Model

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I think it makes sense that we use a model that's similar to vector-borne disease models because trematodes have two hosts: snails and ducks. Ignoring spatial features and sexuality, we're going to use the following model:

$$X'_{S} = b - \beta_{X} X_{S} Y_{I} / N_{X} - \mu_{X} X_{S} - a X_{S} Y / N_{X}$$

$$X'_{I} = \beta_{X} X_{S} Y_{I} / N_{X} - (\mu_{X} + \alpha) X_{S} - a X_{I} Y / N_{X}$$

$$Y'_{S} = \mu_{Y} N_{Y} - \beta_{Y} Y_{S} X_{I} / N_{Y} - \mu_{Y} Y_{S}$$

$$Y'_{I} = \beta_{Y} Y_{S} X_{I} / N_{Y} - \mu_{Y} Y_{S}$$

Here, X and Y represent snail and duck population and subscripts represent the infection status. We keep the size of the duck population constant for simplicity. Snail population is hunted by the duck population and suffer from natural mortality.

Now, let's separate asexual and sexual reproduction. Sexual snails are diploids and asexual snails are triploid clones but we assume diploidy for both for simplicity. Following Ashby and Gupta (2014); Ashby and King (2015), we let birth rates for asexual and sexual population be

$$b_{ij}^{A} = r(X_{ij,S}^{A} + f \sum_{j} X_{ij,I}^{A})(1 - h_b N)$$

$$b_{ij}^{S} = c_b r \frac{(2 - \delta_{ij})X_i^S X_j^S}{2\sum_k X_k^S} (1 - h_b N),$$

where X_i is the number of gamete i in sexual population following recombination. We write it as $X_i = u + fv$, following Ashby and Gupta (2014). As Gibson et al. (2017) showed that there the snail system is consistent with the two fold cost of sex, we let $c_b = 1$.

We want to model outcrossing and recombination of parasies. We can define P_{ij} be number of parasite genotypes in ducks after sexual recombination. Simply, we can write

$$P_{ij} = \frac{(2 - \delta_{ij})Y_i^S Y_j^S}{\sum_k Y_k^S}$$

Then, our model becomes:

$$X_{ij,S}' = b_{ij} - \beta_X X_{ij,S} Z_{ij} / N_X - \mu_X X_{ij,S} - a X_{ij,S} Y / N_X$$

$$X_{ij,I}' = \beta_X X_{ij,S} Z_{ij} / N_X - (\mu_X + \alpha) X_{ij,S} - a X_{ij,S} Y / N_X$$

$$Y_S' = \mu_Y N_Y - \beta_Y Y_S \left(\sum_{i,j} X_{ij,I} \right) - \mu_Y Y_S$$

$$Y_{ij,I}' = \beta_Y Y_S X_{ij,I} - \mu_Y Y_{ij,I}$$

References

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