-sigma-1-mm-3D_glcM_InverseVariance	t2_log-sigma-1-mm-3D_glcM_ClusterShade	Age
t1_log-sigma-3-mm-3D_firstorder_Skewness	t2_log-sigma-5-mm-3D_glszm_GrayLevelNonUniformityNormalized	
t1_log-sigma-3-mm-3D_glcM_InverseVariance	t2_original_glcM_Imc2	
t1_log-sigma-5-mm-3D_glszm_LargeAreaEmphasis	t2_original_glrIm_LongRunHighGrayLevelEmphasis	
t1_original_glrIm_LongRunHighGrayLevelEmphasis	t2_wavelet-HHH_glcM_Idmn	
t1_original_shape_Flatness	t2_wavelet-HHH_glrIm_LongRunLowGrayLevelEmphasis	

t1_wavelet-HHH_glszm_GrayLevelNonUniformityNormalized	t2_wavelet-HHH_glszm_GrayLevelNonUniformityNormalized	
t1_wavelet-HHH_glszm_SmallAreaEmphasis	t2_wavelet-HHH_glszm_SmallAreaEmphasis	
t1_wavelet-HHL_firstorder_Mean	t2_wavelet-HHL_firstorder_Median	
t1_wavelet-HHL_glcm_ClusterShade	t2_wavelet-HHL_firstorder_Skewness	
t1_wavelet-HHL_glszm_ZoneEntropy	t2_wavelet-HHL_glszm_GrayLevelNonUniformityNormalized	
t1_wavelet-HLH_firstorder_Mean	t2_wavelet-HLH_firstorder_Skewness	
t1_wavelet-HLH_glcm_Imc1	t2_wavelet-HLH_glcm_MCC	
t1_wavelet-HLL_glrIm_RunLengthNonUniformity	t2_wavelet-HLH_glszm_SizeZoneNonUniformityNormalized	
t1_wavelet-LHL_glcm_MCC	t2_wavelet-HLL_glcm_Idmn	
t1_wavelet-LLH_firstorder_Kurtosis	t2_wavelet-HLL_glcm_InverseVariance	
	t2_wavelet-HLL_glszm_GrayLevelNonUniformityNormalized	
	t2_wavelet-LLH_firstorder_Skewness	

Supplemental Table 4b. Number of features retained by the final classifier model for each binary pairing.

Embryonal vs HGG	
Age	1

1 st order	6
shape	2
glcm	8
glszm	5
glrlm	1

Embryonal vs EP

1 st order	3
shape	0
glcm	1
glszm	0
glrlm	0

EP vs HGG

1 st order	8
shape	1
glcm	11
glszm	10
glrlm	4

Supplemental Table 5. Listing of the top three features for each binary pairing.

Feature	Interpretation	Higher Group
Embryonal-HGG		
age	Age	HGG
t2_log-sigma-1-mm-3D_glcm_ClusterShade	A measure of the skewness and uniformity Higher cluster shade implies greater asymmetry about the mean.	HGG
t1_wavelet-HLH_firstorder_Mean	The Mean gray level intensity within the ROI	Embryonal
Embryonal-EP		
t2_wavelet-LLL_firstorder_Kurtosis	The 'peakedness' of the distribution of values Higher signifies greater mass distribution in tail; lower signifies concentration toward peak/mean	Embryonal
t1_log-sigma-1-mm-3D_glcm_Imc2	The correlation between probability distributions l and j quantifying the complexity of the texture Range (0 – 1): value of 0 representing two independent distributions (no mutual information) and value of 1 representing two fully dependent/uniform distribution (maximal mutual information)	EP
t1_wavelet- HLL_firstorder_Skewness	The asymmetry of the distribution about the Mean Positive is longer right tail. → higher Skewness in EP → EP darker	Embryonal

EP-HGG		
t1_wavelet-HLH_firstorder_Mean	The Mean gray level intensity within the ROI	EP
t1_wavelet-HHL_glcm_ClusterShade	A measure of the skewness and uniformity Higher cluster shade implies greater asymmetry about the mean.	EP
t2_wavelet-HLH_glcm_MCC	MCC: complexity of the texture, with range $0 \leq \text{MCC} \leq 1$. Greater MCC is with right shifted probability curve a “brighter (less homogenously gray-toned)”	HGG

Supplemental Table 6a. Area under-the-curve (AUC) of the 6 classifiers trialed in each of the binary classifiers. (Also, below is a brief description the unique aspects of the six models used.)

	micro-averaged AUC		
	Embryonal - HGG	Embryonal - EP	EP - HGG
SVM	0.9	0.78	0.71
LR	0.98	0.81	0.94
KNN	0.75	0.66	0.81
RF	0.92	0.8	0.85
XGB	0.95	0.82	0.84
NN	0.92	0.74	0.96

SVM: Support vector machine models identify an optimal separating line (or hyperplane) between predicted classes.

LR: Logistic regression interprets a generalized linear function such that the outcome variable is interpreted as the probability of given outcomes.

KNN: K-nearest neighbors evaluates the K-training points closest to a given datapoint to predict its classification.

RF: Random Forest aggregates the scoring from multiple decision trees to produce a classification for data points based on features.

XGB: XGBoost is another model of multiple decision trees (learners) and retroactively aims to learn from incorrectly identified datapoints at the potential cost of overfitting.

NN: Neural Networks are constructed with layers of nodes, where each node consists of a linear combination and a non-linear activation function, that collectively yield a final prediction. Excess layers can also lead to overfitting with small datasets.

Supplemental Table 6b. Area under-the-curve (AUC) of the 6 classifiers trialed in a 3-way classifier.

	micro-averaged AUC
	Embryonal - HGG - EP
SVM	0.75
LR	0.71
KNN	0.67
RF	0.7
XGB	0.75
NN	0.77

Supplemental Appendix 1. Configuration files for radiomic feature extraction.

setting:

normalize: true

normalizeScale: 100

binWidth: 10

label: 1

interpolator: 'sitkBSpline' # This is an enumerated value, here None is not allowed

resampledPixelSpacing: [1,1,1] # This disables resampling, as it is interpreted as None, to enable it, specify spacing in x, y, z as [x, y, z]

weightingNorm: # If no value is specified, it is interpreted as None

geometryTolerance: 0.0001

correctMask: True

imageType:

Original: {} # for dictionaries / mappings, None values are not allowed, '{}' is interpreted as an empty dictionary

LoG: {'sigma': [5,3,1]}

Wavelet: {}

featureClass:

shape: ['VoxelVolume',

'MeshVolume',

'SurfaceArea',

'SurfaceVolumeRatio',

'Sphericity',
'SphericalDisproportion',
'Maximum3DDiameter',
'Maximum2DDiameterSlice',
'Maximum2DDiameterColumn',
'Maximum2DDiameterRow',
'Elongation',
'Flatness'] # Only enable these shape descriptors (disables redundant Compactness 1 and Compactness 2)

firstorder: [] # specifying an empty list has the same effect as specifying nothing.

glcm: # for lists none values are allowed, in this case, all features are enabled

glrlm:

glszm:

Supplemental Appendix 2. Parameters for image pre-processing, feature extraction and feature reduction.

Image Pre-Processing

Prior to feature extraction, we normalized (normalize scale = 100) and resampled images to isotropic 1-mm voxels. Below is the link regarding the exact method: <https://pyradiomics.readthedocs.io/en/latest/radiomics.html#radiomics.imageoperations.normalizeImage>.

A bin width of 10 was used for grey-level discretization in both normalized MR images.

Feature Classes

Extracted features classes included *First Order* statistics, *2D/3D Shape*, *Gray Level Co-occurrence Matrix (GLCM)*, *Gray Level Run Length Matrix (GLRLM)*, and *Gray Level Size Zone Matrix (GLSZM)*.

Image Filters

Features were computed on original, wavelet filtered, and Laplacian of Gaussian (LoG) filtered images. Wavelet filters included high band-pass (H) and low-band pass filters (L) in the x, y, and z direction resulting in 8 different combinations of decompositions.

Feature Reduction by Least Absolute Shrinkage and Selection Operator

Training was performed with 10-fold cross validation and repeated for 1000 cycles. The mean squared error was calculated for 100 lambdas in each cycle or until a minimum was achieved. The optimal lambda was identified as the lowest mean squared error value and used for feature reduction and coefficient calculations. Features represented in $\geq 80\%$ of the cycles were retained for subsequent classifier optimization.

Supplemental Appendix 3. Final hyperparameters following grid search for six classifiers evaluated binary classifiers.

Classifier	Optimal Algorithm	Parameters
Embryonal – EP	XGB	{'learning_rate': 0.5, 'max_depth': 6}
Embryonal - HGG	LR	{'C': 1, 'penalty': 'l2'}
EP - HGG	NN	{'hidden_layer_sizes': (100, 100, 50), 'learning_rate': 'constant'}