ReadMe - V_rodents_snaptrapping_abundance_intensive

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1 Protocol

Snap trapping of small rodents has been conducted using the small quadrat method following the COAT protocol 'protocol snap trapping of rodents'.

1.1 Timing of sampling and changes in that

The trapping is conducted over two trappinghts (i.e. two checks) twice a year (spring and fall).

From 2005 to 2008 spring-trapping was conducted in late July (21.07.-28.07.) and fall-trapping in late August/early September (26.08.-08.09.).

From 2009 on, spring-trapping is conducted in early July (02.07.-04.07.) and fall-trapping in early September (02.09.-04.09.) with some exceptions. Trapping during the summer at Sandfjorddalen is usually done the three days preceding other trapping, due to logistic reasons.

Summer trapping in 2017 was done in mid-July 14.07.-16.07, due to late snowmelt.

1.2 Spatial layout of sampling and changes in that

From 2005 to 2008 the trapping was conducted at three watershed areas (Ifjordfjellet, Komagdalen and Vestre Jakobselv). Within these, three (Vestre Jakobselv and Komagdalen) or four (Ifjordfjellet) spatially separate sections were included in the design, either as sections of river valley/ separate river valleys. The sections were some kilometers from the nearest neighbouring section.

In each section, sampling quadrats are placed as pairs including a quadrat in each of the two habitat types: "thicket" (i.e. meadow and thicket mosaics in riparian habitats, the sampling quadrats being in the meadow habitat with one edge aligned with a thicket edge) and "heath" (i.e. dwarf shrub heath).

Between one and five quadrat pairs per section were included in the design.

From 2009 onwards, four sections (sections Gurrojohka and Suolojavri at Ifjordfjellet, section Jakobselv in Vestre Jakobselv and section Komagdalen_nedre in Komagdalen) were excluded from the design.

From 2013 onwards, three new sections were included in the design (sections Iesjohka and Giksjohka at Ifjordfjelle and section Sandfjorddalen in Komagdalen).

In 2017, three additional sites (ko_ko_m_new1, ko_sa_m_new1 and ko_sa_m_new2) were included in the study design because the regular sites were inaccessible due to late snowmelt.

In 2020, some of the heath sites were moved.

More detailed information about which sites were included in the study design can be found in the auxialiary file 'V rodents snaptrapping abundance intensive aux.txt'.

1.3 Changes in the sampling protocol

Species: Presence of shrews (*Sorex sp.*) and birds was recorded systematically from 2017 onwards (including 2017). Earlier records exist, but these are not complete.

2 Description of the dataset

The dataset includes three different types of files and all files are saved as ;-separated txt-files:

- One data file per year (_YEAR.txt)
- One coordinate file with coordinates of all sites (_coordinates.txt)
- One auxiliary file with information about which sites are included in the study design (_aux.txt)

2.1 V_rodents_snaptrapping_abundance_intensive_YEAR.txt

These files contain abundance data of trapped animals per site. The following species and species groups are included in the dataset:

```
[1] "Aves"
[2] "Cricetidae"
[3] "Lemmus lemmus (Norwegian lemming)"
[4] "Microtus oeconomus (Tundra vole)"
[5] "Myodes rufocanus (Grey-sided vole)"
[6] "Myodes rutilus (Northern-red-backed vole)"
[7] "Sorex araneus (Common shrew)"
[8] "Sorex caecutiens (Masked shrew)"
[9] "Sorex sp (Shrew)"
```

Example of the first rows of the data files:

```
sn_region
              sn_locality sn_section
                                         sn_site sc_type_of_sites_ecological
1 varanger ifjordfjellet eastordalen if_ea_hn_a
                                                                   heath near
2 varanger ifjordfjellet eastordalen if_ea_hn_a
                                                                   heath_near
  varanger ifjordfjellet eastordalen if_ea_hn_a
                                                                   heath near
      t_date t_season t_session v_species v_abundance v_observer v_comment
1 2005-08-30
                 fall
                              1
                                  mic oec
                                                     0
                                                             <NA>
2 2005-08-30
                                                             <NA>
                 fall
                                  myo_ruf
                                                     1
                                                                         NA
                              1
3 2005-08-30
                 fall
                                  myo_rut
                                                             <NA>
                                                                         NA
```

Description of the columns included in the data files:

Column name	Description	Possible values
sn_region	Study region	varanger
sn_locality	Locality (within region)	ifjordfjellet, komagdalen, vestre_jakobselv
sn_section	Section (within locality)	eastordalen, gurrojohka, storelva, suolojavri, komagdalen_midtre, komagdalen_nedre, komagdalen_ovre, bearalveaijohka, torvhaugdalen, jakobselv, giksjohka, iesjohka, sandfjorddalen
sn_site	Unique Site ID	e.g. if_ea_hn_a, if_ea_hn_e, if_ea_m_e, if_st_hn_c, if_st_m_d, ko_km_hn_a, ko_km_m_a, ko_kn_hn_a, ko_ko_hn_b, ko_ko_m_b
$sc_type_of_sites_ecological$	Habitat type	heath_near, meadow
t_date	Sampling date	YYYY-MM-DD
t _season	Sampling season	fall, spring
$t_session$	Trapping night	1, 2
$v_species$	Species (or larger taxa if species cannot be identified)	mic_oec, myo_ruf, myo_rut, lem_lem, sor_sp, cricetidae, aves, sor_ara, sor_cae
$v_abundance$	Number of trapped individuals	[numeric]
v_observer	Initials of observer	e.g. es (Eeva Soininen)
v_comment	Comments	[character]

$2.2 \quad V_rodents_snaptrapping_abundance_intensive_coordinates.txt$

This file contains the coordinates of all sites included in the study desgin. Coordinates are given in decimal degrees and UTM 33 (WGS 84).

Example of the first rows of coordinate files:

```
sn_site e_dd n_dd e_utm33 n_utm33
1 if_ea_hn_a 27.36533 70.40762 959977.6 7858316
2 if_ea_m_a 27.36179 70.40552 959895.0 7858059
3 if_ea_hn_b 27.34243 70.42363 958774.6 7859895
```

${\bf 2.3 \quad V_rodents_snaptrapping_abundance_intensive_aux.txt}$

This file contains further information about the dataset such as old site names (for example used in raw data files before 2019) and the years when sites were first included in the study design and when sites were excluded from the study design.

Example of the first rows of auxiliary files:

```
sn_region
              sn_locality sn_section
                                         sn_site sn_site_old year_first
1 varanger ifjordfjellet eastordalen if ea hn a
                                                         ae1h
  varanger ifjordfjellet eastordalen if_ea_hn_b
                                                         ae2h
                                                                    2005
3 varanger ifjordfjellet eastordalen if_ea_hn_c
                                                                    2005
                                                         ae3h
  year_last v_comment
       2016
2
       2016
                   NA
3
       2016
                   NA
```

3 Data cleaning and formatting

Rawdata from 2005 to 2018 has been cleaned and formatted according to the requirements of the COAT dataportal by Eeva Soininen and Hanna Boehner.

From 2019 onwards, rawdata is cleaned and formatted in two steps.

1. Data cleaning: Rawdata (individual data and trapping dates) are entered in excel-templates and are quality checked using the script 01_check_raw_data_snap_trapping_intesive_COAT.R . The script checks that all entries are correct (e.g. site_id, species names), possible errors are corrected in the script and the cleaned data is saved as a txt-file.

In particular the script checks for:

- Correct spelling of all variables.
- t_date: Correct format (yyyy-mm-dd), dates in other formats are reformatted.
- v embryos and v weight: Outliers in number of embryos and weight.
- v_observer: Correct format (initials and lowercase letters), observer is reformatted e.g. if full names were used.
- **v_comment:** Correct spelling and format (lowercase letters and english). Comments are edited or translated if necessary.
- Matching number of observations in the different columns.
- The columns are renamed according to the the requirements of the COAT dataportal.

All corrections that go beyond simple typing mistakes and lead to differences between rawdata and cleaned data are double-checked in the fieldbooks and a comment is added.

2. Data formatting: The script

02_produce_reformatted_snap_trapping_data_files_individual_plotbased_trapstatus_COAT.R is used to produce the three datasets that are derived from snaptrapping data. The files will be formatted according to the requirements of the COAT data portal, will be saved as ;-separated txt-files and will be uploaded to the COAT data portal:

- V_rodents_snaptrapping_individuals_intensive_YEAR.txt
- V_rodents_snaptrapping_abundance_intensive_YEAR.txt
- V_rodents_snaptrapping_trapstatus_intensive_YEAR.txt

^{*} year last is NA if the site is still included in the study design