Longitudinal automated assessment of paramagnetic rim lesions in multiple sclerosis using RimNet

Maxence Wynen,¹ Francesco La Rosa,²,³,⁴ Amina Sellimi,⁵ Germán Barquero,²,³,⁴ Gaetano Perrotta,⁶ Valentina Lolli,² Vincent Van Pesch,⁵ Cristina Granziera,⁰, Tobias Kober,¹⁰ Pascal Sati,¹¹,¹², Benoit Macq,¹³ Daniel S. Reich,¹¹ Martina Absinta,¹¹,¹⁴, Meritxell Bach Cuadra,²,³,⁴,⁺ Pietro Maggi⁵,¹⁵,⁺

SYNOPSIS (100 words)

The automatic assessment of paramagnetic rim lesions in multiple sclerosis is important, and a deep learning-based algorithm called RimNet has recently been proposed. This work evaluates the generalizability of RimNet and its longitudinal performance on MRI data acquired at different clinical centers. We found that RimNet's performance was nearly as good on totally unseen data as in the original paper (receiver-operating-characteristic area-under-the-curve (AUC) 0.88 vs. 0.94, precision-recall AUC 0.69 vs. 0.70), and it made consistent predictions on longitudinal data (binary consistency 82%, probability consistency 93%).

SUMMARY OF THE MAIN FINDINGS

RimNet, a deep-learning-based segmentation algorithm for chronic active multiple sclerosis lesions, is effective for unseen MRI data acquired on a scanner from a different vendor, and makes consistent predictions on longitudinal data.

¹ Ecole Polytechnique de Louvain, Université Catholique de Louvain, Louvain-la-Neuve, Belgium.

² Signal Processing Laboratory (LTS5), École Polytechnique Fédérale de Lausanne, Switzerland.

³ Medical Image Analysis Laboratory, Center for Biomedical Imaging (CIBM), University of Lausanne, Switzerland.

⁴ Radiology Department, Lausanne University Hospital and University of Lausanne, Switzerland.

⁵ Department of Neurology, Cliniques Universitaires Saint-Luc, Université Catholique de Louvain, Brussels, Belgium.

⁶ Department of Neurology and ⁷ Radiology, Erasme University Hospital, Université Libre de Bruxelles, Brussels, Belgium.

⁸ Neurologic Clinic and Policlinic, Departments of Medicine, Clinical Research and Biomedical Engineering, University Hospital Basel and University of Basel, Switzerland.

⁹ Translational Imaging in Neurology (ThINk) Basel, Department of Medicine and Biomedical Engineering, University Hospital Basel and University of Basel, Basel, Switzerland.

¹⁰ Advanced Clinical Imaging Technology, Siemens Healthcare AG, Lausanne, Switzerland.

¹¹ Translational Neuroradiology Section, National Institute of Neurological Disorders and Stroke, National Institutes of Health, Bethesda, MD, USA.

¹² Department of Neurology, Cedars-Sinai Medical Center, Los Angeles, CA, USA.

¹³ ICTEAM Institute, Université catholique de Louvain, Louvain-la-Neuve, Belgium.

¹⁴ Department of Neurology, Johns Hopkins University, Baltimore, MD, USA

¹⁵ Department of Neurology, Lausanne University Hospital and University of Lausanne, Lausanne, Switzerland.

^{*} the last two authors contributed equally

INTRODUCTION

In multiple sclerosis (MS), some brain white matter lesions are more destructive and feature an edge of chronic/smoldering inflammation, which corresponds to a paramagnetic rim on susceptibility-based MRI. Higher paramagnetic rim lesion (PRL) burden is associated with more aggressive disease course, however the longitudinal effect of available MS disease modifying treatments (DMT) on PRL is still largely unknown. Considering that manual rim classification is tedious and associated with inter-observer variability, a multimodal convolutional neural network for automated detection (RimNet) has been recently developed In this work, we tested RimNet (i) on data acquired in two centers using MRI scanners from two different vendors (including one not used for training in the original paper); and (ii) longitudinally on MRI data acquired in primary and secondary progressive MS patients (PMS) before and after DMT.

METHODS

Two sets of MRI images were acquired from 13 PMS patients [mean ± SD age: 44 ± 10; female/male: 4/9; median EDSS (range): 4.5 (3.0–6.5)] who started a DMT (ocrelizumab or siponimod) at 2 different hospitals. The first dataset (baseline) was acquired before DMT administration, and the second at a median time of 13 months (range: 6–18) after DMT administration (follow-up). The protocol included 3D Fluid Attenuated Inversion Recovery (FLAIR) and 3D T2*-weighted echo planar imaging (T2*-EPI), providing T2* magnitude and phase images (which we simply refer to as T2* and phase, respectively). FLAIR* images were generated by voxel-wise multiplication of the FLAIR and T2*-EPI images, as previously described. The scanners and acquisition parameters are reported in Table 1.

Two experts manually annotated the PRL by examining phase images by consensus. The first pre-processing step consisted of motion-correction of the FLAIR images using Freesurfer⁶, and then automated lesion segmentation using SAMSEG, ⁷ yielding a total of 704 lesions. The next steps consisted of rigid registration of the FLAIR and lesion mask to the T2*-EPI space using ANTs,8 manual correction of the lesion mask (including splitting of confluent lesions), and rigid coregistration of baseline and follow-up images for all subjects. Next, all images were skull-stripped using FSL,9 and the intensities of FLAIR and T2* images were normalized using histogram matching¹⁰ to two reference images (T2* and FLAIR) used during the training. Finally, 3D patches of size 28x28x28 voxels were extracted for all lesions.⁴ The resulting lesions were then tested using two bimodal networks of RimNet⁴ pre-trained on different datasets. The first model, trained with phase and FLAIR (phase+FLAIR model), was tested with the same images. The other model, which was trained using phase and T2*-EPI, was tested on two image sets: one including phase and T2*-EPI (phase+T2*) and the other phase and FLAIR* (phase+FLAIR*). We evaluated PRL classification performance after fixing a false positive rate (FPR) of 0.10. In addition to commonly used metrics (accuracy, F1-score, receiver-operating-characteristic (ROC) and precision-recall (PR) curves per session), we also calculated a consistency measure "c" to evaluate the similarity between the baseline and follow-up predictions. This was computed for each PRL using the following formula:

$$c = 1-(|p1-p2|)$$

where p1 and p2 are, respectively, the binary or probability predictions of RimNet for the baseline (p1) and follow-up (p2) sessions. For the consistency analysis, we tested RimNet using the baseline

PRL patch as lesion mask in both sessions in order to eliminate possible differences due to the lesion segmentation method.

RESULTS

After experts' consensus, 54 PRL were found in each session (Figure 1). RimNet achieved an overall accuracy for phase+FLAIR*, phase+T2*, and phase+FLAIR of 87%, 85%, and 82%, respectively. The complete metrics for each separate session are shown in Table 2. The ROC and PR curve, and the relative AUC for the various models, are reported in Figure 2. The phase+FLAIR* model (AUC=0.88) clearly outperformed phase+T2* (AUC=0.83) and phase+FLAIR (AUC=0.72). Of 54 PRL, the phase+FLAIR* model gave the same binary prediction in both sessions for 44 (binary consistency 82%) and a probability consistency (before applying threshold) of 93%. The detailed results per patient of this analysis can be found in Figure 3.

DISCUSSION & CONCLUSION

In this work, we not only evaluated RimNet's generalizability on images acquired on a scanner from a different vendor (compared to the original training⁴) but also its consistency to make predictions on longitudinal PRL detection. Interestingly, the model trained on phase+T2* but tested on phase+FLAIR* outperformed the model trained and tested on phase+T2* (AUC=0.88 vs AUC=0.83), possibly because of a higher signal-to-noise ratio of FLAIR* vs. T2* images in the testing dataset. The overall results for the best model (phase+FLAIR*) had lower ROC AUC than reported in the original RimNet paper (0.88 vs. 0.94), but a similar PR AUC (0.69 vs. 0.70). This slight underperformance can be explained by the different scanners and acquisition parameters used, but RimNet's ability to generalize well to an unseen dataset is nevertheless promising. In the longitudinal assessment, the relatively high binary and probability base consistency of >80% and >90%, respectively, indicates that RimNet makes consistent predictions on longitudinal data, at least for two consecutive timepoints. Future work should confirm these hypotheses by studying a larger cohort of patients at more timepoints.

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FIGURES

Table 1: MRI acquisition parameters of the protocol used.

N. of image sets	24		2			
Magnet strength	3T		3T			
Manufacturer	Philip	s	Siemens			
Model	Ingeni	ia	Prisma/Skyra			
	3D T2*-EPI	FLAIR	3D T2*-EPI	FLAIR		
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	3D T2*-EPI	FLAIR	3D T2*-EPI	FLAIR	
Resolution (mm)	0.55x0.55x0.55	1.0x1.0x1.0	0.65x0.65x0.65	1.0x1.0x1.0	
Repetition time (TR, ms)	53	4800	64	5000	
Echo Time (TE, ms)	29	373	35	386	
Inversion time (TI, ms)	-	1600	-	1800	
Flip angle (degrees)	10	90	10	120	
Acquisition time	4'40"	6'00"	6'20"	4'47"	

Figure 1: Representative 3D FLAIR* and Phase images from one PMS patient showing the same PRL a) at baseline and b) at the 13 months post-DMT follow-up.

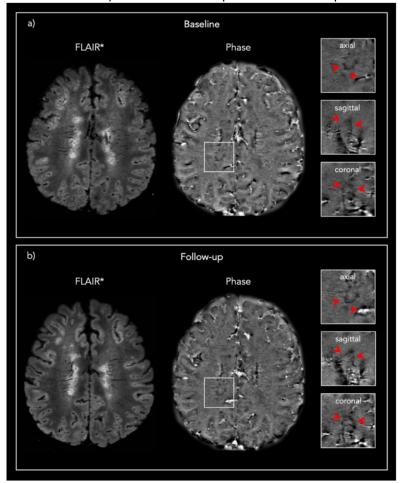


Figure 2: Combined (Figure 2a) and session-based (baseline Figure 2b and follow-up Figure 2c) receiver-operating characteristic (ROC) and precision-recall (PR) curves per model tested. AUC: area under the curve.

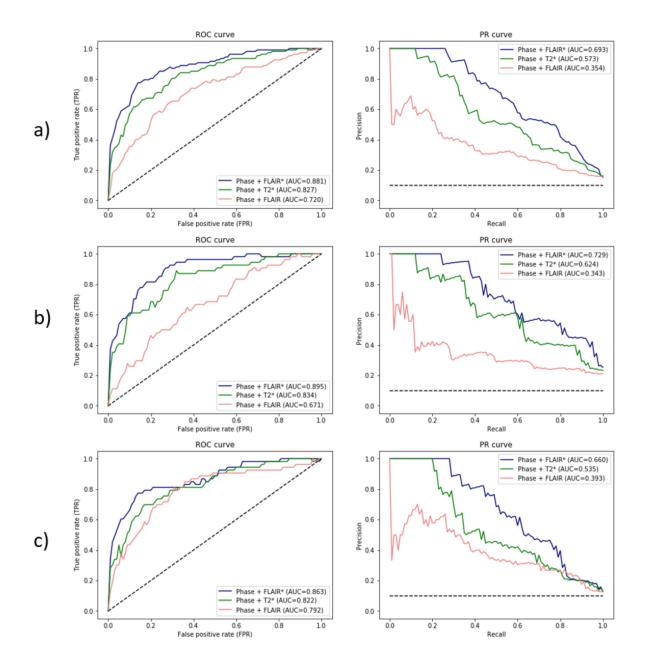


Figure 3: Summary of the lesion-wise results for each patient with paramagnetic rim lesions (n=11 of 13 total patients studied, 85%). a) Number of rims detected in session 1 (baseline) and session 2 (follow-up) for each patient, overlaid on the total number of rims adjudicated in the consensus manual reading. b) Number of matching predictions between baseline and follow-up. Abbreviations: Consistency_p, probability-based consistency; Consistency_b, binary-based consistency.

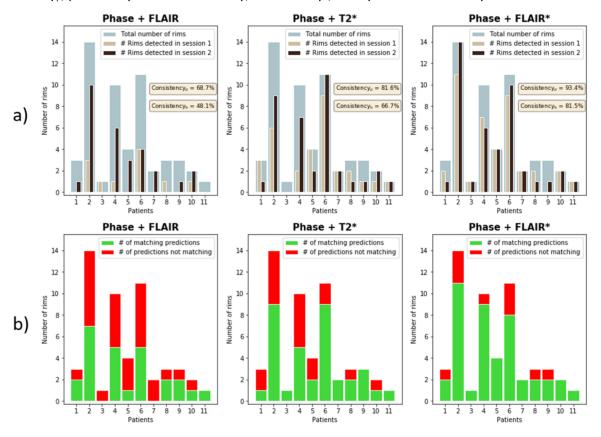


Table 2: Complete metrics per session for each model tested. Abbreviations: TN, True Negatives; FP, False Positives; FN, False Negatives; TP, True Positives; TPR, True Positive Rate; FPR, False Positive Rate; PPV, Positive Predictive Value; NPV, Negative Predictive Value.

Model	Session	TN	FP	FN	TP	Accuracy	F1	TPR	FPR	PPV	NPV
PHASE + FLAIR*	1	198	26	14	40	85.6	66.6	74.0	11.0	60.0	93.0
THASE + FLAIR	2	337	35	16	37	88.0	59.2	69.8	9.4	51.3	95.4
PHASE + FLAIR	1	205	19	41	13	78.4	30.2	24.0	8.5	40.6	83.3
I HASE + PLAIR	2	332	40	26	27	84.5	45.0	50.9	10.7	40.2	92.7
PHASE + T2*	1	203	21	21	33	84.8	61.1	61.1	9.3	61.1	90.6
FHASE + 12	2	334	38	22	31	85.8	50.8	58.5	10.2	44.9	93.8