

Study of pulse-coupled neural networks for glioma segmentation.

Brad Niepceron, Ahmed Nait-Sidi-Moh and Filippo Grassia. University of Picardie Jules Vernes

Speaker : Brad Niepceron. brad.niepceron@etud.u-picardie.fr

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CONTEXT

Towards lighter
diagnosis solutions

High interest for **computer aided diagnosis**

Deep Learning as a solution for medical image analysis

- Training models is **time** consuming
- Need for expensive dedicated **hardware**
- High energy cost

Rise of **biologically plausible** solutions

Image processing using **Pulse-coupled neural networks**

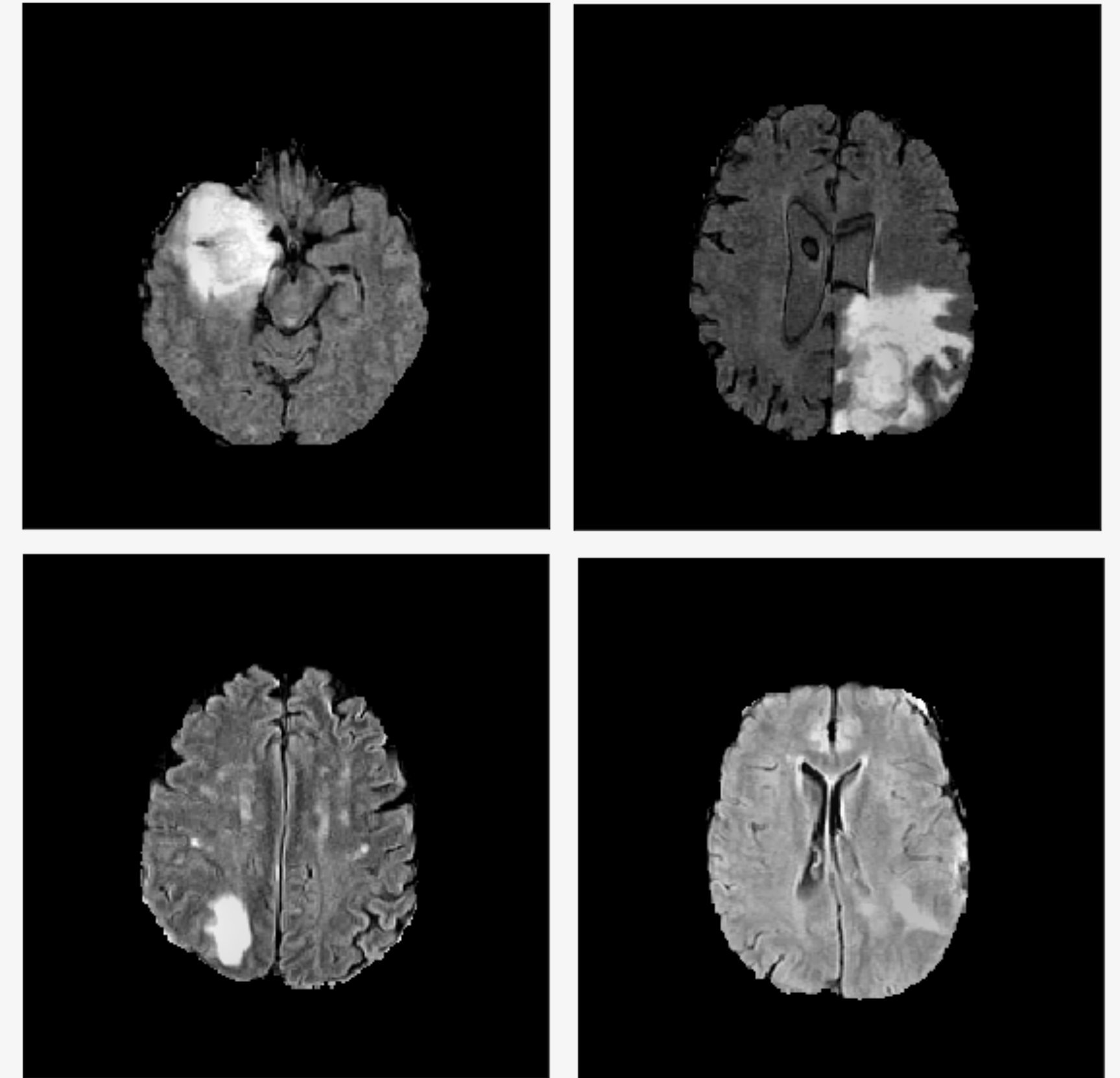


Fig 1. Example of MRI

Dataset info :

- Pre-operative MRI scans
- Manually** segmented by 1-4 raters
- Four MRI sequences : **Flair, T2, T1, T1C**
- Contains **3** unhealthy cell labels
- Focused on **segmentation** and **overall survival**

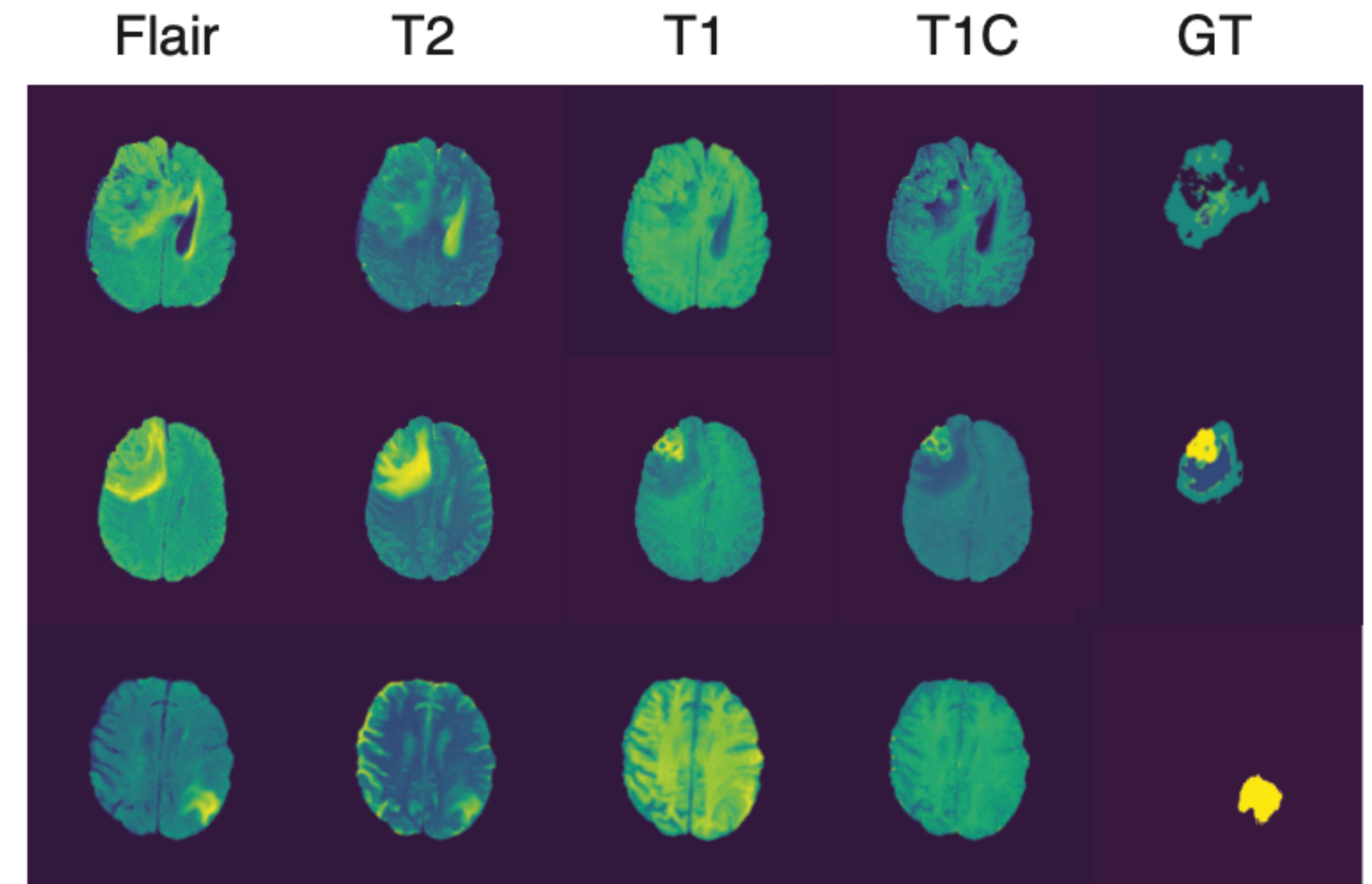


Fig 2. MRI cases taken from the BRaTS Dataset

Pixel intensity normalization and histogram matching applied

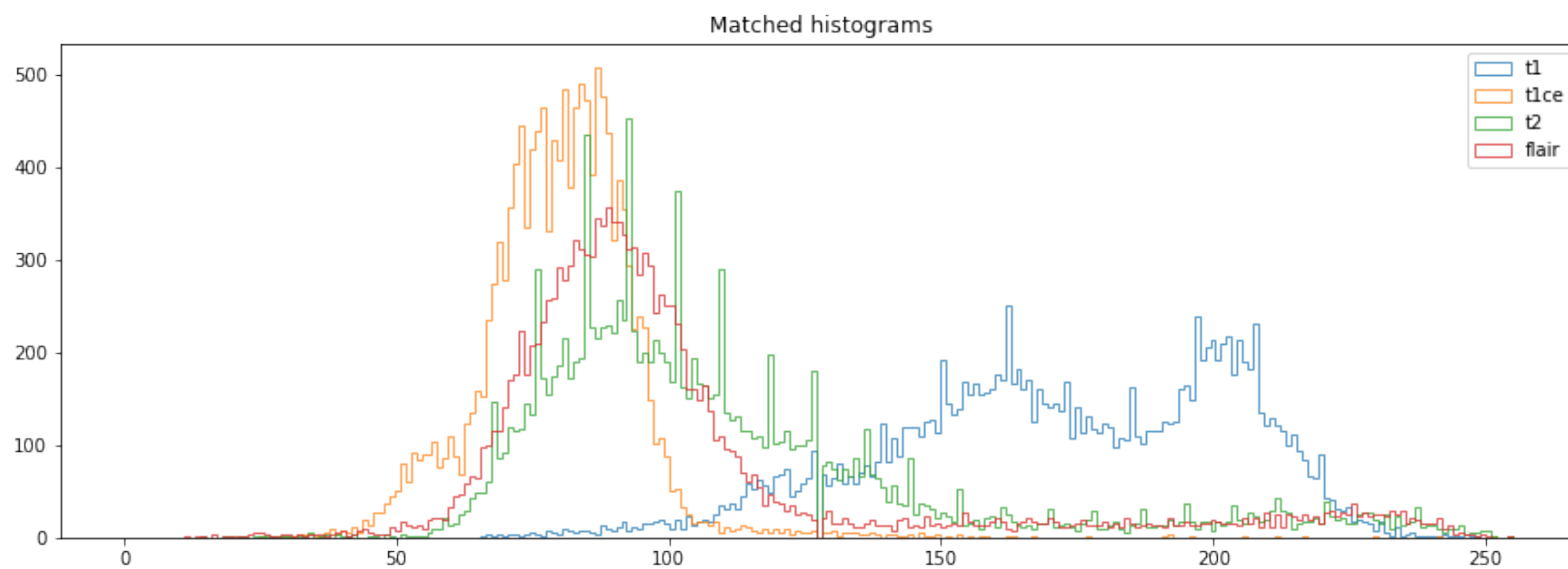


Fig 3. MRI sequence histogram matching

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METHODS

BRaTS 2020 Dataset

 Fusion strategy using Discrete Wavelet Transform

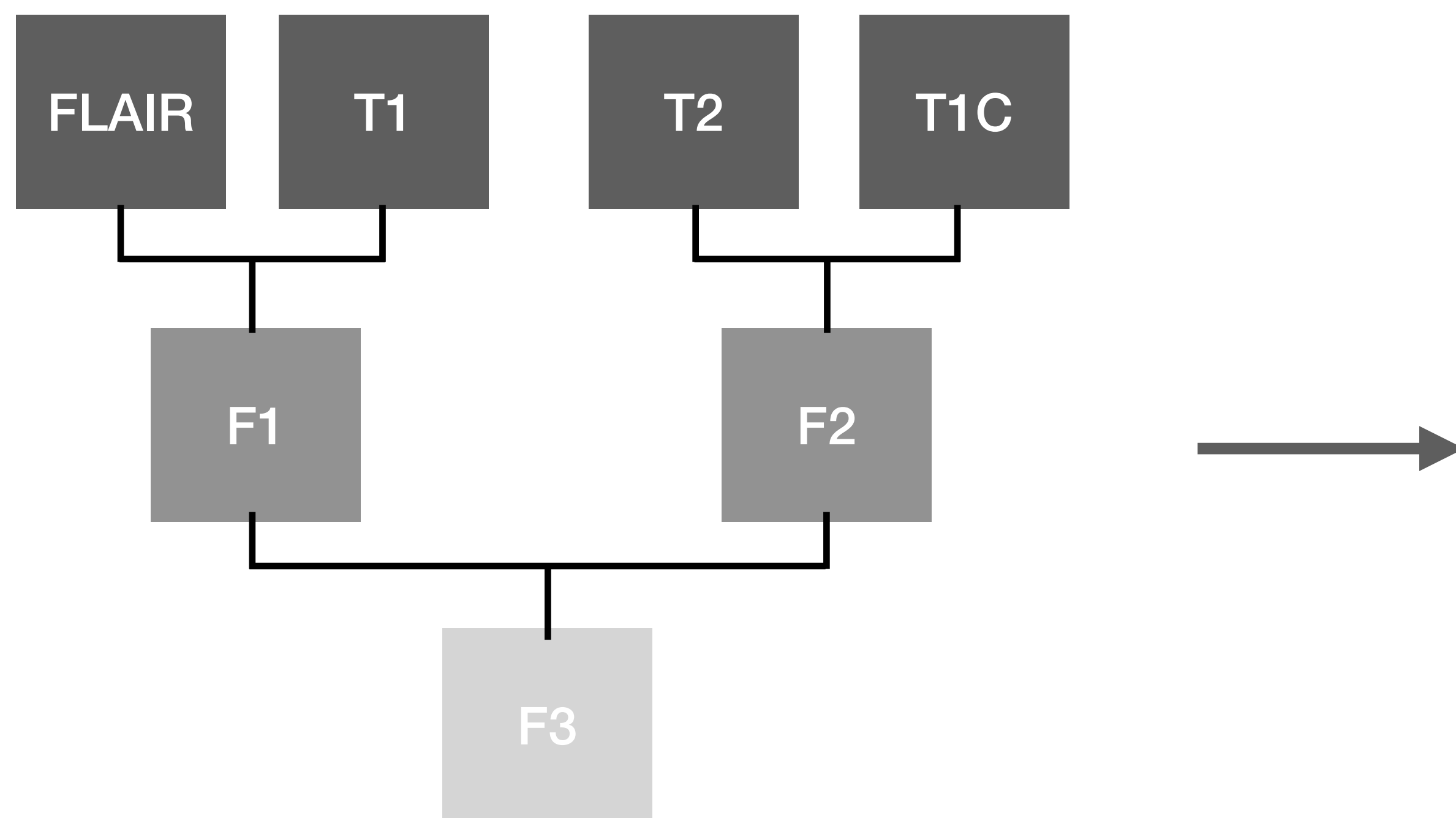


Fig 4. DWT computation tree

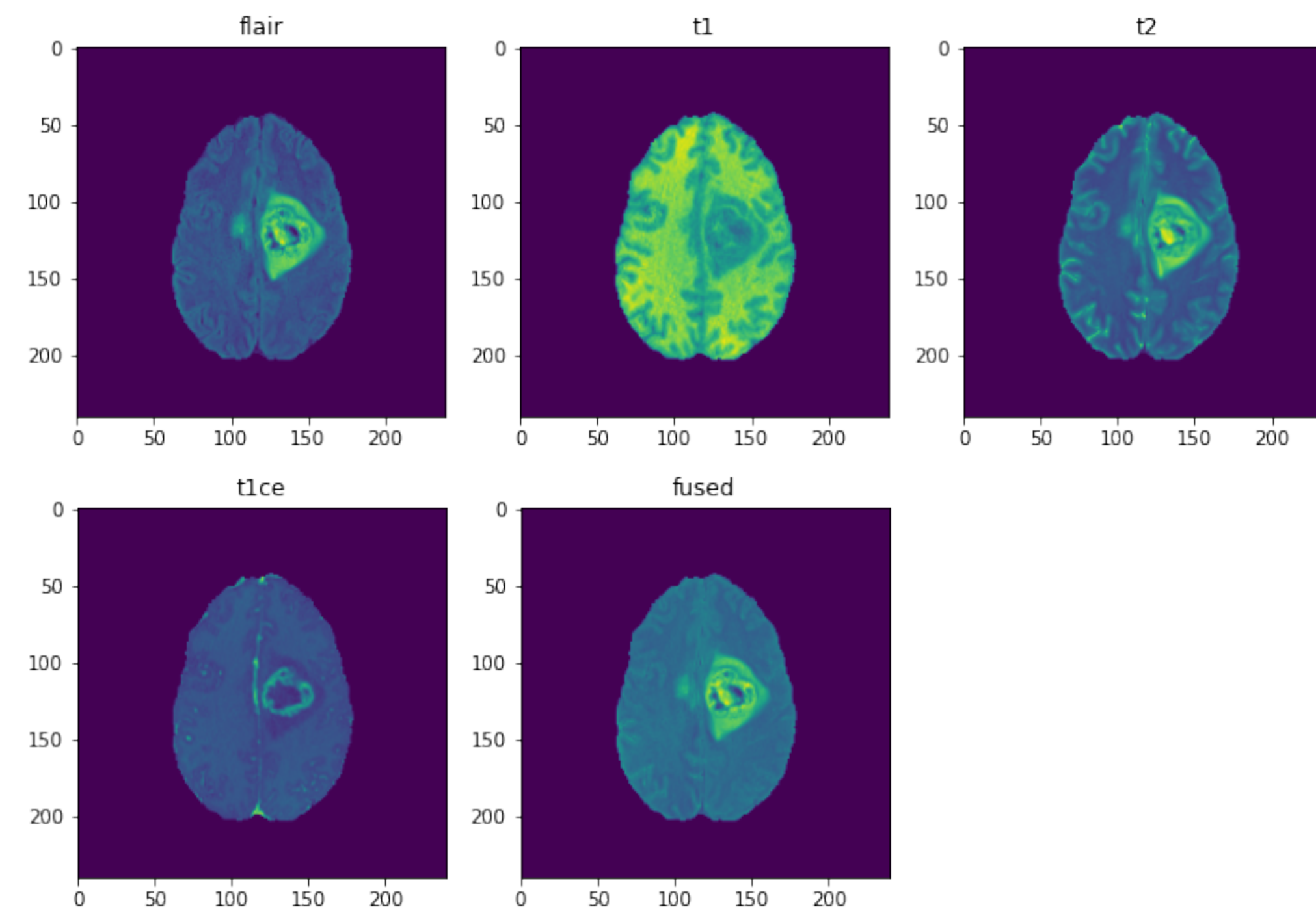


Fig 5. Sequences and fused image

Standard PCNN :

- Laterally-connected neurons
- 2D input image
- Feeding** : Computes voltage with input stimulus
- Linking** : Updates neuron's internal activity
- Pulse** : Fire if membrane potential exceeds threshold

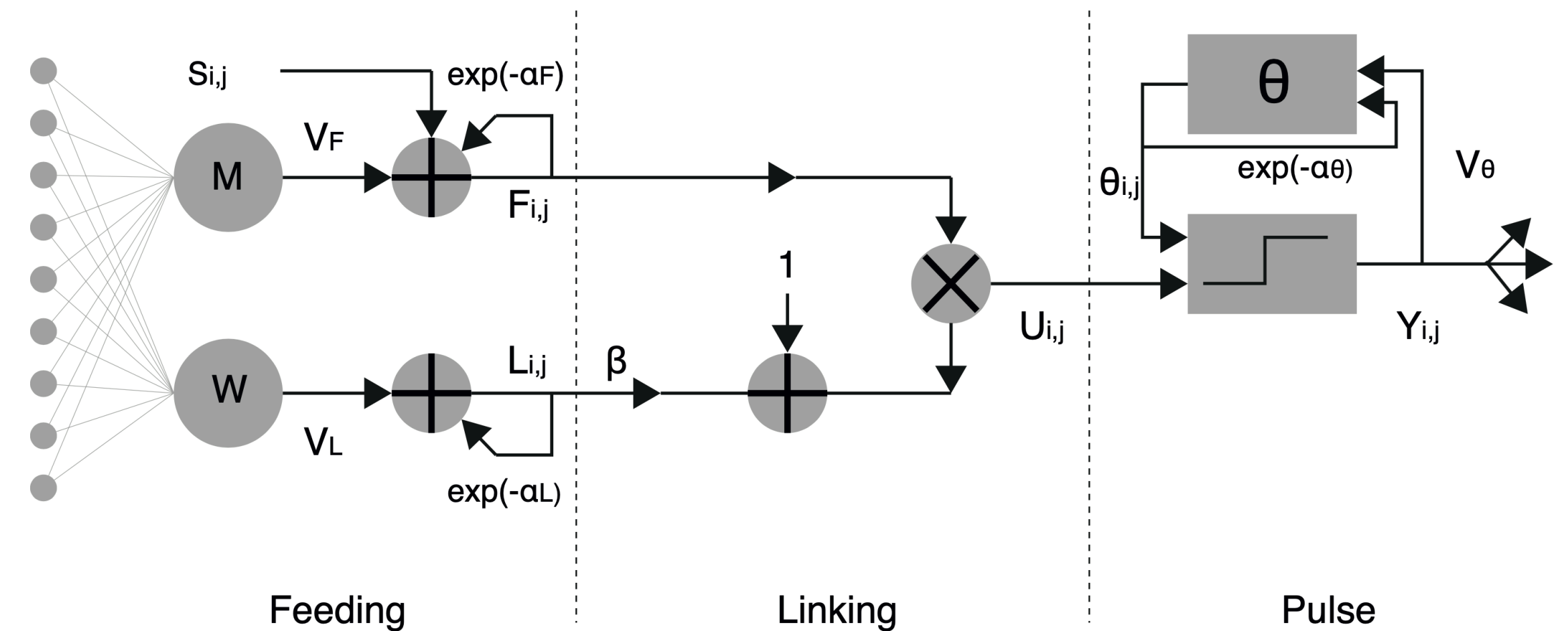


Fig 6. Standard PCNN

Unit-linking

- Simplified version of PCNN
- Reduced computational cost
- Reduced parameters

Changes

- **Feeding :** Now equal to the intensity of a pixel
- **Linking :** Allows a neuron to fire when one or more neighbors fired

Fast-linking

- Uses Spiking cortical neurons
- Fast linking synapses
- Neurons fire faster

Changes

- **Feeding :** As for Unit-linking
- **Pulse :** Combines stimulus and synaptic modulation to charge the membrane

Parameters settings :

- Differential Evolution for optimization
- Dice Score as loss function $\frac{2|Y \cap \hat{Y}|}{|Y| + |\hat{Y}|}$
- Iteration number constrained to 10

Model	Standard PCNN	ULPCNN	FLSCM
β	0.47	0.47	0.44
α_{θ}	0.0125	0.015	-
α_F	0.96	-	-
α_L	0.81	-	-
V_{θ}	20	20	20
V_F	0.21	-	-
V_L	0.36	-	-
W	3x3 gaussian kernel	-	-
M	W	-	-
α_U	-	-	0.49

Table 1. Parameters used for our experiments

RESULTS

Segmentation Evaluation

Segmentation evaluated with Dice Score

Iteration can be stopped if best dice score has been found.

Fast-linking average running time : **17 seconds**

Unit-linking average running time : **17 seconds**

PCNN average running time : **60 seconds**

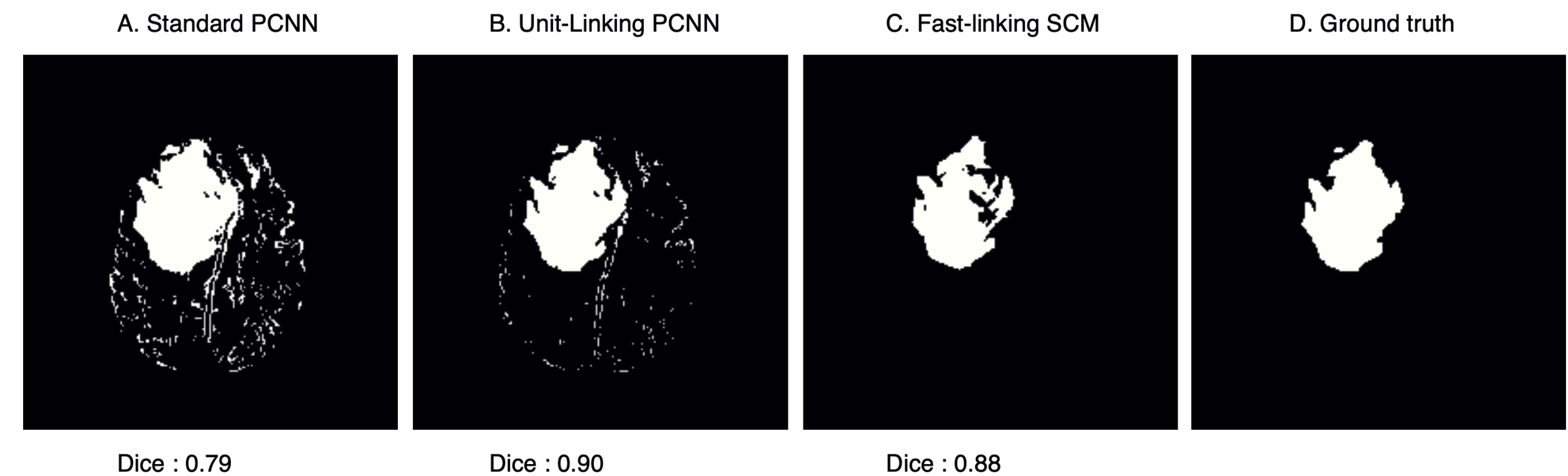



Fig7. Results of PCNN based segmentation



Experiments proved the efficiency of PCNN models



Fast computation makes PCNN a **perfect fit for diagnosis tasks**



Differential Evolution coupled with **Dice Loss** appeared efficient for segmentation parameters optimization



Multi-channel versions could be use for **semantic segmentation**



PCNN can be coupled with a **spiking classifier**

Thank you.

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