

Aims

- Understand what deterministic and stochastic modelling are
- Be able to build stochastic models based on given parameters
- Appreciate the added complexity of drawing from random distributions rather than applying rates and averages
- Be able to run stochastic simulations on a cluster
- Be able to process and visualise stochastic simulations
- Appreciate where deterministic and stochastic modelling substantially differ

Population modelling

- 1. Population modelling overview and stochastic vs deterministic
- 2. Demographic population models and matrix population models
- 3. Stochastic adaptation

Population modelling

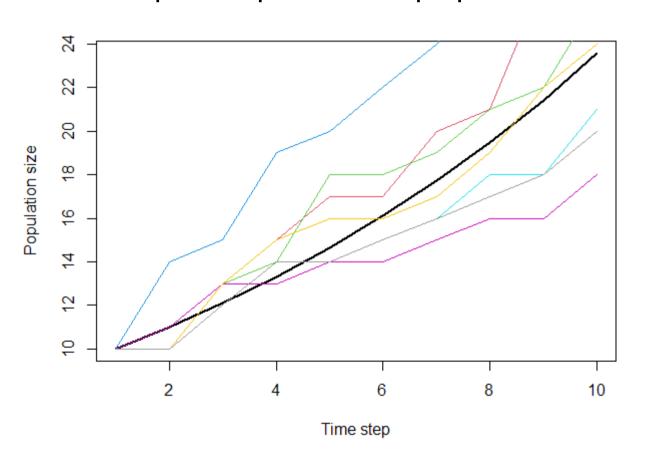
- 1. Population modelling overview and stochastic vs deterministic
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What is a population model?

- Models how a population (or populations) changes over time
- We might be interested in...
 - The effect of different parameters on the population dynamics
 - Identifying an appropriate control or conservation strategy
 - The likelihood of extinction or recovery
- We can make different assumptions:
 - Homogeneous every individual is the same
 - Heterogeneous individuals can be different
 - Compartmental there are different compartments and every individual in the same compartment is the same
 - Individual-based we explicitly model every single individual and all their traits

Stochastic vs deterministic modelling

- Deterministic = determined
 - Assumes that outcomes follow determined paths
 - Can be simulated once to observe modelled behaviour
- Stochastic = random
 - Assumes that outcomes are drawn from random distributions
 - Must be simulated many times to observe modelled behaviour, since every simulation can be different
 - Typically closer to reality as life is inherently random!



Deterministic: (black line)

Every time step, the bacterial population multiplies by 1.1

$$p_{t+1} = 1.1p_t$$

Stochastic: (coloured lines)

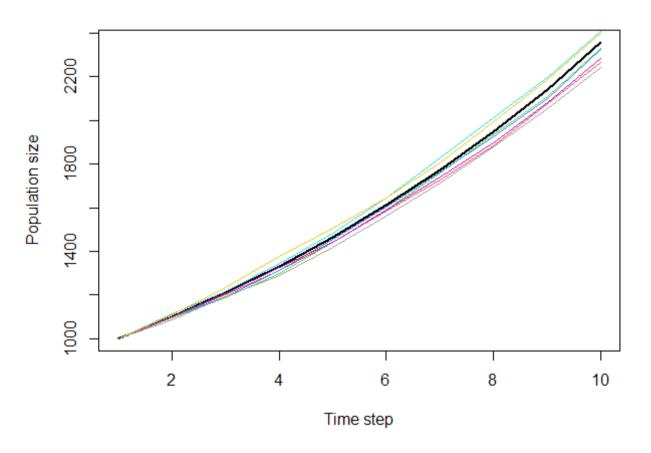
Every time step, there is a 10% chance of each individual bacterium reproducing

$$p_{t+1} = p_t + N_{t+1}$$

$$N_{t+1} \sim \text{Bin}(p_t, 0.1)$$

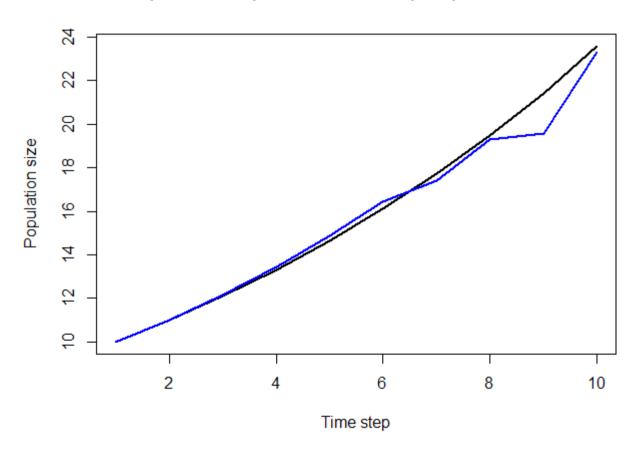
model_type	simulation_number	time_0	$time_1$	time_2	time_3	time_4	time_5	time_6	time_7	time_8	time_9
deterministic	<na></na>	10	11	12.1	13.31	14.64	16.11	17.72	19.49	21.44	23.58
stochastic	1	10	12	13	14	17	19	20	22	24	26
stochastic	2	10	11	11	14	15	15	16	20	24	29
stochastic	3	10	11	12	15	16	19	20	22	25	27
stochastic	4	10	11	13	14	16	19	20	22	24	26
stochastic	5	10	11	13	13	15	17	18	19	20	22
stochastic	6	10	11	12	12	12	12	12	12	12	14
stochastic	7	10	10	11	12	13	14	16	18	18	19

- Note that the deterministic simulation is based on averages and therefore results in continuous – not discrete – outputs
- If we were modelling number of bacteria, this would be biologically unrealistic
- In this particular situation, the deterministic simulations reflects the average behaviour of the stochastic model



In some situations, deterministic and stochastic modelling may produce very similar outcomes – the deterministic model is a **good approximation** to the stochastic model.

E.g., in this example the initial population size was set at 1000 rather than the previous value of 10.



In some situations, the deterministic model reflects the **average behaviour** of the stochastic model.

E.g., the blue line in this plot shows the average population size at each time point across the 7 stochastic simulations (with initial population size of 10) (the black line is still the deterministic simulation)

Stochastic modelling

- Enables us to include realistic probability distributions (rather than assuming that everything follows the "average" outcome)
- Can enable us to handle discrete outcomes
 - E.g., whole number of organisms; not 16.11 bacteria
- Most important in situations where:
 - Stochastic events can make a substantial difference
 - Future changes are sensitive to the current value
 - We want some certainty bounds around our results
- Should be run many times in order to get an overall sense of model behaviour this is where cluster computing can be useful
- Can present a trade-off: biological realism vs computational effort

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Demographic population models

- A demographic population model describes how a population changes in terms of, and in response to, a demographic structure in the population.
- The demographic structure would typically be a set of traits which affect life history processes (i.e., birth and death).
- A simple demographic trait to structure a population model is age
- Other traits could include body weight, sex, or sexual maturity
- Demographic population models can also incorporate evolutionary processes ("evolutionary demography") which have been used to explore evolution of life history "strategies"

Matrix population models

- The term matrix population model usually refers to an age- or stage-structured matrix population model
- In this format, a population is divided into discrete life stages or age classes
- Each stage/class has a set of parameters to describe demographic processes (i.e., survival, maturation, and recruitment)
- The population is represented by a state vector containing the number of individuals in each stage/class
- The parameters are put into a **projection matrix** which can be applied to the state vector to project the population forwards in time
- Matrix population models are typically deterministic

Matrix population models

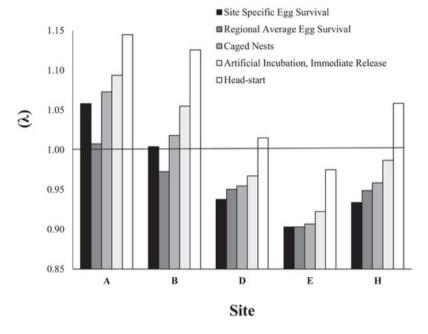
- Allows us to model population dynamics, which are hard to observe, based on individual-level processes, which are easier to observe
- The parameters can be inferred by field studies such as capture-mark-recapture
- Online database of matrices, so there is field-based evidence to support the modelling of many different populations



- Version 6.22.5.0
- 760 Taxonomic Species
- 643 Publications
- 8851 Matrix Population Models



- Version 4.21.8.0
- 415 Taxonomic Species
- 395 Publications
- 3317 Matrix Population Models

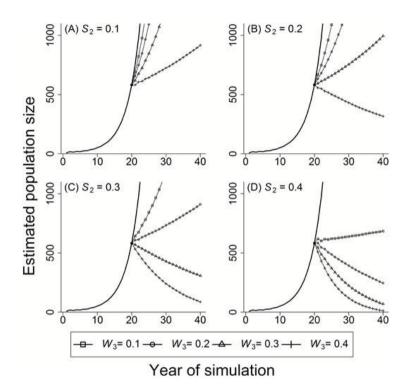


Bog turtle conservation in North Carolina, USA.

Knoerr, M.D., Tutterow, A.M., Graeter, G.J., Pittman, S.E. and Barrett, K. (2022), Population models reveal the importance of early life-stages for population stability of an imperiled turtle species. Anim. Conserv., 25: 53-

64. https://doi.org/10.1111/acv.12718

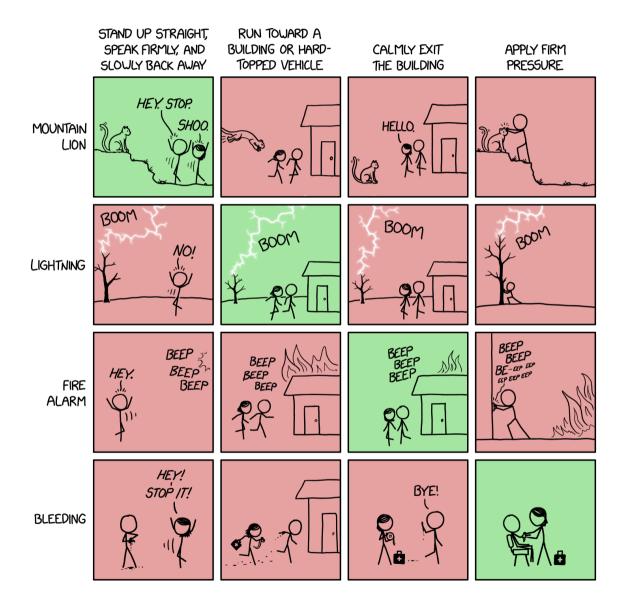




Control of the invasive Chinese mitten crab in North America and Europe.

Zhang, Z., Yokota, M., Strüssmann, C.A. (2019), A periodic matrix population model to predict growth potential of the invasive Chinese mitten crab *Eriocheir sinensis* (H. Milne Edwards, 1853) (Decapoda: Brachyura: Varunidae), *J. Crustac. Biol.*, 39 (1):28–35. https://doi.org/10.1093/jcbiol/ruy090





Matrix multiplication

$$A.B = ?$$

$$\begin{pmatrix} a_{11} & a_{12} & a_{13} \\ a_{21} & a_{22} & a_{23} \\ a_{31} & a_{32} & a_{34} \end{pmatrix} \begin{pmatrix} b_{11} & b_{12} \\ b_{21} & b_{22} \\ b_{31} & b_{32} \end{pmatrix} = \begin{pmatrix} ? & ? \\ ? & ? \\ ? & ? \end{pmatrix}$$

Matrix multiplication

$$A.B = ?$$

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$$a_{21} b_{12} + a_{22} b_{22} + a_{23} b_{32}$$

Matrix multiplication

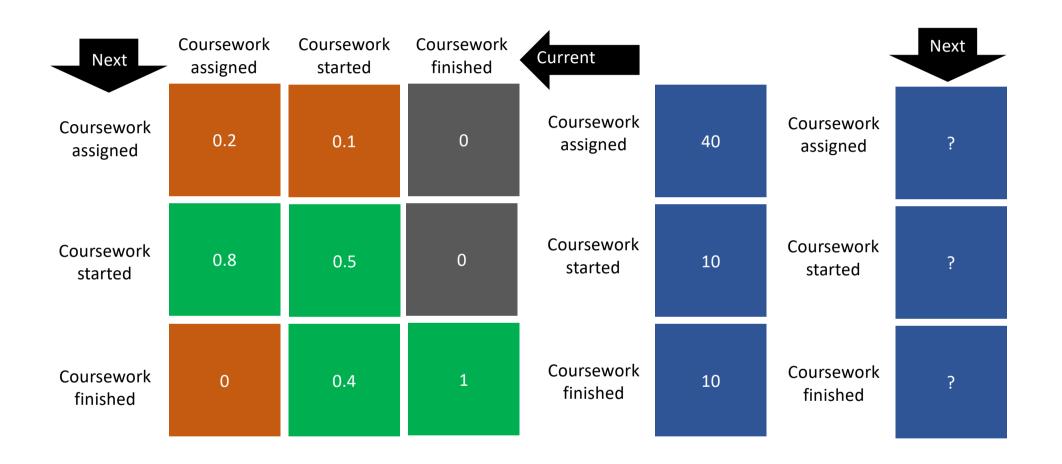
$$A.B = ?$$

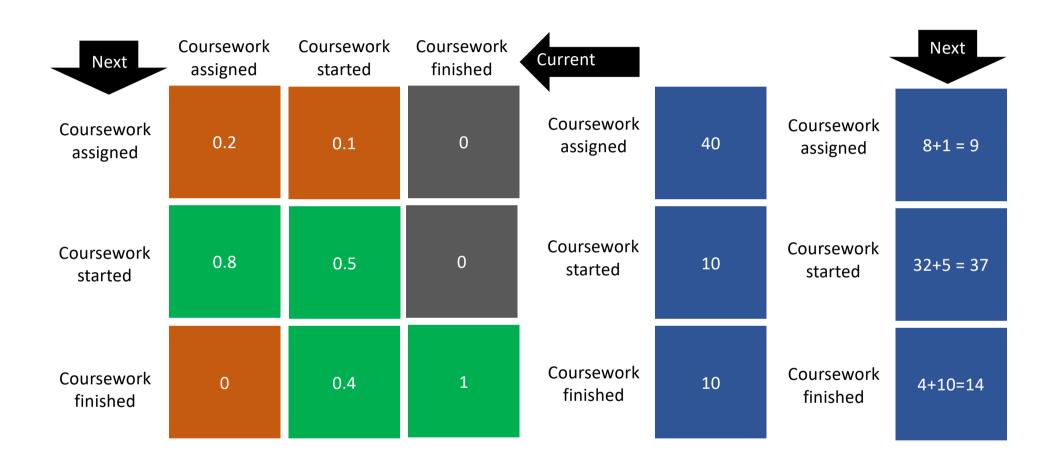
$$\begin{pmatrix} a_{11} & a_{12} & a_{13} \\ a_{21} & a_{22} & a_{23} \\ a_{31} & a_{32} & a_{34} \end{pmatrix} \begin{pmatrix} b_{11} \\ b_{21} \\ b_{31} \end{pmatrix} = \begin{pmatrix} ? \\ ? \\ ? \end{pmatrix}$$

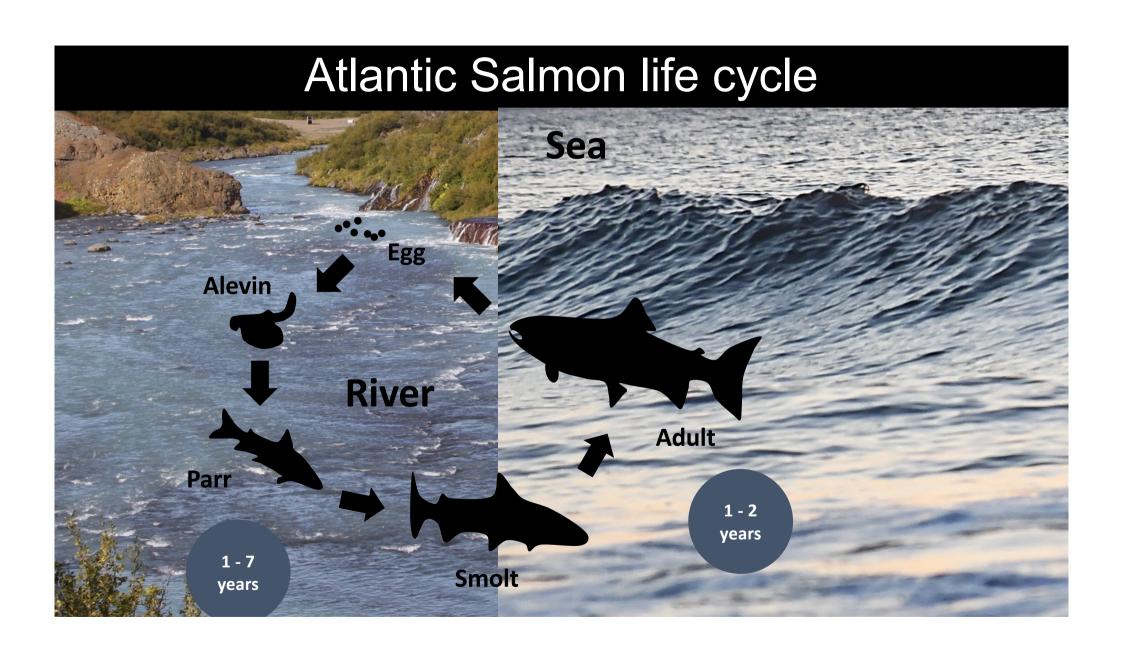
$$a_{21} b_{11} + a_{22} b_{21} + a_{23} b_{31}$$

Next state	Coursework assigned	Coursework started	Coursework finished	Current state
Coursework assigned	Laziness or too busy	Data loss disaster without a backup	Impossible	
Coursework started	Do work	Keep working but not done yet	Impossible	
Coursework finished	Cheating	Do work	Celebrate! it's all done	

Next state	Coursework assigned	Coursework started	Coursework finished	Current state
Coursework assigned	Laziness or too busy 0.2	Data loss disaster without a backup 0.1	Impossible 0	
Coursework started	Do work 0.8	Keep working but not done yet 0.5	Impossible 0	
Coursework finished	Cheating 0	Do work 0.4	Celebrate! it's all done 1	







Atlantic Salmon toy matrix population model

Atlantic Salmon toy matrix population model

Matrix population models

- In a **deterministic** application of a matrix population model, we assume that population changes will follow the "average" (which are the parameters in the projection matrix)
- We can compute these changes by multiplying the **state vector** by the **projection matrix** using matrix multiplication

Matrix population models

- The matrix formulation is helpful because there are some mathematical shortcuts we can use:
 - The population growth rate is the **eigenvalue** of the matrix
 - The eigenvector represents the eventual proportions of individuals in each stage
- Limitations:
 - random outcomes such as random extinction cannot occur
 - technically the population state can never reach zero (it can only get closer and closer to zero) so we would have to define a threshold for extinction if we were modelling this

Matrix population models – no extinction?

The projection matrix used here is making the population smaller and smaller, but the population size actually never reaches zero.

8.97725984

5.02864527

2.81678401

1.57781332

0.49506226

0.27730769

0.15533310

0.08700939

0.04873806

8.34983123

4.67723943

2.61993959

1.46755160

0.46046601

0.25792871

0.14447802

0.08092895

0.04533211

population_size_time_series

1.82381210

0.57224801

0.32054307

0.17955127

0.10057513

0.05633687

5.40646068

3.02841747

1.69635935

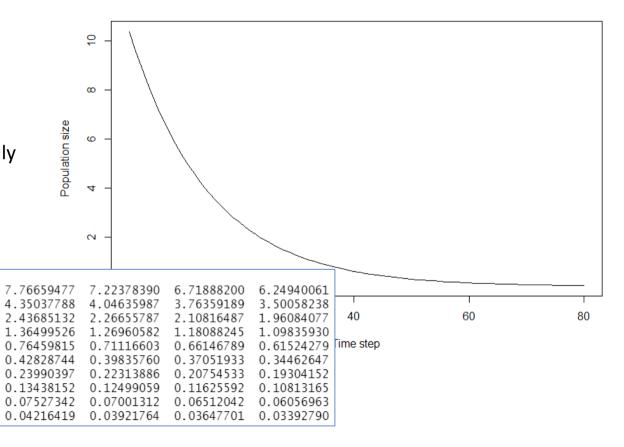
0.53225783

0.29814268

0.16700376

0.09354667

0.05239990

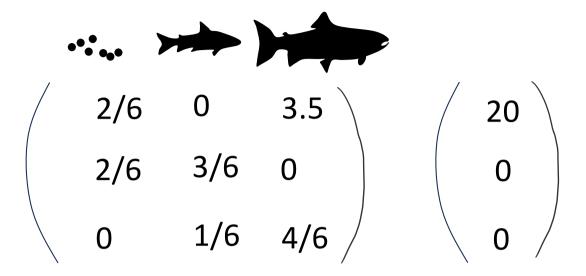


Population modelling

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Stochastic version

- As discussed already, deterministic models do not always reflect reality
- We will carry out the same kind of age-structured population model except this time we will not assume determinism – processes will be stochastic
- This will allow us to compare the deterministic and stochastic equivalents
- How can we convert biological parameters inferred from field studies into a stochastic formulation?
- Think about every process involved/assumed in the model and what probability distribution it is drawn from



Keeper of the eggs

Roll your dice

1 or 2 means they die (return to bag)

3 or 4 means you keep them

5 or 6 means put aside to mature you give these dice to the keeper of juveniles at the end of this turn

Keeper of the juveniles

Roll your dice

1 or 2 means they die (return to bag)

3 - 5 means you keep them

6 means put aside to mature you give these dice to the keeper of adults at the end of this turn

Keeper of the adults

Roll your dice

sum the numbers of all your dice (whether they die or not) and give this number of dice (from the bag) to the keeper of eggs at the end of this turn

Also ...

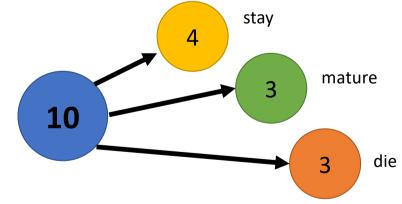
1 or 2 means they die (return to bag)

Making a deterministic matrix population model stochastic: Survival and maturation

Suppose there are 10 individuals in stage 2. The projection matrix tells us that at the next time step **we expect** a proportion 0.4 to stay in stage 2, and **we expect** a proportion 0.3 to have matured into stage 3. (We expect the remaining proportion 0.3 to have died.)

$$\begin{pmatrix} 0.1 & 0 & 3.5 \\ 0.5 & 0.4 & 0 \\ 0 & 0.3 & 0.6 \end{pmatrix}$$

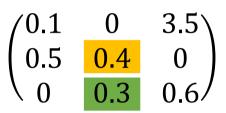


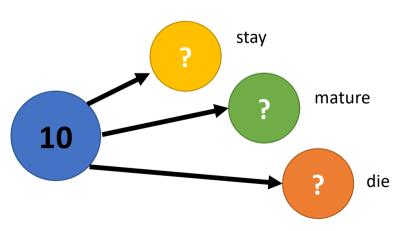


Making a deterministic matrix population model stochastic: Survival and maturation

Suppose there are 10 individuals in stage 2. The projection matrix tells us that at the next time step **we expect** a proportion 0.4 to still be in stage 2, and **we expect** a proportion 0.3 to have matured into stage 3. (We expect the remaining proportion 0.3 to have died.)

In a stochastic formulation, we consider the **probability** of each of these three events (staying in the same life stage, maturing into the next life stage, and death) for every individual.



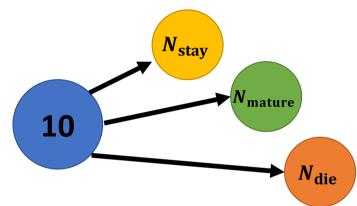


Making a deterministic matrix population model stochastic: Survival and maturation

For each individual in stage 2,

$$P(\text{stay in stage 2}) = 0.4$$

 $P(\text{mature into stage 3}) = 0.3$
 $P(\text{die}) = 0.3$



So if there were 10 individuals in stage 2, then we could draw the outcomes as

$$(N_{\text{stay}}, N_{\text{mature}}, N_{\text{die}}) = \text{Multinomial}(n = 10, p = (0.4, 0.3, 0.3))$$

• When looking at **life stage 1**, we need to find out $N_{\rm stay}$ and $N_{\rm mature}$ for this stage. These will correspond to stage 1 and stage 2 respectively. Suppose that for life stage 1, $N_{\rm stay}=5$ and $N_{\rm mature}=27$.

$\sqrt{0.1}$	0	3.5
0.5	0.4	0
/ 0	0.3	0.6/

• When looking at **life stage 1**, we need to find out $N_{\rm stay}$ and $N_{\rm mature}$ for this stage. These will correspond to stage 1 and stage 2 respectively. Suppose that for life stage 1, $N_{\rm stay}=5$ and $N_{\rm mature}=27$.

 $\begin{pmatrix} 0.1 & 0 & 3.5 \\ 0.5 & 0.4 & 0 \\ 0 & 0.3 & 0.6 \end{pmatrix}$

• We need to do the same for **life stage 2**. Suppose that $N_{\text{stay}} = 10$ and $N_{\text{mature}} = 8$.

- When looking at **life stage 1**, we need to find out $N_{\rm stay}$ and $N_{\rm mature}$ for this stage. These will correspond to stage 1 and stage 2 respectively. Suppose that for life stage 1, $N_{\rm stay} = 5$ and $N_{\rm mature} = 27$.
- We need to do the same for **life stage 2**. Suppose that $N_{\text{stay}} = 10$ and $N_{\text{mature}} = 8$.
- For **life stage 3**, individuals can only survive or die (not mature). Suppose that $N_{\text{stay}} = 14$

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- We need to do the same for **life stage 2**. Suppose that $N_{\rm stay}=10$ and $N_{\rm mature}=8$.
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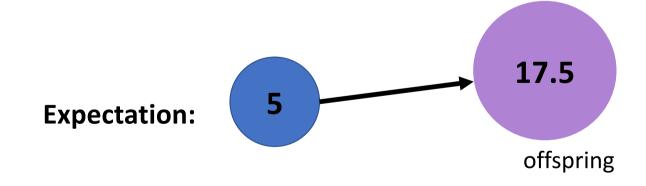
Then to obtain the population state at the next time step, we need to add these together in the correct entries of the life state vector:

$$\begin{pmatrix}
5 \\
27 + 10 \\
8 + 14
\end{pmatrix}$$

But this doesn't include recruitment – we ignored the 3.5 entry of the vector since this did not reflect survival or maturation.

Suppose there are 5 individuals in stage 3. The projection matrix tells us that at the next time step **we expect** each individual to have recruiting 3.5 new offspring, which will be in stage 1.

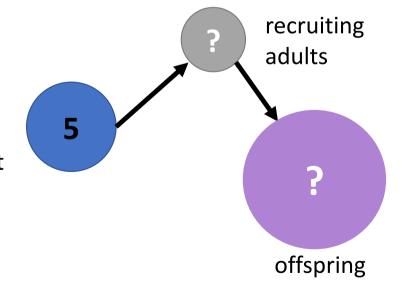
$$\begin{pmatrix} 0.1 & 0 & 3.5 \\ 0.5 & 0.4 & 0 \\ 0 & 0.3 & 0.6 \end{pmatrix}$$



Suppose there are 5 individuals in stage 2. The projection matrix tells us that at the next time step **we expect** each individual to have recruiting 3.5 new offspring.

$$\begin{pmatrix} 0.1 & 0 & 3.5 \\ 0.5 & 0.4 & 0 \\ 0 & 0.3 & 0.6 \end{pmatrix}$$

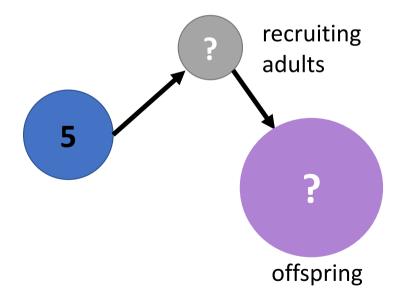
In a stochastic formulation, we consider the **probability** of each individual recruiting (either they recruit or they don't) and also the **probability** of the litter/clutch size taking different values.



We have a problem here because the **probability** of a given adult recruiting and the **probability** of each clutch/litter size taking different values have been conflated into one value, 3.5, which is the "recruitment rate".

We need to separately know the recruitment probability and the probability distribution for clutch/litter size.

$$\begin{pmatrix} 0.1 & 0 & 3.5 \\ 0.5 & 0.4 & 0 \\ 0 & 0.3 & 0.6 \end{pmatrix}$$



Suppose from field studies we also knew the probability distribution for clutch/litter sizes looked like this:

(0.06, 0.08, 0.13, 0.15, 0.16, 0.18, 0.15, 0.06, 0.03)

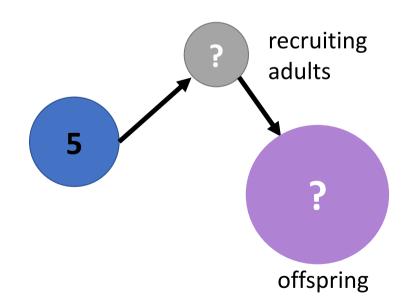
$$\begin{pmatrix} 0.1 & 0 & 3.5 \\ 0.5 & 0.4 & 0 \\ 0 & 0.3 & 0.6 \end{pmatrix}$$

Then we could compute the **expected clutch/litter size** using the probability distribution as

$$E(\text{clutch size}) = \sum_{n} nP(\text{clutch size} = n)$$

And we can then compute the individual **recruitment probability** as

$$P(\text{recruit}) = \frac{3.5}{E(\text{clutch size})}$$



Suppose that the expected clutch size was

$$E(\text{clutch size}) = 7$$
 so $P(\text{recruit}) = 0.5$

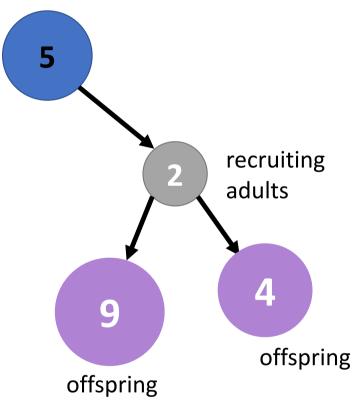
Using these probabilities, we can now draw the number of individuals which recruit:

$$N_{\text{recruit}} = \text{Bin}(n = 5, p = 0.5)$$

 \rightarrow Suppose the number drawn was $N_{\text{recruit}} = 2$

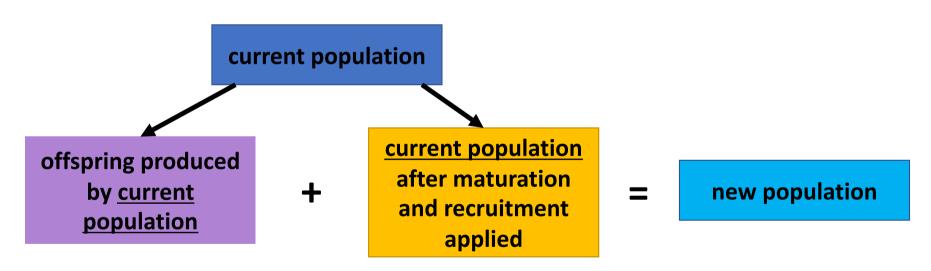
For each clutch, we could then draw the clutch size based on our clutch size distribution.

Supposing the numbers drawn were 9 and 4 for each clutch respectively, then the total number of offspring would be 13.



Making a deterministic matrix population model stochastic: Applying processes in turn

One of the challenges with applying a stochastic simulation is that different processes have to be applied one-by-one, which seems counterintuitive since in a deterministic model everything happens at the same time.



The resulting model – a stochastic demographic population model

Stochastic because processes are drawn from random distributions

Demographic because the population is structured by life stage (a demographic trait)

Pros:

- In general, it is a better reflection of reality than the deterministic model (as long as we run it many times to observe outcomes)
- We can observe random extinction and the population can reach zero, so we can compute extinction probability

Cons:

- It needs to be run many times to see a representative sample of outcomes
- It was more awkward to code than the deterministic matrix population model
- It required additional data to simulate correctly (clutch size distribution)