

# DeepLabCut AI Residency

## Day 1 Session 3:

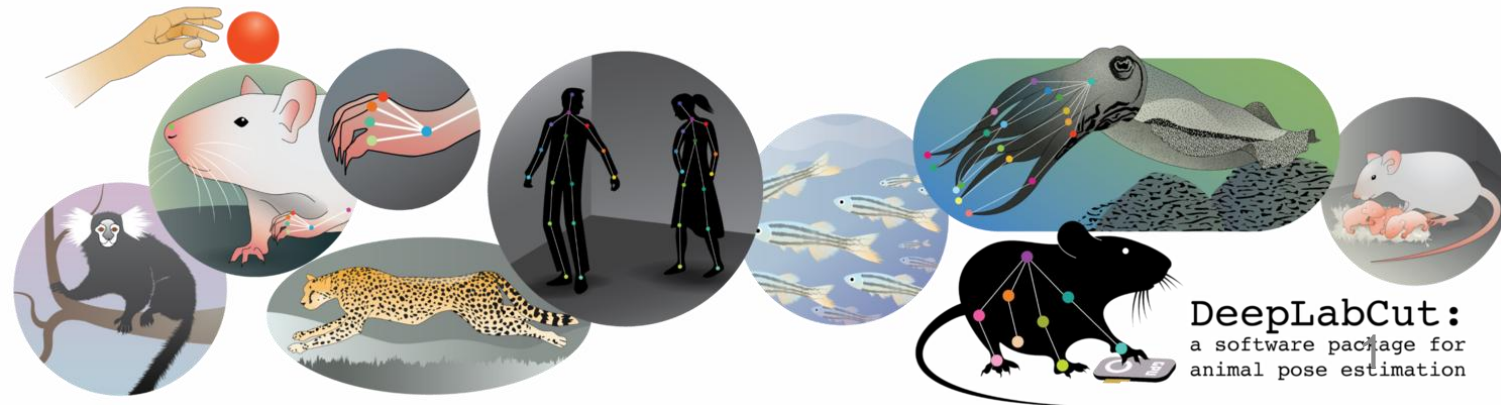
### Project creation, labeling, and data management

July 30 & August 1, 2025  
McGill University, Montreal

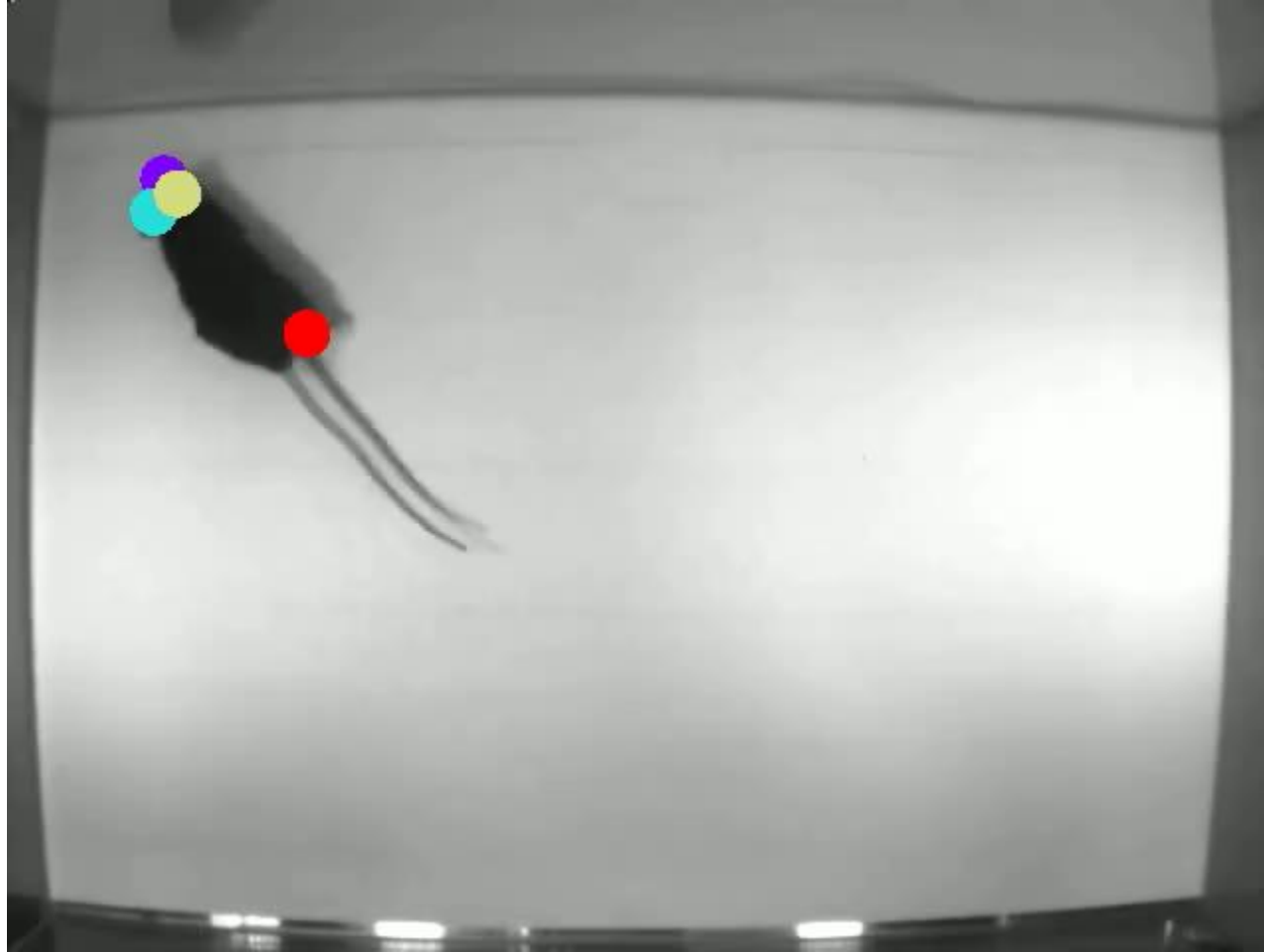
Jiayue Yang  
Vic Shao-Chinh Chiang



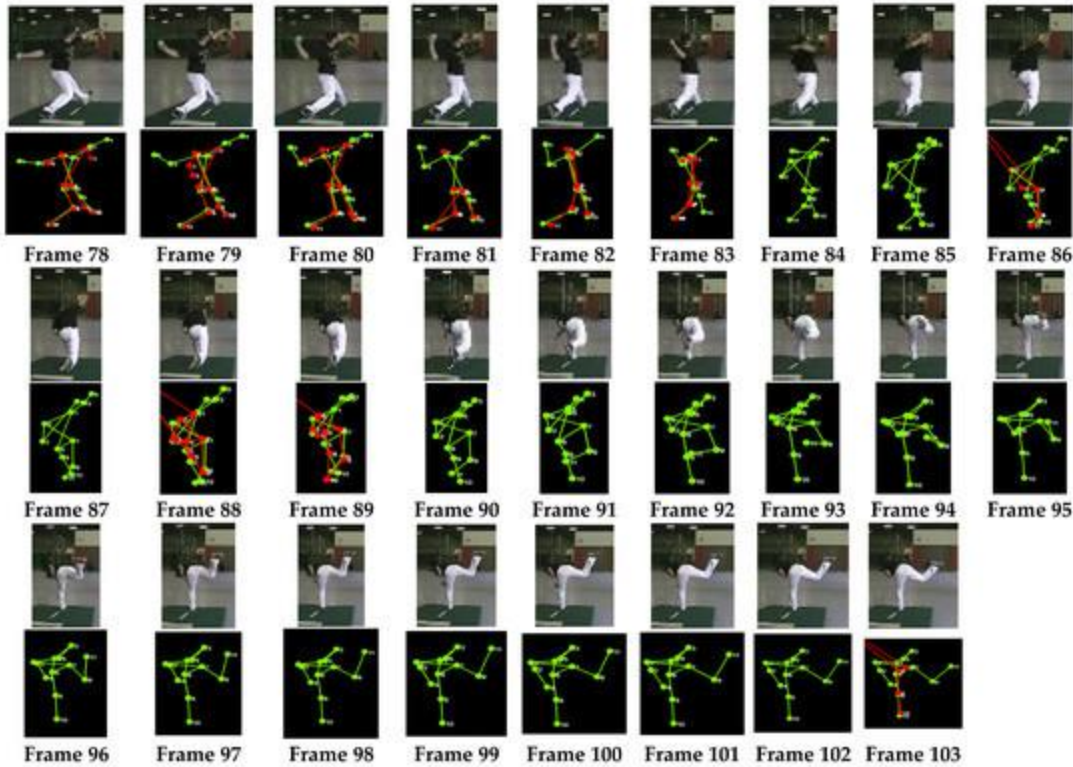
**McGill**



# Example of DeepLabCut

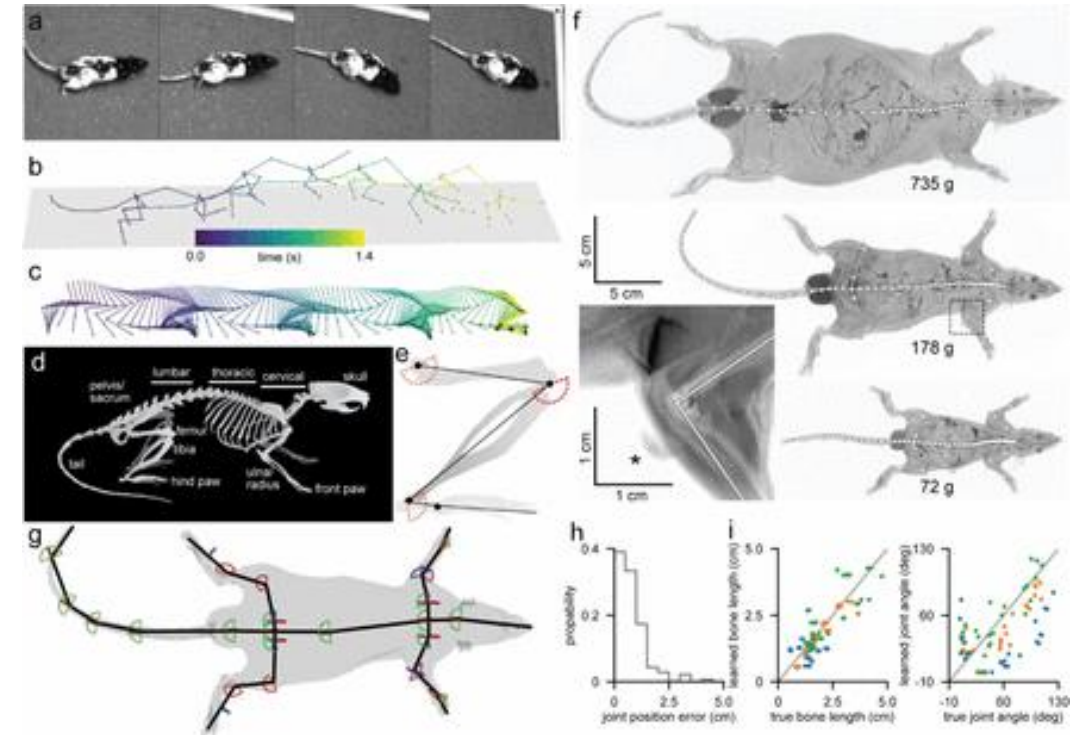


# Pose estimation: bodyparts & skeleton



OnePose detection example (humans).

Chung et al. (2022). Future Internet.



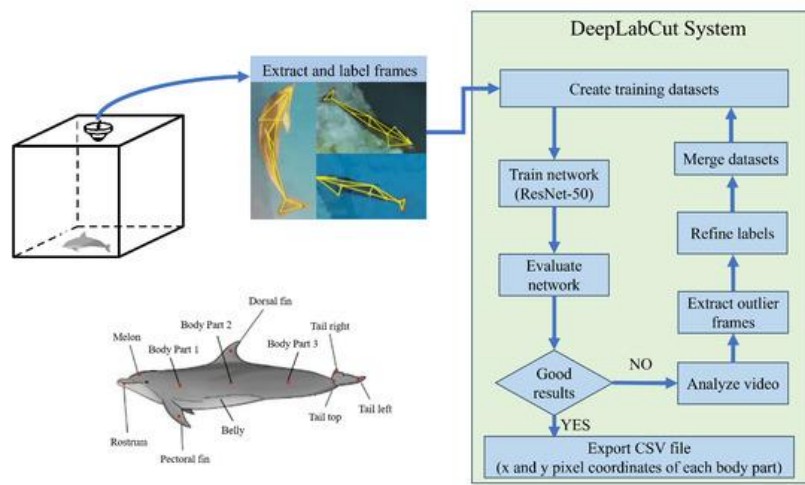
Anatomically constrained skeleton model (ACM) example (mice).

Monsees et al. (2021). Biorxiv.

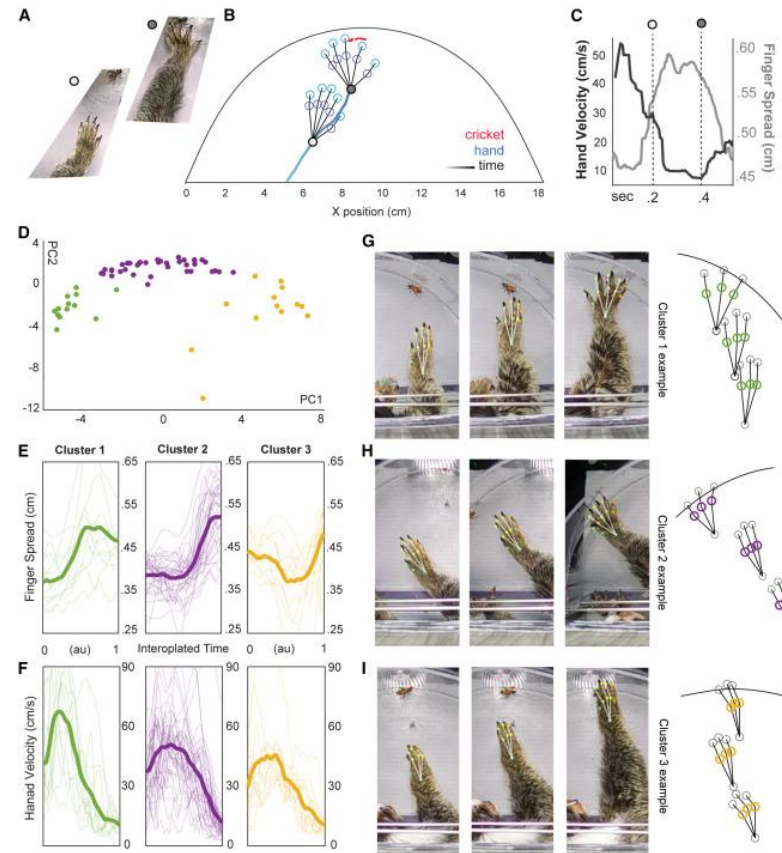
# Which bodyparts can be detected?

- Anything you want to focus on your animal(s), joints, hands, fingers, snout/tail base, ears, whiskers...
- But **doesn't** need to be a moving object (i.e. light, box, toy, etc. also works)

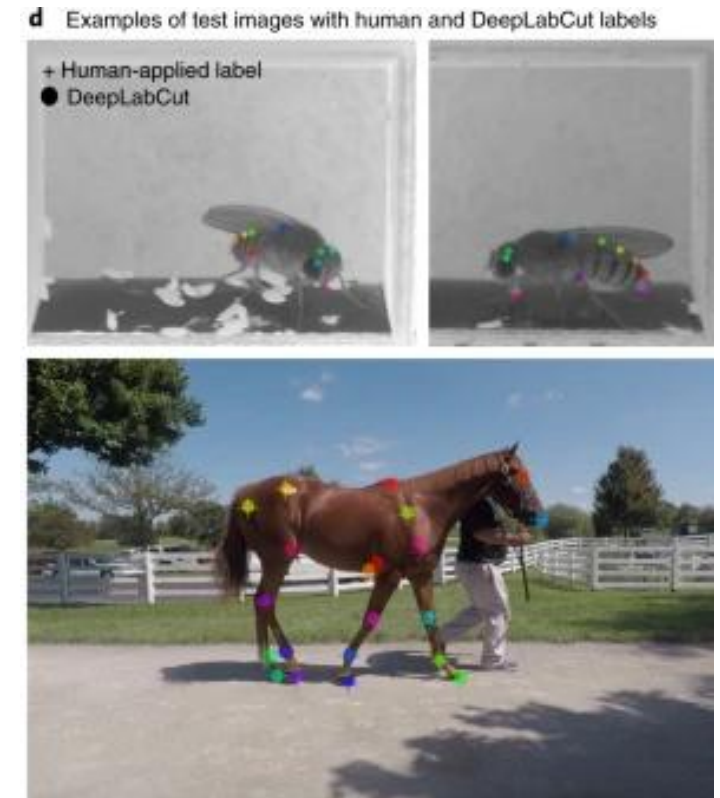
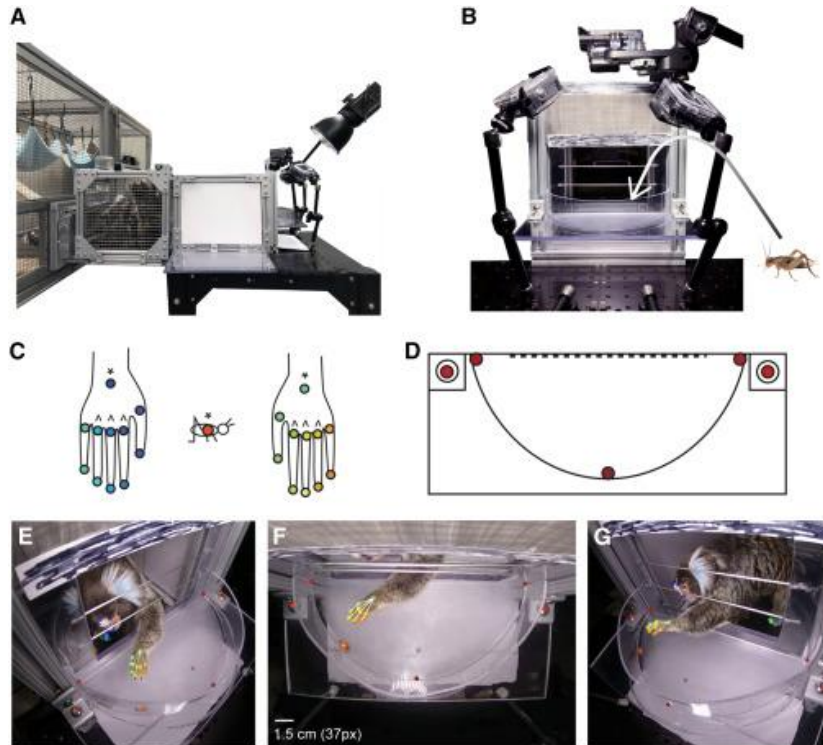




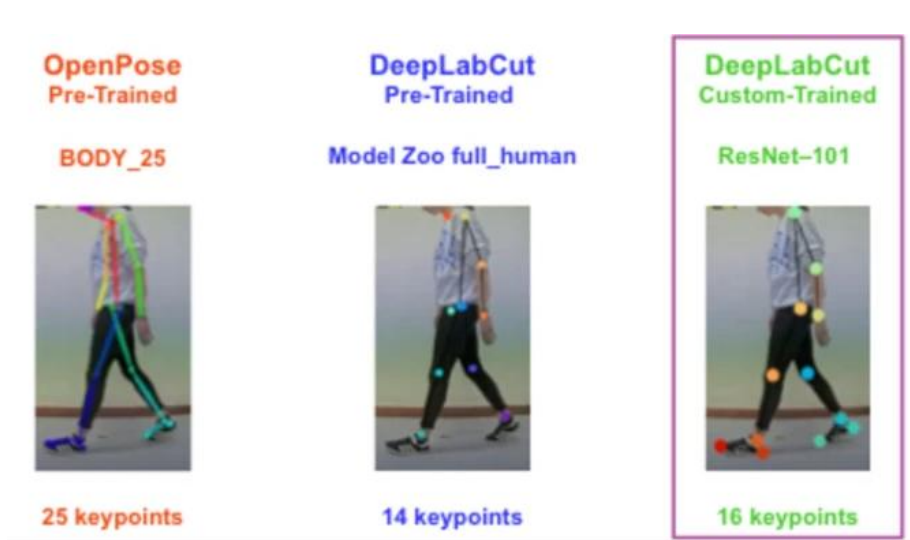
Tseng, et al. (2024). J. Mar. Sci. Eng.



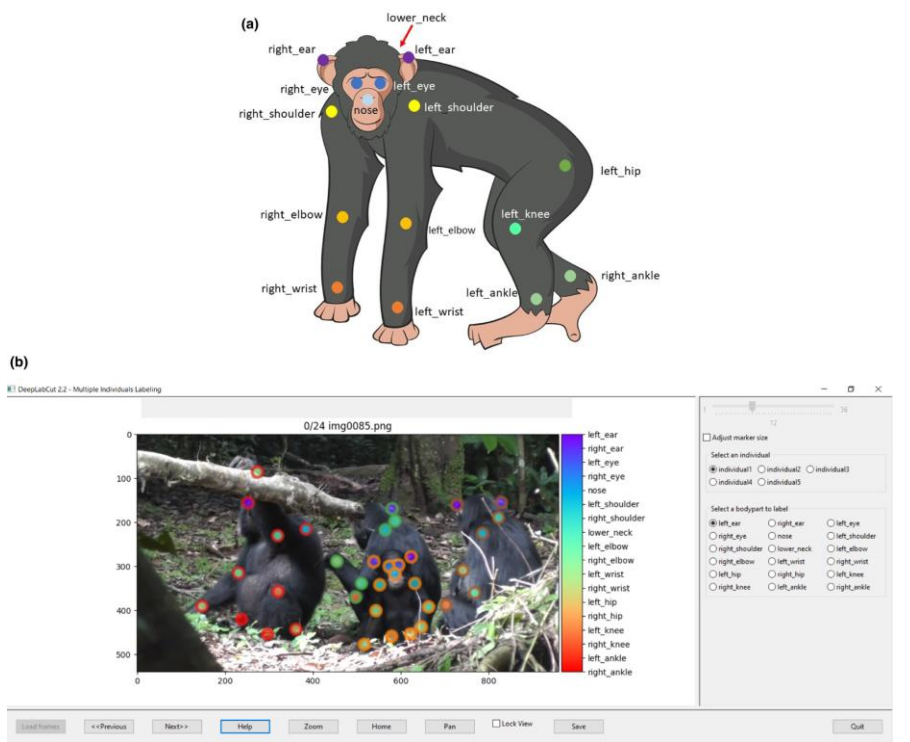
Shaw, et al. (2023). Current Biology.



Nath, et al. (2019). Nature Protocols.

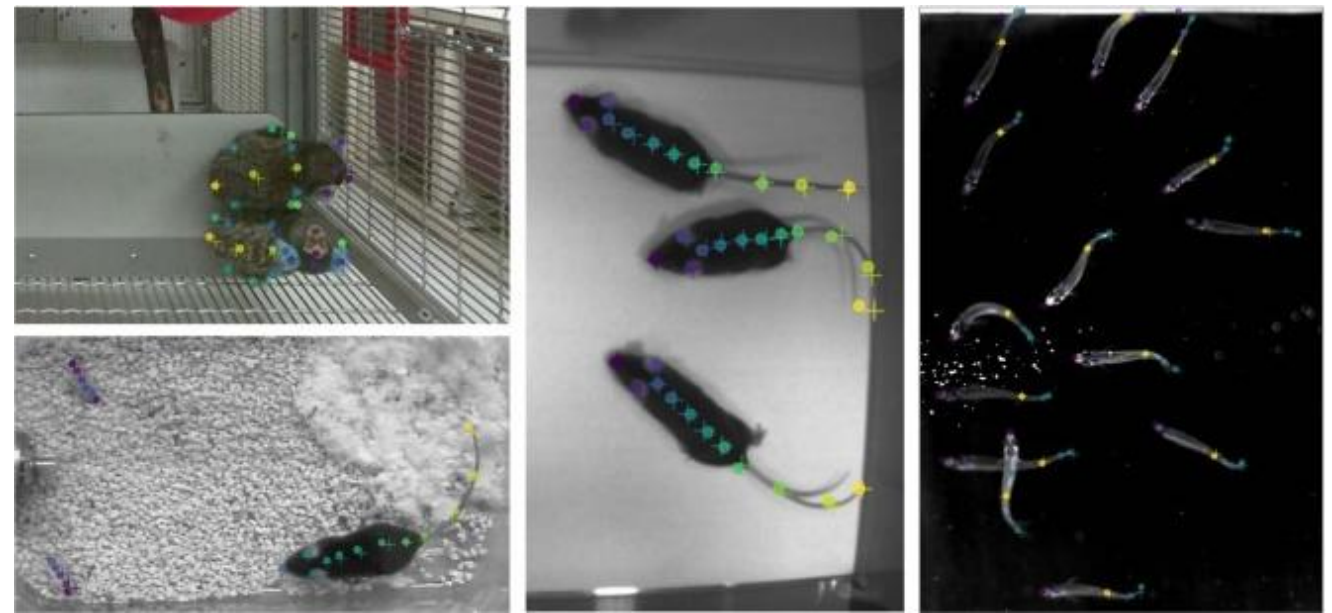
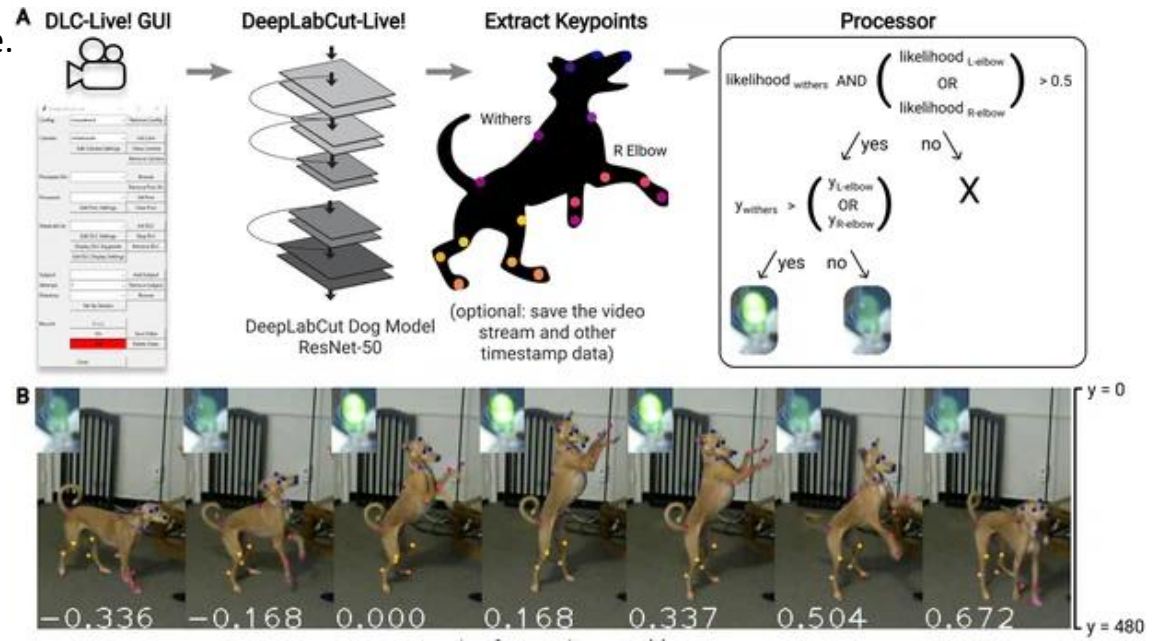


Panconi et al. (2025). Scientific Reports.



Wiltshire, et al. (2023). Journal of Animal Ecology.

Kane, et al. (2020). eLife.



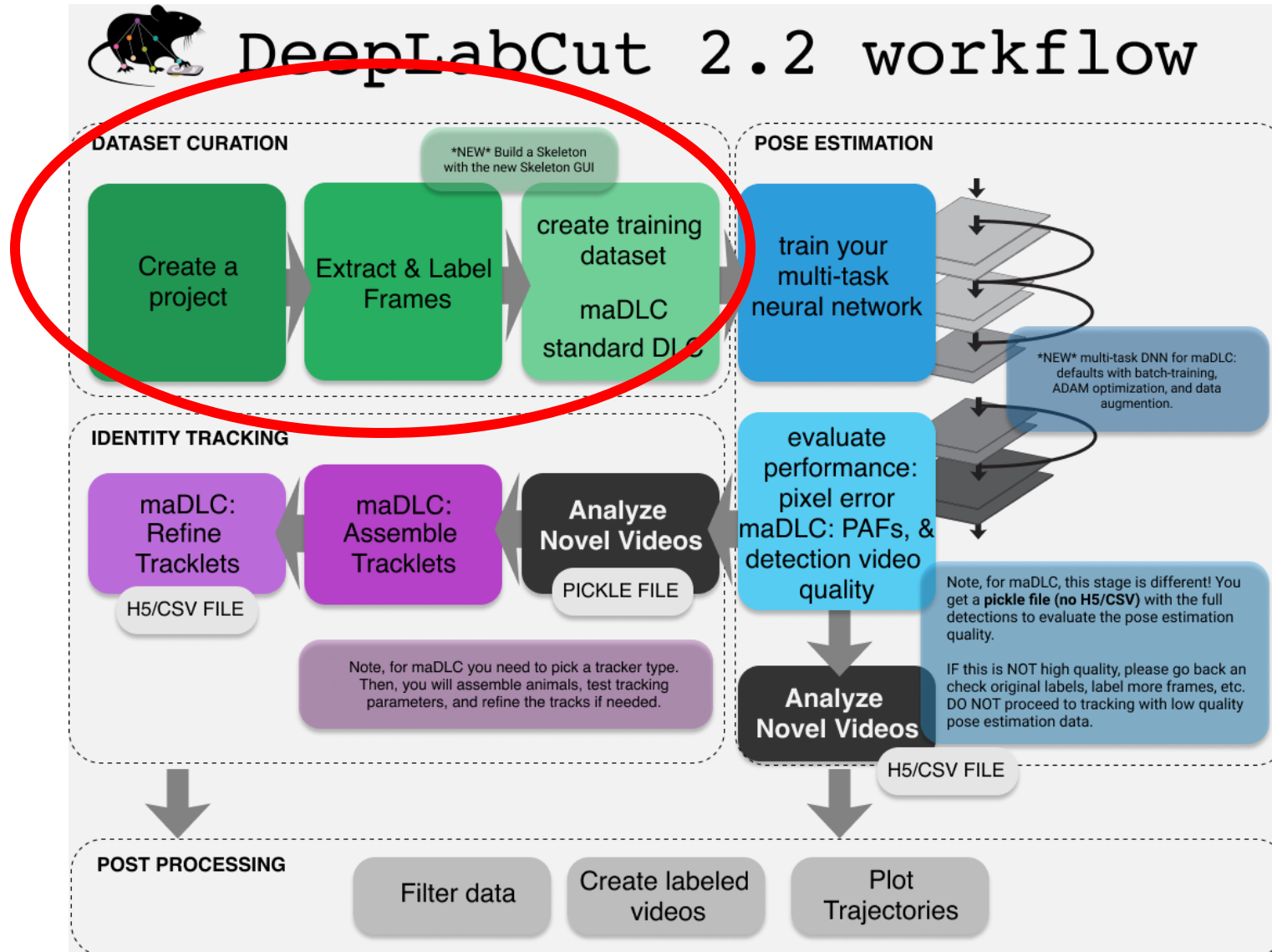
Lauer, et al. (2022). Nature Methods.

# DLC workflow

1. Opening a python session,
2. Importing deeplabcut,
3. Creating a project,
  - single animal vs. multiple animals
4. Selecting frames,
5. Labeling frames,
6. Then training a network...



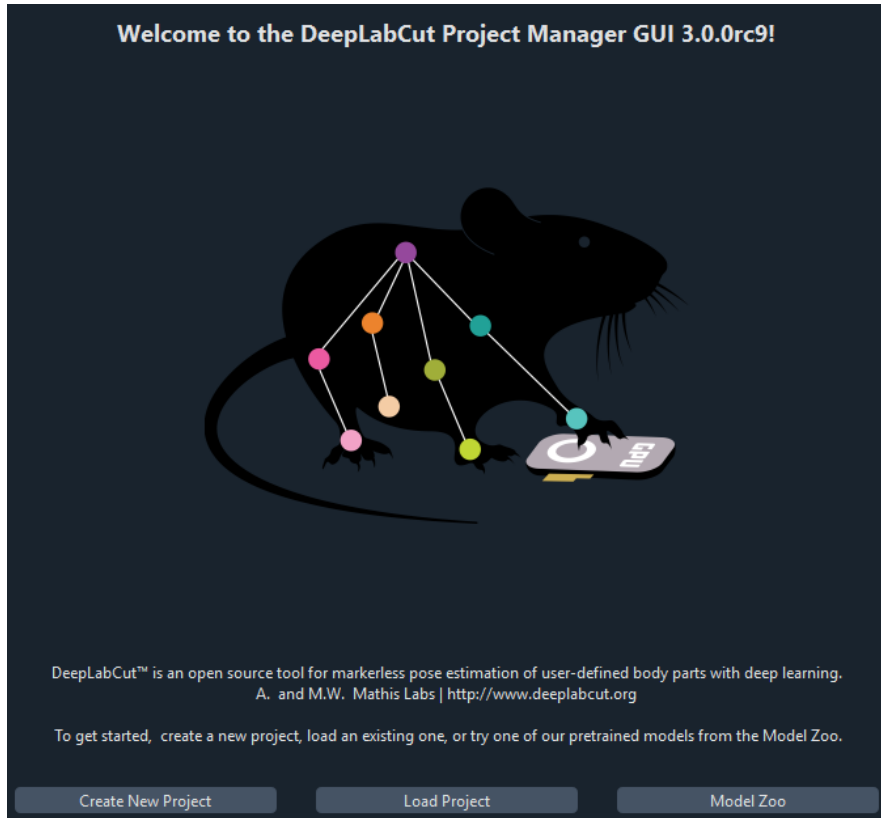
# DLC workflow – integration of annotation, training and inference



Lauer et al. (2022). Nature Methods.  
Nath et al. (2019). Nature Protocols.



# DLC GUI vs Google colab



- Simple user interface
- All analysis steps could be done here
- Could also use Prompt commands



- Access to powerful GPUs on Google's servers for faster training
- Session timeout
- Larger datasets or more complex models

But...

- Session timeout
- No Persistent Environment

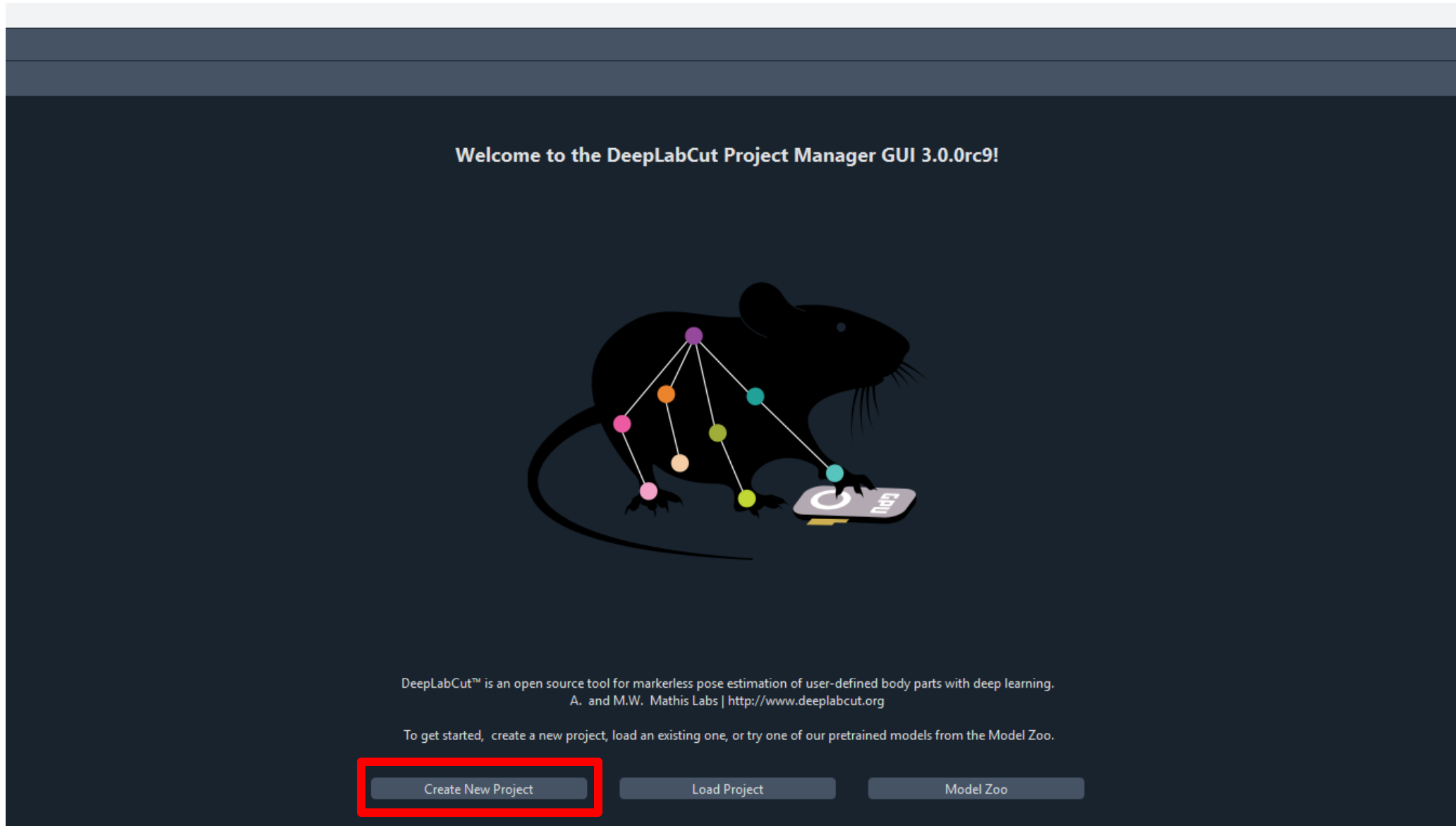
# Summary of DLC commands

Table 1   Summary of commands	
Operation	Command
Open IPython and import DeepLabCut (Step 1)	<code>ipython</code> <code>import deeplabcut</code>
Create a new project (Step 2)	<code>deeplabcut.create_new_project('project_name', 'experimenter', ['path of video 1', 'path of video2', ...])</code>
Set a <code>config_path</code> variable for ease of use (Step 3)	<code>config_path = '/yourdirectory/project_name/config.yaml'</code>
Extract frames (Step 4)	<code>deeplabcut.extract_frames(config_path)</code>
Label frames (Steps 5 and 6)	<code>deeplabcut.label_frames(config_path)</code>
Check labels (optional)(Step 7)	<code>deeplabcut.check_labels(config_path)</code>
Create training dataset (Step 8)	<code>deeplabcut.create_training_dataset(config_path)</code>
Train the network (Step 9)	<code>deeplabcut.train_network(config_path)</code>
Evaluate the trained network (Step 11)	<code>deeplabcut.evaluate_network(config_path)</code>
Video analysis and plotting results (Step 11)	<code>deeplabcut.analyze_videos(config_path, ['path of video 1 or folder', 'path of video2', ...])</code>
Video analysis and plotting results (Step 12)	<code>deeplabcut.plot_trajectories(config_path, ['path of video 1', 'path of video2', ...])</code>
Video analysis and plotting results (Step 13)	<code>deeplabcut.create_labeled_video(config_path, ['path of video 1', 'path of video2', ...])</code>
Refinement: extract outlier frames (Step 14)	<code>deeplabcut.extract_outlier_frames(config_path, ['path of video 1', 'path of video 2'])</code>
Refine labels (Step 15)	<code>deeplabcut.refine_labels(config_path)</code>
Combine datasets (Step 16)	<code>deeplabcut.merge_datasets(config_path)</code>

Nath et al. (2019). Nature Protocols.

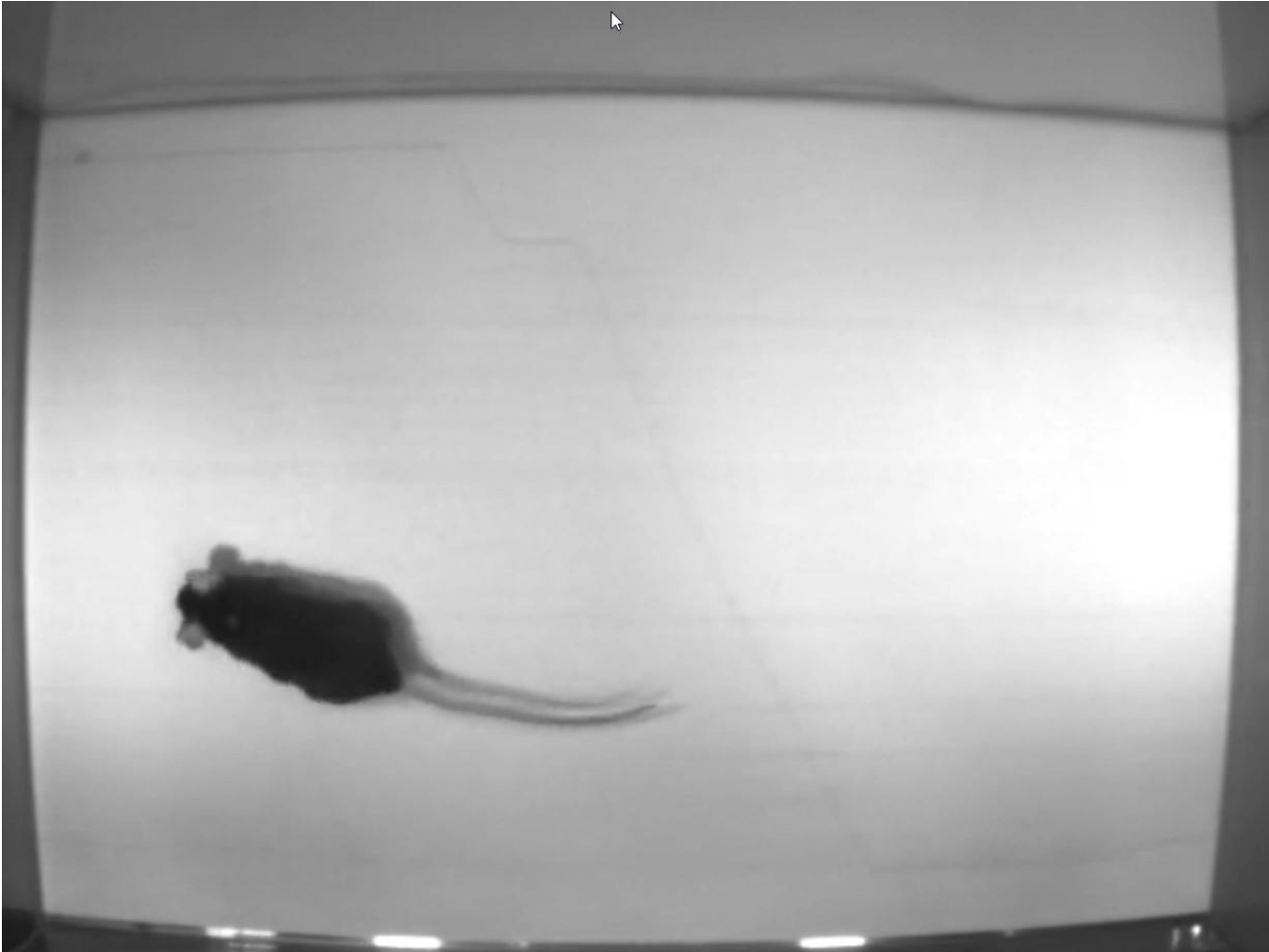
<https://www.nature.com/articles/s41596-019-0176-0/tables/1>

# Have a play with DeepLabCut GUI



# Test video

(reference: <https://github.com/DeepLabCut/DeepLabCut/tree/main/examples/openfield-Pranav-2018-10-30>)



Mathis et al. (2018). Nature Neuroscience.  
Nath, Mathis et al. (2019). Nature Protocols.  
Lauer et al. (2022). Nature Methods.  
Ye et al. (2024). Nature Communications.



New Project

Project: test

Experimenter: jiayue

Location: C:\Users\jyang291\Desktop\test-jiayue-2025-07-25

No

Do you want to create a 3D pose estimation project?

(What is needed for a 3D project?)

No

Are there multiple individuals in your videos?

(Why does this matter?)

Bodyparts to track

- snout
- leftear
- rightear
- tailbase
-

☒ 1 file selected

☒ E:/DLC\_residency\_slides\m3v1mp4.mp4

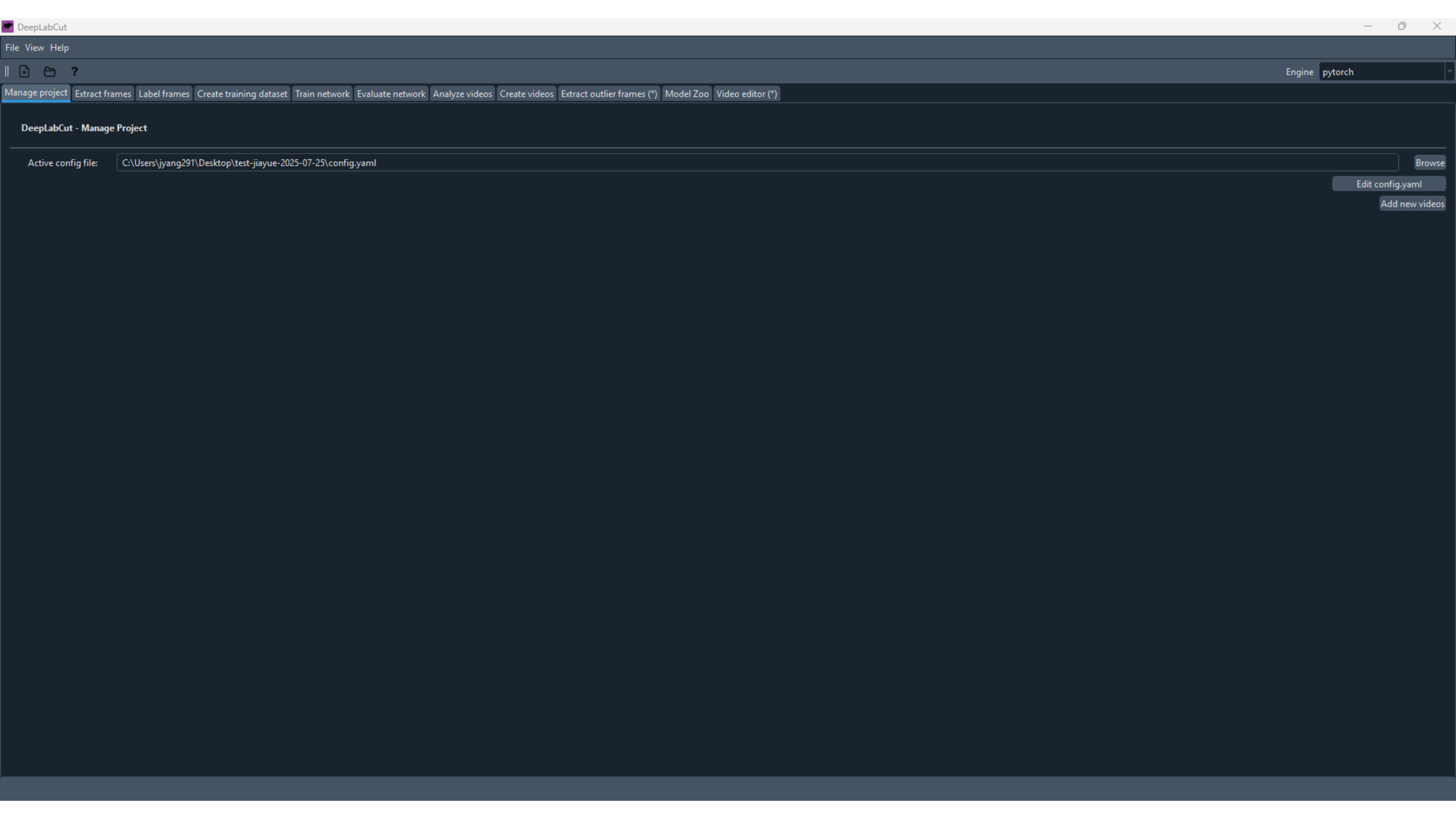
Browse folders for videos

Clear

☒ Copy videos to project folder

Create

Select if 3D or multi-animal videos



DeepLabCut - Manage Project

Active config file: C:\Users\jyang291\Desktop\test-jiayue-2025-07-25\config.yaml Browse

Edit config.yaml

Add new videos

**Box 1 | Glossary of parameters in the project configuration file (*config.yaml*)**

The *config.yaml* file sets the various parameters for generation of the training set file and evaluation of results. The meaning of these parameters is defined here, as well as referenced in the relevant steps.

**Parameters set during the project creation**

- **task:** Name of the project (e.g., mouse-reaching). (Set in Step 1; do not edit.)
- **scorer:** Name of the experimenter (set in Step 1; do not edit).
- **date:** Date of creation of the project. (Set in Step 1; do not edit).
- **project\_path:** Full path of the project, which is set in Step 1; edit this if you need to move the project to a cluster/server/another computer or a different directory on your computer.
- **video\_sets:** A dictionary with the keys as the full path of the video file and the values, *crop* as the cropping parameters used during frame extraction. (Step 1; use the function *add\_new\_videos* to add more videos to the project; if necessary, the paths can be edited manually, and the *crop* can be edited manually).

**Important parameters to edit after project creation**

- **bodyparts:** List containing names of the points to be tracked. The default is set to *bodypart1*, *bodypart2*, *bodypart3*, *objectA*. Do not change after labeling frames (and saving labels). You can add additional labels later, if needed.
- **numframes2pick:** This is an integer that specifies the number of frames to be extracted from a video or a segment of video. The default is set to 20.
- **colormap:** This specifies the colormap used for plotting the labels in images or videos in many steps. Matplotlib colormaps are possible ([https://matplotlib.org/examples/color/colormaps\\_reference.html](https://matplotlib.org/examples/color/colormaps_reference.html)).
- **dotsize:** Specifies the marker size when plotting the labels in images or videos. The default is set to 12.
- **alphavalue:** Specifies the transparency of the plotted labels. The default is set to 0.5.
- **iteration:** This keeps the count of the number of refinement iterations used to create the training dataset. The first iteration starts with 0 and thus the default value is 0. This will auto-increment once you merge a dataset (after the optional refinement stage).

**If you are extracting frames from long videos**

- **start:** Start point of interval to sample frames from when extracting frames. Value in relative terms of video length, i.e., [*start*=0, *stop*=1] represents the full video. The default is 0.
- **stop:** Same as *start*, but specifies the end of the interval. Default is 1.

**Related to the Neural Network Training**

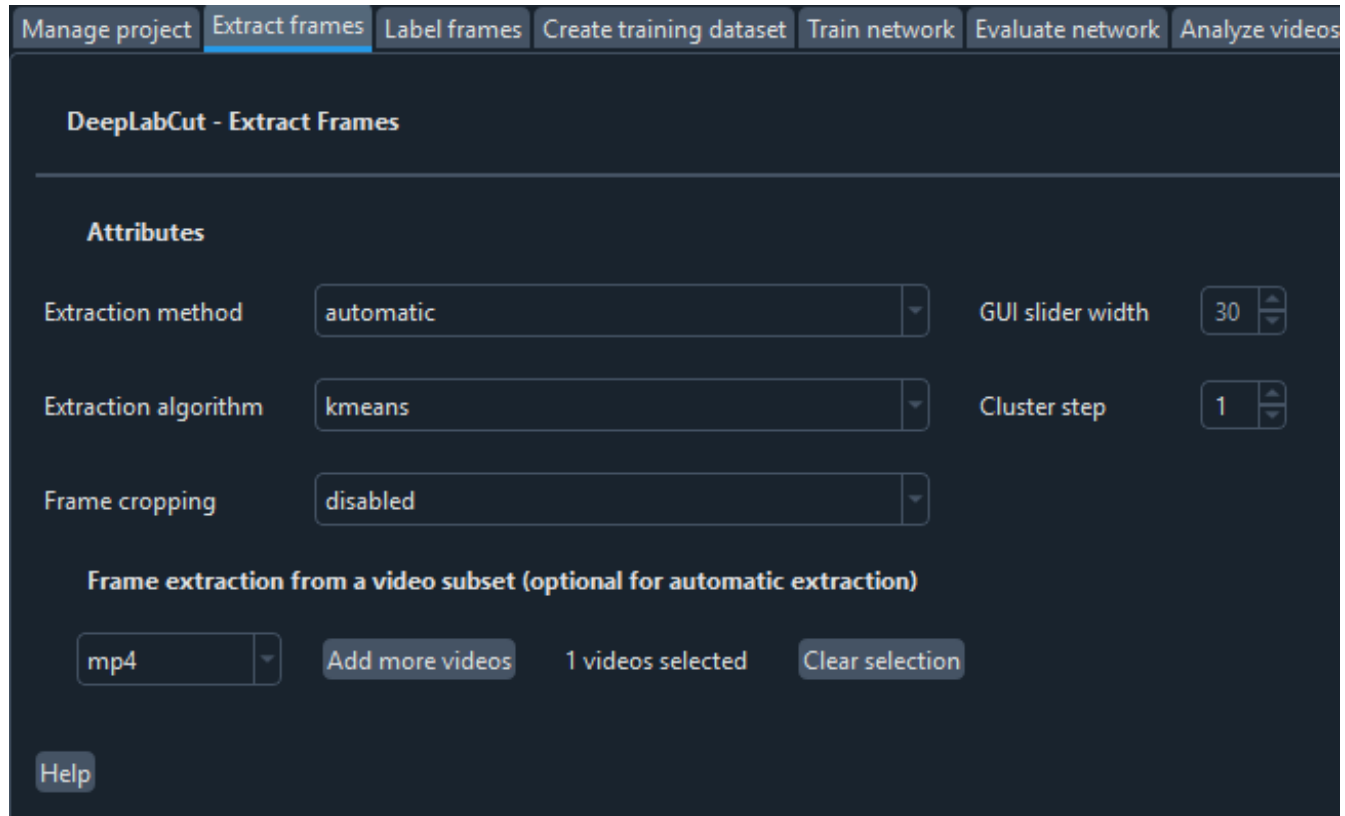
- **TrainingFraction:** This is a two-digit floating-point number in the range [0–1] used to split the dataset into training and testing datasets. The default is 0.95.
- **resnet:** This specifies which pre-trained model to use. The default is 50 (user can choose 50 or 101; see also Mathis et al.<sup>12</sup>).

**Used during video analysis (Step 13)**

- **batch\_size:** This specifies how many frames to process at once during inference (for tuning of this parameter, see Mathis and Warren<sup>27</sup>).
- **snapshotindex:** This specifies which checkpoint to use to evaluate the network. The default is –1. Use *all* to evaluate all the checkpoints. Snapshots refer to the stored TensorFlow configuration, which holds the weights of the feature detectors.
- **pcutoff:** This specifies the threshold of the likelihood and helps to distinguish likely body parts from uncertain ones. The default is 0.1.
- **cropping:** This specifies whether the analysis video needs to be cropped (in Step 13). The default is *False*.
- **x1, x2, y1, y2:** These are the cropping parameters used for cropping novel video(s). The default is set to the frame size of the video.

**Used during refinement steps**

- **move2corner:** In some (rare) cases, the predictions from DeepLabCut will be outside of the image (because of the location refinement shifts). This binary parameter ensures that those points are mapped to a user-defined point within the image so that the label can be manually moved to the correct location. The default is *True*.
- **corner2move2:** This is the target location, if *move2corner* is *True*. The default is set to (50, 50).



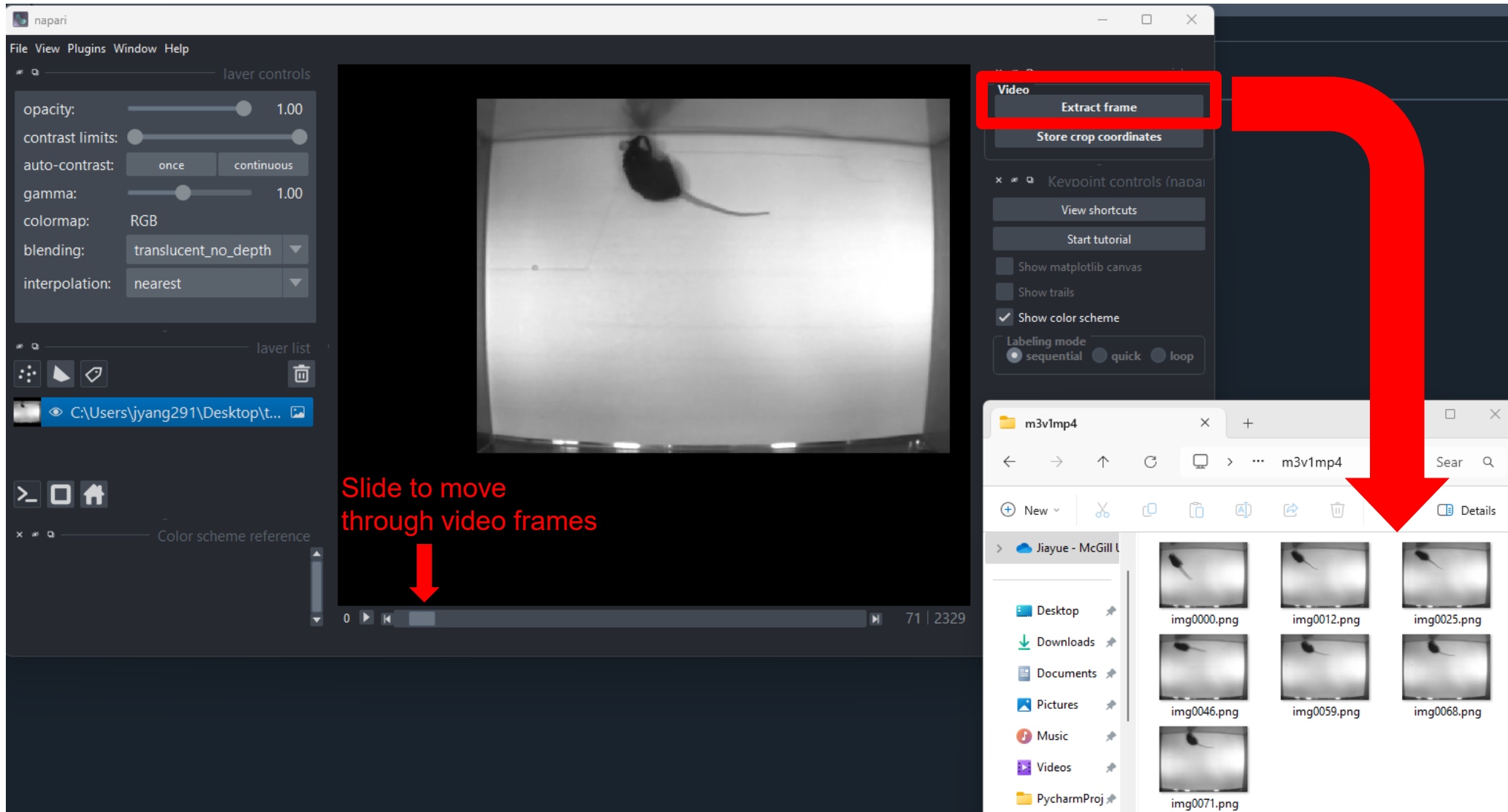
With **automatic** extraction:

→ directly proceed to the frame labeling

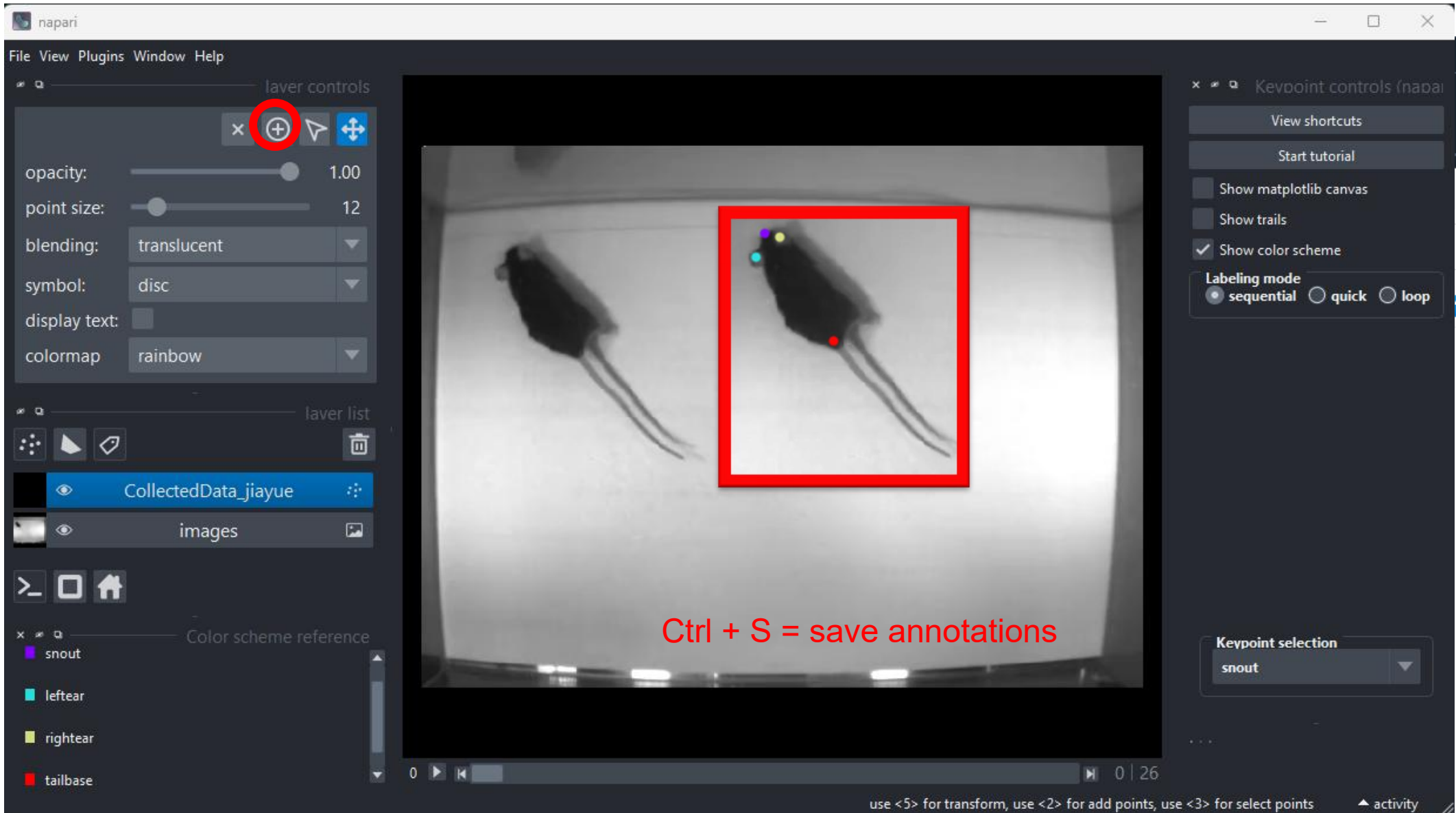
→ Console: “successfully extracted frames.”



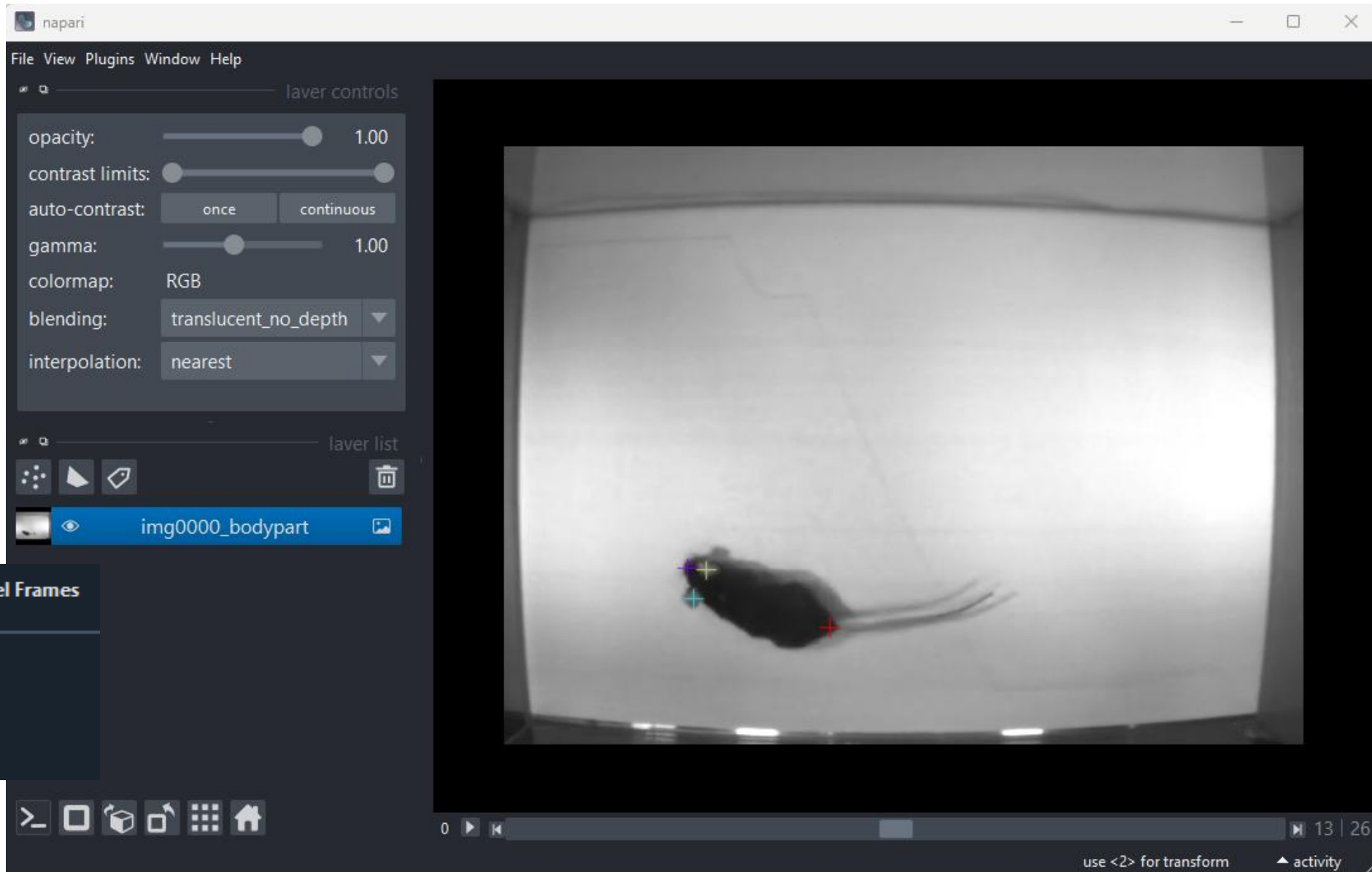
# If manual extraction:



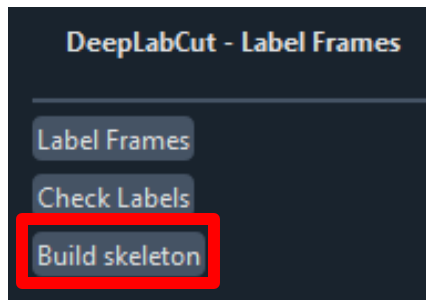
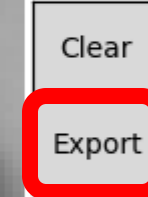
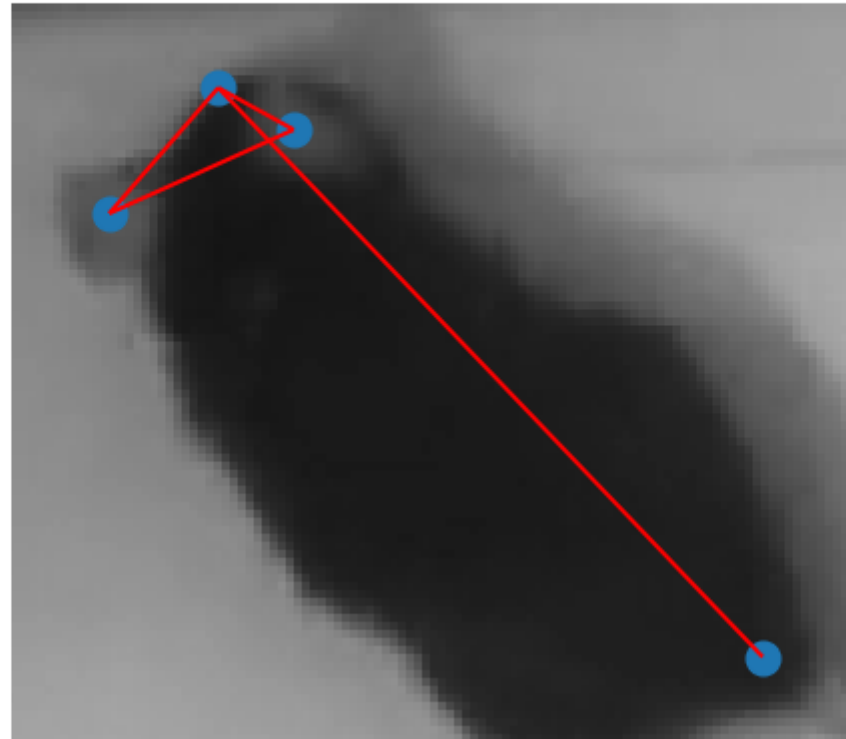
# Label frames



# Label frames



# Label frames





# What if multiple animals?

New Project

Project: multi-test

Experimenter: jiyue

Location: C:\Users\jyang291\Desktop\multi-test-jiyue-2025-07-28

☐ No Do you want to create a 3D pose estimation project? [\(What is needed for a 3D project?\)](#)

☒ Yes Are there multiple individuals in your videos? [\(Why does this matter?\)](#)

☐ No Do you have unique bodyparts in your video? [\(What are unique bodyparts?\)](#)

☐ No Label with identity? [\(What is labeling with identity?\)](#)

**Unique objects (appear only once per frame)**

**Any marker (collar beads, tail marking) on animal?**

Bodyparts to track

1.

Individual names

1.

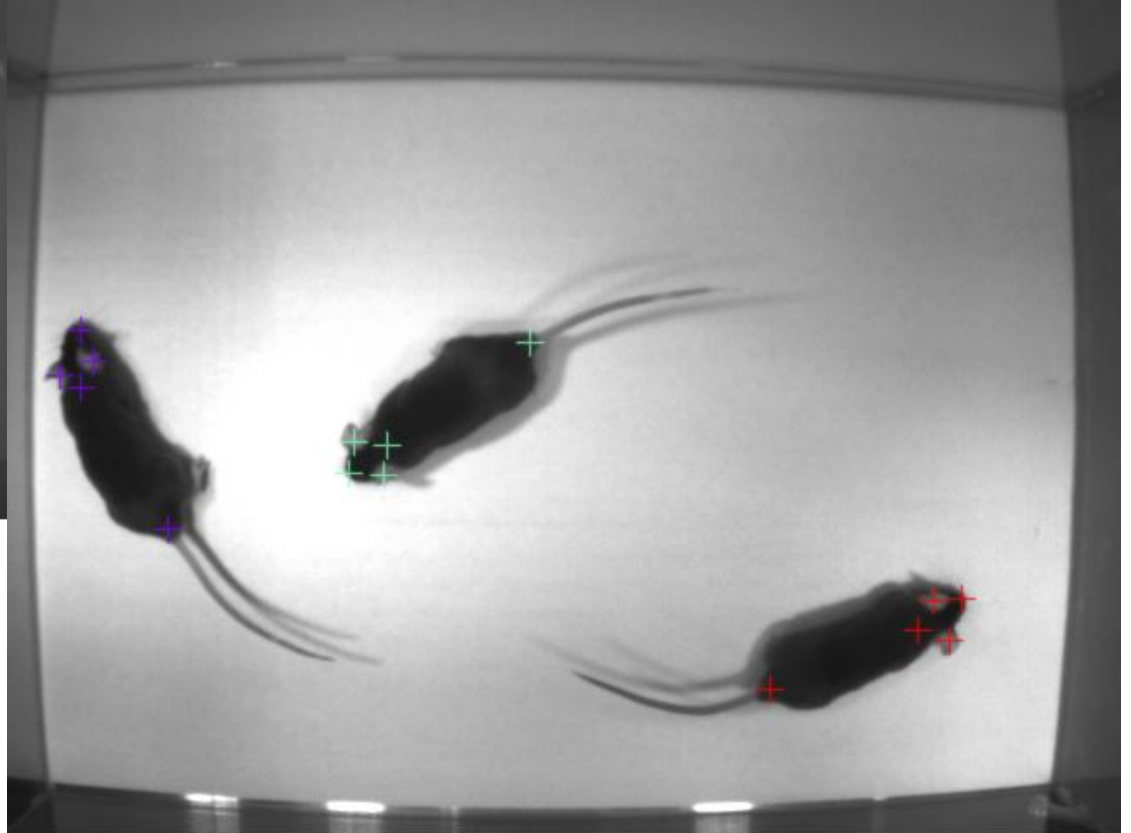
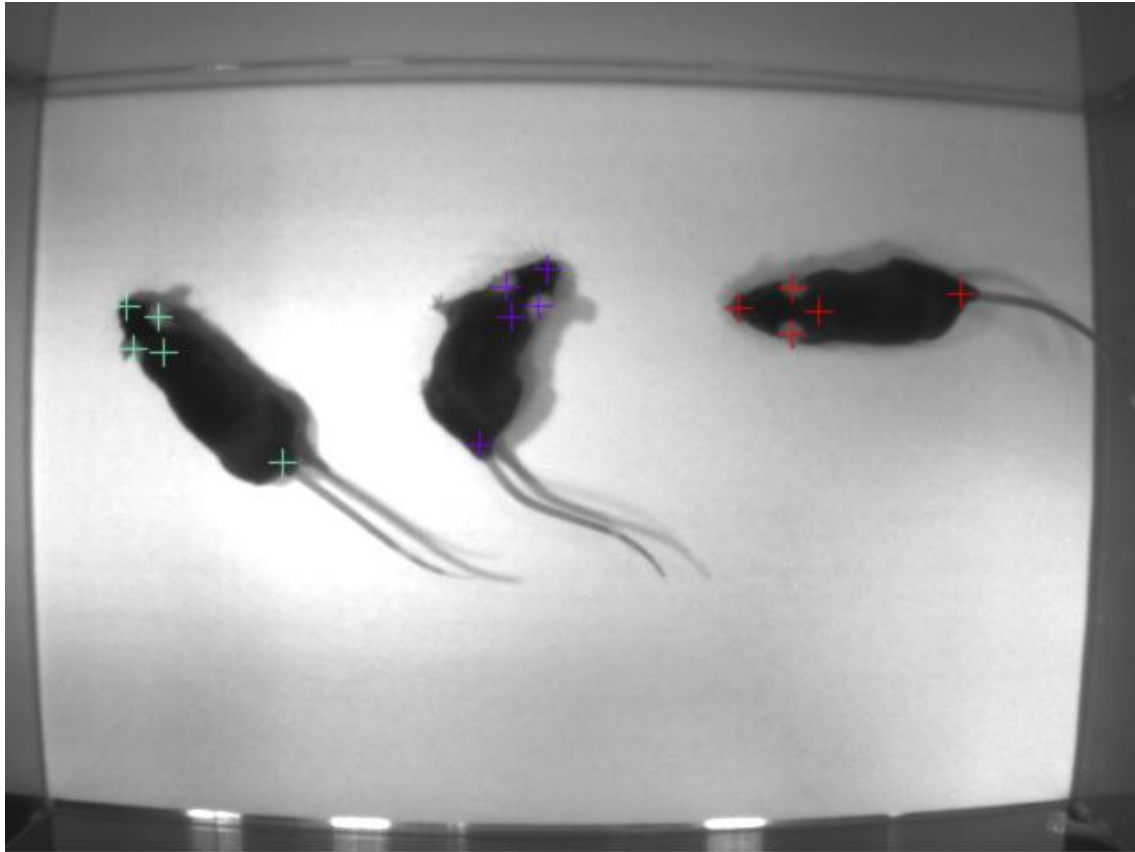
☒ Files

Browse folders for videos Clear

☐ Copy videos to project folder

Create

## Example: mice



napari

File View Plugins Window Help

layer controls

×

+

↶

↷

opacity: 1.00

point size: 12

blending: translucent

symbol: disc

display text: ☐

colormap: rainbow

layer list

⋮

◀

▶

🗑

👁

CollectedData\_jiayue

⋮

🖼

👁

images

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⏪

⏩

🏠

Color scheme reference

🟡

 head


🟢

 neck

🔴

 tailbase

Ready



Keypoint controls (napari)

View shortcuts

Start tutorial

Show matplotlib canvas

Show trails

☒ Show color scheme

Labeling mode

☒ sequential ☐ quick ☐ loop

Keypoint coloring mode

☒ bodypart ☐ individual

Keypoint selection

hippo1

head

0 | 15

use <4> for pan/zoom, use <5> for transform, use <3> for select points

activity

Switch between individuals &amp; bodyparts here

layer controls

opacity: 1.00  
point size: 40  
blending: translucent  
symbol: disc  
display text: ☐  
colormap: rainbow

layer list

CollectedData\_jiayue  
images

Color scheme reference

- head
- neck
- tailbase

Individuals differ by colors



Keypoint controls (napari)

View shortcuts

Start tutorial

- Show matplotlib canvas
- Show trails
- ☒ Show color scheme

Labeling mode

☒ sequential ☐ quick ☐ loop

Keypoint coloring mode

☒ bodypart ☐ individual

Keypoint selection

hippo3

head

0 5 | 15

use &lt;4&gt; for pan/zoom, use &lt;5&gt; for transform, use &lt;3&gt; for select points

activity

Last saved at 10:55:37



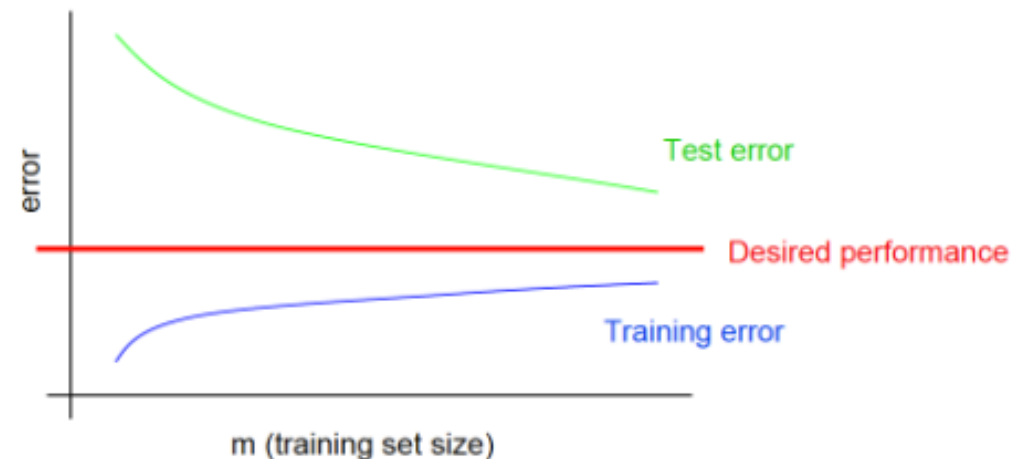
# Data management:

## How should a training dataset look like?

A good training set should look like:

- Covers the range of **variability** expected in your videos, with **clear and consistently** labeled frames showing **diverse poses, conditions, and subjects**.

Typical learning curve for high variance:



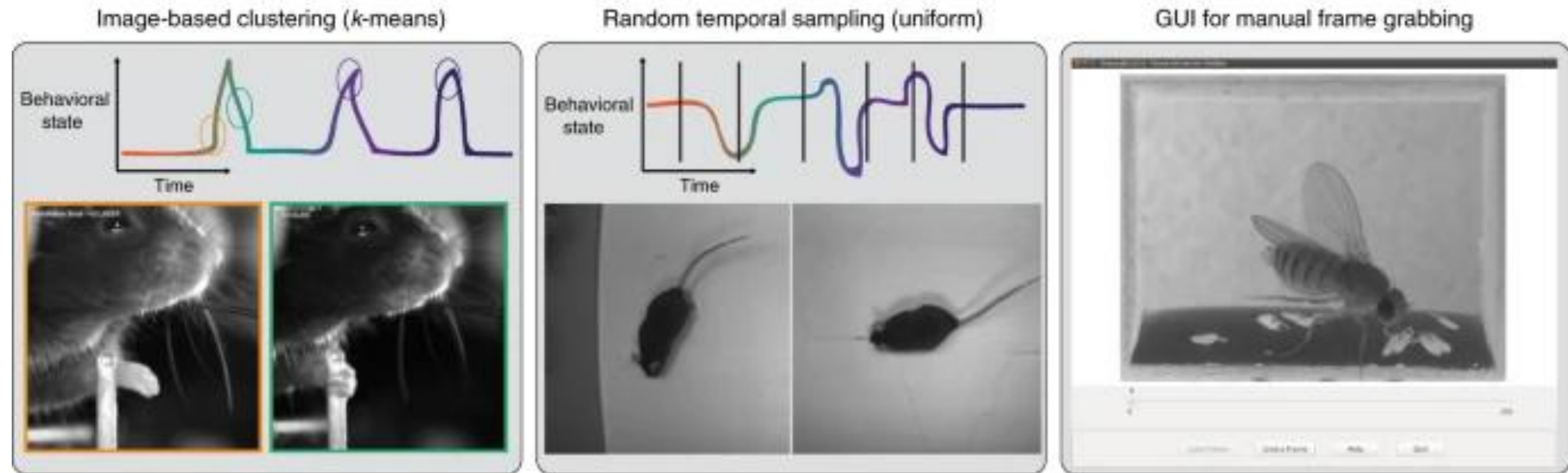
# Frame extraction

Select videos from which to grab frames:

Use videos with images from

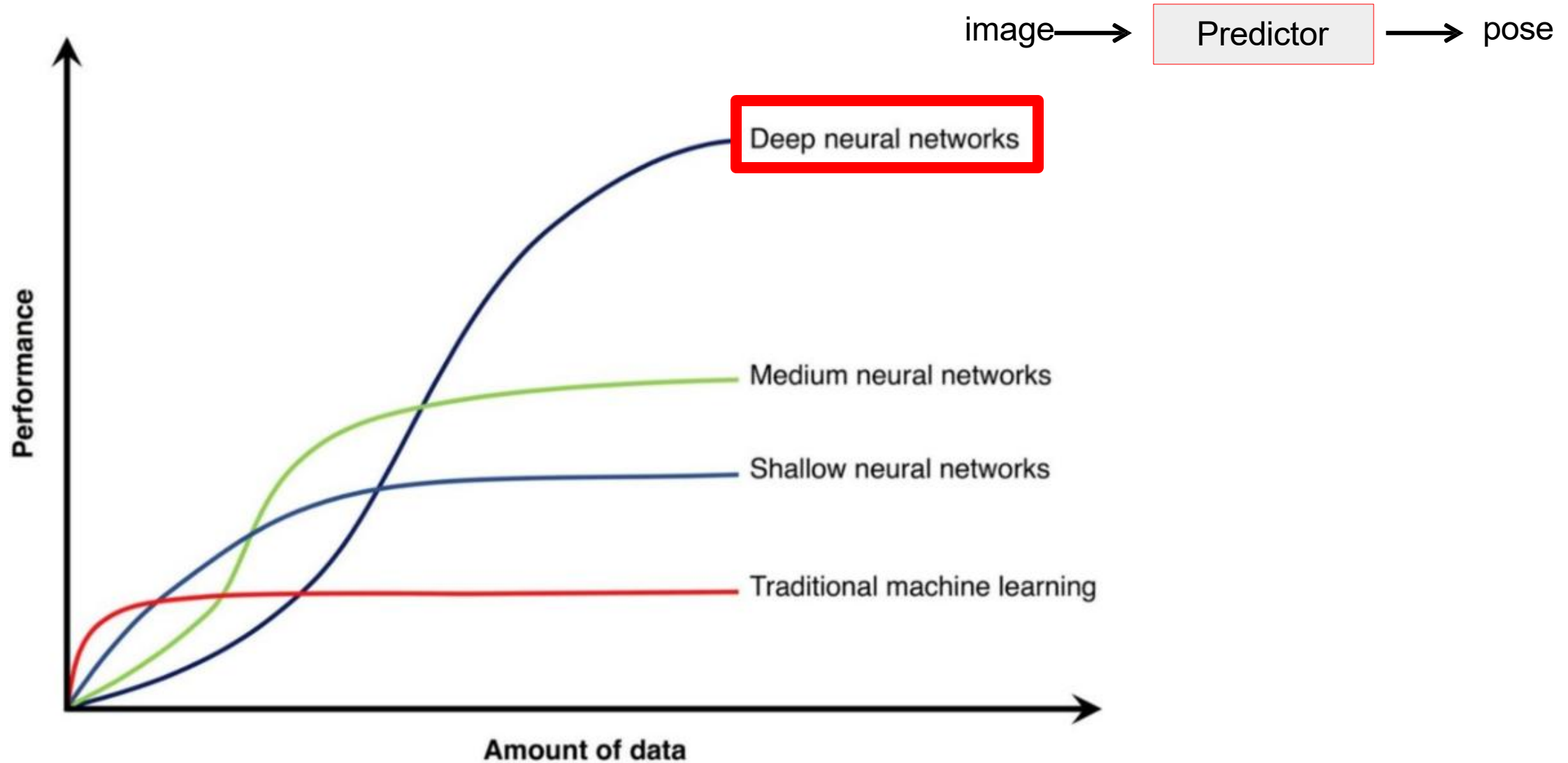
- Different sessions reflecting (if the case) varying light conditions, backgrounds, setups, and camera angles
- Different individuals, especially if they look different (i.e., brown and black mice)

3 methods for frame extraction to create a labeled train/test set



Nath et al. (2019). Nature Protocols.

# Data amount vs. performance



# Labeling on GUI (napari):

- Zooming
- Only need to label visible features/bodyparts
- The bodypart label (names) are defined in the config file
  - could be edited or added (if need to add extra)
- The labels and their meanings could be defined by users

# Labeling on GUI (napari):

- **Consistent** labeling
  - **Same locations** across frames
- Please avoid label error, especially:
  - If **small number** of images
  - **Left** and **right** labels, avoid flipping when labeling
- **Check labels** by plotting them on all figures (before training)
- Can reopen GUI to **correct labels** if mistakes