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GENERAL INFORMATION

1. Data from 'Evolution of thermal physiology alters projected distributions of threespine stickleback under climate change'

2. Author information
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3. Date of data collection: Sept 2014 – July 2018

4. Geographic location of data collection: Bamfield, BC and Calgary, AB, CANADA

DATA & FILE OVERVIEW

1. File list
    a. dryad_f1.csv
    b. dryad_f2.csv
    c. dryad_wild.csv

METHODOLOGICAL INFORMATION

1. Description of methods used for collection/generation of data
    Adult threespine stickleback were run through a standard critical thermal minimum (CTmin) experiment, allowed to recover for at least three days, run through a standard critical thermal maximum (CTmax) experiment, and allowed to recover for an additional three days before being placed in a Shuttlebox holding a static temperature gradient. Mass measurements were taken at time of tagging and sex was determined by polymerase chain reaction (PCR) for the sex-specific *idh* locus.
2. Instrument- or software-specific information needed to interpret data
    Specific packages/libraries for data processing can be found in the manuscript methods section
3. People involved with sample collection, processing, analysis, or submission
    a. Sample collection:
        i. Sara JS Wuitchik
        ii. Tegan N Barry
        iii. Jori Harrison
    b. Sample processing:
        i. Sara JS Wuitchik
        ii. Tegan N Barry
        iii. Antoine Paccard
    c. Sample analysis:
        i. Sara JS Wuitchik
    d. Submission:
        i. Sara JS Wuitchik

DATA-SPECIFIC INFORMATION FOR DRYAD_F1.CSV

1. Number of variables: 10
2. Number of rows: 125 (including header)
3. Variable list
    a. Habitat: which marine or freshwater habitat the individual is from (M, marine, or FW, freshwater)
    b. Source Pop: which population the individual is from
    c. Indiv: the individual identifier by cross and number
    d. Tag: the unique visible implant elastomer tag each individual was tattooed with

        i. Either R (right) or L (left)
        ii. Either D (dorsal) or C (caudal)
        iii. Either S (single tag) or D (double tag)
        iv. Pk (pink), R (red), B (blue), G (green), O (orange), W (white), Y (yellow)
    e. ctmin: the CTmin value at loss of equilibrium (LOE) (in °C)
    f. ctmax: the CTmax value at LOE (in °C)
    g. pref: the average preferred ambient temperature from the Shuttlebox experiment (in °C)
    h. coretemp: the average estimated core body temperature from the Shuttlebox experiment (in °C)
    i. mass: mass at time of tagging (in g)
    j. sex: genetic sex determination (M, male, or F, female)

4. Missing data codes
    a. Empty cells mean the individual did not complete that experiment

DATA-SPECIFIC INFORMATION FOR DRYAD_F2.CSV

1. Number of variables: 8
2. Number of rows: 104
3. Variable list:
    a. Cross: source populations of cross
    b. Indiv: the individual identifier by cross and number
    c. Tag: the unique visible implant elastomer tag each individual was tattooed with
        i. Either R (right) or L (left)
        ii. Either D (dorsal) or C (caudal)
        iii. Either S (single tag) or D (double tag)
        iv. Pk (pink), R (red), B (blue), G (green), O (orange), Y (yellow), Br (brown), Pu (purple)
    d. ctmin: the CTmin value at loss of equilibrium (LOE) (in °C)
    e. ctmax: the CTmax value at LOE (in °C)
    f. pref: the average preferred ambient temperature from the Shuttlebox experiment (in °C)
    g. coretemp: the average estimated core body temperature from the Shuttlebox experiment (in °C)
    h. mass: mass at time of tagging (in g)
4. Missing data codes:
    a. Empty cells mean the individual did not complete that experiment

DATA-SPECIFIC INFORMATION FOR DRYAD_WILD.CSV

1. Number of variables
2. Number of rows
3. Variable list:

a. habitat: which marine or freshwater habitat the individual is from (M, marine, or FW, freshwater)
b. pop: which population the individual is from
c. Individual: the individual identifier by cross and number
d. Tag: the unique visible implant elastomer tag each individual was tattooed with
    i. Either R (right) or L (left)
    ii. Either D (dorsal) or C (caudal)
    iii. Either S (single tag) or D (double tag)
    iv. Pk (pink), R (red), B (blue), G (green), O (orange), W (white), Y (yellow)
e. ctmin: the CTmin value at loss of equilibrium (LOE) (in °C)
f. ctmax: the CTmax value at LOE (in °C)
g. pref: the average preferred ambient temperature from the Shuttlebox experiment (in °C)
h. coretemp: the average estimated core body temperature from the Shuttlebox experiment (in °C)
i. mass: mass at time of tagging (in g)
j. sex: genetic sex determination (M, male, or F, female)

4. Missing data codes:
    a. Empty cells mean the individual did not complete that experiment