**Description of data fields (columns) in the frog susceptibility dataset: Knapp\_lab\_experiment\_data.csv**

**frogid**: unique identifier for each frog, consisting of tank\_frognum

**tank**: number of tank (1 through 16)

**frognum**: number assigned to first or second frog in each tank (1 or 2)

**day**: number of days since the start of the experiment (i.e. number of days since the frogs were exposed to Bd and became infected)

**week**: number of weeks since the start of the experiment

**ZE**: number of Bd zoospore equivalents on a skin swab (measured by qPCR)

**weight**: weight of the frog in grams

**svl**: snout-to-vent length of the frog in millimeters

**sex**: sex of frog (M or F)

**frogsource**: name of population from which the frog was collected

**bdsource**: name of population from which Bd was collected

**frogpersistent**: frog from persistent population (1), frog from naïve population (0)

**bdpersistent**: Bd from persistent population (1), Bd from die-off population (0)

**logZE1:** log of Zoospore Equivalents + 1 (log10(ZE+1))