Automatic Analysis of Fungal-Infected Tissue using Deep Learning

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1. Experimental setup

- Combine optical image analysis with MALDI images to detect fungal infected regions
- Learn one model for each staining method:
 - Gomori (Gömöri trichrome stain used on muscle tissue)
 - PAS (Periodic acid-Schiff reaction)





 Classify mass spectrum (MS) according to respective fungus species



2. Optical image segmentation

Deep Learning of Convolutional Neural Networks (CNN)



• Network – loss-value on test images: $L(y, \tilde{y}) = \sum_{j=0}^{\infty} \sum_{i=0}^{\infty} (y_{ij} * log(\tilde{y}_{ij})) + DC = 0.3067$



(A) Test region of manually annotated fungus-infected tissue

(B) Network prediction: white regions correspond to high probability P of fungal-infected tissue Scalebar: P(fungal-infected tissue)

(C) Complete tissue sample: green regions represent the annotated fungus

(D) Network inference for the entire sample image

3. Interface for optical image – MALDI image



4. Classification of MALDI-imaging



5. Conclusions

- Deep learning of neural networks allows to identify fungal-infected tissue
- Novel approach for the registration of optical tissue image and MALDI image
- Machine learning models allow to associate fungal-infected tissue with MS-spectra

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