Preface

Stroke is the second most frequent cause of death and a major cause of disability in industrial countries. In patients who survive, stroke is generally associated with high socioeconomic costs due to persistent disability. Its most frequent manifestation is the ischemic stroke, whose diagnosis often involves the acquisition of brain magnetic resonance (MR) scans to assess the stroke lesion's presence, location, extent, evolution and other factors. An automated method to locate, segment and quantify the lesion area would support clinicians and researchers alike, rendering their findings more robust and reproducible.

New methods for stroke segmentation are regularly proposed. But, more often than desirable, it is difficult to compare their fitness, as the reported results are obtained on private datasets. Challenges aim to overcome these shortcomings by providing (1) a public dataset that reflects the diversity of the problem and (2) a platform for a fair and direct comparison of methods with suitable evaluation measures. Thus, the scientific progress is promoted.

With ISLES, we provide such a challenge covering ischemic stroke lesion segmentation in multi-spectral MRI data. The task is backed by a well established clinical and research motivation and a large number of already existing methods. Each team may participate in either one or both of two sub-tasks:

**SISS** Automatic segmentation of ischemic stroke lesion volumes from multi-spectral MRI sequences acquired in the sub-acute stroke development stage.

**SPES** Automatic segmentation of acute ischemic stroke lesion volumes from multi-spectral MRI sequences for stroke outcome prediction.

The participants downloaded a set of training cases with associated expert segmentations of the stroke lesions to train and evaluate their approach, then submitted a short paper describing their method. After reviewing by the organizers, a total of 17 articles were accepted and compiled into this volume. At the day of the challenge, each teams' results as obtained on an independent test set of cases will be revealed and a ranking of methods established.

For the final ranking and more information, visit [WWW.ISLES-CHALLENGE.ORG](http://WWW.ISLES-CHALLENGE.ORG).

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Random forests for acute stroke penumbra estimation

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Abstract. Ischemic stroke treatment decisions are time-critical and depend largely on the potentially salvageable tissue. This rises the need for accurate, reproducible and reliable segmentation of acute ischemic stroke lesions from brain MR scans. This article details a contribution to the Acute Stroke Penumbra ESTimation (SPES) sub-task of the Ischemic Stroke Lesion Segmentations Challenge (ISLES), organized in conjunction with the MICCAI 2015. The proposed method bases on previous works, which showed the approach to handle the tasks well and to be applicable to potentially flawed data acquired in clinical routine. The method is described in detail and all chosen parameter values are disclosed. Preliminary results on the training data places the approach among the highest ranking contributions.

Keywords: acute ischemic stroke, lesion segmentation, penumbra estimation, magnetic resonance imaging, brain MRI, random forest, RDF

1 Introduction

Ischemic stroke is caused by an obstruction of the blood supply to the brain and the subsequent death of brain tissue. Its diagnosis often involves the acquisition of brain magnetic resonance (MR) scans to assess the strokes presence, location, extent, evolution and other factors. If diagnosed early, part of the under-perfused tissue could still be salvaged by re-establishing the blood flow. Since the available treatment options are not risk-free and can e.g. lead to inter-cranial bleeding, the decision has to be made individually, depending on the potential gain and under great time restriction. An automated method to distinguish the already necrotic from the potentially salvageable tissue (furthermore termed penumbra, although this term is disputed) would be highly beneficial for the clinical routine and reduce incorrect decisions. The ISLES 2015 challenge offers the first platform for researchers to compare their methods directly and fair. Our contribution draws its base from a previously published method targeted towards sub-acute ischemic stroke lesions [3], which showed good results. It is based on carefully selected features extracted from the MR sequences and used to train a random forest (RF).
2 Method

The challenge’s training data consists of multi-spectral (T1c, T2, DWI, CBF, CBV, TTP, Tmax) scans of 30 patients displaying acute ischemic stroke. For training the manual segmentations of a single expert rater has been provided. More details on the data can be found on www.isles-challenge.org.

2.1 Pre-processing

The image data is provided with a 1 mm isotropic resolution, already co-registered and skull-stripped. Nevertheless, the training cases of the challenge display high intensity differences, a normal occurrence for MRI, where intensity ranges are not standardized. With a learning based intensity standardization method implemented in MedPy [2] and based on [4] we harmonize each T1c, T2 and DWI sequences intensity profile without a prior bias-correction step. The Tmax sequence, which is considered the most discriminative for penumbra estimation, is cut-off at an upper value of 100, which corresponds to 10 s.

2.2 Forest classifier

We employ the RF classifier implemented in [5], which is similar to the propositions made by [1]. The classification of brain lesions in MRI is a complex task with high levels of noise [3], hence a sufficiently large number of trees must be trained.

2.3 Features

The primary distinction criteria for identifying pathological tissue of stroke lesions is the MR intensity in the different sequences. The bulk of our voxel-wise features therefore bases on the intensity values.

intensity First feature is the voxel’s intensity value.

gaussian Due to the often low signal-to-noise ratio in MR scans and intensity inhomogeneities of the tissue types, we furthermore regard each voxel’s value after a smoothing of the volume with a 3D Gaussian kernel at three sizes: \( \sigma = 3, 5, 7 \) mm.

hemispheric difference Gliomas mostly affect a single hemisphere, therefore we extract the hemispheric difference (in intensities) after a Gaussian smoothing of \( \sigma = 1, 3, 5 \) mm to account for noise. For simplicity, the central line of the saggital view is taken as sufficiently close approximation of the sagittal midline.
Finally, we extract the distance to the image center (assumed here to coincide roughly with the brain’s center of mass) in mm as final feature. Note that this is not intensity based, but rather discloses each voxel’s rough location inside the brain.

All features are extracted from each of the MR sequence, hence in total we obtain 52 values per multi-spectral voxel. Note that all of these features are implemented in MedPy [2].

2.4 Post-processing

After thresholding the a-posteriori class probability maps for a crisp segmentation, all but the largest connected component are removed. No other post-processing steps are employed.

3 Experiments

3.1 Training choices and parameter values

For training our RF, we sample 1,000,000 voxels randomly from all training cases. The ratios between classes in each case are kept intact (i.e. stroke class samples will be highly under-represented). A total of 100 trees are trained for the forest. As split criteria the Gini impurity is employed, a maximum of $\sqrt{52}$ features is considered at each node. No growth restrictions are imposed. The a-posteriori class probabilities produced by the forest are thresholded at a value of 0.35 to counter under-segmentation.

3.2 Preliminary results

Online evaluation is provided with the Dice’s coefficient (DC), the average symmetric surface distance (ASSD) and the Hausdorff distance (HD) as quality metrics. Using a leave-one-out evaluation scheme, we have obtained the scores presented in Tab. 1.

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<tr>
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<th>DC</th>
<th>ASSD</th>
<th>HD</th>
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<tbody>
<tr>
<td>30 cases</td>
<td>0.83 ± 0.06</td>
<td>1.38 ± 0.66</td>
<td>23 ± 13</td>
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4 Discussion and conclusion

After the challenge date, the final results can be found on the challenge webpage www.isles-challenge.org. Preliminary results rank our method among the best performing. An advantage of our approach is its flexible design, that allows an application for a large number of brain lesion segmentation tasks (see e.g. [3]). Slightly adapted versions have been handed in to the sibling challenges ISLES 2015: SISS and BRATS 2015.

By employing RFs, we have a powerful classifier at our hand that is robust against uninformative features, generalizes well and produces good results for a wide range of parameters. Mixing widely used with specially designed features, we can successfully learn to discriminate between the acute stroke area and healthy brain tissue.

On the downside, they suffer from the same drawbacks as all other machine learning based methods: The training set must be carefully chosen and types of cases not present in the training data can not be processed.

References