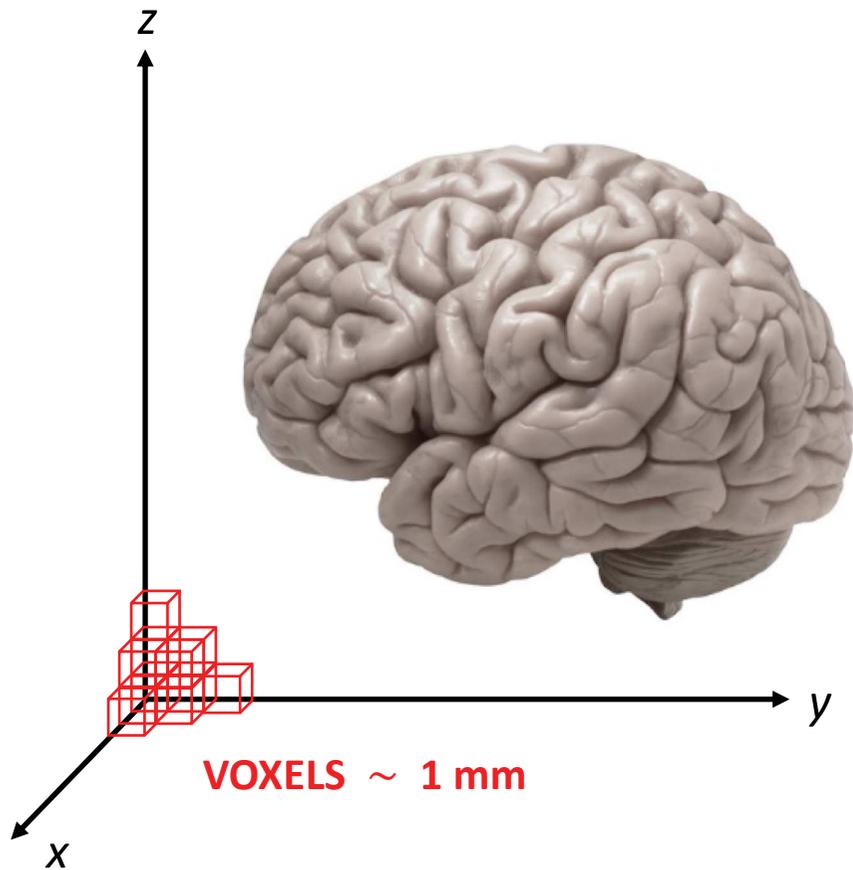


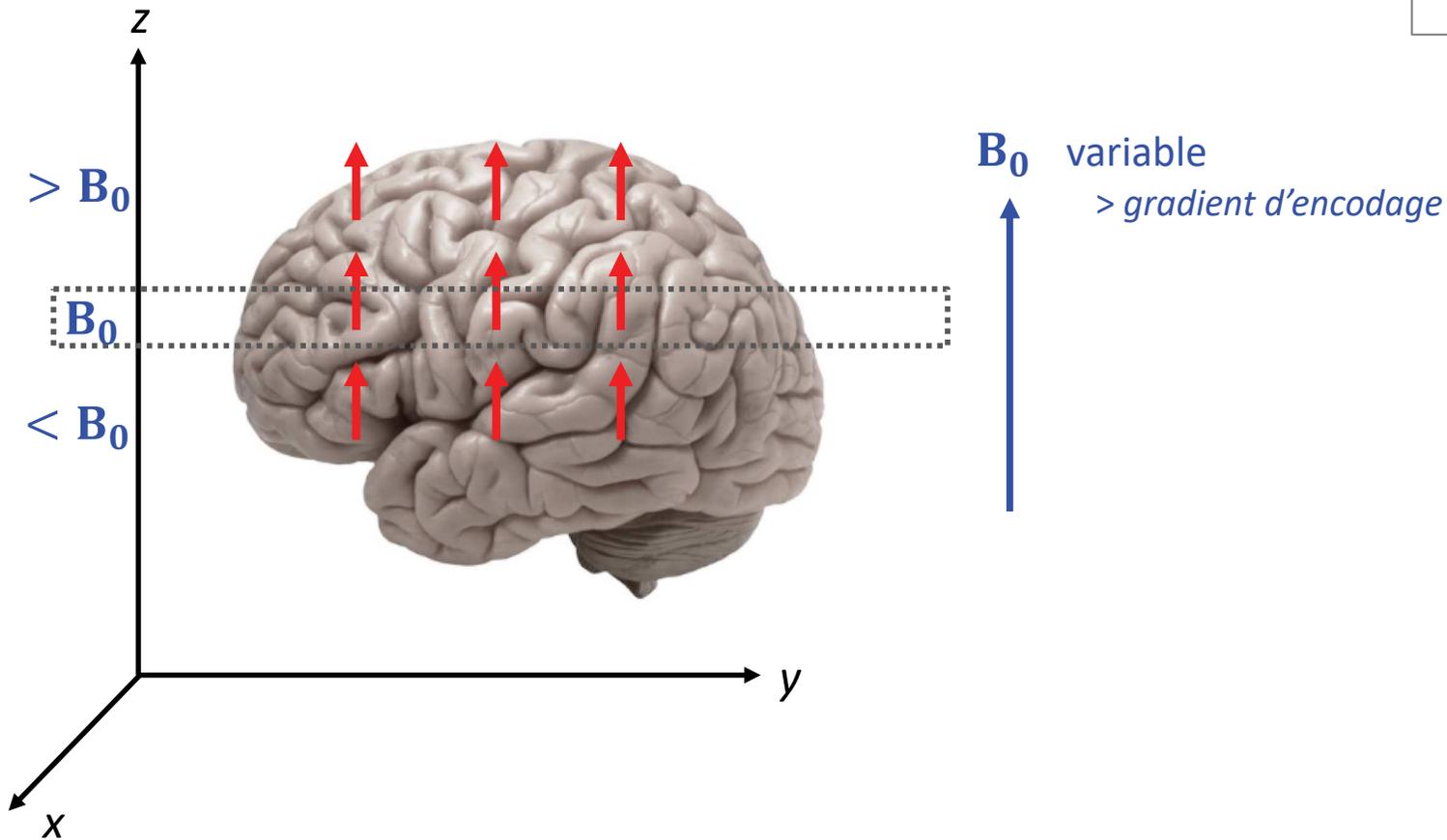
Applications : IRM / SRM

Imagerie : IRM



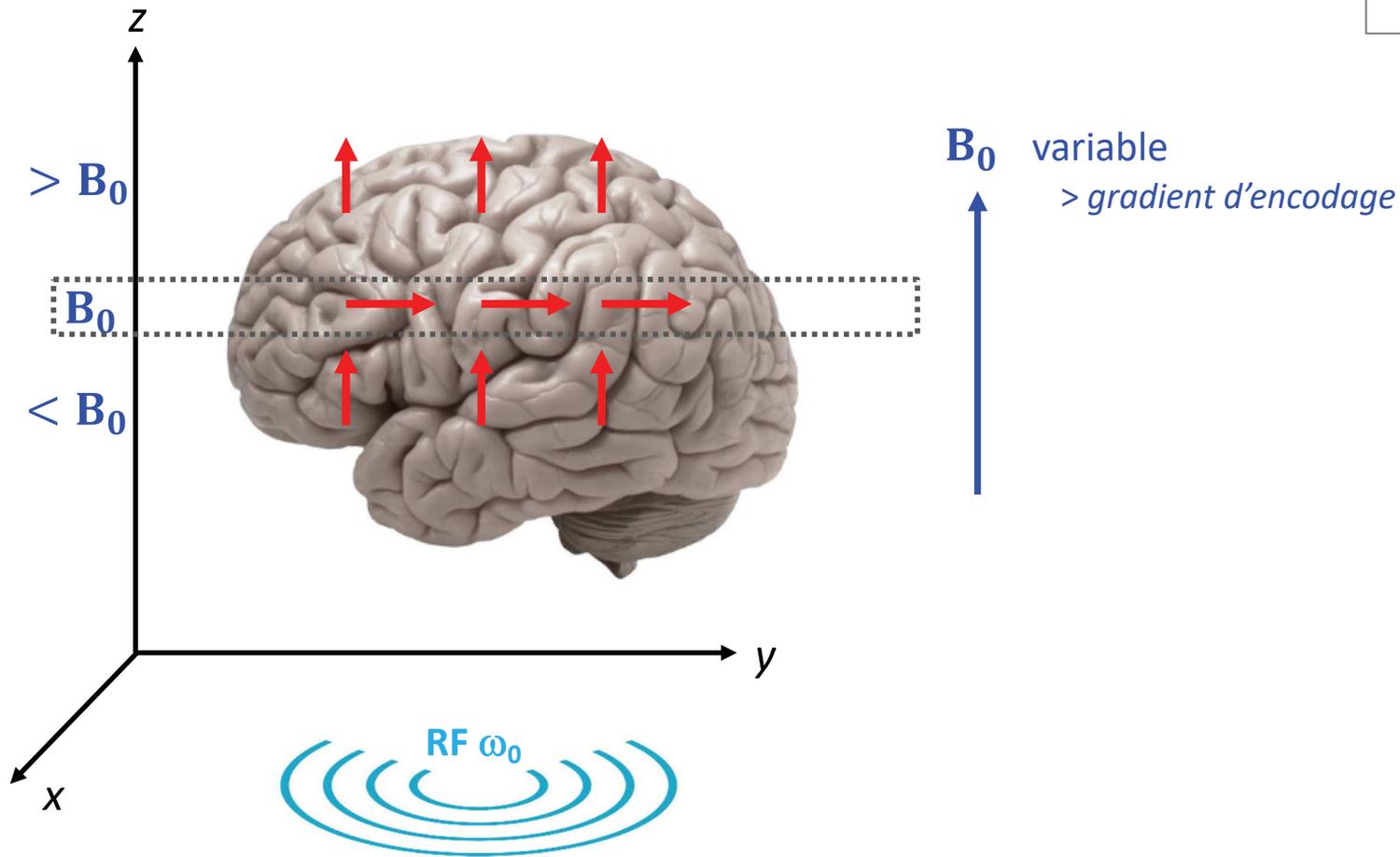
Applications : IRM / SRM

Imagerie : IRM



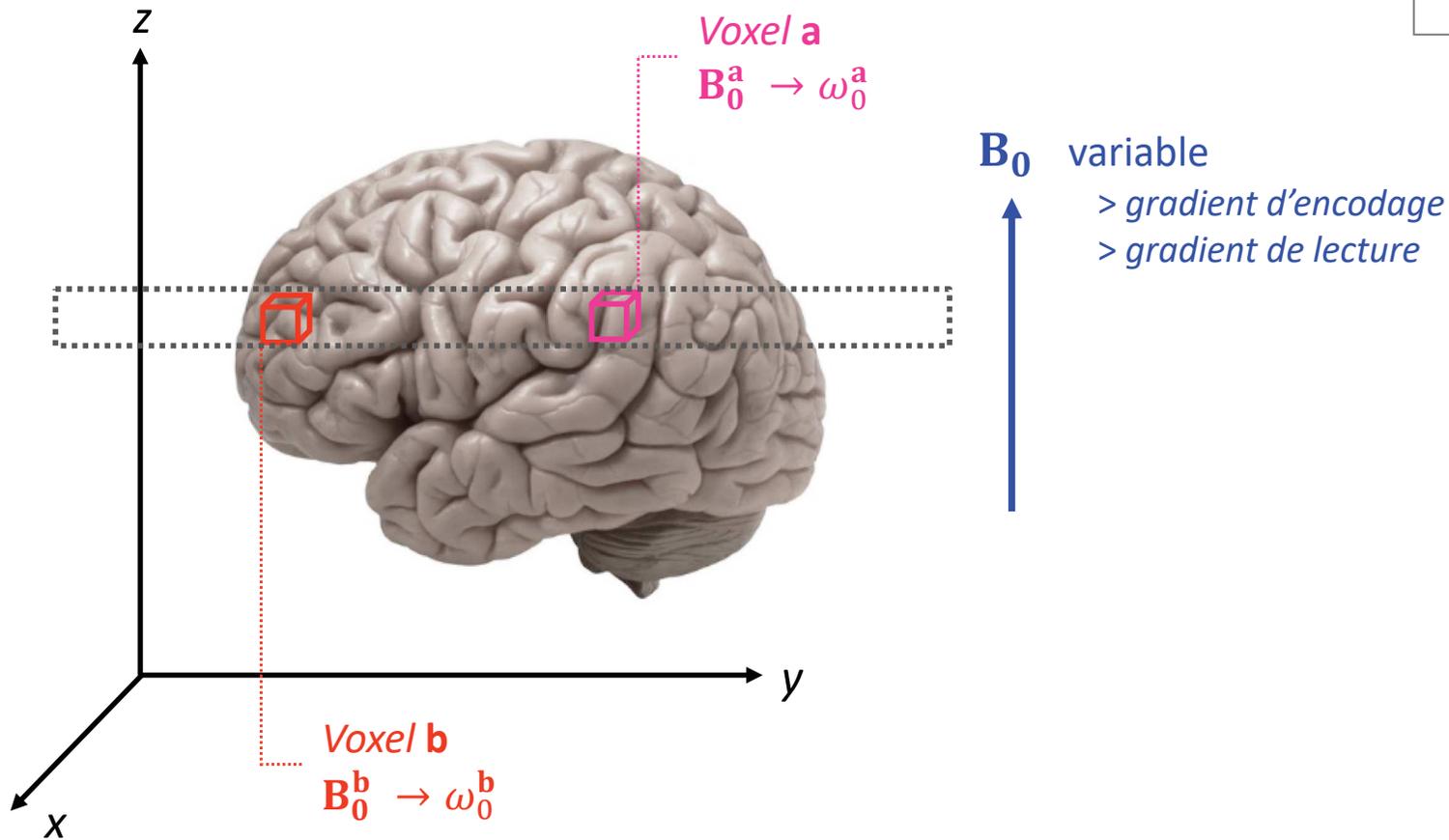
Applications : IRM / SRM

Imagerie : IRM



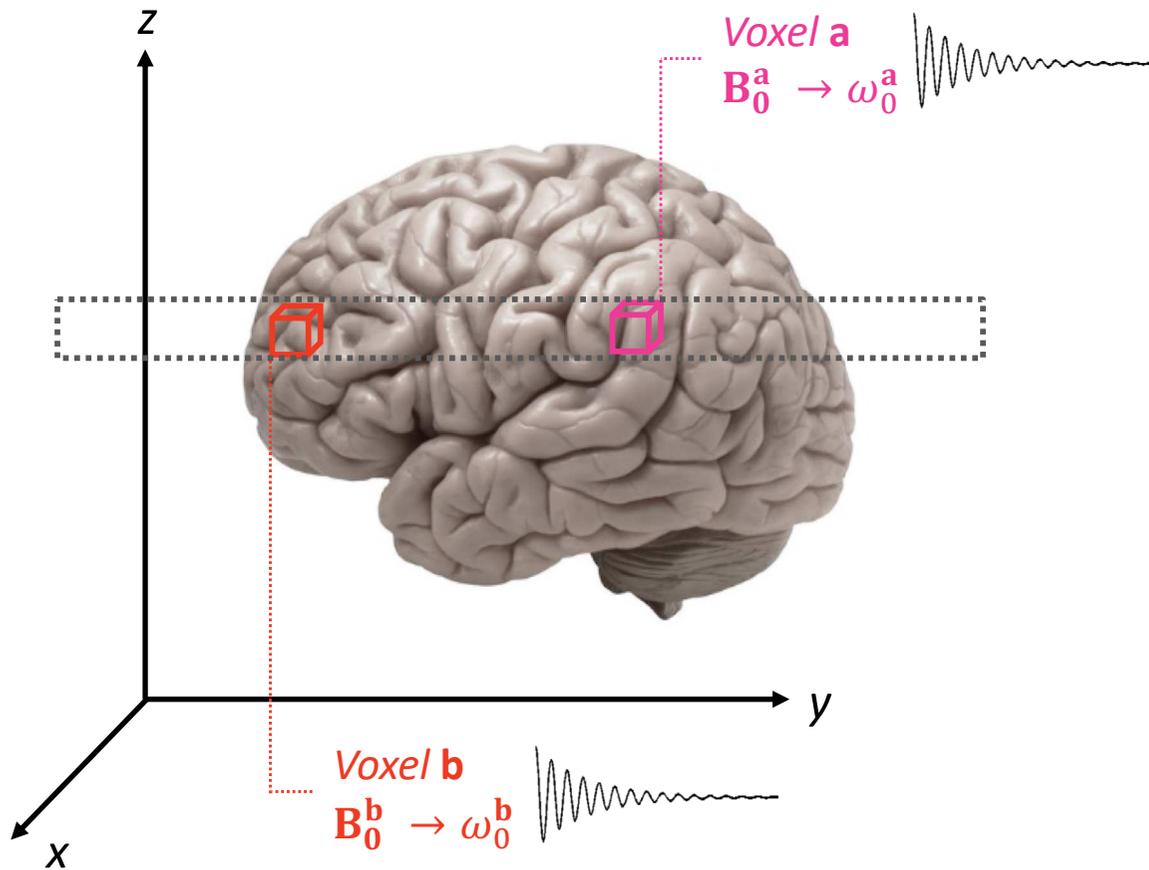
Applications : IRM / SRM

Imagerie : IRM

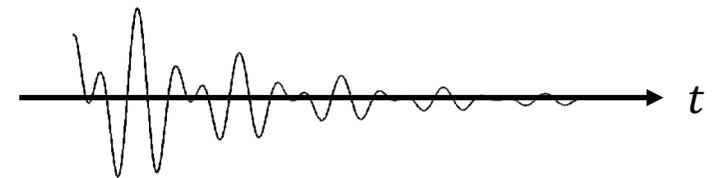


Applications : IRM / SRM

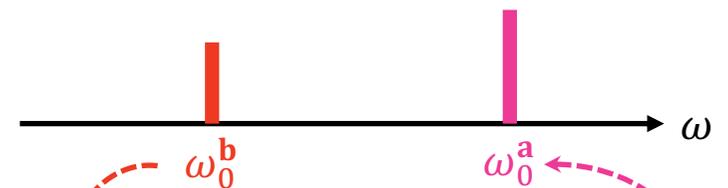
Imagerie : IRM



FID = signal (a) + signal(b)



Spectre (FID)

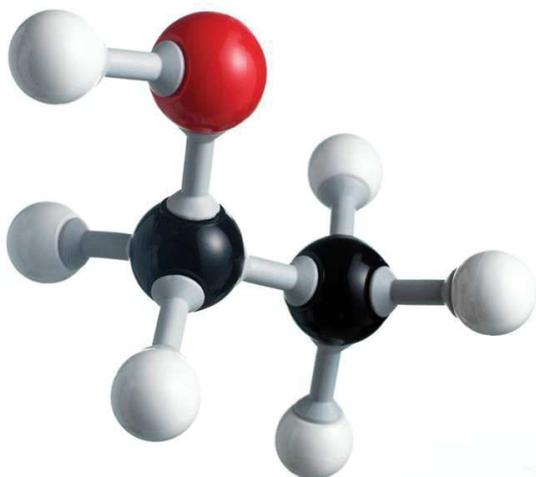


Applications : IRM / SRM

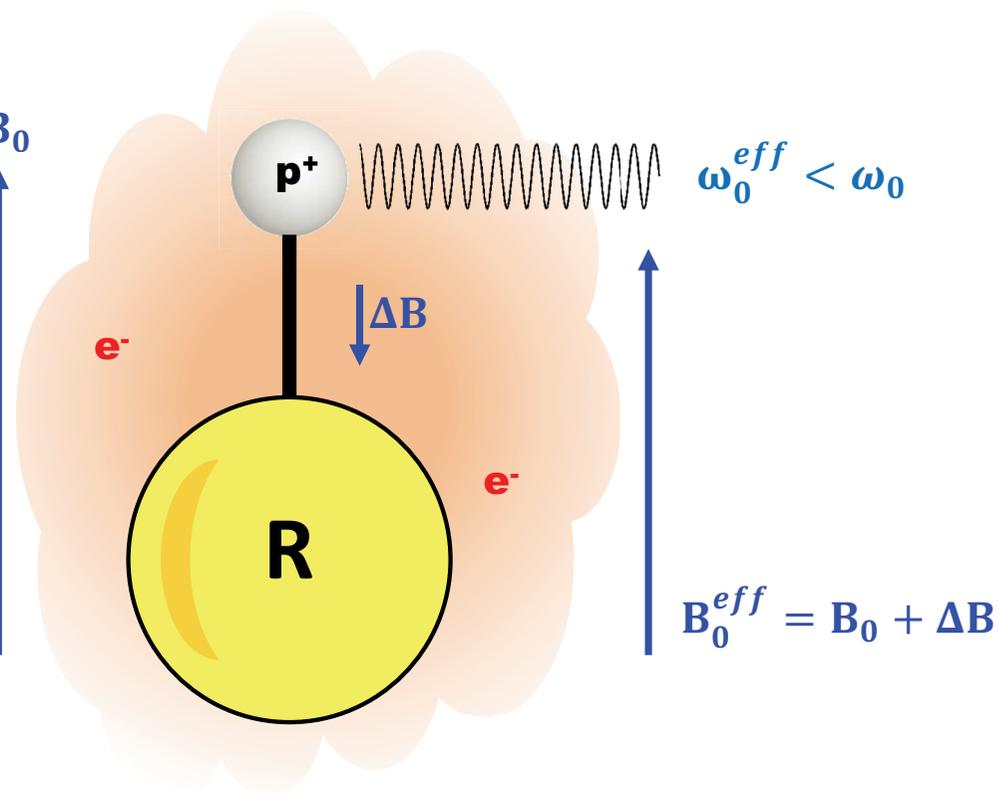
Spectroscopie : SRM

Ethanol $\text{CH}_3\text{-CH}_2\text{-OH}$

B_0



B_0



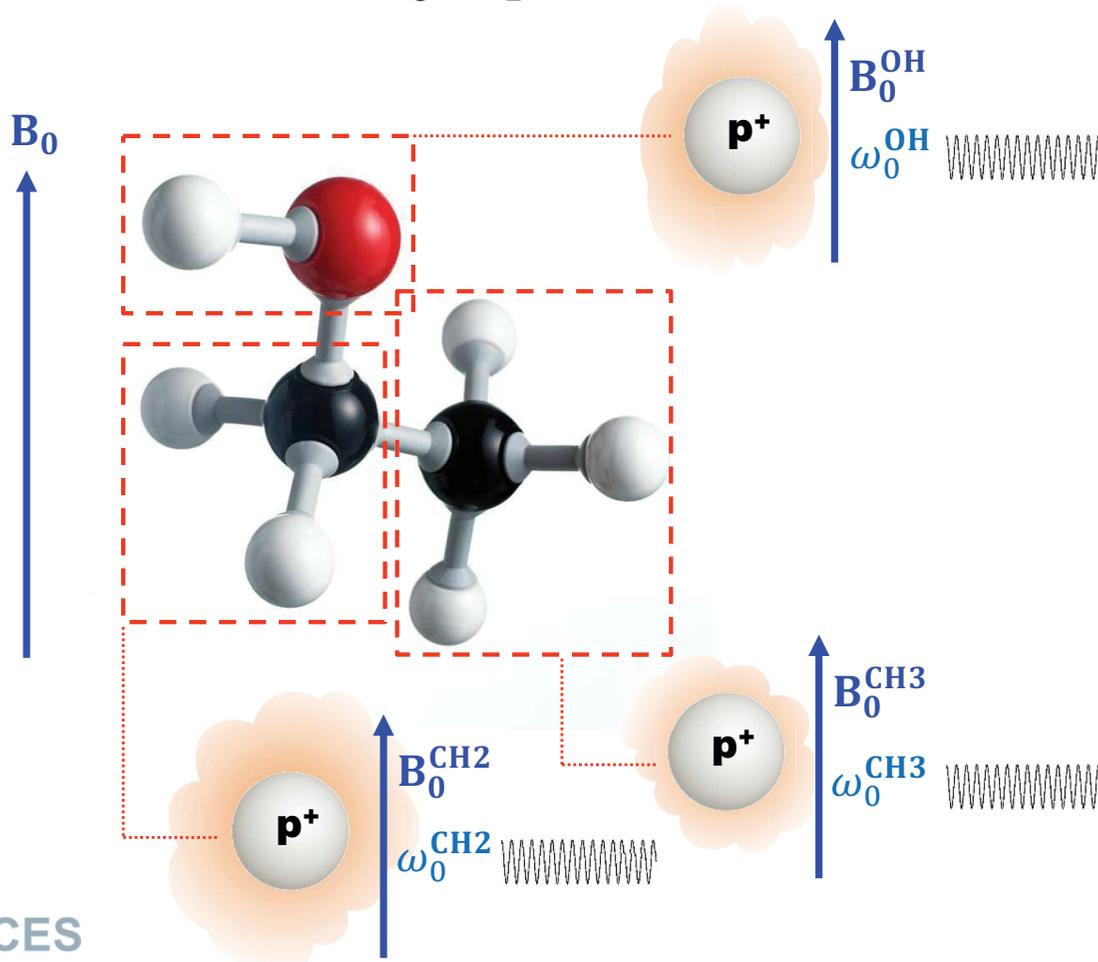
$$\text{Déplacement chimique : } \delta = 10^6 \frac{\omega_0^{eff} - \omega_0^{ref}}{\omega_0} \text{ [ppm]}$$



Applications : IRM / SRM

Spectroscopie : SRM

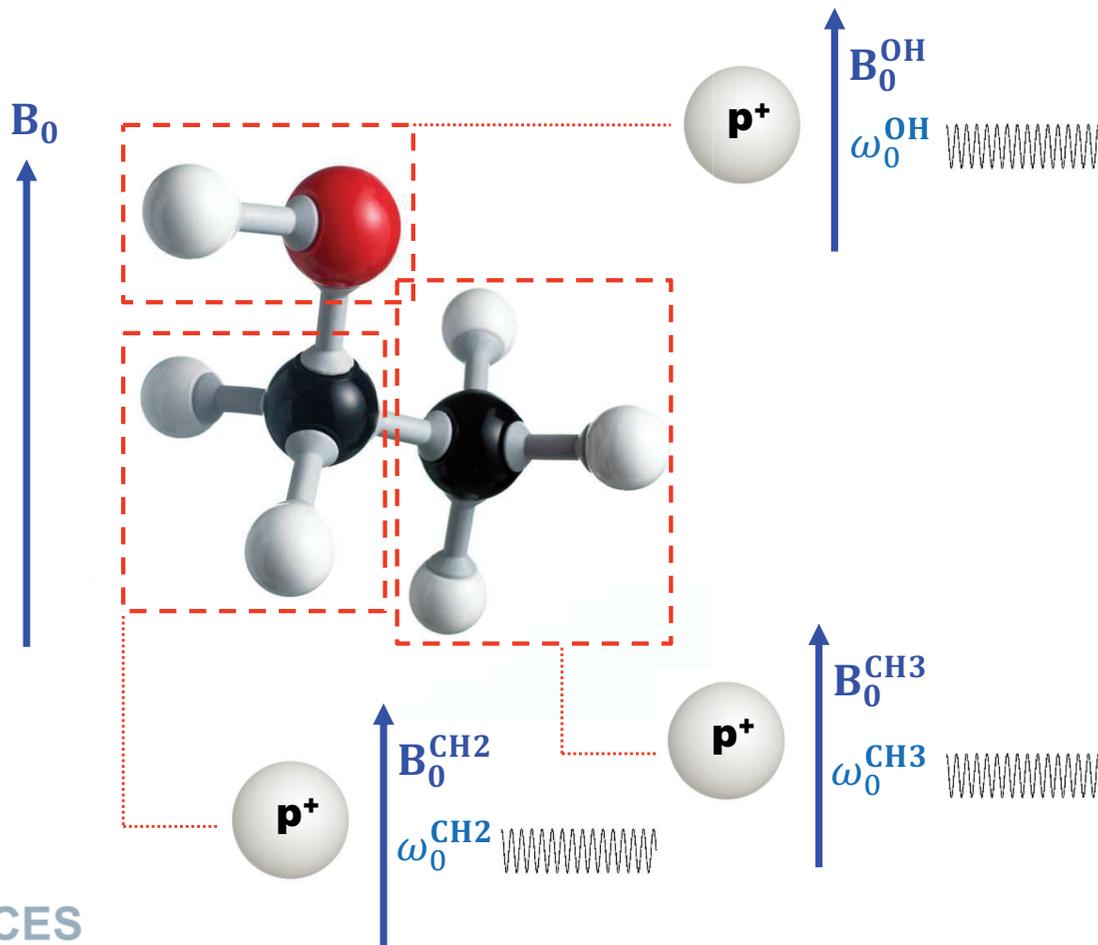
Ethanol $\text{CH}_3\text{-CH}_2\text{-OH}$



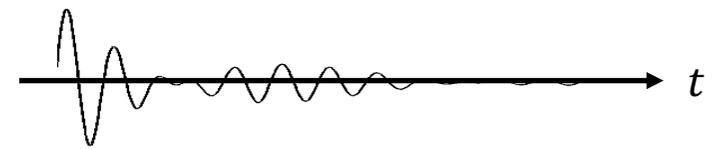
Applications : IRM / SRM

Spectroscopie : SRM

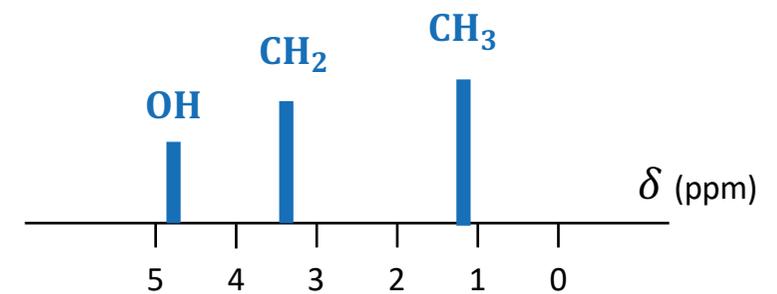
Ethanol $\text{CH}_3\text{-CH}_2\text{-OH}$



FID = signal (OH) + signal(CH₂) + signal(CH₃)

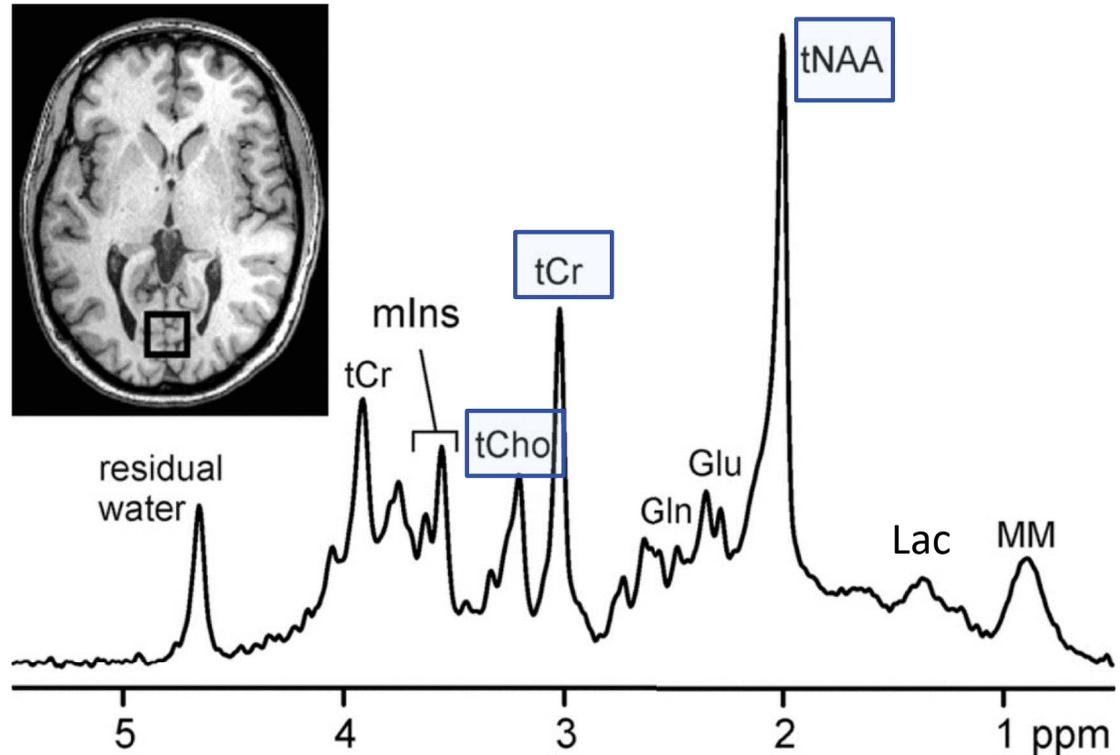
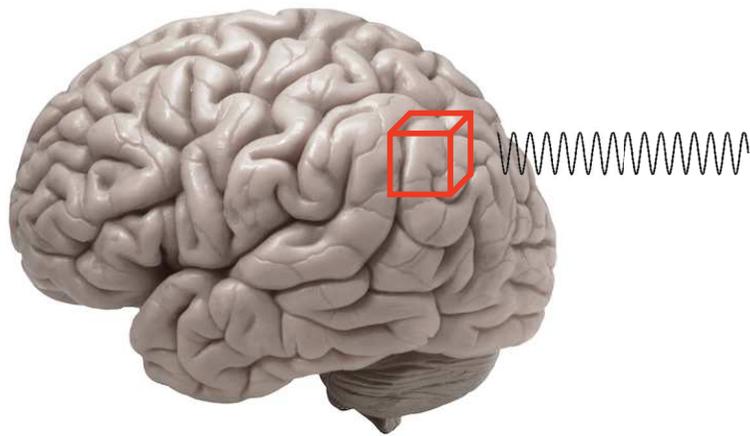


Spectre (FID)



Applications : IRM / SRM

Spectroscopie : SRM



Cho : choline - catabolisme membranaire

Cr : créatine - métabolisme aérobie

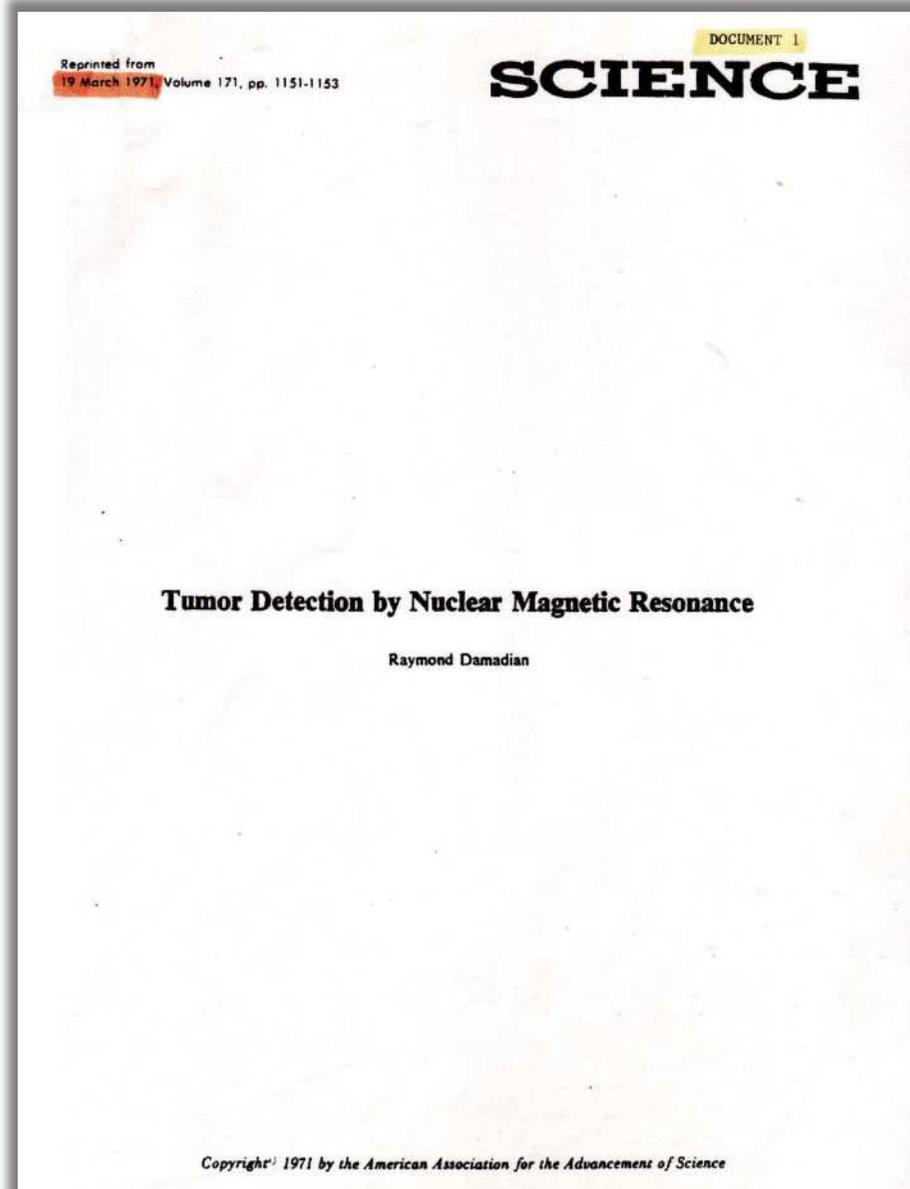
NAA : N-acétylaspartate - viabilité cellulaire

Lac : lactates - métabolisme anaérobie

MM : macro-molécules

Gülin et al. Radiology 2014.

IRM – exemples



IRM – exemples

Table 1. Spin-lattice (T_1) and spin-spin (T_2) relaxation times (in seconds) in tumors.

Rectus muscle		Liver	
T_1	T_2	T_1	T_2
0.493	0.050	0.286	0.050
.548	.050	.322	.060
.541	.050	.241	.050
.576 (0.600)*	.070	.306 (0.287)*	.048
.531		.300	
Mean and S.E.		Mean and S.E.	
0.538 ± 0.015	0.055 ± 0.005	0.293 ± 0.010	0.052 ± 0.005

relaxation time after the specimen stood overnight at room temperature.

Table 2. Spin-lattice (T_1) and spin-spin (T_2) relaxation times (in seconds) in tumors.

Rat No.	Weight (g)	T_1	T_2
<i>Walker sarcoma</i>			
6	156	0.700	0.100
7	150	.750	.100
8	495	.794 (0.794)*	.100
9	233	.688	
10	255	.750	
Mean and S.E.		0.736 ± 0.022	.100
P		< .01	
<i>Novikoff hepatoma</i>			
11	155	0.798	0.120
12	160	.852	.120
13	231	.877	.115
Mean and S.E.		0.826 ± 0.013	0.118 ± 0.002
P		< .01	
<i>Fibroadenoma (benign)</i>			
14		0.448	
15		.537	
Mean		.492	
<i>Distilled water</i>			
		2.691	
		2.690	
		2.640	
Mean and S.E.		2.677 ± 0.021	

* Spin-lattice relaxation time after the specimen stood overnight at room temperature. † The P values are the probability estimates of the significance of the difference in the means of T_1 for the malignant tumor and for brain.

Kidney T_1	Brain T_1
	0.573
	.573
	.596
41 (0.530)*	520 (0.614)
89	512
480 ± 0.026	0.595 ± 0.007

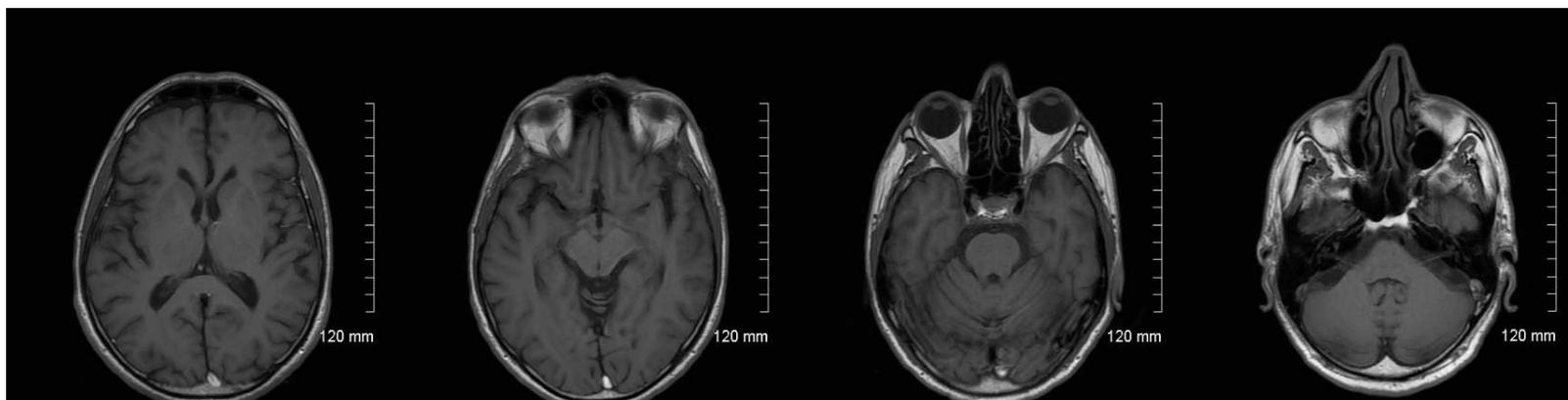


IRM – exemples

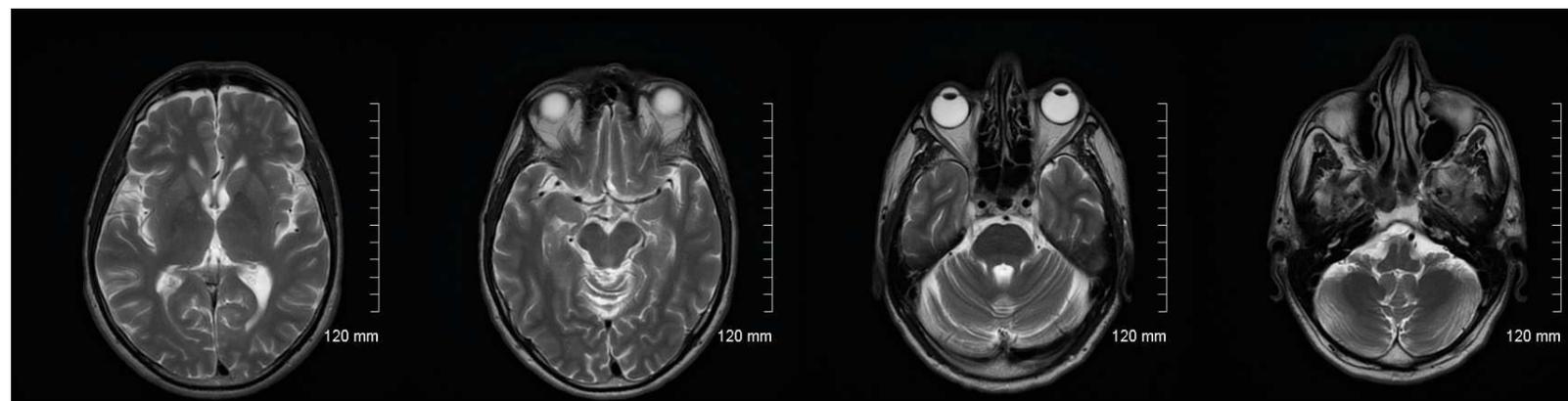
T_1 : contraste anatomique

T_2 : contraste « inversé »

T_1



T_2

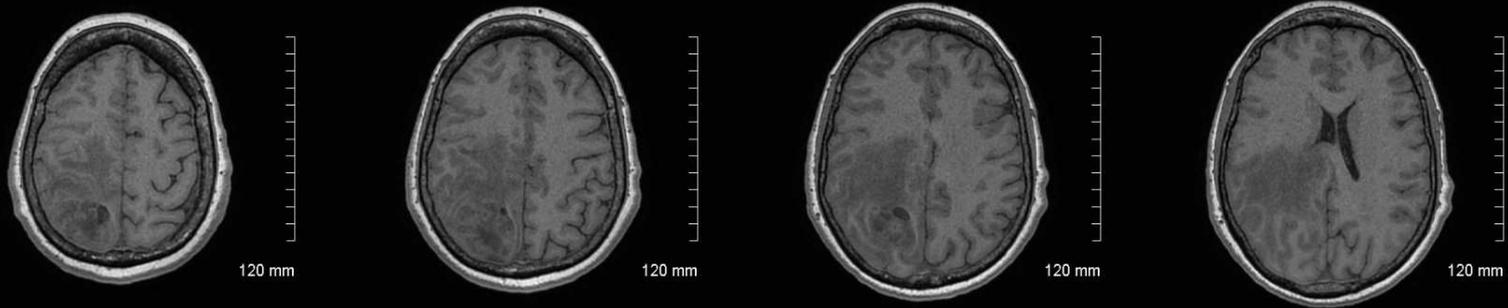


IRM – exemples

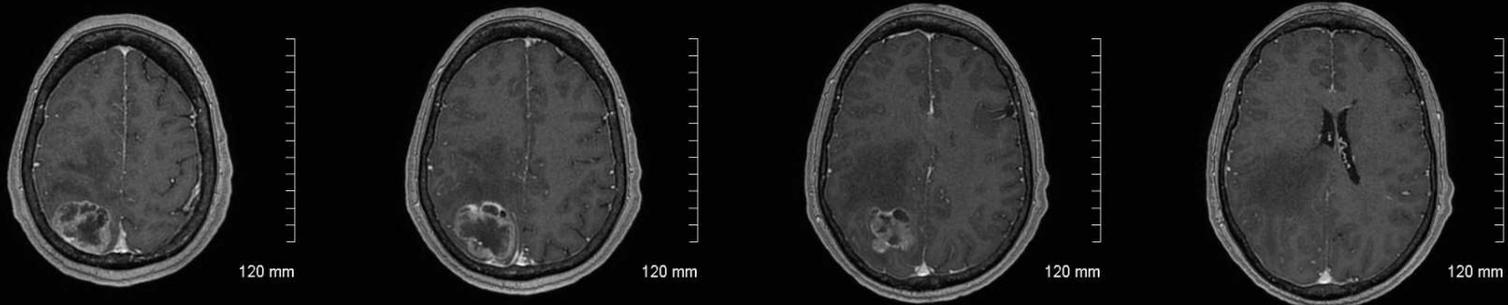
Métastase cérébrale

- Rupture BHE
- Effet de masse
- Œdème

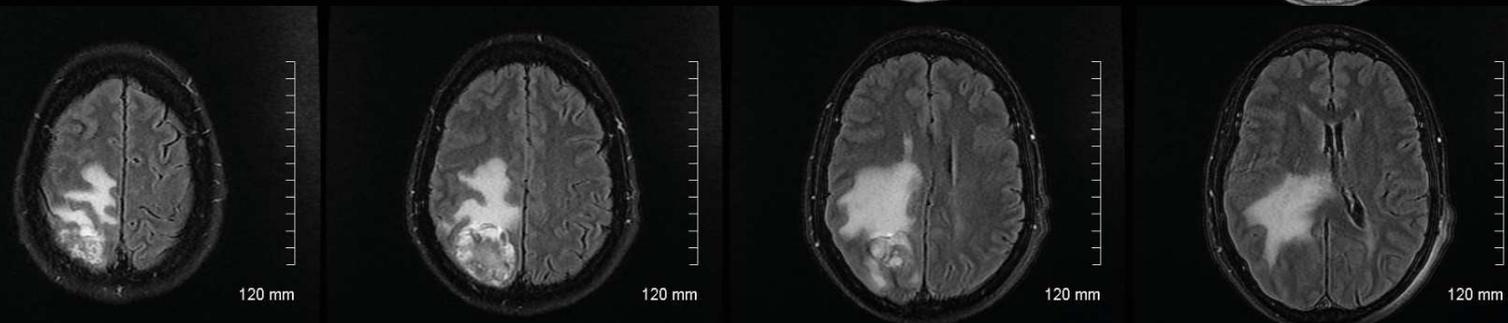
T_1



T_1 gado.



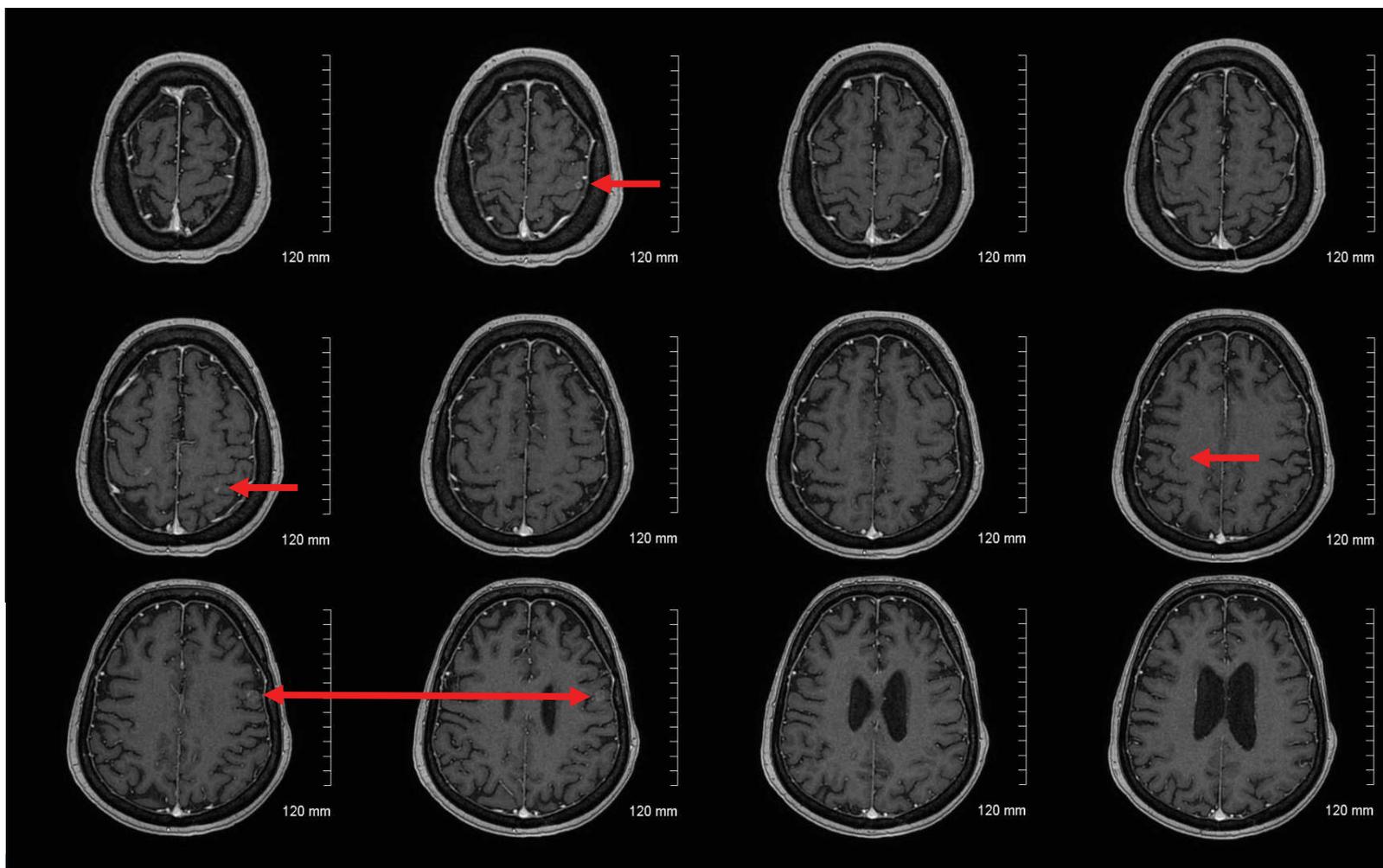
T_2 FLAIR



IRM – exemples

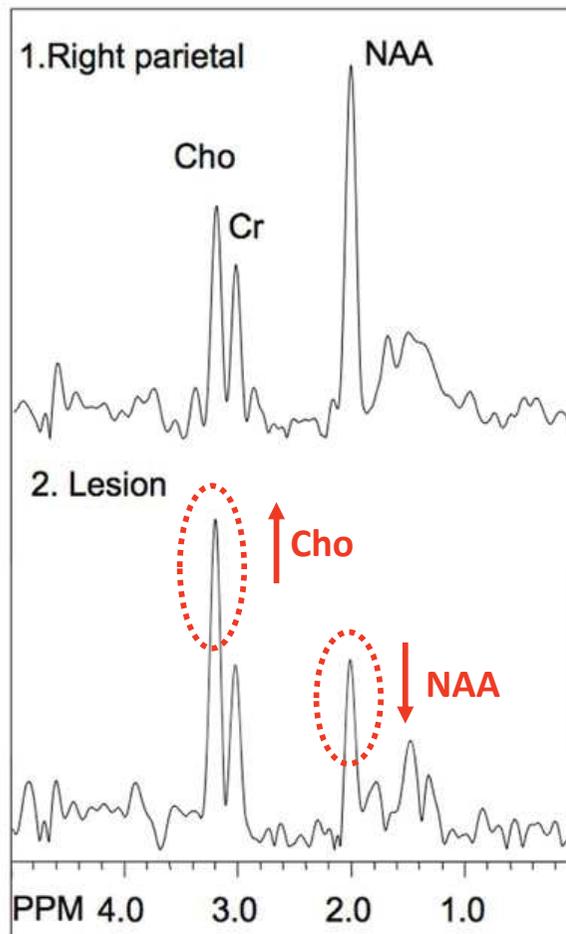
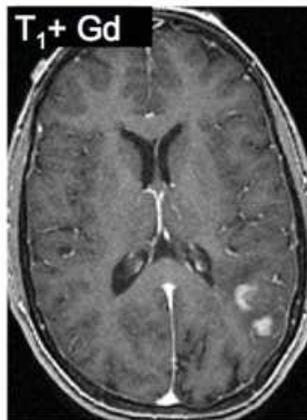
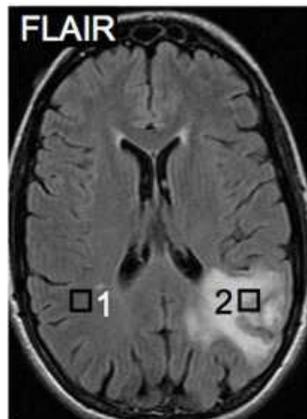
Bilan initial ADK pulmonaire

T₁ gado.

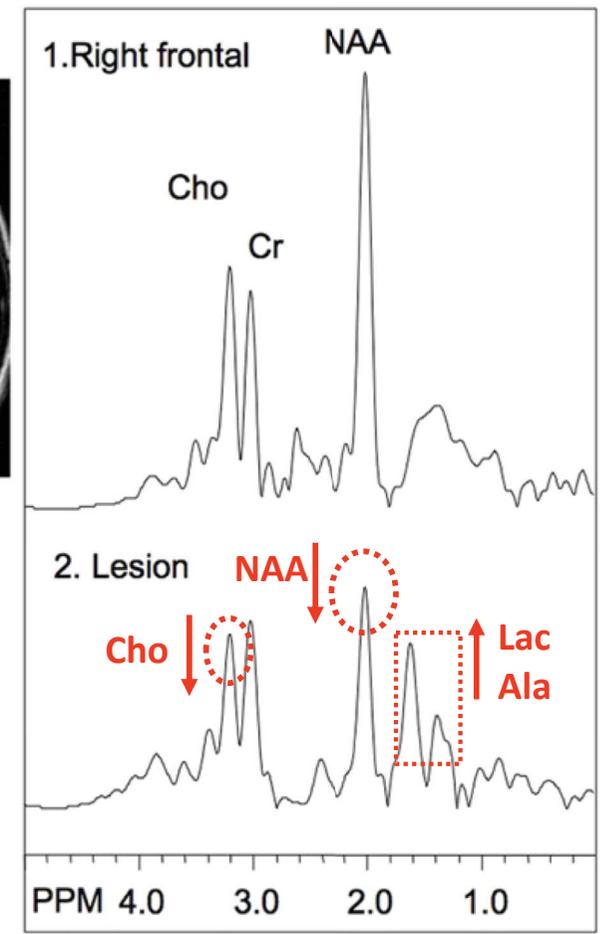
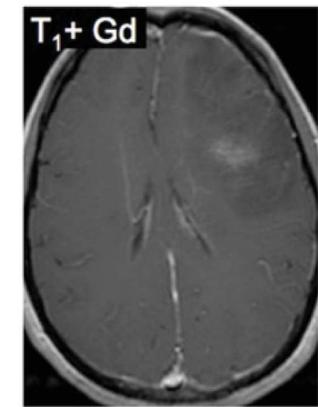
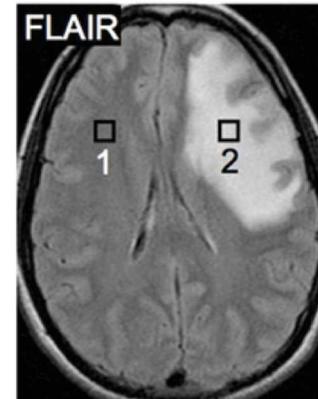


IRM – exemples

Lymphome cérébral

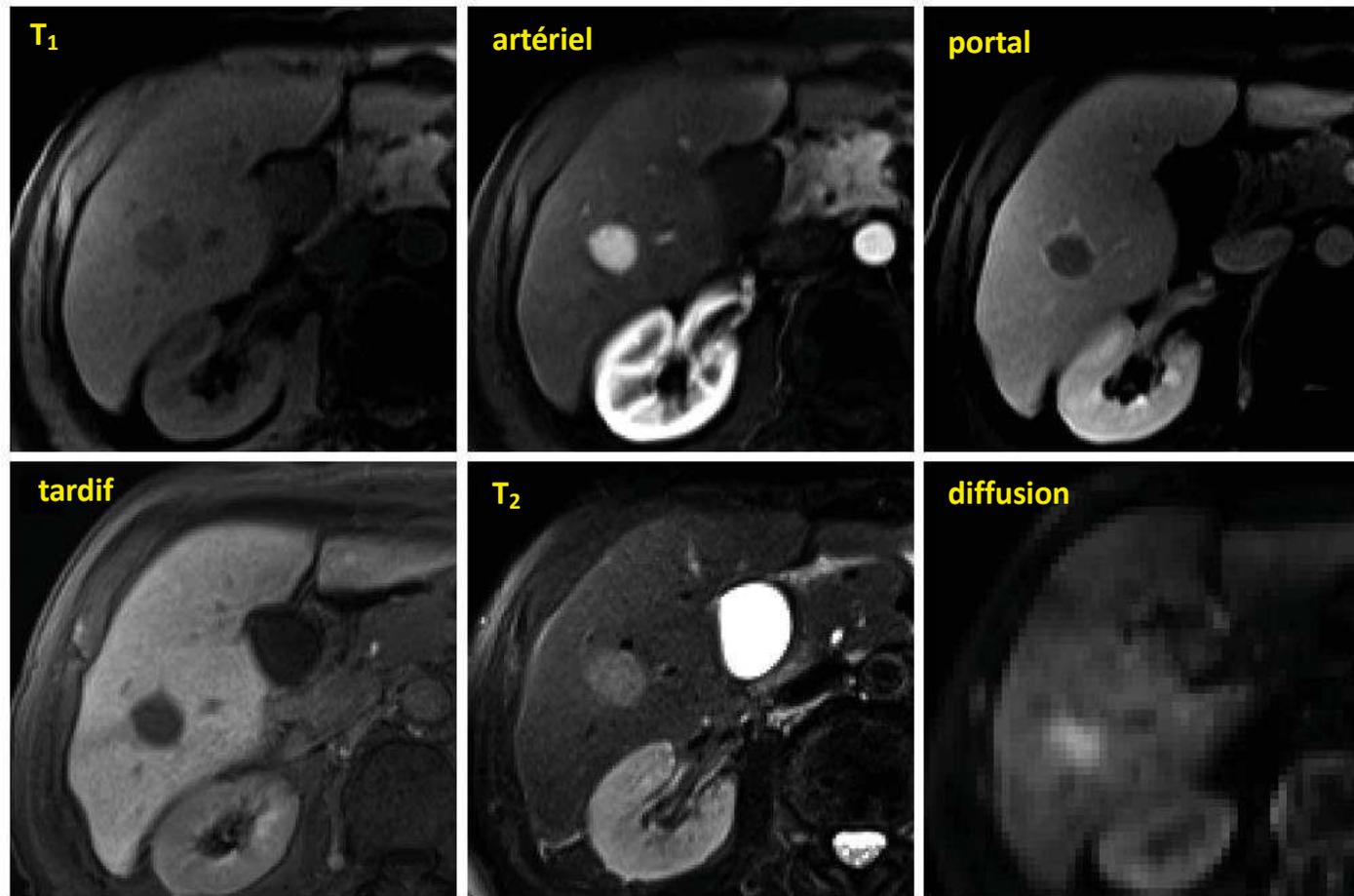


Méningoencéphalite



IRM – exemples

Carcinome hépatocellulaire



Park et al. World J Gastroenterol 2016.

IRM – exemples

Angiome hépatique

