

# Exploitable Results by Third Parties

14001 3DPathology

---

## Project details

Project leader:	Dominique Segers (Barco)
Email:	dominique.segers@barco.com
Website:	www.barco.com

Name: DCGMM histopathology stain-color normalization		
Input(s):	Main feature(s)	Output(s):
DCGMM Source Code	DCGMM is applied for histopathology stain-color normalization	Color-normalized images
Unique Selling Proposition(s):	Fully-unsupervised generative model learns the color distribution among the images and transforms it to resemble the color distribution in an arbitrary template image.	
Integration constraint(s):	<ul style="list-style-type: none"> <li>▪ Python 2.7</li> <li>▪ TensorFlow 1.5.0</li> </ul>	
Intended user(s):	<ul style="list-style-type: none"> <li>▪ Application developers or research engineers in computational pathology that have interest to exploit GPU capabilities. In histopathology CAD system developer, this module can be used as a pre-processing stage prior to further image analysis such as tumor detection or metastasis grading .</li> </ul>	
Provider:	<ul style="list-style-type: none"> <li>▪ Eindhoven University of Technology – VCA research group.</li> <li>▪ <a href="https://github.com/FarhadZanjani/Histopathology-Stain-Color-Normalization">https://github.com/FarhadZanjani/Histopathology-Stain-Color-Normalization</a></li> </ul>	
Contact point:	<ul style="list-style-type: none"> <li>▪ Farhad G. Zanjani – f.ghazvinian.zanjani@tue.nl</li> </ul>	
Condition(s) for reuse:	<ul style="list-style-type: none"> <li>▪ Free license</li> </ul>	
<i>Latest update: 25 June 2018</i>		

Name: 3D Reconstructor for 2D stack images		
Input(s):	Main feature(s)	Output(s):
<ul style="list-style-type: none"> <li>2D Mass Spec Imaging Data</li> </ul>	<ul style="list-style-type: none"> <li>Recombining rotated, skewed, stretched 2D images to one 3D data block ready for 3D viewer</li> </ul>	<ul style="list-style-type: none"> <li>3D Mass Spec Imaging Dataset</li> </ul>
Unique Selling Proposition(s):	<ul style="list-style-type: none"> <li>Automated reconstruction of 3D block image from 2D stack image, making use of optimized number of channels</li> <li>Working with different image data types</li> </ul>	
Integration constraint(s):	<ul style="list-style-type: none"> <li>Versatile Interfacing</li> <li>No special OS requirements</li> <li>Algorithms can run on instrument's computer</li> <li>Ongoing development with MSI hardware / resolution improvement</li> </ul>	
Intended user(s):	<ul style="list-style-type: none"> <li>MSI Instrument Suppliers, Research Organizations</li> <li>2D-from-3D imaging developers and suppliers</li> </ul>	
Provider:	<ul style="list-style-type: none"> <li>Target Holding B.V.</li> </ul>	
Contact point:	<ul style="list-style-type: none"> <li>dr. Rolf Neubert, rolf.neubert@target-holding.nl</li> </ul>	
Condition(s) for reuse:	<ul style="list-style-type: none"> <li>Commercial license on annual basis to be negotiated</li> <li>University license to be negotiated for special rates</li> </ul>	
<i>Latest update: 25 June 2018</i>		

Name: VISPAT 3D Viewer (digital pathology web-based player)		
Input(s):	Main feature(s):	Output(s):
<ul style="list-style-type: none"> <li>▪ Digital pathology images</li> <li>▪ HPC</li> <li>▪ Big Data</li> </ul>	<ul style="list-style-type: none"> <li>▪ displaying 2D/3D medical image/models based on very large files</li> <li>▪ pan, zoom, rotate the 2D/3D images</li> <li>▪ take screenshots for both 2D/3D images</li> <li>▪ use filters (contrast, brightness, erosion, dilation, invert, grey scale, convolution) that can be applied to each image or model, for any zoom level in 2D/3D mode;</li> <li>▪ download sample/test files;</li> <li>▪ share models between users through internal unique codes</li> <li>▪ dynamically model generation based on atlas files</li> <li>▪ dynamically generation of atlas files based on provided image files</li> <li>▪ 2d perspectives on each of 3D axes (X, Y, Z)</li> </ul>	<ul style="list-style-type: none"> <li>▪ Analysis of relevant areas from medical digital images</li> <li>▪ Provides web-based visualization of medical data</li> <li>▪ Suitable for different platforms</li> </ul>
Unique Selling Proposition(s):	<ul style="list-style-type: none"> <li>▪ Dynamic application for big 2D/3D images visualization</li> <li>▪ Source to source tool to generate fast access and displaying big images</li> <li>▪ Auto adaptive library for different types of data/images</li> </ul>	
Integration constraint(s):	<ul style="list-style-type: none"> <li>▪ Suited for heterogeneous platforms (CPU/GPU on HPC)</li> <li>▪ Node.js</li> <li>▪ Three.js</li> <li>▪ X3dom</li> <li>▪ Caman.js</li> <li>▪ Openseadragon.js</li> <li>▪ Bootstrap</li> <li>▪ Openseadragonfiltering.js</li> <li>▪ Require.js</li> <li>▪ Upload.js</li> <li>▪ DZI (deep zoom images)</li> <li>▪ Connector.js</li> <li>▪ Download.js</li> <li>▪ make2D.js</li> <li>▪ make3D.js</li> </ul>	

Name: VISPAT 3D Viewer (digital pathology web-based player)	
Intended user(s):	<ul style="list-style-type: none"> <li>▪ pathologists</li> <li>▪ clinicians</li> <li>▪ researchers in digital pathology</li> <li>▪ students</li> <li>▪ web 3d practitioners</li> <li>▪ doctors</li> </ul>
Provider:	<ul style="list-style-type: none"> <li>▪ ALTFACOR SRL, Romania, <a href="http://www.altfactor.ro">www.altfactor.ro</a> , <a href="http://www.3dpathology.ro">www.3dpathology.ro</a></li> </ul>
Contact point:	<ul style="list-style-type: none"> <li>▪ Marius Petrut IVANOV, <a href="mailto:marius.ivanov@altfactor.ro">marius.ivanov@altfactor.ro</a></li> </ul>
Condition(s) for reuse:	<ul style="list-style-type: none"> <li>▪ End-User License Agreement (EULA)</li> </ul>

*Latest update: 25 June 2018*

Name: Mobile 3D Pathology Viewer (pathology viewer based on mobile technologies)		
Input(s):	Main feature(s):	Output(s):
<ul style="list-style-type: none"> <li>▪ Digital images</li> <li>▪ Big data images</li> </ul>	<ul style="list-style-type: none"> <li>▪ displaying 2D/3D images from different modalities on mobile devices</li> <li>▪ pan and zoom the 2D/3D images</li> <li>▪ Rapid pre-viewing of 2D/3D Pathology high-resolution datasets. This volume can be moved, rotated, zoomed, etc.;</li> <li>▪ Fast performance and accessibility from everywhere;</li> <li>▪ Intuitive and interactive user interface;</li> <li>▪ Flexibility with both high and low-level control;</li> <li>▪ Interactive remote visualization of multi-modal 3D image</li> </ul>	<ul style="list-style-type: none"> <li>▪ Viewing of relevant areas from medical digital images</li> <li>▪ Provides mobile based technologies for medical data visualization</li> <li>▪ Suitable for different mobile devices</li> </ul>
Unique Selling Proposition(s):	<ul style="list-style-type: none"> <li>▪ Mobile application for big images visualisation</li> <li>▪ Fast integration on different platforms</li> <li>▪ Accessible for any mobile device regardless of the device's performance: PC, laptops, mobile devices</li> </ul>	
Integration constraint(s):	<ul style="list-style-type: none"> <li>▪ Client-Side Storage</li> <li>▪ Slower processor speeds</li> <li>▪ Wireless network connections with latency</li> <li>▪ Smaller screen that can display a limited area of the medical images</li> </ul>	
Intended user(s):	<ul style="list-style-type: none"> <li>▪ pathologists</li> <li>▪ clinicians</li> <li>▪ researchers in digital pathology</li> </ul>	
Provider:	<ul style="list-style-type: none"> <li>▪ SIVCO, Romania; <a href="http://www.sivco.ro">www.sivco.ro</a>; <a href="http://www.3dpathology.ro">www.3dpathology.ro</a></li> </ul>	
Contact point:	<ul style="list-style-type: none"> <li>▪ Dorothea Caraman, <a href="mailto:dorothea.caraman@sivco.ro">dorothea.caraman@sivco.ro</a></li> </ul>	
Condition(s) for reuse:	<ul style="list-style-type: none"> <li>▪ Mobile Application End-User License Agreement (EULA)</li> </ul>	

Latest update: 25 June 2018