

IT ecosystem based on machine learning methods and data analysis technologies for radiobiological research

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On behalf of the **BIOHLIT** group



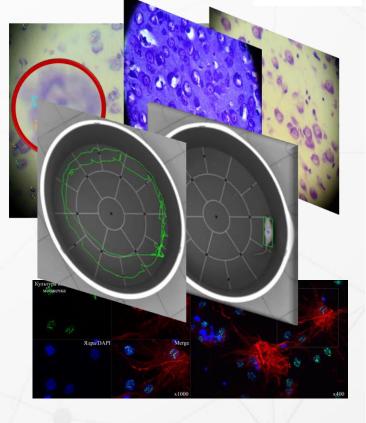
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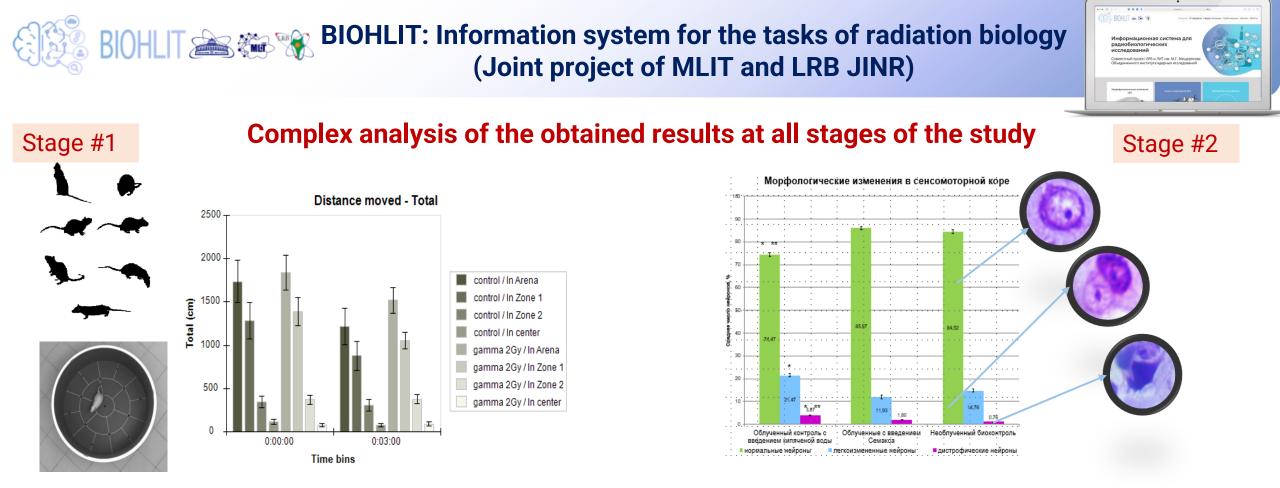


In collaboration with

University of Belgrade

Coordinator: Dr Marko Ćosić

In collaboration with Germany: Streltsov Alexei I. Gromov E.



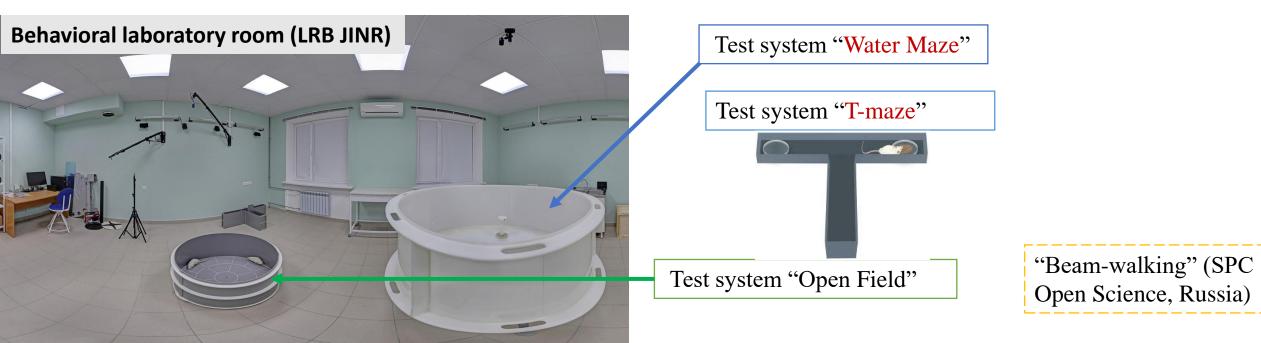
The IS is necessary for:

- **Storing** the entire array of experimental data (photo and video materials, pdf, excel, doc files with information about the experiment) and minimizing the risk of information loss.
- Convenient access for all members of the research group and ability to conduct both step-by-step and comprehensive data analysis in a unified information space.
- **Reducing** the time spent on data processing, increasing the speed of obtaining qualitative results and reducing the subjectivity of the experimental data processing approach.

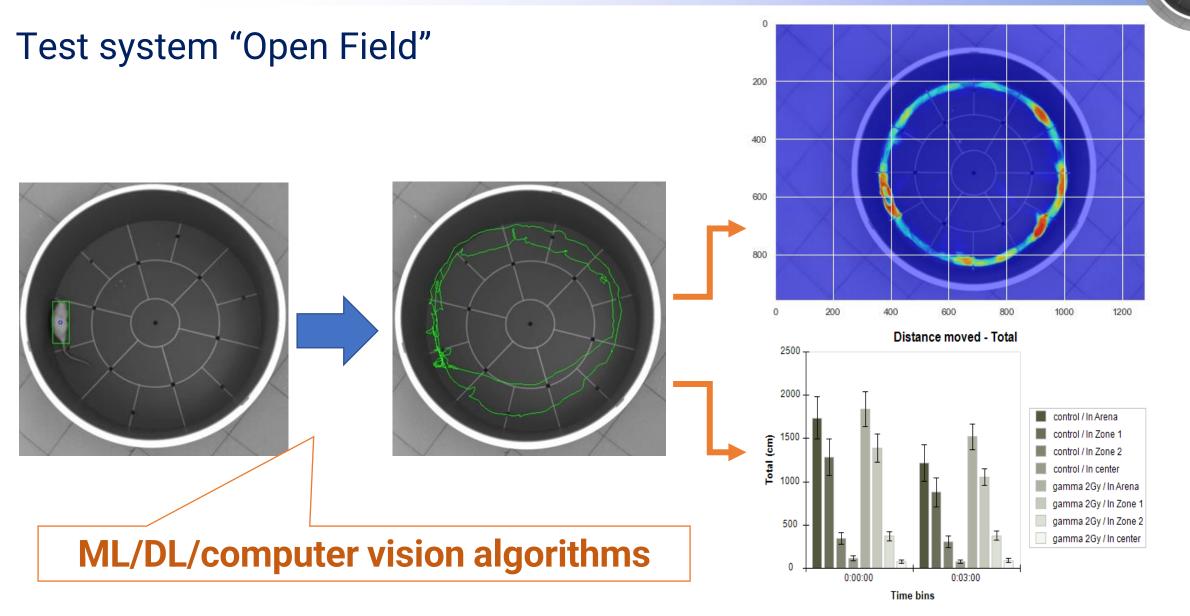
BIOHLIT AND Stage #1: analysis of behavioral reactions of laboratory animals

The first stage is related to the analysis of behavioral reactions of laboratory animals exposed to ionizing radiation, as well as of the possible effect on the behavior of pharmacological preparations.

From the point of view of data analysis, this stage is connected with the collection and analysis of **video data** reflecting behavioral reactions of laboratory animals during experimental tests using specialized stands.



BIOHLIT & Trage #1: analysis of behavioral reactions of laboratory animals



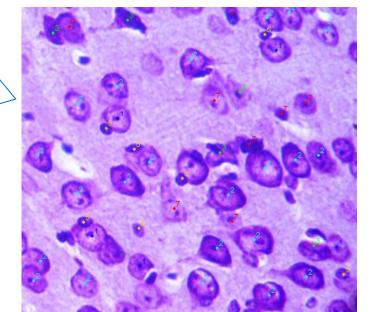
BIOHLIT AN Stage #2: studying morphological changes in the cells of the central nervous system

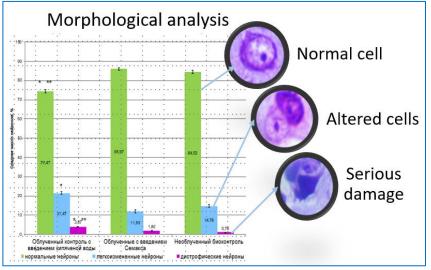
The **second stage** is aimed at studying morphological changes in the cells of the central nervous system to establish a connection with the behavioral reactions of laboratory animals obtained during the first stage of investigations.

The histological examination of other organs of laboratory animals is also carried out.

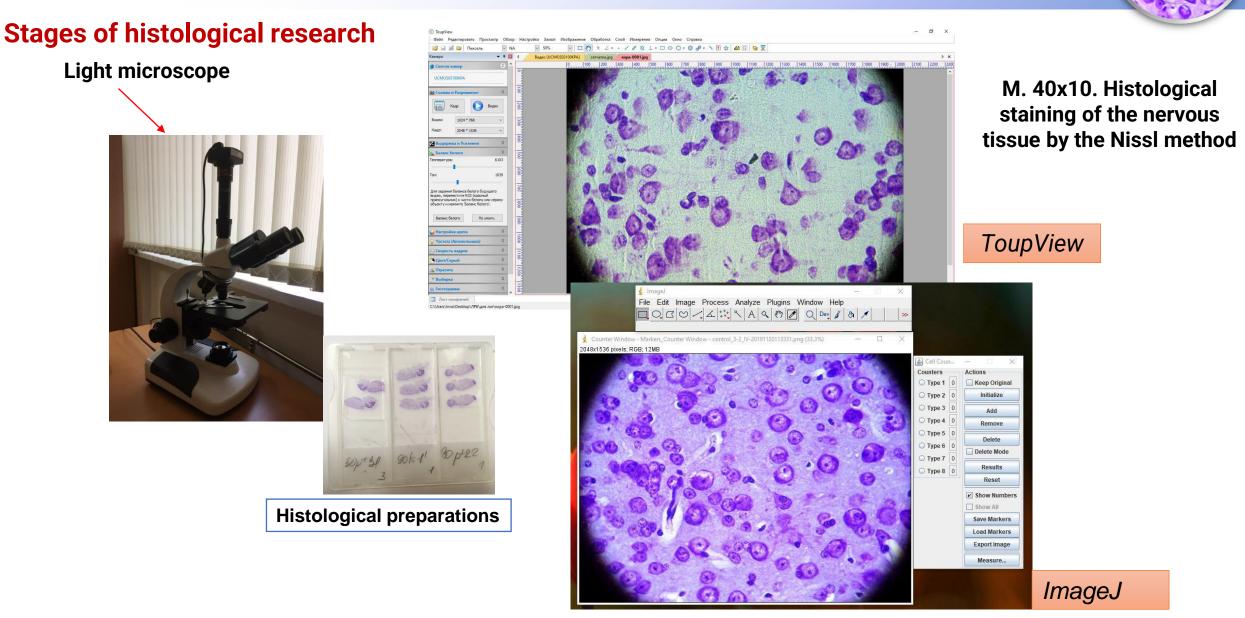
From the point of view of data analysis, this stage is connected with the analysis of **images of slices** of the nervous tissue of different parts of the brain obtained using a light microscope, a camera and programs that allow counting cells with certain changes.

The problem of automating the morphological analysis of histological preparations is solved within the project by implementing algorithms based on the neural network approach and computer vision methods.





BIOHLIT A Stage #2: studying morphological changes in the cells of the central nervous system



BIOHLIT AN WE'R Stage #2: studying morphological changes in the cells of the central nervous system

Used classification of brain cells:

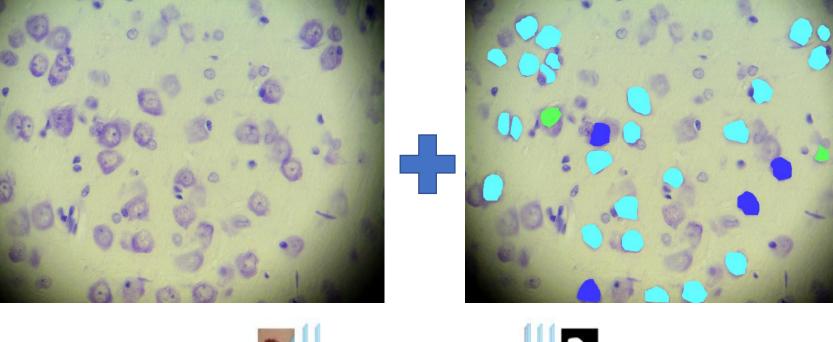
□ normal (1);

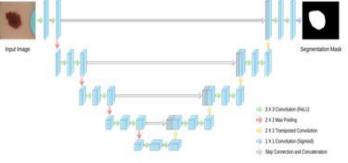
- easily modified (2): morphofunctional and compensatoryadaptive;
- \Box degenerative (3);

□glia (5);

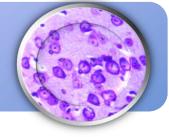
elements of the nervous tissue that are difficult to identify (7)

For the segmentation task, the U-net neural network architecture is well suited





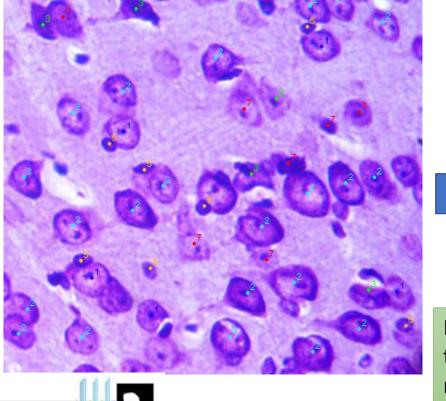
BIOHLIT A Stage #2: studying morphological changes in the cells of the central nervous system

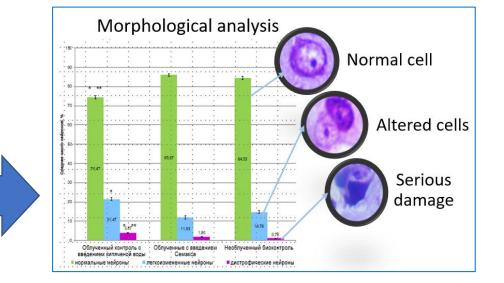


Stages of histological research

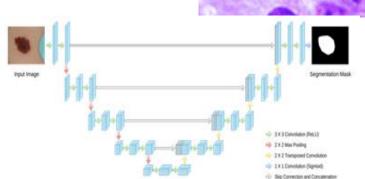
Automation options:

- automation of the morphological analysis of brain cells of experimental animals using photos of micropreparations
- storage of photo images before and after their analysis by an expert





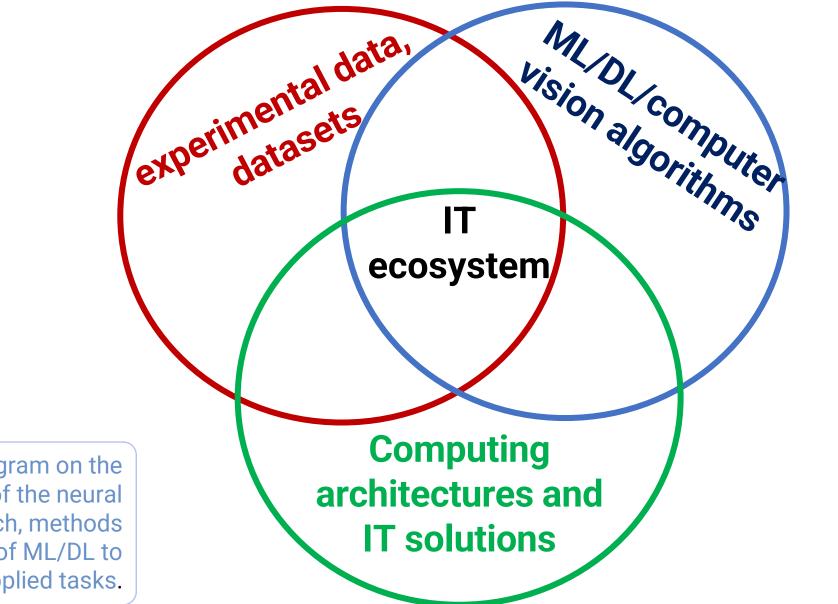
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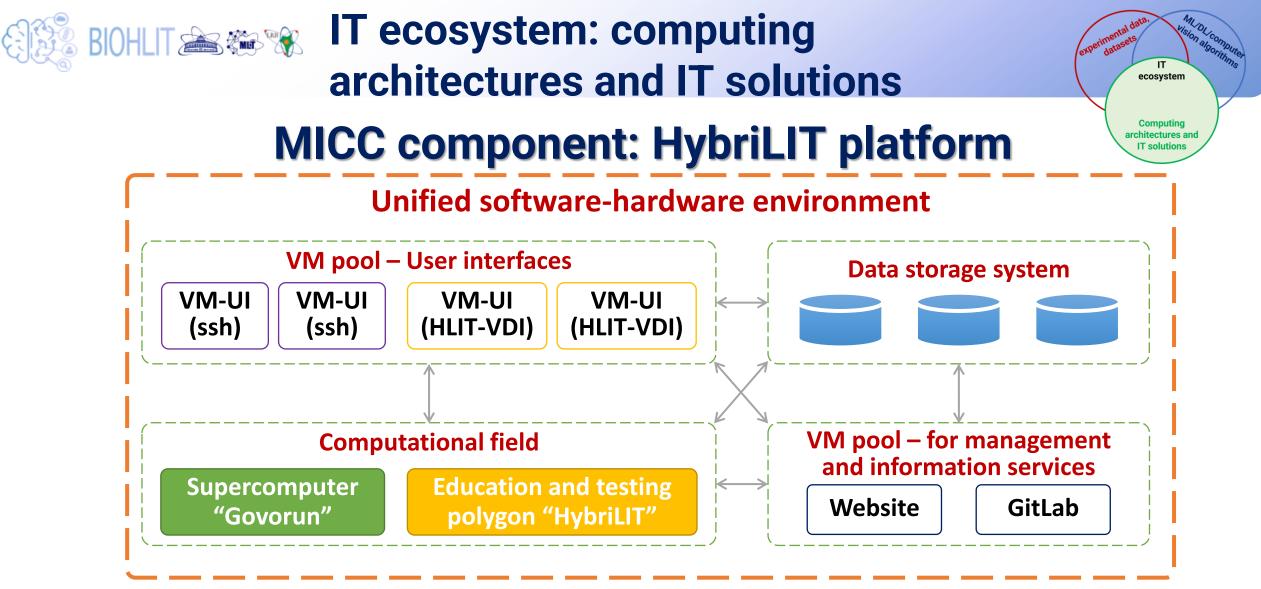








Venn diagram on the implementation of the neural network approach, methods and algorithms of ML/DL to solve applied tasks.



The unified software and information environment of the HybriLIT platform allows users to use the education and testing polygon aimed at exploring the possibilities of novel computing architectures and IT solutions, to develop and debug their applications, furthermore, to carry out calculations on the supercomputer, which enables to effectively use the supercomputer resources.







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Full stack deep learning software preinstalled eplaces 400 traditional dual CPU servers on DL applications

GPU component based on NVIDIA DGX-1 Volta





"Govorun" Supercomputer

The "Govorun" supercomputer is a hyper-converged software-defined system, ranked 17th in the current edition of the **IO500** list (July 2020). For the high-speed data storage system, RSC Group received the prestigious Russian DC Award 2020 in "The Best IT Solution for Data Centers" nomination at the awards ceremony held on 10 December 2020 in Moscow.

CPU component based on the latest Intel architectures: Intel Xeon Phi gen.2 and Intel CascadeLake processors Total peak performance: **1.7** PFLOPS SP 860 TFLOPS DP **300** Gb/s Data IO rate



NVIDIA.

IT ecosystem: computing architectures and IT solutions





The GPU component consists of **5 NVIDIA DGX-1 servers**. Each server has **8 GPUs NVIDIA Tesla V100** based on the latest architecture NVIDIA Volta. Moreover, one server NVIDIA DGX-1 has **40,960** CUDA cores, which are equivalent to 800 high-performance central processors. A whole number of novel technologies are used in DGX-1, including the NVLink 2.0 wire with a bandwidth of up to 300 Gb/s.



The GPU component gives supercomputer users the ability to perform massively parallel computing for general-purpose tasks using such technologies as CUDA and OpenCL, as well as to use applications already adapted for this architecture. In addition, the GPU component allows using machine learning and deep learning algorithms to solve applied problems by a neural network approach.

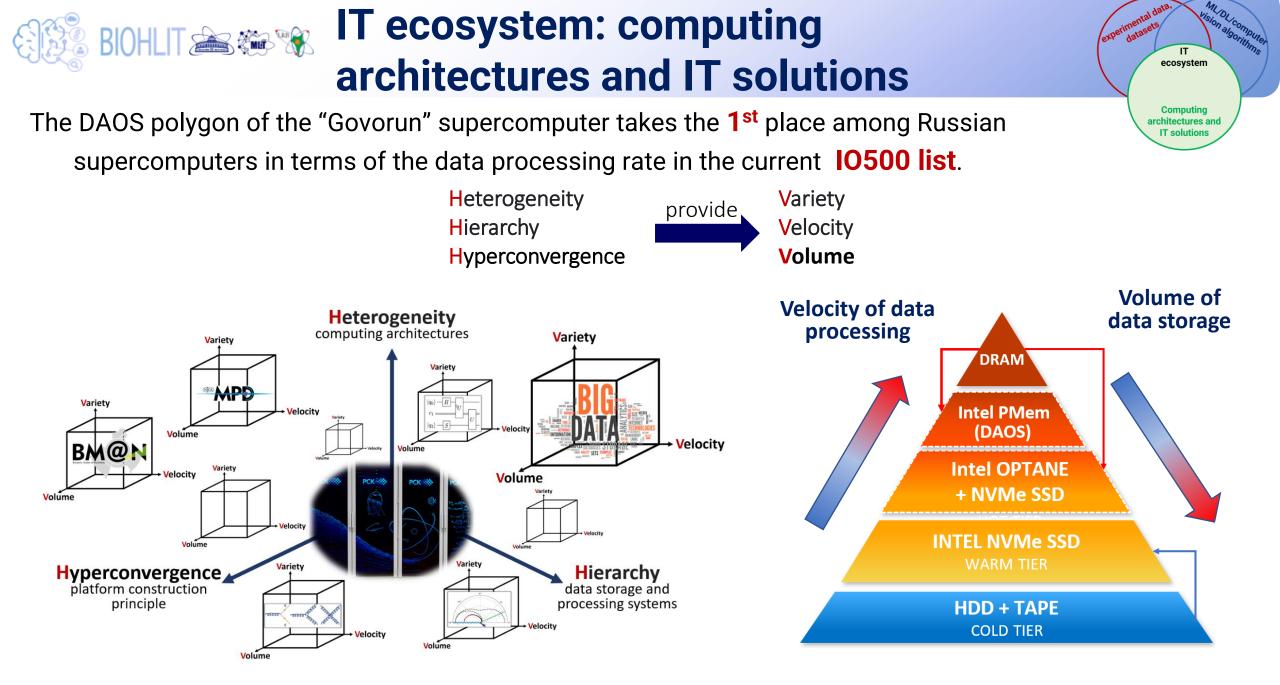


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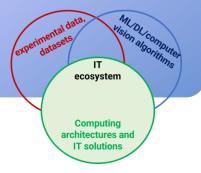
Caffe

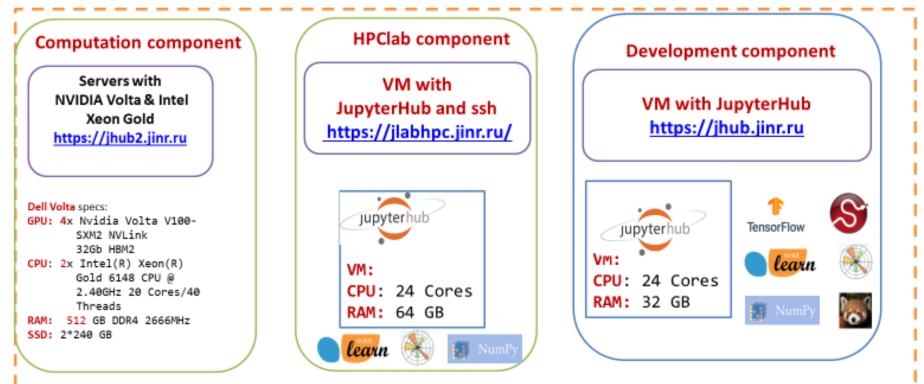




IT ecosystem: computing architectures and IT solutions

Environment for development





Ecosystem for HPC and ML/DL tasks

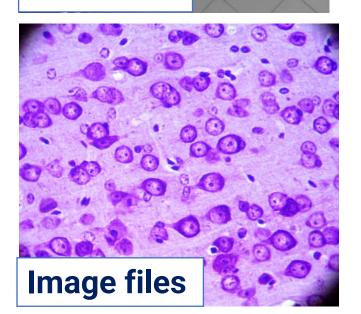


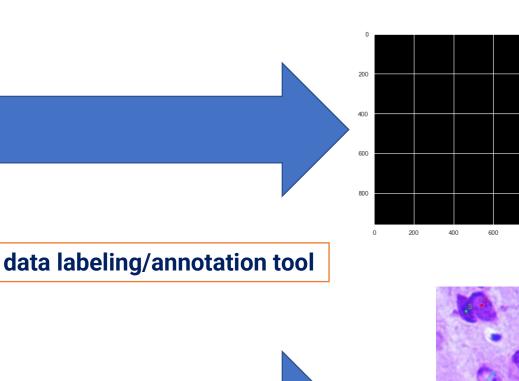
IT ecosystem: experimental data, datasets



experimental data

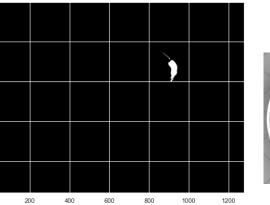






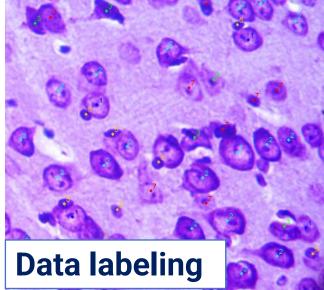
data labeling/annotation tool





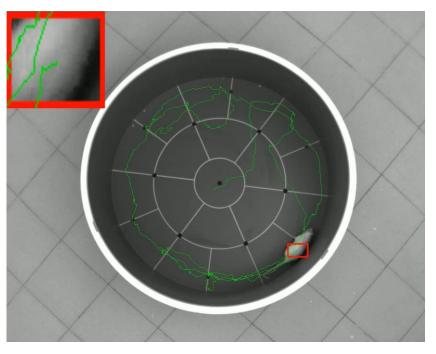


Action recognition



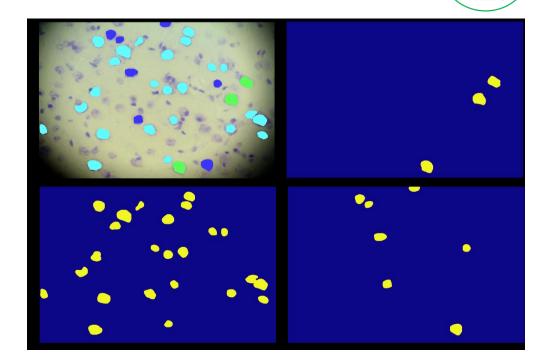
labeled/ annotated data

IT ecosystem: ML/DL/computer vision BIOHLIT 🚵 🐲 😵 algorithms ecosystem



Tracking a laboratory animal: Approaches:

- Tracking using a correlation filter
- Tracking with object-background Ο segmentation assessment Best: CSRT tracker
- GOTURN
- MOSSE Ο



IT

architectures and

Neural networks for the task of neuron segmentation on brain slice images

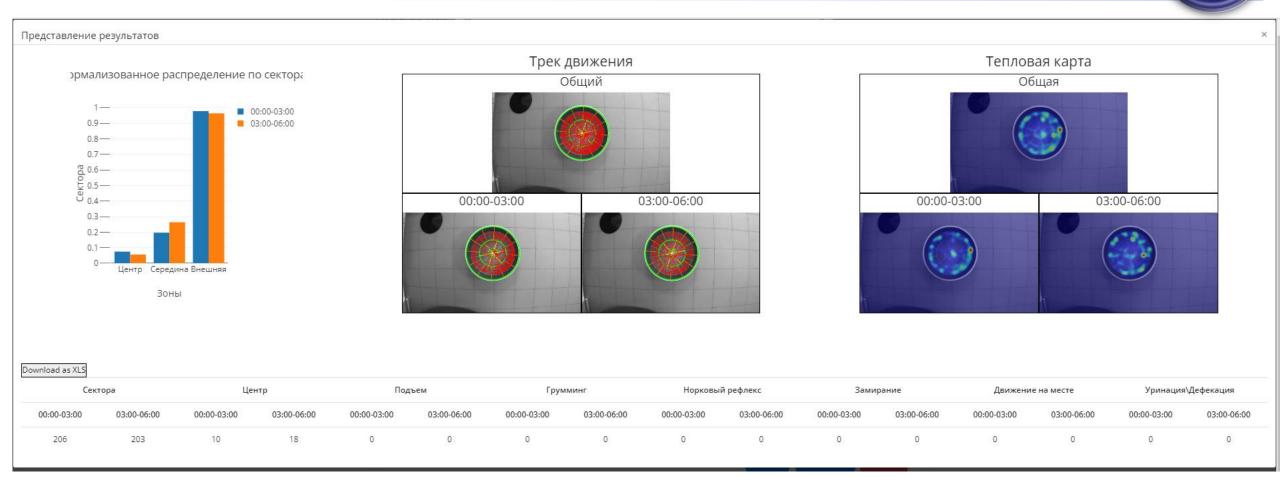
IT ecosystem: web service

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Эксперимент:		• Редактировать	Поведени	е живот	НЫХ						
вьетнам			Открытое поле	Моррис	Т-лабиринт						
Описание:			Загрузить								
лечение			Всего видео:7								
Дата забоя:	Тип животных:	Пол животных:	Выбраны:								
28-06-2022	mouse										A
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			□ 3	Ссылка	Control_mouse3 10-38-39.avi	FAIL	2022-06-28 11:27:16	2022-06-28 11:27:33	Анализ	Результаты	Удалить файл

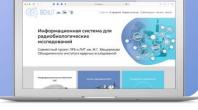
IT ecosystem: web service

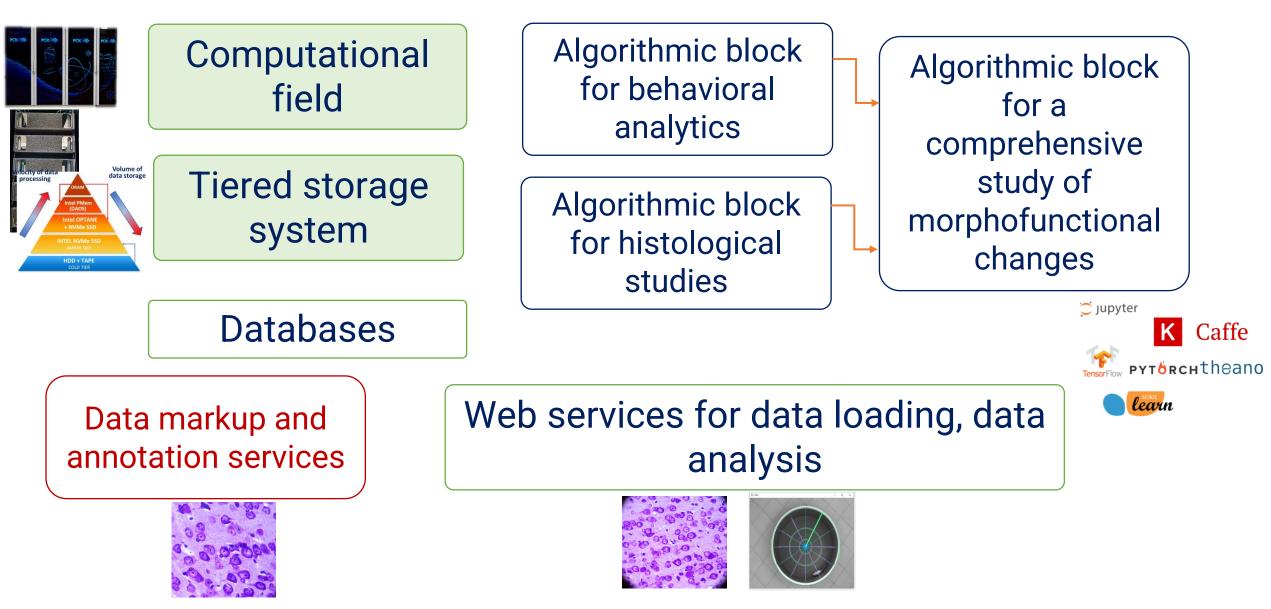
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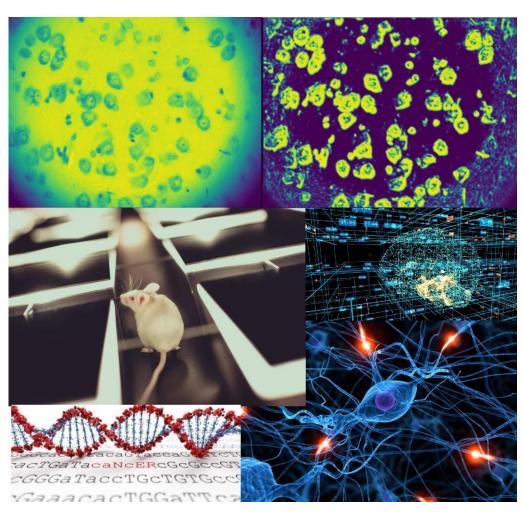
IT ecosystem







BIOHLIT A BIOHLIT: Information system for the tasks of radiation biology (Joint project of MLIT and LRB JINR)



The studies are carried out using all the capabilities of the HybriLIT platform

The joint project of MLIT and LRB is focused on creating an Information System (IS) for analyzing behavioral and pathomorphological changes in the central nervous system when studying the effects of ionizing radiation and other factors on biological objects.

The **IS** is based on:

- computer vision algorithms on top of machine and deep learning technologies (ML/DL);
- modern IT solutions for data storage, processing and visualization.

The **IS** will allow one to simplify and accelerate:

- experimental data processing by automating the morphological classification of neural cells:
- data analysis techniques using the latest neural network algorithms based on ML/DL;
- work with experimental data for different research groups;
- experimental data systematization and development of effective methods for preventing and countering the negative effect of ionizing radiation.





Thank you for your attention!



