

Population: a set of individuals of the same species co-occurring in space and time.

Characteristics of populations

Total number of individuals or population 'density' Age/Stage/Size structure

Sex ratio

Spatial distribution

$$dN/dt = rN$$

$$N_t = N_0 e^{rt}$$

$$ln(2)/r = t_{double}$$

$$dN/dt = rN\left(1 - \frac{N}{K}\right)$$

$$dN/dt = rN\left(\frac{K-N}{K}\right)$$

$$N_{t} = N_{0} \frac{K}{(K - N_{0})e^{-rt} + N_{0}}$$

Four important assumptions

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- 1. Per capita growth rate is a linear function of *N*.
- 2. Population growth rate responds instantly to changes in N (no time lags).
- 3. The external environment has no influence on population growth.
- 4. All individuals are equal (i.e., no age or size effects).

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These models work for 'simple' organisms like single-celled bacteria or protozoa.



But for most plants and animals, birth and death rates depe on age.



Northern leopard frog, Rana pipiens



A newborn elephant cannot reproduce immediately, but instead must wait ~10 years.





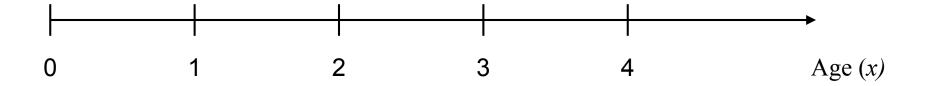
Hatchlings and seedlings often have higher mortality rates than adults.



A population that consisted of only old, post-reproductive chimpanzees will soon go extinct.



A population of only tadpoles cannot begin to grow until tadpoles become adults and can reproduce.







When using age classes, we treat all individuals in the same class as equivalents.

The fecundity schedule: the average number of female offspring born per unit time to an individual female of a particular age.

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$$b(6) = 3$$
 — A female of age 6 has, on average, 3 female offspring

x	S(x)	b(x)	l(x) = S(x)/S(0)	g(x) = l(x+1)/ $l(x)$	l(x)b(x)	l(x)b(x)x	Initial estimate $e^{-rx}l(x)b(x)$
0	500	0					
1	400	2					
2	200	3					
3	50	1					
4	0	0					

The survivorship schedule

x	S(x)	b(x)	l(x) = S(x)/S(0)	g(x) $= l(x+1)/$ $l(x)$	l(x)b(x)	l(x)b(x)x	Initial estimate $e^{-rx}l(x)b(x)$
0	500	0					
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The proportion of the original cohort that survives to the start of that age class

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0	500	0					
1	400	2					
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x	S(x)	b(x)	l(x) = S(x)/S(0)	g(x) = l(x+1)/ $l(x)$	l(x)b(x)	l(x)b(x)x	Initial estimate $e^{-rx}l(x)b(x)$
0	500	0	1				
1	400	2	0.8				
2	200	3	0.4				
3	50	1	0.1				
4	0	0	0				

The proportion of the original cohort that survives to the start of that age class The probability of surviving to that age class

The survival probability

x	S(x)	b(x)	l(x) = S(x)/S(0)	g(x) = l(x+1)/ $l(x)$	l(x)b(x)	l(x)b(x)x	Initial estimate $e^{-rx}l(x)b(x)$
0	500	0	1				
1	400	2	0.8				
2	200	3	0.4				
3	50	1	0.1				
4	0	0	0				

The probability that an individual of age x survives to age x + 1

The survival probability

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0	500	0	1	0.80			
1	400	2	0.8	0.50			
2	200	3	0.4	0.25			
3	50	1	0.1	0			
4	0	0	0				

The probability that an individual of age x survives to age x + 1

x	S(x)	b(x)	l(x) = S(x)/S(0)	g(x) = l(x+1)/ $l(x)$	l(x)b(x)	l(x)b(x)x	Initial estimate $e^{-rx}l(x)b(x)$
0	500	0	1	0.80			
1	400	2	0.8	0.50			
2	200	3	0.4	0.25			
3	50	1	0.1	0			
4	0	0	0				

The mean number of female offspring produced per female over her lifetime.

x	S(x)	b(x)	l(x) = S(x)/S(0)	g(x) = l(x+1)/ $l(x)$	l(x)b(x)	l(x)b(x)x	Initial estimate $e^{-rx}l(x)b(x)$
0	500	0	1	0.80	0		
1	400	2	0.8	0.50	1.6		
2	200	3	0.4	0.25	1.2		
3	50	1	0.1	0	0.1		
4	0	0	0				

The mean number of female offspring produced per female over her lifetime. The reproductive potential adjusted by the mortality schedule.

x	S(x)	b(x)	l(x) = S(x)/S(0)	g(x) = l(x+1)/ $l(x)$	l(x)b(x)	l(x)b(x)x	Initial estimate $e^{-rx}l(x)b(x)$
0	500	0	1	0.80	0		
1	400	2	0.8	0.50	1.6		
2	200	3	0.4	0.25	1.2		
3	50	1	0.1	0	0.1		
4	0	0	0 R	$C_0 = \sum l(x)b(x)$	(x) = 2.9 off	fspring	

If $R_0 > 1$, there is a net surplus of offspring produced each generation, and the population increases exponentially.

x	S(x)	b(x)	l(x) = S(x)/S(0)	g(x) = l(x+1)/ $l(x)$	l(x)b(x)	l(x)b(x)x	Initial estimate $e^{-rx}l(x)b(x)$
0	500	0	1	0.80	0		
1	400	2	0.8	0.50	1.6		
2	200	3	0.4	0.25	1.2		
3	50	1	0.1	0	0.1		
4	0	0	0 R	$c_0 = \sum l(x)b(x)$	x) = 2.9 of	fspring	

If $R_0 < I$, the population declines

x	S(x)	b(x)	l(x) = S(x)/S(0)	g(x) = l(x+1)/ $l(x)$	l(x)b(x)	l(x)b(x)x	Initial estimate $e^{-rx}l(x)b(x)$
0	500	0	1	0.80	0		
1	400	2	0.8	0.50	1.6		
2	200	3	0.4	0.25	1.2		
3	50	1	0.1	0	0.1		
4	0	0	0 R	$R_0 = \sum l(x)b(x)$	x) = 2.9 off	fspring	

If $R_0 = I$, the population is stable.

Generation time, G

X	S(x)	b(x)	$l(x) = S(x)/S(\theta)$	g(x) = l(x+1)/ $l(x)$	l(x)b(x)	l(x)b(x)x	Initial estimate $e^{-rx}l(x)b(x)$
0	500	0	1	0.80	0		
1	400	2	0.8	0.50	1.6		
2	200	3	0.4	0.25	1.2		
3	50	1	0.1	0	0.1		
4	0	0	0				

Average age of the parents of all the offspring produced by a single cohort

Generation time, G

$$\frac{\sum l(x)b(x)x}{\sum l(x)b(x)}$$

x	S(x)	b(x)	$l(x) = S(x)/S(\theta)$	g(x) $= l(x+1)/$ $l(x)$	l(x)b(x)	l(x)b(x)x	Initial estimate $e^{-rx}l(x)b(x)$
0	500	0	1	0.80	0	0	
1	400	2	0.8	0.50	1.6	1.6	
2	200	3	0.4	0.25	1.2	2.4	
3	50	1	0.1	0	0.1	0.3	
4	0	0	0			$\Sigma = 4.3$	

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$$\frac{\sum l(x)b(x)x}{\sum l(x)b(x)}$$

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0	500	0	1	0.80	0	0	
1	400	2	8.0	0.50	1.6	1.6	
2	200	3	0.4	0.25	1.2	2.4	
3	50	1	0.1	0	0.1	0.3	
4	0	0	0	G = 4	.3/2.9 = 1.4	$\sum = 4.3$ 483 years	

Average age of the parents of all the offspring produced by a single cohort

$$rpproxrac{\ln(R_{ heta})}{G}$$

$$r \approx \frac{\ln(R_0)}{G}$$
 $r \approx \frac{\ln(2.9)}{1.483} \approx 0.718 \text{ individuals}$
per individual per year

$$R_0 = \sum l(x)b(x) = 2.9$$
 offspring $G = 4.3/2.9 = 1.483$ years

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As generation time increases, rate of increase decreases; as number of offspring per female increases, rate of increase increases.

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This is just an estimate, but a close estimate (usually within ~10%).

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0	500	0	1	0.80	0	0	0
1	400	2	0.8	0.50	1.6	1.6	0.780
2	200	3	0.4	0.25	1.2	2.4	0.285
3	50	1	0.1	0	0.1	0.3	0.012
4	0	0	0				$\Sigma = 1.077$

The Euler equation (pronounced "Oiler")

$$1 = \sum_{e^{-rx}l(x)b(x)}$$

Unfortunately, there is no way to solve this equation except to plug in different values of *r*.

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0	500	0	1	0.80	0	0	0
1	400	2	0.8	0.50	1.6	1.6	0.780
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			$r \approx 0.7$	718			

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4	0	0	0				$\Sigma = 1.077$
			$r \approx 0.7$	718	<i>P</i> (Euler)	= 0.77	76

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$$\mathbf{n}(t) = \begin{array}{c} n_2(t) \\ \vdots \\ n_k(t) \end{array}$$

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Using the mortality and fertility schedules from the life table, we can prediction how the age structure changes from one time period $\mathbf{n}(t)$ to the next, $\mathbf{n}(t+1)$

Calculating survival probabilities for age classes (Pi)

x	S(x)	b(x)	$I(x) = S(x)/S(\theta)$	g(x) = l(x+1)/ $l(x)$	I(x)b(x)	l(x)b(x)x	Initial estimate e ^{-rz} l(x)b(x)
0	500	0	1	0.80	0	0	0
1	400	2	0.8	0.50	1.6	1.6	0.780
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$$n_1(t+1) = \sum_{i=1}^k F_i n_i(t)$$

Once the Fi is known for each age class, we multiply these fertilities by the number of individuals in each age class. This product is then summed over all age classes to calculate the number of new offspring.

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A Leslie matrix describes the changes in population size due to mortality and reproduction.

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A Leslie matrix describes the changes in population size due to mortality and reproduction.

If there are k age classes, the Leslie matrix is a $k \times k$ square matrix.

$$\mathbf{A} = \begin{bmatrix} F_1 & F_2 & F_3 & F_4 \\ P_1 & 0 & 0 & 0 \\ 0 & P_2 & 0 & 0 \\ 0 & 0 & P_3 & 0 \end{bmatrix}$$

$$n_1(t+1) = F_1 n_1(t) + F_2 n_2(t) + F_3 n_3(t) + F_4 n_4(t)$$

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Each **column is the age at time** *t*Each **row is the age at time** *t* **+1**The first row are the fertilities - contributions to newborn of each age class.

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$$n_1(t+1) = F_1 n_1(t) + F_2 n_2(t) + F_3 n_3(t) + F_4 n_4(t)$$

 $n_2(t+1) = P_1 n_1(t)$
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The survival probabilities are always in the subdiagonal. They represent transitions from one age class to the next.

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$$n_1(t+1) = F_1 n_1(t) + F_2 n_2(t) + F_3 n_3(t) + F_4 n_4(t)$$

 $n_2(t+1) = P_1 n_1(t)$
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Each column is the age at time t Each row is the age at time t +1

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The survival probabilities are always in the subdiagonal. They represent transitions from one age class to the next. 0's: those transitions aren't possible.

$$\mathbf{A} = \begin{bmatrix} F_1 & F_2 & F_3 & F_4 \\ P_1 & 0 & 0 & 0 \\ 0 & P_2 & 0 & 0 \\ 0 & 0 & P_3 & 0 \end{bmatrix}$$

x	i (age class)	S(x)	b(x)	l(x)	Pi = l(i)/l(i-1)	Fi = b(i)Pi
0		500	0	1		
1	1	400	2	0.8	0.80	1.6
2	2	200	3	0.4	0.50	1.5
3	3	50	1	0.1	0.25	0.25
4	4	0	0	0	0	0

х	i (age class)	S(x)	b(x)	I(x)	Pi = l(i)/l(i-1)	Fi = b(i)Pi
0		500	0	1		
-1	1	400	2	0.8	0.80	1.6
2	2	200	3	0.4	0.50	1.5
3	3	50	1	0.1	0.25	0.25
4	4	0	0	0	0	0

$$\mathbf{A} = \begin{bmatrix} 1.6 & 1.5 & 0.25 & 0 \\ 0.8 & 0 & 0 & 0 \\ 0 & 0.5 & 0 & 0 \\ 0 & 0 & 0.25 & 0 \end{bmatrix}$$

$$\mathbf{A} = \begin{bmatrix} 1.6 & 1.5 & 0.25 & 0 \\ 0.8 & 0 & 0 & 0 \\ 0 & 0.5 & 0 & 0 \\ 0 & 0 & 0.25 & 0 \end{bmatrix}$$

Let's compare two populations.

$$\mathbf{A} = \begin{bmatrix} 1.6 & 1.5 & 0.25 & 0 \\ 0.8 & 0 & 0 & 0 \\ 0 & 0.5 & 0 & 0 \\ 0 & 0 & 0.25 & 0 \end{bmatrix} \qquad \mathbf{n}_0 = \begin{pmatrix} 200 \\ 0 \\ 0 \\ 0 \end{pmatrix}$$

$$\mathbf{n}_0 = \begin{pmatrix} 200 \\ 0 \\ 0 \\ 0 \end{pmatrix}$$

$$\mathbf{A} = \begin{bmatrix} 1.6 & 1.5 & 0.25 & 0 \\ 0.8 & 0 & 0 & 0 \\ 0 & 0.5 & 0 & 0 \\ 0 & 0 & 0.25 & 0 \end{bmatrix} \qquad \mathbf{n}_0 = \begin{pmatrix} 50 \\ 50 \\ 50 \\ 50 \end{pmatrix}$$

$$\mathbf{n}_0 = \begin{pmatrix} 50 \\ 50 \\ 50 \\ 50 \end{pmatrix}$$

APPENDIX: BASIC MATRIX MANIPULATIONS

Matrix Multiplication

A matrix is a table of numbers, and the mathematical manipulation of matrices derives from analysis of systems of linear equations. Thus, for example, if we have the system

$$Y_1 = a_1X_1 + b_1X_2 + c_1X_3,$$
 (A1a)

$$Y_2 = a_2 X_1 + b_2 X_2 + c_2 X_3,$$
 (A1b)

and

$$Y_3 = a_3X_1 + b_3X_2 + c_3X_3,$$
 (A1c)

simply as a matter of convenience we can group the X_i s to the right and more easily visualize the structure of the system as follows:

$$\begin{vmatrix} Y_1 \\ Y_2 \\ Y_3 \end{vmatrix} = \begin{vmatrix} a_1 & b_1 & c_1 \\ a_2 & b_2 & c_2 \\ a_3 & b_3 & c_3 \end{vmatrix} \begin{vmatrix} X_1 \\ X_2 \\ X_3 \end{vmatrix}.$$

Here we have three matrices: first a matrix with a single column (the Ys), second a square matrix with three columns (sometimes referred to as the detached coefficient matrix), and third a matrix with a single column (the Xs). Sometimes the various matrices are simply referred to using a single letter, but it is customary when speaking of matrices to put them in boldface type, so the above equation could be

$$Y = AX$$
. (A2)

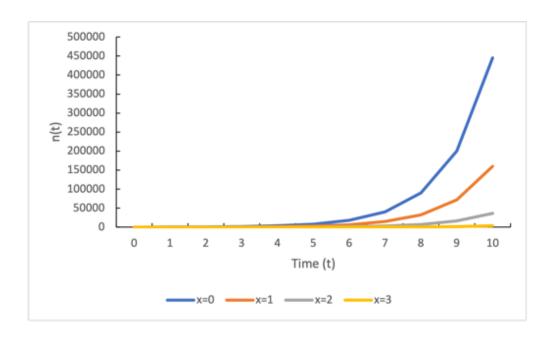
Let's compare two populations.

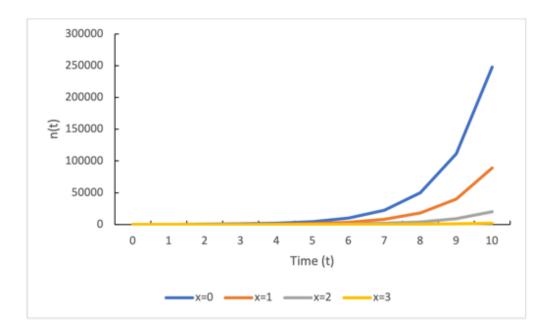
$$\mathbf{A} = \begin{bmatrix} 1.6 & 1.5 & 0.25 & 0 \\ 0.8 & 0 & 0 & 0 \\ 0 & 0.5 & 0 & 0 \\ 0 & 0 & 0.25 & 0 \end{bmatrix}$$

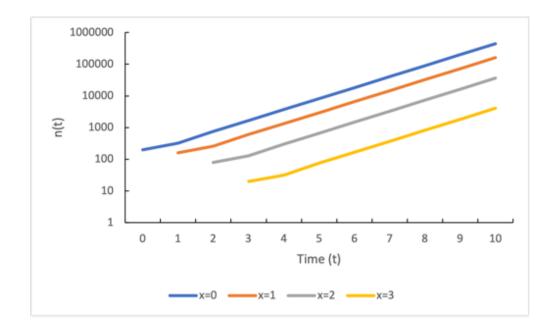
$$\mathbf{A} = \begin{bmatrix} 1.6 & 1.5 & 0.25 & 0 \\ 0.8 & 0 & 0 & 0 \\ 0 & 0.5 & 0 & 0 \\ 0 & 0 & 0.25 & 0 \end{bmatrix} \qquad \mathbf{n}_0 = \begin{pmatrix} 200 \\ 0 \\ 0 \\ 0 \end{pmatrix} \begin{array}{l} 1.6*200 + 1.5*0 + 0.25*0 + 0*0 = 320 \\ 0.8*200 + 0*0 + 0*0 + 0*0 = 160 \\ 0*200 + 0.5*0 + 0*0 + 0*0 = 0 \\ 0*200 + 0*0 + 0.25*0 + 0*0 = 0 \end{bmatrix}$$

$$\mathbf{A} = \begin{bmatrix} 1.6 & 1.5 & 0.25 & 0 \\ 0.8 & 0 & 0 & 0 \\ 0 & 0.5 & 0 & 0 \\ 0 & 0 & 0.25 & 0 \end{bmatrix}$$

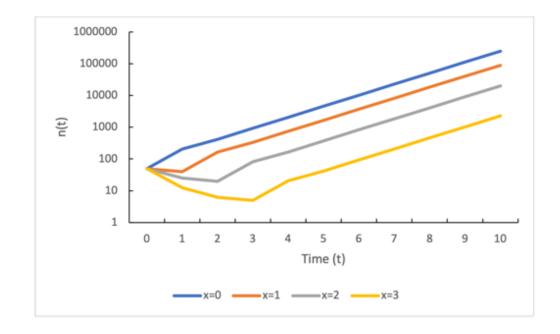
$$\mathbf{A} = \begin{bmatrix} 1.6 & 1.5 & 0.25 & 0 \\ 0.8 & 0 & 0 & 0 \\ 0 & 0.5 & 0 & 0 \\ 0 & 0 & 0.25 & 0 \end{bmatrix} \qquad \mathbf{n}_0 = \begin{pmatrix} 50 \\ 50 \\ 50 \\ 50 \end{pmatrix} \begin{cases} 1.6*50 + 1.5*50 + 0.25*50 + 0*50 = 205 \\ 0.8*50 + 0*50 + 0*50 + 0*50 = 40 \\ 0*50 + 0.5*50 + 0*50 + 0*50 = 25 \\ 0*50 + 0.5*50 + 0.25*5 + 0*50 = 12.5 \end{cases}$$







If a population is growing with constant birth and death rates, it will quickly converge on a stable age distribution, with relative numbers in each age class staying about the same.



Stage- and size-structured population growth

Sometimes size or life stage is more important than age.





Remember each column represents the stage at time t and each row represents the stage at time t + 1.

	egg	larva	adult
egg	Γ0	0	F_{ae}
larva	P_{el}	P_{ll}	0
adult	Lο	P_{la}	P_{aa}

Remember each column represents the stage at time t and each row represents the stage at time t + 1.

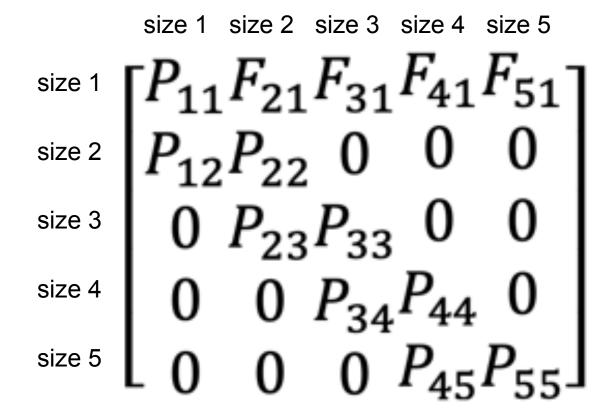
	egg	larva	adult
egg	Γ0	0	F_{ae}
larva	P_{el}	P_{ll}	0
adult		P_{la}	P_{aa}

The first row represents the fertilities.

Remember each column represents the stage at time t and each row represents the stage at time t + 1.

The first row represents the fertilities.

The other entries are *transition probabilities*.



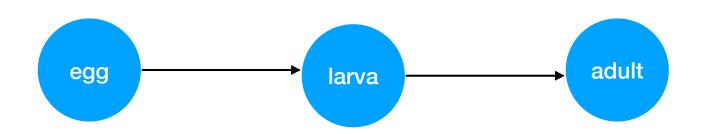
$$\begin{array}{c|cccc} & \text{egg} & \text{larva} & \text{adult} \\ \hline \\ \text{egg} & \begin{bmatrix} 0 & 0 & F_{ae} \\ P_{el} & P_{ll} & 0 \\ 0 & P_{la} & P_{aa} \\ \end{bmatrix} \\ \text{adult} \end{array}$$

egg

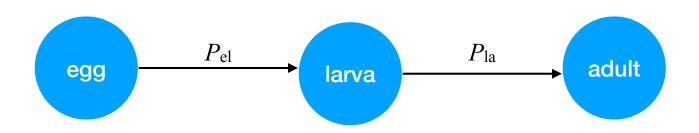
larva

adult

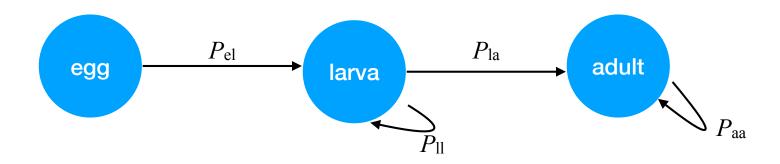
$$\begin{array}{c|cccc} & \text{egg} & \text{larva} & \text{adult} \\ \hline \\ \text{egg} & \begin{bmatrix} 0 & 0 & F_{ae} \\ P_{el} & P_{ll} & 0 \\ 0 & P_{la} & P_{aa} \\ \end{bmatrix} \\ \text{adult} \end{array}$$



$$\begin{array}{c|cccc} & \text{egg} & \text{larva} & \text{adult} \\ \hline \\ \text{egg} & \begin{bmatrix} 0 & 0 & F_{ae} \\ P_{el} & P_{ll} & 0 \\ 0 & P_{la} & P_{aa} \\ \end{bmatrix} \\ \text{adult} \end{array}$$



$$\begin{array}{c|cccc} & \text{egg} & \text{larva} & \text{adult} \\ \hline \\ \textbf{egg} & \begin{bmatrix} 0 & 0 & F_{ae} \\ P_{el} & P_{ll} & 0 \\ 0 & P_{la} & P_{aa} \\ \end{bmatrix} \\ \textbf{adult} \end{array}$$



$$\begin{array}{c|cccc} & \text{egg} & \text{larva} & \text{adult} \\ \hline \\ \textbf{egg} & \begin{bmatrix} 0 & 0 & F_{ae} \\ P_{el} & P_{ll} & 0 \\ 0 & P_{la} & P_{aa} \\ \end{bmatrix} \\ \textbf{adult} \end{array}$$

