SLEAP into Behavioral Analysis: A Deep Learning Approach

M.S. Project

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Motivation

- Understanding how mice use sniffing behaviors to communicate can uncover fundamental principles of social interaction.
- Differentiating sniffing types provides insight into how age, sex, and individual traits influence social behaviors.
- Linking sniffing patterns to neural circuitry helps identify the neurological basis of social interaction.
- Analyzing temporal patterns of social behaviors offers a deeper understanding of how interactions evolve over time.
- This research can advance neuroscience by revealing mechanisms behind behavioral differences in animal models.
- Machine learning enables efficient and precise analysis of complex social behaviors, paving the way for innovative approaches to behavior research.

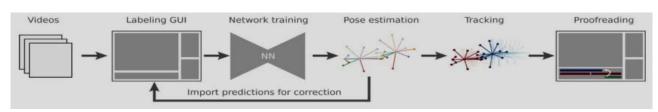
Problem Statement

- Develop methods to predict specific social behaviors (e.g., ano-genital sniffing, nose-to-nose, nose-to-neck, nose-to-left ear, nose-to-right ear) across individual videos and group categories, such as sex and age groups.
- Quantify the frequency of behavioral events, such as the number of occurrences of ano-genital sniffing.
- Measure the duration of each behavioral event to understand its temporal characteristics.
- Calculate the latency to the first occurrence of each behavior within observed interactions.
- Analyze behavior patterns observed during the initial three minutes of interaction, including corresponding latencies.
- Automate these analyses using machine learning to eliminate the need for manual tracking and annotation, improving efficiency and accuracy.

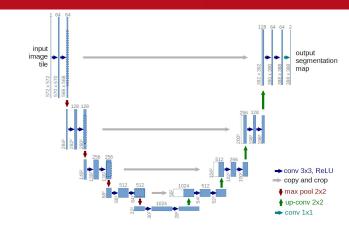


Proposed Solution

- Utilize transfer learning with pre-trained models such as U-Net, ResNet152, Hourglass, Pretrained Encoder and many more models to accurately capture and analyze complex social behaviors.
- Leverage open-source tools, including SLEAP and DeepLabCut, renowned for their effectiveness in animal pose estimation, to extract behavioral parameters.
- Implement SLEAP for automated behavioral tracking to streamline data collection and analysis.
- Employ U-Net as the backbone model to achieve precise identification and classification of specific behaviors.







U-NET Architecture



SLEAP for APT





Approach

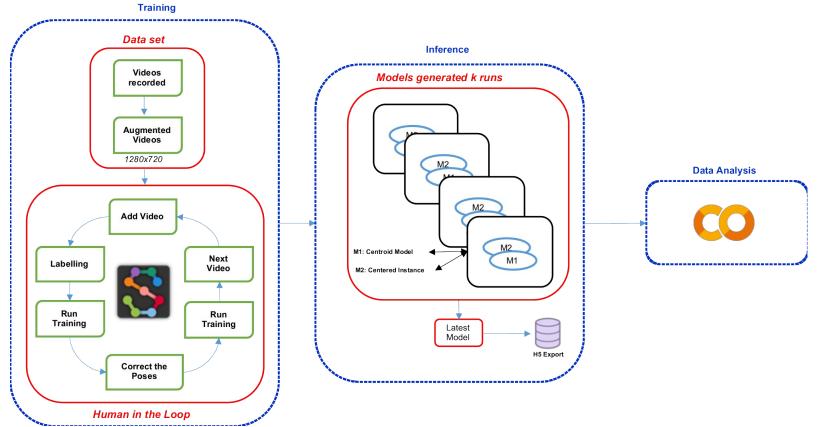




SLEAP – Why?

- Deep-learning-based framework for precise multi-animal pose tracking and behavioral analysis.
- Enables training and utilization of models to automatically track movements across diverse animal types.
- Provides a user-friendly interface for labeling, training, and proofreading pose estimation.
- Supports single- and multi-animal tracking with both top-down and bottom-up training strategies.
- Offers state-of-the-art pretrained and customizable neural network architectures, requiring minimal labeled data.
- Ensures fast training and inference, facilitating efficient analysis of recorded videos.
- Supports remote workflows and includes a flexible developer API for integration and customization.

Design Chart







Training





Data Labelling - Before





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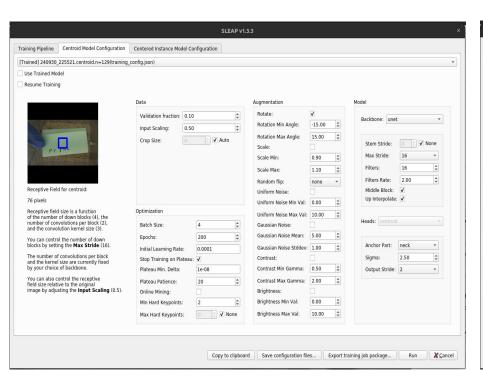
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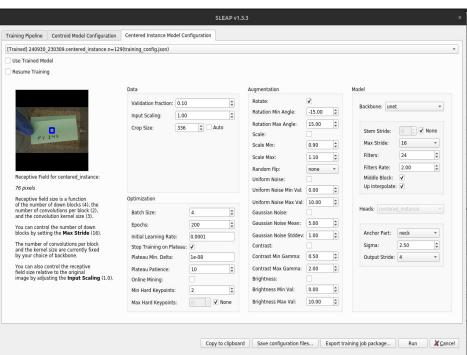
Data Labelling – After Correction





Model training Parameters



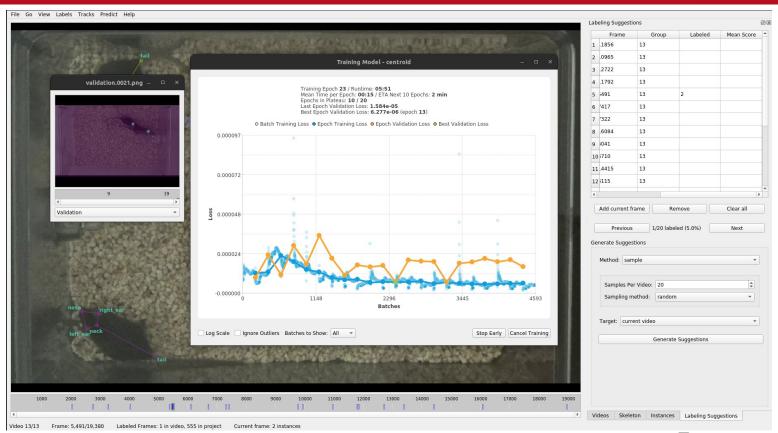


Centroid Model

Centered Instance

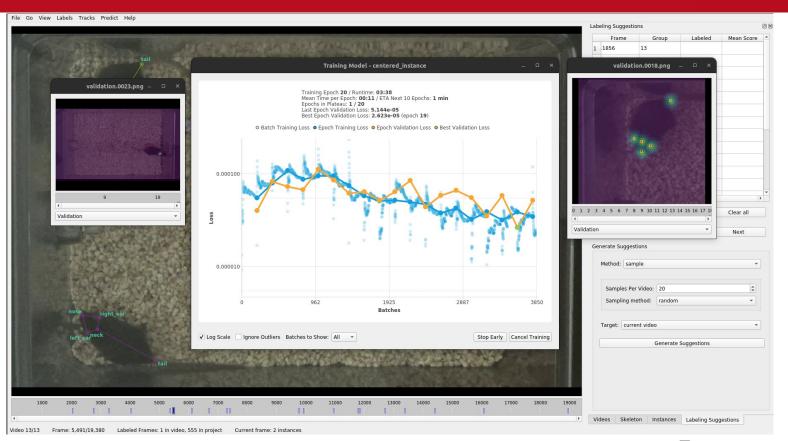


Centroid Loss





Centered Instance Loss



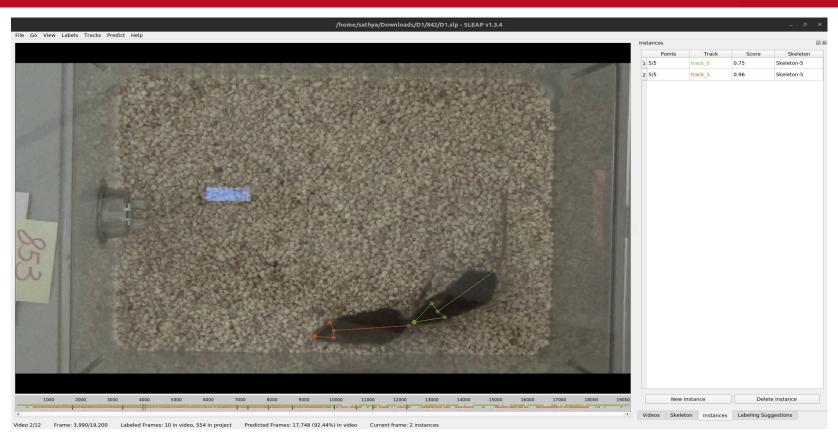


Inference





Prediction and Export H5 file





Analysis



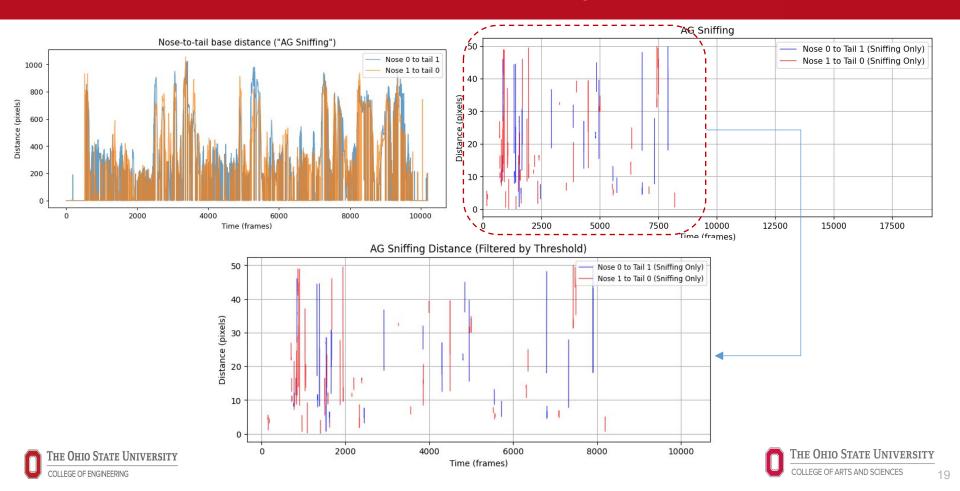
Results

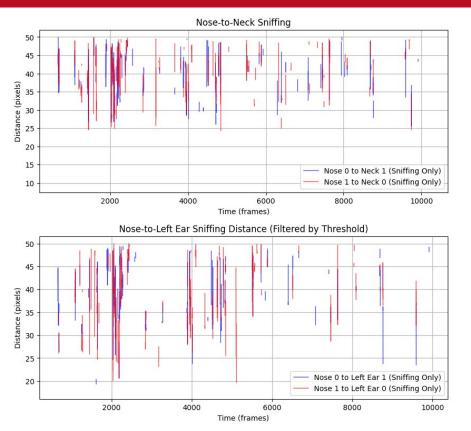
- Analyze the following behavioral parameters for both individual videos and groups of five videos:
 - Nose-to-tail sniffing (ano-genital sniffing).
 - Nose-to-nose sniffing.
 - Nose-to-right-ear sniffing.
 - Nose-to-left-ear sniffing.
 - Nose-to-neck sniffing.
- Determine the first occurrence of each behavior for one video
- Quantify the total occurrences of each behavior for one video
- Examine each behavior and its latency during the first three minutes of the video.

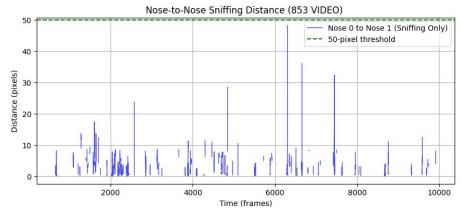
```
with h5py.File("/content/drive/MyDrive/Neuroscience/H5/Female\
     adult C/853.000 853 interaction.analysis.h5", "r") as f:
        print("Available datasets:")
        def printname(name):
            print(name)
        f.visit(printname) # This will print all dataset names
→ Available datasets:
    edge_inds
    edge names
    instance scores
    labels path
    node names
    point_scores
    provenance
    track names
    track_occupancy
    tracking scores
    tracks
    video_ind
    video path
```

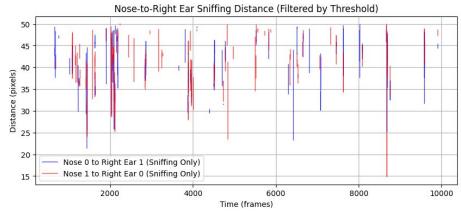
Structure of H5 file













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Number of AG sniffing events (Nose 0 to Tail 1): 146 Number of AG sniffing events (Nose 1 to Tail 0): 198 Total number of AG sniffing events: 344

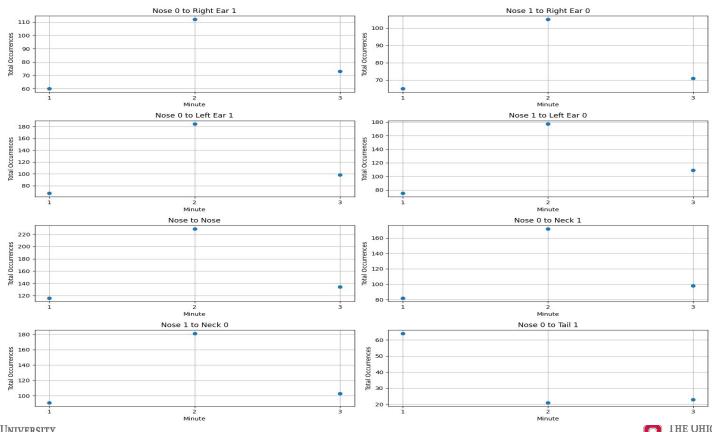
Nose-to-Neck sniffing events (Nose 0 to Neck 1): 392 Nose-to-Neck sniffing events (Nose 1 to Neck 0): 418 Total Nose-to-Neck sniffing events: 810

Total Nose-to-Nose sniffing events: 406

Nose-to-Left Ear sniffing events (Nose 0 to Left Ear 1): 352 Nose-to-Left Ear sniffing events (Nose 1 to Left Ear 0): 342 Total Nose-to-Left Ear sniffing events: 694

Nose-to-Right Ear sniffing events (Nose 0 to Right Ear 1): 270 Nose-to-Right Ear sniffing events (Nose 1 to Right Ear 0): 304 Total Nose-to-Right Ear sniffing events: 574

First occurrence of Nose 0 to Right Ear 1: Frame 663, Time 22.10 seconds First occurrence of Nose 0 to Left Ear 1: Frame 663. Time 22.10 seconds First occurrence of Nose to Nose: Frame 663. Time 22.10 seconds First occurrence of Nose 0 to Neck 1: Frame 663. Time 22.10 seconds First occurrence of Nose 1 to Neck 0: Frame 665. Time 22.17 seconds First occurrence of Nose 1 to Right Ear 0: Frame 679, Time 22.63 seconds First occurrence of Nose 1 to Left Ear 0: Frame 679. Time 22.63 seconds First occurrence of Nose 0 to Tail 1: Frame 690, Time 23.00 seconds



Observation of each behavior and its latency in the first 3 minutes of the video

Nose 0 to Right Ear 1:

- •Minute 1: 60 occurrences, Total Time: 2.00 seconds
- Minute 2: 112 occurrences, Total Time: 3.73 seconds
- •Minute 3: 73 occurrences, Total Time: 2.43 seconds

Nose 1 to Right Ear 0:

- •Minute 1: 65 occurrences, Total Time: 2.17 seconds
- Minute 2: 105 occurrences, Total Time: 3.50 seconds
- •Minute 3: 71 occurrences, Total Time: 2.37 seconds

Nose 0 to Left Far 1:

- •Minute 1: 67 occurrences. Total Time: 2.23 seconds
- •Minute 2: 184 occurrences. Total Time: 6.13 seconds
- •Minute 3: 98 occurrences, Total Time: 3.27 seconds

Nose 1 to Left Ear 0:

- •Minute 1: 75 occurrences. Total Time: 2.50 seconds
- •Minute 2: 177 occurrences, Total Time: 5.90 seconds
- •Minute 3: 109 occurrences, Total Time: 3.63 seconds

Nose to Nose:

- •Minute 1: 116 occurrences, Total Time: 3.87 seconds
- •Minute 2: 229 occurrences, Total Time: 7.63 seconds
- •Minute 3: 134 occurrences, Total Time: 4.47 seconds

Nose 0 to Neck 1:

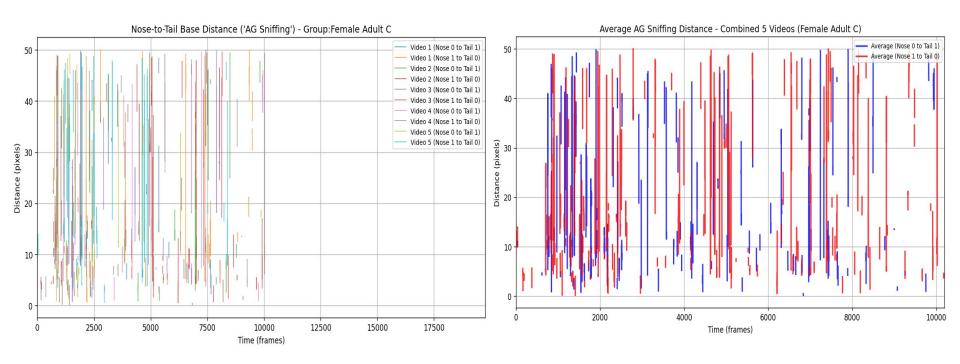
- •Minute 1: 82 occurrences, Total Time: 2.73 seconds
- •Minute 2: 172 occurrences, Total Time: 5.73 seconds
- •Minute 3: 98 occurrences, Total Time: 3.27 seconds

Nose 1 to Neck 0:

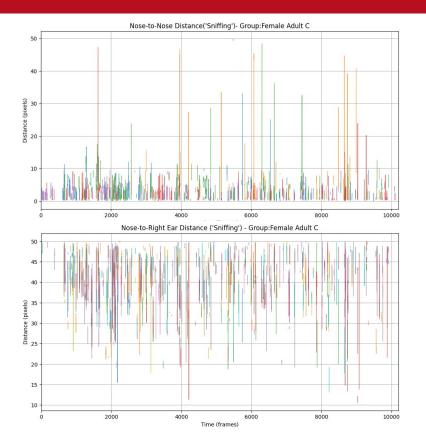
- Minute 1: 91 occurrences, Total Time: 3.03 seconds
- •Minute 2: 181 occurrences, Total Time: 6.03 seconds
- •Minute 3: 103 occurrences, Total Time: 3.43 seconds

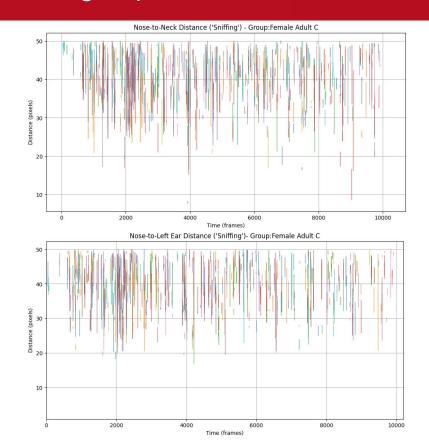
Nose 0 to Tail 1:

- Minute 1: 64 occurrences, Total Time: 2.13 seconds
- •Minute 2: 21 occurrences, Total Time: 0.70 seconds
- Minute 3: 23 occurrences, Total Time: 0.77 seconds











Conclusion and Future Work

The study successfully utilized SLEAP for automated pose tracking to analyze mouse social behaviors, quantifying sniffing interactions with high accuracy. Key findings highlight patterns in behavior duration, frequency, and latency, providing insights into social and neural interactions. This methodology improves efficiency over manual tracking and lays a foundation for scalable behavioral neuroscience studies.

FUTURE WORK : MOSEQ (MOtion SEQuencing)

Learning Stereotyped Patterns: MoSeq identifies and learns repetitive, consistent movement patterns ("stereotyped behaviors") from keypoint tracking data, enabling automatic behavioral classification.

Temporal Analysis: MoSeq determines when these learned behaviors occur over time, providing insights into the sequence and timing of actions during social interactions.

Comprehensive Toolkit: MoSeq offers tools to fit its probabilistic model to keypoint tracking data (e.g., from SLEAP) and analyze the results for behavioral studies.

REFERENCES:

https://sleap.ai/tutorials/tutorial.html

https://www.nature.com/articles/s41592-022-01426-1

https://keypoint-moseq.readthedocs.io/en/latest/





Questions?

Thank You



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