

Multiparametric microvascular MRI: a cluster approach to characterize glioma

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For solid tumors, several microvascular factors are relevant to:

- tumor diagnosis
- follow-up

...

- response to treatment

Some parameters can be obtained with MRI:

- apparent diffusion constant (ADC)
- cerebral blood volume (CBV)
- cerebral blood flow (CBF),
- integrity of the vascular wall:
 - area under curve after Gd-DOTA injection (AUC_{Gd-DOTA})
- tissue oxygen saturation (StO₂)
- cerebral metabolic rate of oxygen (CMRO₂)

Parameter maps accumulation yields however some difficulty for interpretation

cluster-based approach to highlight the tissue structures that exhibit similar physiological characteristic in two rat models of glioma



Methods – experimental design

Two orthotopic rat models of glioma:

C6 / Wistar rats	- n:	13 rats
	- inoculation:	10 ⁵ cells, 5 μL, caudate nucleus
	- days of imaging:	21-23 days
	- mean tumor size +/- SD:	85.1 +/- 46.4 mm ³
F98 / Fischer rats	- n:	13 rats
	- inoculation:	10 ³ cells , 5 μL, caudate nucleus
	- day of imaging:	22-24 days
	- mean tumor size +/-SD:	104.9 +/- 28.0 mm ³
Physiological control	- mechanical ventilation	
	- blood samples in femoral	vein and artery before and after MRI
	- similar pO_2 and pCO_2 betw	veen models
Multiparametric MRI		
	Magnet 4.7 T (Avance III co	onsole; Bruker)
	Three regions of interest (F	ROIs):
	- tumor bulk	(n=3023 voxels; Wistar-C6=1500, Fischer-F98=1523)
	- contralatoral striatum	(n=E70 vovole) Wistor CE-21E Fischer E09-264)

- contralateral striatum (n=579 voxels; Wistar-C6=315, Fischer-F98=264)
- contralateral cortex (n=811 voxels; Wistar-C6=417, Fischer-F98=394)



Six MRI parameters:

Apparent diffusion coefficient (ADC)	TR/TE = 3000/28.6 ms, b=900 s/mm ²			
Vascular integrity:	DCE imaging approach: - multiple T1-weighted images; n = 60, 15.0 - bolus of Gd-DOTA (200µmol/kg)	Lemasson et al., Radiology 2010 6 s per image; TR/TE: 800/4.2 ms		
	-> Area Under Curve of Gd-DUTA	(AUC _{Gd-DOTA})		
Cerebral blood flow (CBF)	continous arterial spin labeling (CASL) - spin-echo EPI, TE=17.2 ms	Clément Debacker		
	 labeling duration = 4 sec, postlabeling de 	lay = 0.2 sec, 20 pairs		
Cerebral blood volume (CBV)	Steady state approach: - multiple gradient echo sequence; 26 echo - bolus of P904 (200µmol/kg, Guerbet)	<i>Valable et al., NMR Biomed 2008</i> os TR/TE: 6000/3.5-78.5 ms		
Tissue oxygen saturation (StO ₂)	Computation with: - CBV map	Lemasson et al., Radiology 2012		
	 - 3D multiple gradient echo sequence; 25 echos TR/TE: 100/3-87 ms - multiple spin echo sequence; 28 echos TR/TE: 2300/12-336 ms 			
Cerebral metabolic rate of oxygen (CM	$(1RO_2)$ CMRO ₂ = CBF x (1	– StO ₂ /100)		



Multiparametric MRI

Classical ROI-based analysis



-Similarities or heterogeneities between Wistar/Fischers rats, C6/F98 tumor models -Mean measurement do not reflect disparities within ROI -Loss of spatial information



Mclust-based Clustering approach

Cluster-based analysis C. Fraley and A. E. Raftery (2002). Journal of the American Statistical Association

Normal mixture modeling: Mclust Values standardization for scale effect minimization between parameters Allows prediction of new data set from built model Model choice / optimal number of clusters > Bayesian Information Criterion (BIC)

Cluster number determination

5 clusters are sufficient to explain variability of the data set





Mclust-based Clustering approach

Clusters separation

Data reduction after clustering Total of 6 directions Cluster distribution in the 4 first directions are depicted



> Clusters are well grouped> Good clusters separation

Dir. 2



Healthy			Pathological			
	mMRI parameters	in each cluster (mea	n +/- SD of absolute	values)		
cluster	1	2	3	4	5	
voxel number	875	837	758	805	637	
ADC (µm ² /s)	797.7 +/- 56.7	925.2 +/- 143.5	1188.2 +/- 242.0	1012.6 +/- 148.9	1102.3 +/- 216.7	
AUC _{Gd-DOTA} (10 ⁵ a.u).	1.2 +/- 2.7	8.7 +/- 10.6	177.8 +/- 85.9	65.5 +/- 58.9	131.4 +/- 92.8	
CBV (%)	4.0 +/- 1.2	3.3 +/- 0.9	5.0 +/- 2.5	6.1 +/- 2.6	3.4 +/-1.6	
CBF (mL/min/100g)	197.3 +/- 54.1	151.0 +/- 45.1	36.8 +/- 19.4	170.2 +/- 57.6	78.8 +/- 47.5	
StO _{2 (%)}	83.0 +/- 6.8	64.1 +/- 11.7	65.4 +/- 13.9	81.0 +/- 8.4	36.6 +/- 20.2	
CMRO ₂ (mL/min/100g)	33.8 +/- 17.4	54.5 +/- 24.7	12.6 +/- 7.7	31.9 +/- 17.5	47.0 +/- 33.0	



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mMRI parameters in each cluster (mean of standardized values)





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mMRI parameters in each cluster (mean of standardized values)



ROIs distribution in each cluster ?



	Healthy Pathological			Excluded			
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	mMRI parameters	in each cluster (mea	n of standardized val	ues)			
	AUC Gebota StO2 CBF CBV ADC CBV ADC	AUC Geboora	AUC Gebora Sto2 CMRo2	AUC StO2 CBF CBV CBV ADC CBV ADC CBV ADC CBV ADC CBV ADC CBV ADC CBV ADC CBV ADC CBV ADC CBV ADC CBV ADC CDV ADC CDV CDV ADC CDV CDV CDV CDV CDV CDV CDV CDV CDV C	AUC GebOOTA StO2 CBF CBV AUC CMRO2	c	
	ROI distribution in	each cluster					
striatum cortex tumor 1000 voxels							



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	AUC Ge-DOTA StO2 CMRO2	AUC Gebota SIO2 CBV CBV AUC CBV AUC CBV AUC CBV AUC CBV AUC CBV AUC CBV AUC CBV AUC CBV AUC CBV AUC CBV CBV AUC CBV CBV CBV	AUC GHOOTA StO2 CBF CBV CBV ADC CBV CDC CBV ADC CBV CDC CBV CDC CBV CDC CBV CDC CBV CDC CBV CDC CBV CDC CBV CDC CBV CDC CBV CDC CBV CDC CDC CDC CBV CDC CDC CDC CDC CDC CDC CDC CDC CDC CD	AUC SHOOTA	AUC GODTA StO2 CBF CBV CBV CBV CBV CBV CBV CBV CBV CBV CBV	0
	ROI distribution in	each cluster				
striatum cortex tumor 1000 voxels 750 500 250						
	Rat stem of in striat	distribution tum and (ROIs?	Tumor ı ir	models dist i tumor RO	ribution ?	

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	AUC Gebota Sto2 CMRO2	AUC HODIA STO2 CBV STO2 CMRO2	AUC Gebora Sto2 CBF Gebora CBV ADC CBV ADC CBV ADC CBV ADC CBV ADC CBV ADC CBV ADC CBV ADC CBV ADC CBV	AUC SHOTA	AUC GebOTA StO2 CBF CBV CBV ADC CBV ADC CBV CBV ADC		
	ROI distribution in e	each cluster					
striatum cortex tumor 1000 voxels 750 500						6	
	striatum and cortex	:	tumor				
C6-Wistar F98-Fischer							



Cluster maps comparison



Clusters

- 1: Healthy spécific
- 2: Healthy/Wistar specific
- 3: Tumor/F98 specific
- 4: Tumor specific (tendency toward C6 specificity)
- 5: Tumor specific

-Rat stem differences highlighted. Cluster 2 (orange) mostly presents in Wistar rat striatum -Some healthy clusters present in drawn tumor ROI: partial volume, false positive tumor voxels -Spatial organization of cluster in tumor



Conclusion/Perspectives

Conclusion

Normal Mixture model-based clustering:

- separates tumor tissue from healthy tissue
- shows cluster composition differences between C6 and F98 glioma models
- explain rat brain variability between rat stem
- highlight the spatial distribution of clusters within tumor

Perspectives

Preclinical research: effect of treatments on cluster composition ...detection of responsive/non responsive clusters

Clinical research (Julien Bouvier, PhD student):

tumor localization
 diagnosis
 grades

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UNIVERSITE JOSEPH FOURIER SCIENCES TECHNOLOGIE SANTÉ

Cluster-based approach validation on StO2 map vs Pimonidazol histology





Physiologic controls

