

Set-up as in grooming_mouseSubsetSelection

```
1 from database.database import Database
  from data.constants import dbDetails, dbUser_Krista
  from models.experiments import Experiment
  from models.reviewers import Reviewer
  from database.update_database.update_from_data_dirs import update_from_data_dirs

  Database.initialize(**dbDetails, **dbUser_Krista)
  experiment_name = 'grooming'
  reviewer_name = 'Krista K'

  experiment = Experiment.from_db(experiment_name=experiment_name)
  reviewer = Reviewer.from_db(reviewer_fullname=reviewer_name)

  update_from_data_dirs(experiment)

  No mouse in the database with mouse number 7044
  No mouse in the database with mouse number 7068
  No mouse in the database with mouse number 7025
  No mouse in the database with mouse number 7076
  No mouse in the database with mouse number 747
  No mouse in the database with mouse number 7035
  No mouse in the database with mouse number 750
```

In order to select the sequences, I'll have to know how the session data is stored either:

1. All videos are present and no prior processing has been completed; or
2. Session folders have 'cut' folders inside of them, containing trials.

In the case of 2, we can easily collect a list of all trials and then sample one per session.

In the case of 1, we can randomly sample a numbers 1-30, and this provides the starting point for looking for a grooming sequence.

Our list of mice for analysis is: 7169, 7043, 7014, 745, 7061, 7062, 7063, 7064, 7065, 7166

First, the mouse details will be loaded from the ParticipantDetails folder in order to obtain a list of all sessions for this animal.

```
6 from random import randrange
  from pathlib import Path

  from models.mouse import Mouse
  from models.participant_details import ParticipantDetails
  from models.sessions import Session

  all_eartags = [7169, 7043, 7014, 745, 7061, 7062, 7063, 7064, 7065, 7166]

  all_selected_sessions = dict()
  for eartag in all_eartags:
      mouse = Mouse.from_db(eartag)
```

```

mouse_details = ParticipantDetails.from_db(eartag, experiment_name)
all_sessions = Session.list_all_sessions(mouse, experiment)
pre_sr_sessions = [session for session in all_sessions if 'G4' not in session.session_dir]
post_sr_sessions = [session for session in all_sessions if 'G4' in session.session_dir]
session_timePoints_preSR = list()
session_timePoints_postSR = list()
for session in pre_sr_sessions:
    cut_folders = list(Path(session.session_dir).glob('*_cut/'))
    if len(cut_folders) > 0:
        selected_cut_folder = cut_folders.pop()
        trials_in_folders = list(Path(selected_cut_folder).glob('*'))
        selected_timepoint = trials_in_folders.pop()
    else:
        selected_timepoint = randrange(1,30)
    session_timePoints_preSR.append((session.session_dir, selected_timepoint))
for session in post_sr_sessions:
    cut_folders = list(Path(session.session_dir).glob('*_cut/'))
    if len(cut_folders) > 0:
        selected_cut_folder = cut_folders.pop()
        trials_in_folders = list(Path(selected_cut_folder).glob('*'))
        selected_timepoint = trials_in_folders.pop()
    else:
        selected_timepoint = randrange(1,30)
    session_timePoints_postSR.append((session.session_dir, selected_timepoint))
if len(post_sr_sessions) == 0:
    selected_timepoint = randrange(1,30)
    session_timePoints_postSR.append(('G4', selected_timepoint))
all_selected_sessions[eartag] = {'preSR': session_timePoints_preSR, 'postSR': session_time

```

```

7 import pprint
pprint.pprint(all_selected_sessions)

```

```

{745: {'postSR': [{'/Volumes/SharedX/Neuro-Leventhal/data/mouseGrooming/et745/et745_20190507_C
'preSR': [{'/Volumes/SharedX/Neuro-Leventhal/data/mouseGrooming/et745/et745_20181130_CC
PosixPath('/Volumes/SharedX/Neuro-Leventhal/data/mouseGrooming/et745/et745_2
('/Volumes/SharedX/Neuro-Leventhal/data/mouseGrooming/et745/et745_20181207_CC
PosixPath('/Volumes/SharedX/Neuro-Leventhal/data/mouseGrooming/et745/et745_2
('/Volumes/SharedX/Neuro-Leventhal/data/mouseGrooming/et745/et745_20181214_CC
PosixPath('/Volumes/SharedX/Neuro-Leventhal/data/mouseGrooming/et745/et745_2
7014: {'postSR': [('G4', 12)],
'preSR': [{'/Volumes/SharedX/Neuro-Leventhal/data/mouseGrooming/et7014/et7014_20191111
16),
('/Volumes/SharedX/Neuro-Leventhal/data/mouseGrooming/et7014/et7014_20200602
17),
('/Volumes/SharedX/Neuro-Leventhal/data/mouseGrooming/et7014/et7014_20191119
19)]]},
7043: {'postSR': [{'/Volumes/SharedX/Neuro-Leventhal/data/mouseGrooming/et7043/et7043_2019110
'preSR': [{'/Volumes/SharedX/Neuro-Leventhal/data/mouseGrooming/et7043/et7043_20190822
12),
('/Volumes/SharedX/Neuro-Leventhal/data/mouseGrooming/et7043/et7043_20190809
27),
('/Volumes/SharedX/Neuro-Leventhal/data/mouseGrooming/et7043/et7043_20190819
3)]]},
7061: {'postSR': [{'/Volumes/SharedX/Neuro-Leventhal/data/mouseGrooming/et7061/et7061_2019091
'preSR': [{'/Volumes/SharedX/Neuro-Leventhal/data/mouseGrooming/et7061/et7061_20190527
PosixPath('/Volumes/SharedX/Neuro-Leventhal/data/mouseGrooming/et7061/et706
('/Volumes/SharedX/Neuro-Leventhal/data/mouseGrooming/et7061/et7061_20190502

```

```

PosixPath('/Volumes/SharedX/Neuro-Leventhal/data/mouseGrooming/et7061/et706
('/Volumes/SharedX/Neuro-Leventhal/data/mouseGrooming/et7061/et7061_20190514
PosixPath('/Volumes/SharedX/Neuro-Leventhal/data/mouseGrooming/et7061/et706
7062: {'postSR': [{'/Volumes/SharedX/Neuro-Leventhal/data/mouseGrooming/et7062/et7062_2019091
'preSR': [('/Volumes/SharedX/Neuro-Leventhal/data/mouseGrooming/et7062/et7062_20190515
4),
('/Volumes/SharedX/Neuro-Leventhal/data/mouseGrooming/et7062/et7062_20190502
PosixPath('/Volumes/SharedX/Neuro-Leventhal/data/mouseGrooming/et7062/et706
('/Volumes/SharedX/Neuro-Leventhal/data/mouseGrooming/et7062/et7062_20190527
6)]]},
7063: {'postSR': [{'/Volumes/SharedX/Neuro-Leventhal/data/mouseGrooming/et7063/et7063_2019092
'preSR': [('/Volumes/SharedX/Neuro-Leventhal/data/mouseGrooming/et7063/et7063_20190514
2),
('/Volumes/SharedX/Neuro-Leventhal/data/mouseGrooming/et7063/et7063_20190527
27),
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22)]]},
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4),
('/Volumes/SharedX/Neuro-Leventhal/data/mouseGrooming/et7064/et7064_20190502
1),
('/Volumes/SharedX/Neuro-Leventhal/data/mouseGrooming/et7064/et7064_20190527
28)]]},
7065: {'postSR': [{'/Volumes/SharedX/Neuro-Leventhal/data/mouseGrooming/et7065/et7065_2019092
'preSR': [('/Volumes/SharedX/Neuro-Leventhal/data/mouseGrooming/et7065/et7065_20190502
4),
('/Volumes/SharedX/Neuro-Leventhal/data/mouseGrooming/et7065/et7065_20190527
20),
('/Volumes/SharedX/Neuro-Leventhal/data/mouseGrooming/et7065/et7065_20190515
25)]]},
7166: {'postSR': [('G4', 26)],
'preSR': [('/Volumes/SharedX/Neuro-Leventhal/data/mouseGrooming/et7166/et7166_20191122
18),
('/Volumes/SharedX/Neuro-Leventhal/data/mouseGrooming/et7166/et7166_20200602
9),
('/Volumes/SharedX/Neuro-Leventhal/data/mouseGrooming/et7166/et7166_20191115
6),
('/Volumes/SharedX/Neuro-Leventhal/data/mouseGrooming/et7166/et7166_20191108
3)]]},
7169: {'postSR': [('G4', 14)],
'preSR': [('/Volumes/SharedX/Neuro-Leventhal/data/mouseGrooming/et7169/et7169_20200602
24),
('/Volumes/SharedX/Neuro-Leventhal/data/mouseGrooming/et7169/et7169_20191115
3),
('/Volumes/SharedX/Neuro-Leventhal/data/mouseGrooming/et7169/et7169_20191122
1),
('/Volumes/SharedX/Neuro-Leventhal/data/mouseGrooming/et7169/et7169_20191108
17)]]}

```