

Supplementary Materials

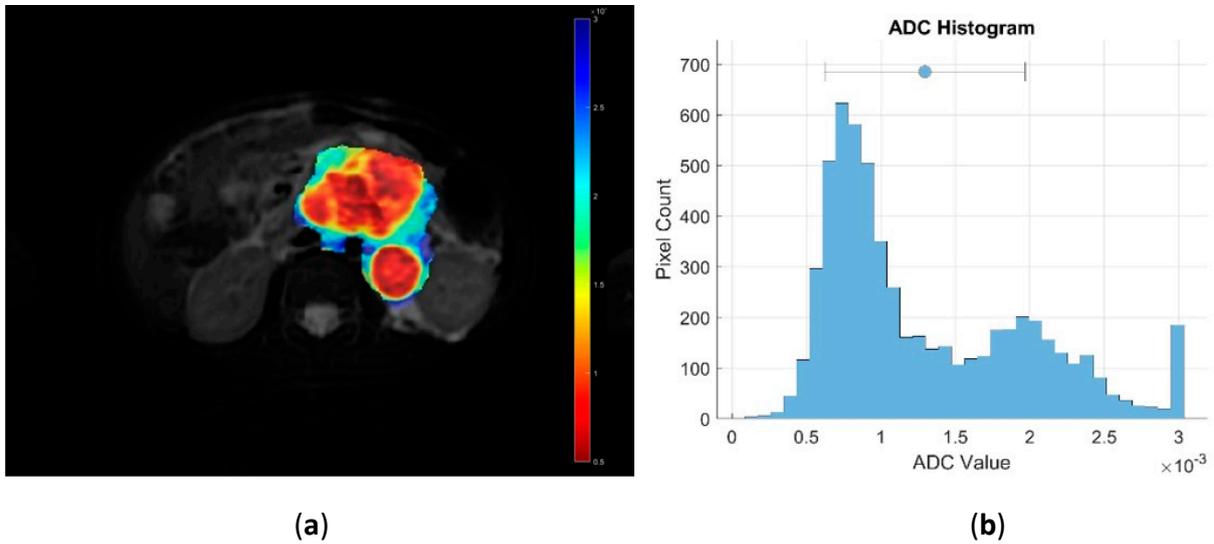


Figure S1. (a) Parametric map and (b) distribution histogram of the apparent diffusion coefficient (ADC, 10^{-3} mm²/s) obtained with the standard voxel-based mono-exponential signal decay model.

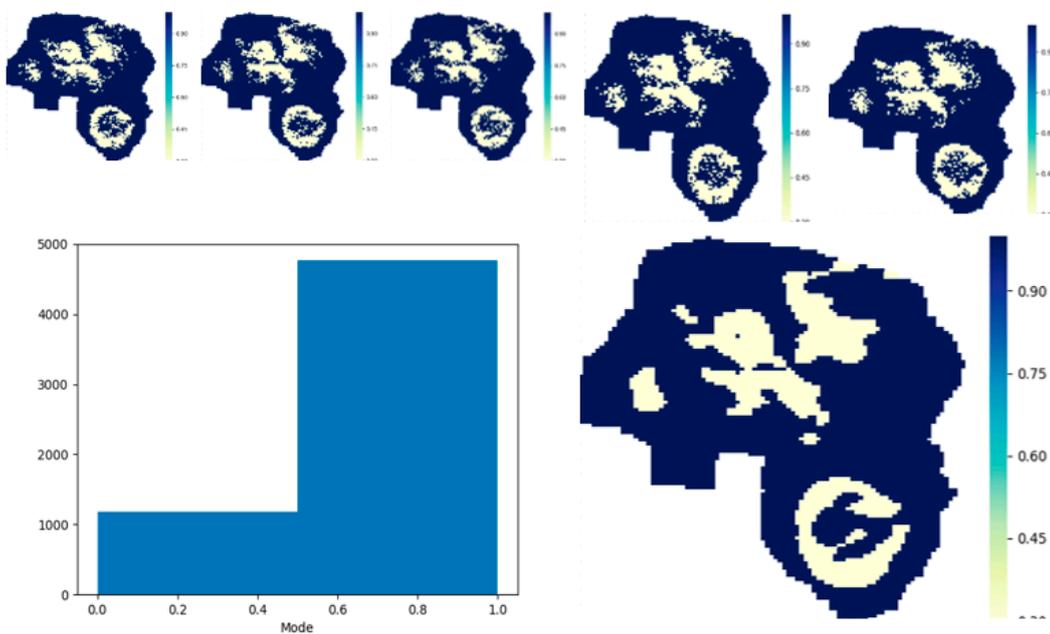


Figure S2. Generation of *in-silico* experiments, smearing of ADC values with a double-sided Crystal Ball function and cluster assignment of voxels. At the bottom, frequency map and frequency distribution of voxel allocation.

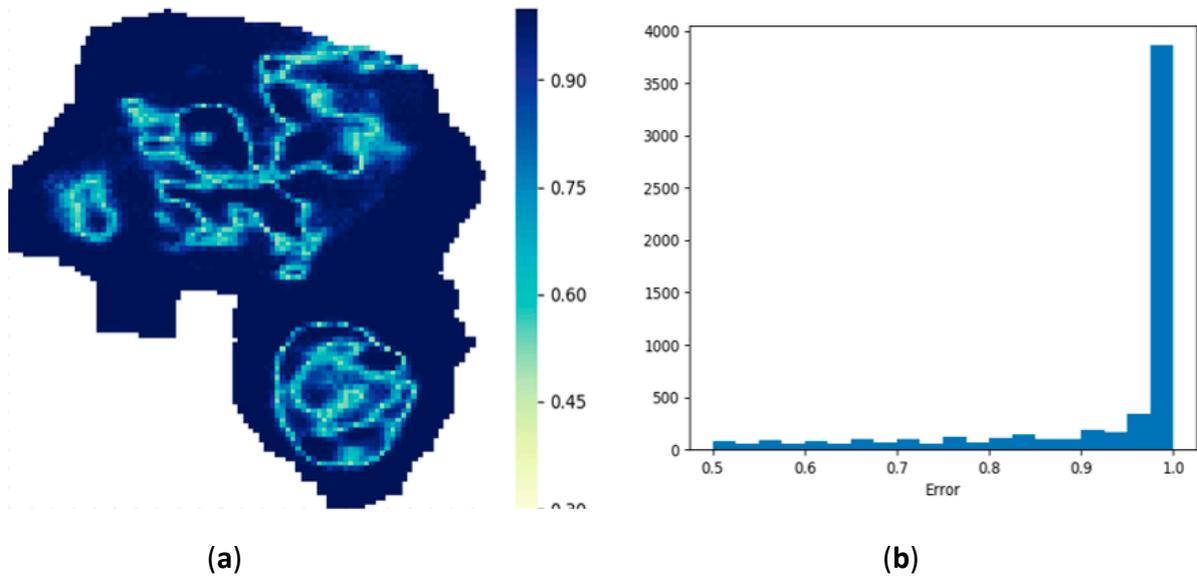


Figure S3. (a) Confidence map obtained from in-silico experiments representing the accuracy of voxel allocation to a clustered habitat. (b) The distribution of the confidence level spectrum per voxel can be used to threshold the voxel groups to exclude uncertainties.

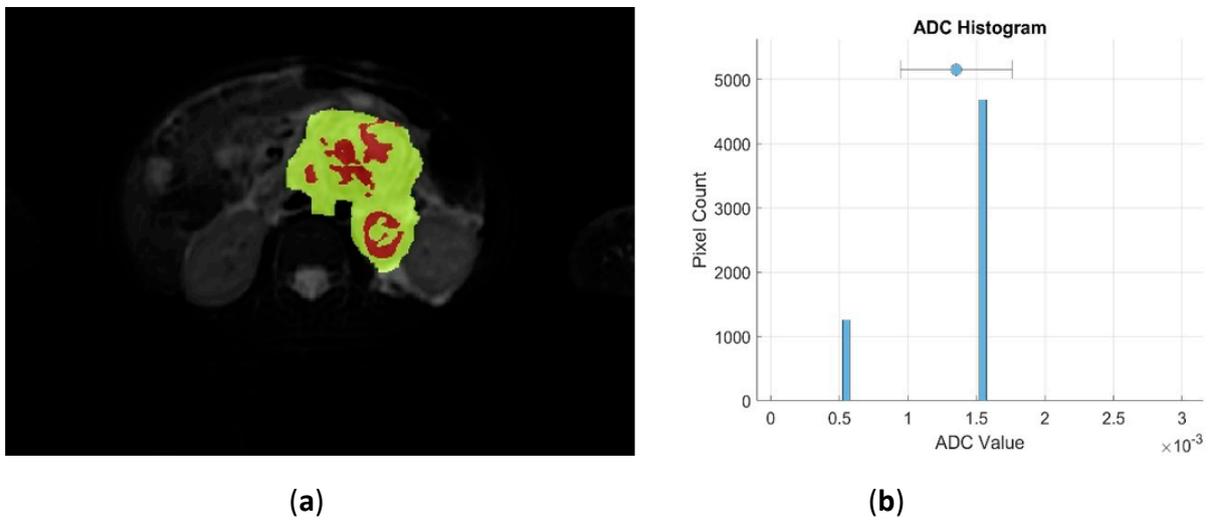


Figure S4. (a) Apparent diffusion coefficient (ADC) map (10^{-3} mm²/s) obtained with the *Fit-Cluster-Fit* method. (b) Distribution of the ADC spectrum performing a voxel-wise clustering into two habitats, with the 5th percentile as the parcellation threshold.

Table S1. Apparent diffusion coefficient (ADC) values per tumor malignancy in voxel-based, cluster-based and *Fit-Cluster-Fit* models with different confidence thresholds to exclude uncertainties (60%, 70%, 80%, 90% and 95%). Clustered habitats were built based on the 10th percentile of the artificially generated ADC distributions.

Method	ADC cut-off value	ADC value (10 ⁻³ mm ² /s)		Sensitivity (%)	Specificity (%)	Accuracy (%)	PPV (%)	NPV (%)	AUC
		Benign	Malignant						
Voxel-based	1.56	1.89±0.55	1.23±0.34	87	78	84	91	70	0.877
Cluster-based	0.80	1.33±0.41	0.60±0.24	87	100	91	100	75	0.966
FCF	0.81	1.36±0.42	0.63±0.24	87	100	91	100	75	0.959
FCF (60%)	0.76	1.33±0.41	0.61±0.24	83	100	88	100	69	0.959
FCF (70%)	0.76	1.27±0.40	0.58±0.24	87	100	91	100	75	0.961
FCF (80%)	0.71	1.18±0.36	0.54±0.21	87	100	91	100	75	0.966
FCF (90%)	0.69	1.11±0.35	0.51±0.20	87	100	91	100	75	0.969
FCF (95%)	0.68	1.06±0.35	0.49±0.19	87	100	91	100	75	0.966

Table S2. Apparent diffusion coefficient (ADC) values per tumor malignancy in voxel-based, cluster-based and *Fit-Cluster-Fit* models with different confidence thresholds to exclude uncertainties (60%, 70%, 80%, 90% and 95%). Clustered habitats were built based on the 15th percentile of the artificially generated ADC distributions.

Method	ADC cut-off value	ADC value (10 ⁻³ mm ² /s)		Sensitivity (%)	Specificity (%)	Accuracy (%)	PPV (%)	NPV (%)	AUC
		Benign	Malignant						
Voxel-based	1.56	1.89±0.55	1.23±0.34	87	78	84	91	70	0.877
Cluster-based	0.83	1.12±0.52	0.60±0.24	87	100	91	100	75	0.966
FCF	0.83	1.13±0.57	0.63±0.24	87	100	91	100	75	0.957
FCF (60%)	0.82	1.10±0.57	0.61±0.24	87	100	91	100	75	0.957
FCF (70%)	0.77	1.05±0.55	0.58±0.24	87	100	91	100	75	0.957
FCF (80%)	0.75	0.96±0.49	0.54±0.21	87	100	91	100	75	0.966
FCF (90%)	0.72	0.91±0.47	0.51±0.20	87	100	91	100	75	0.969
FCF (95%)	0.70	0.87±0.45	0.49±0.19	87	100	91	100	75	0.971

