

Descriptions of Data for “RanavirusTransmissionDataset.csv”

Dataset used in:

Le Sage, M.J., B. D. Towey, J. L. Brunner. 2019. Do scavengers prevent or promote disease transmission? The effect of invertebrate scavenging on *Ranavirus* transmission. *Functional Ecology*, in review.

The dataset tracks each naïve salamander larva in the transmission experiment described in the article above, during the exposure to infected carcasses.

| Column Title | Explanation |
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| Treatment | the treatment for the sample (AS, ANS, NAS, NANS) AS=Access-Scavenger, ANS=Access-No Scavenger, NAS=No Access-Scavenger, NANS=No Access-No Scavenger |
| Replicate | The replicate number for the sample (1-10 for each treatment) |
| ID | The unique ID number given to each salamander larva in a replicate (1-9 for each replicate) |
| LarvaUniqueID | The full Unique Identifier for each salamander larva "Treatment - Replicate# - ID#" |
| BeginDate | The date on which the replicate was begun (for scavenger treatments [S], this is the day that scavenging began; for non scavenger treatments [NS], this is the day that the carcass was introduced into the water) |
| Scav | The family of the scavenger used in this experiment (all were Dytiscidae - larva) |
| ScavSize(mm) | The size of the scavenger used for this replicate in millimeters |
| ScavOrigin | The pond from which the scavenger used in this replicate was collected |
| CarcassSVL(mm) | The Snout-Vent length of the carcass used in this replicate in millimeters |
| CarcassWeight(g) | The weight of the carcass used in this replicate in grams |
| CarcassID | The Unique ID number of the carcass used in this replicate |
| CarcassFV3ExposureTitre(pfu/mL) | The pfu/mL of FV3 that was used in the exposure of the carcass (all were done by injection) |
| ScavDuration(hr) | The duration of scavenging for this replicate (in hours) |
| ExposeDate | The date on which the nine uninfected salamander larvae were introduced to the tank for this replicate |
| EndExposeDate | The date on which the nine uninfected salamander larvae were removed from the tank for this replicate, and then housed separately |

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| ExposeDuration(hr) | The duration of uninfected larva exposure to the conditions in this replicate (in hours) |
| EuthDate | The date on which this larva was euthanized or was found dead in its separate container |
| PostExposeDuration(days) | The duration (in days) that the larva survived post exposure (maximum 15) |
| DissectionDate | The date on which the larva was dissected and the tissue was removed for analysis |
| TissueTubeID | The unique ID on the tissue storage tube for this larva "TreatmentID (A,B,C,D) - Rep# - ID#" |
| SVL(mm) | Snout vent length of the larva in mm |
| weight(g) | weight of the larva in grams |
| Survival(days) | How many days post-exposure to carcass the larva survived (anything < 15d means it died prematurely) |
| UninfectedLarvaComments | comments about the uninfected larva |
| LarvaSymptoms | outward symptoms noticed on the larva before mortality |
| ExtractionDate | the date that the larval tissue was extracted |
| ExtractionType | the type of extraction that was performed on the larval tissue |
| Tissue Type | the type of tissue that was extracted |
| DNAName | The unique ID given to the DNA sample extracted from this larva's tissue |
| ExtractionNeg | The unique ID of the extraction negative used for this larva |
| PCRDate | The date that qPCR was run on this DNA sample |
| PCRName | The name of the qPCR run that contains this sample |
| ExtractionNeg Value | The value 0 / 1 (neg / pos) of the extraction negative's qPCR run |
| PCRNeg Value | The value 0 / 1 (neg / pos) of the PCR plate negative's qPCR run |
| PCRVal1 | The value of the first well replicate of the qPCR run for this sample |
| PCRVal2 | The value of the second well replicate of the qPCR run for this sample |
| PCRVal3 | The value of the third well replicate of the qPCR run for this sample |
| PCRMeanVal | The mean value of all 3 qPCR runs for this sample |
| PCRCall | The call (0 = neg, 1 = pos, or "Rerun" = inconclusive) for this sample |
| PCRRerunDate | The date that qPCR was run on this DNA sample (if a Rerun is needed) |
| PCRRerunName | The name of the qPCR run that contains this sample (if a Rerun is needed) |
| PCRRerunComments | Comments about the qPCR run (if a Rerun is needed) |
| PCRRerunNegValue | The value 0 / 1 (neg / pos) of the PCR plate negative's qPCR run (if a Rerun is needed) |
| PCRRerunVal1 | The value of the first well replicate of the qPCR run for this sample (if a Rerun is needed) |

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| PCRRerunVal2 | The value of the second well replicate of the qPCR run for this sample (if a Rerun is needed) |
| PCRRerunVal3 | The value of the third well replicate of the qPCR run for this sample (if a Rerun is needed) |
| PCRRerunMeanVal | The mean value of all 3 qPCR runs for this sample (if a Rerun is needed) |
| PCRRerunCall | The call (0 = neg, 1 = pos, or "Rerun" = inconclusive) for this sample (if a Rerun is needed) |
| PCRFinalMeanVal | The mean value of qPCR runs for this sample |
| PCRFinalCall | The overall call (0 = neg, 1 = pos, or "Rerun" = inconclusive) for this sample |