




Poster Presentations

Code	Title	Authors
P01	A Python application programming interface for accessing Philips iSyntax whole slide images	Krzysztof Krawczyk, Matthias J. Mickert, Stefan Andersson-Engels, Anders Sjögren
P02	Augmented Reality Microscopy (ARM) Utility for Breast Tumor Measurements	Mustafa Yousif, Liron Pantanowitz
P03	Microscope HD video streaming in University Pathology teaching through YouTube and Twitch	Luis Alfaro, Maria Jose Roca
P04	Analysis of paraffin-embedded slides of esophageal carcinoma after different treatments using QuPath	Benjamin Igbo, Annett Linge, Theresa Suckert, Susanne Frosch, Liane Stolz-Kieslich, Esther G.C. Troost
P05	Instant digital pathology for rapid evaluation of thyroid cytology: a pilot study and molecular test	Martina Verri, S. Scarpino, C. Taffon, A. Palermo, A.M. Naciu, D. Nicoletti, D. Galafate, E. Pillozzi, A. Crescenzi
P06	Physical Color Calibration of Scanners for Deep Learning Based Diagnosis of Prostate Cancer	Xiaoyi Ji, Richard Salmon, Nita Mulliqi, Henrik Olsson, Lars Egevad, Pekka Ruusuuvuori, Martin Eklund, Kimmo Kartasalo
P07	A Python application programming interface for accessing Philips iSyntax whole slide images	Nita Mulliqi, Lars Egevad, Pekka Ruusuuvuori, Martin Eklund, Kimmo Kartasalo
P08	Automatic segmentation of tumor infiltrating lymphocytes in breast histopathology slides	Jakub Gawlik, Agnieszka Łazarczyk, Julita Ciuruś, Michał Okarski, Joanna Szpor
P09	Normal and Neoplastic Salivary Gland Segmentation Using Machine Learning – A pilot study	Ibrahim Alsanie, Eu-Wing Toh, Syed Ali Khurram
P10	An approach to resource saving histology dataset expansion	Artyom Borbat, Inna Yatsenko
P11	Neural network analysis visualization approach for diagnostic pathology	Artyom Borbat, Peter Bondarenko, Sergey Lishchuk
P12	QuPath-based approach to evaluate liver parenchyma status: preliminary data	Artyom Borbat, Alexandr Dushkin, Ilya Serdyuk
P13	Primary and metastatic tumor tissue datasets to train neural network model for metastasis detection	Artyom Borbat, Elena Filatova, Maksim Iaroslavtsev, Tatiana Novikova
P14	Semiautomated workflow for tissue-microarray analysis on tumors of the central nervous system	Teresa San Miguel, Ana Sierra, David Moratal, Miguel Cerdá-Nicolás, Concha López-Ginés, Javier Megías, Lara Navarro, Daniel Monleón
P15	Segmentation of oesophageal cancer lymph nodes within large H&E datasets with explainable AI	Manon Beuque, Avishek Chatterjee, Henry C. Woodruff, Ruth E. Langley, William Allum, Matthew G. Nankivell, David Cunningham, Philippe Lambin, Heike I. Grabsch
P16	Reducing the false negative prostate biopsies using deep-learning assessment of benign biopsies	Bojing Liu, Yinxu Wang, Philippe Weitz, Johan Lindberg, Lars Egevad, Henrik Grönberg, Martin Eklund, Mattias Rantalainen
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P18	The effectiveness of whole slide imaging in assessing the invasive breast carcinoma cases.	Gabriela Baltatescu, M. Aşchie, M. Enciu, G.C. Cozaru, O. Cojocaru, M. Cristian, N. Dobrin, M. Deacu
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P20	Predicting Genetic Intra-tumor Heterogeneity From Digital Histopathology Slides	Mustafa Umit Oner, Hwee Kuan Lee, Wing-Kin Sung, Jianbin Chen, Weiwei Zhai
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P24	New Cytomine open-source software architecture and modules for AI in digital pathology	Raphaël Marée
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P35	AI-Power to the pathologist - IHC guided annotations improved the resolution performance of the algo compared to manual annotated whole slide images.	Lars Bjork, Feria Hikmet, Jonas Gustavsson, Daniel Hägg, Mats Andersson, Filippo Frassetto, Witold Rezner, Mateusz Seliga, Piotr Bobkiewicz, Lex Makkus, Andrey Bychkov, Junya Fukuoka, Kristian Eurén, Stefan Elfving, Cecilia Lindskog



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