

# Comparing mathematical models of NSC Dynamics in Zebrafish against the Mouse

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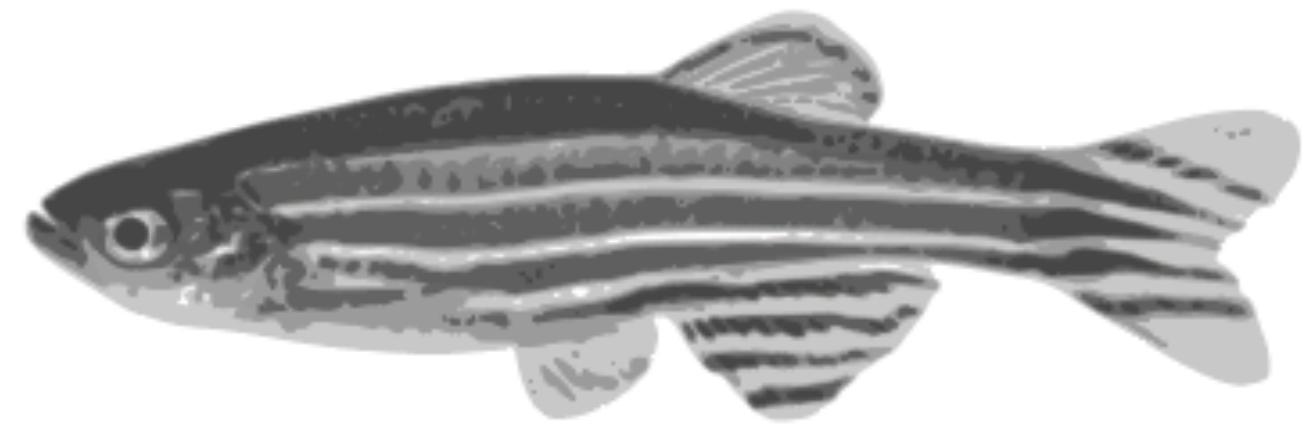
## Observations

### Adult mouse



- NSCs numbers decline with age in the DG and V-SVZ

### Adult zebrafish



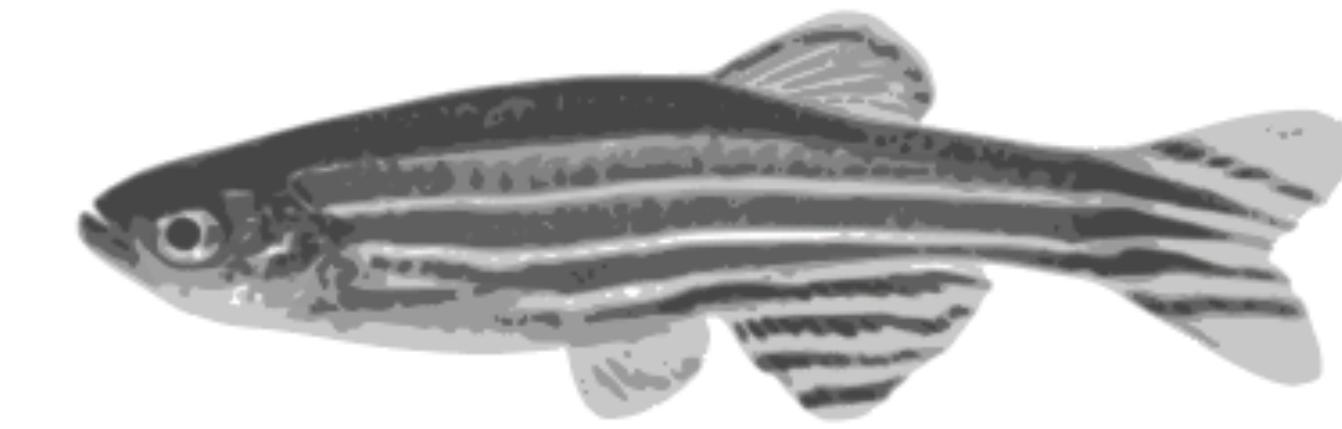
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- NSCs numbers decline with age in the DG and V-SVZ
- Mathematical model of NSC dynamics

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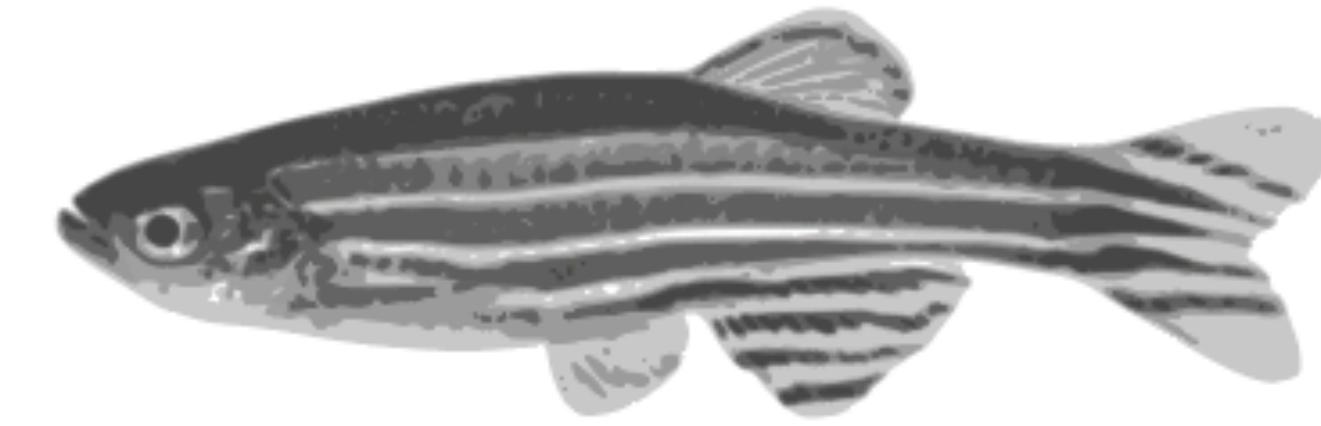
### Unraveling regulatory feedback mechanisms in adult neurogenesis through mathematical modelling

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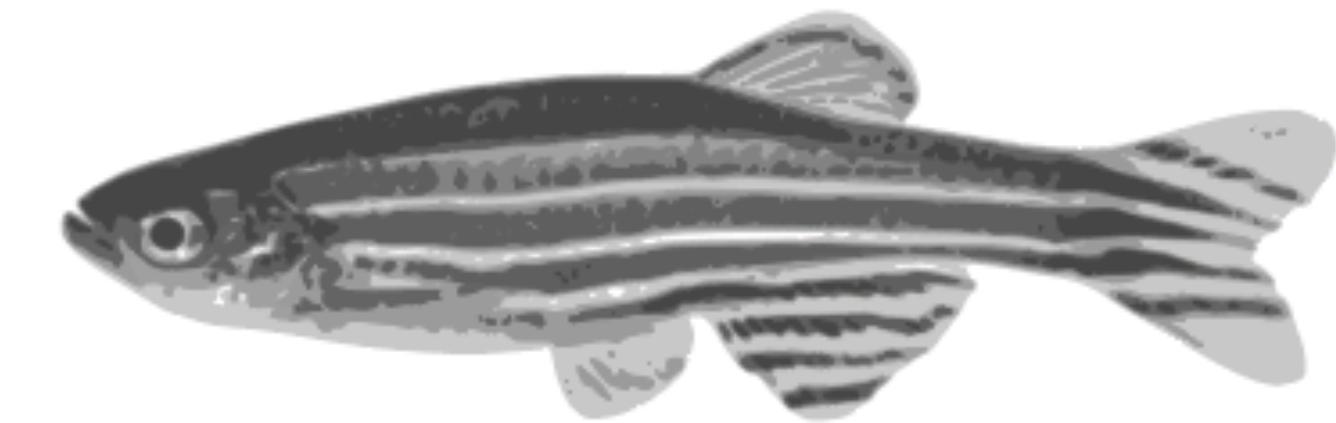
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## Adult zebrafish

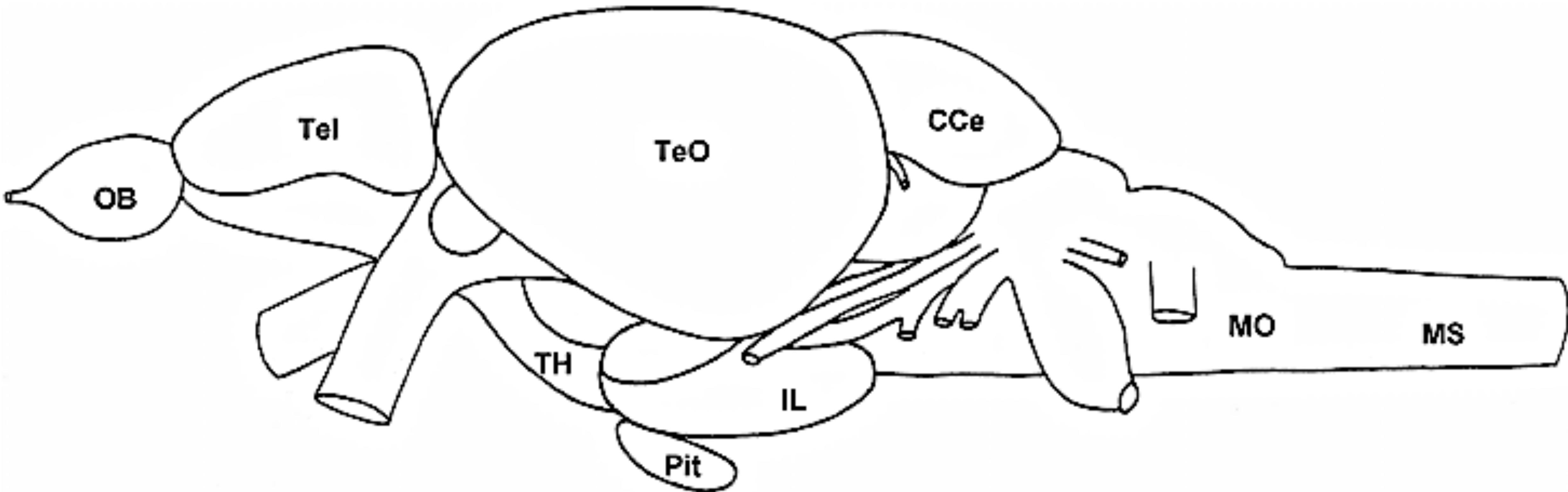


- NSCs in homologous neurogenic territories are maintained in numbers and proportions (homeostasis)



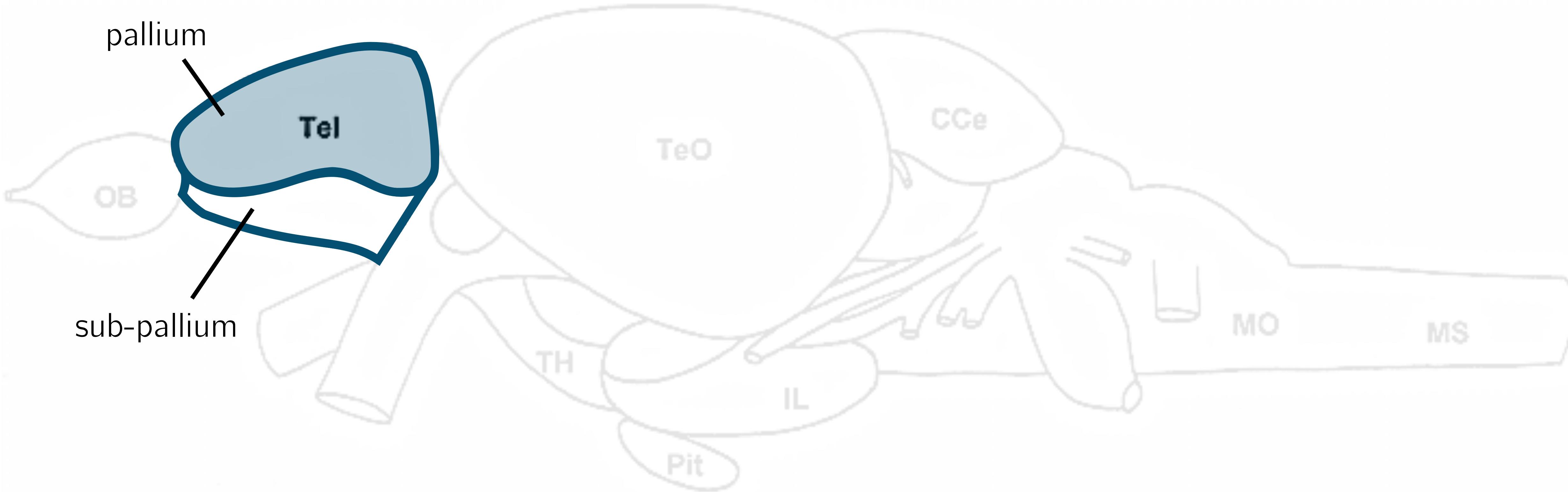
Same?  
Slightly different?  
Completely different?

## Biological Context: NSCs anatomy and lineage progression in the fish brain



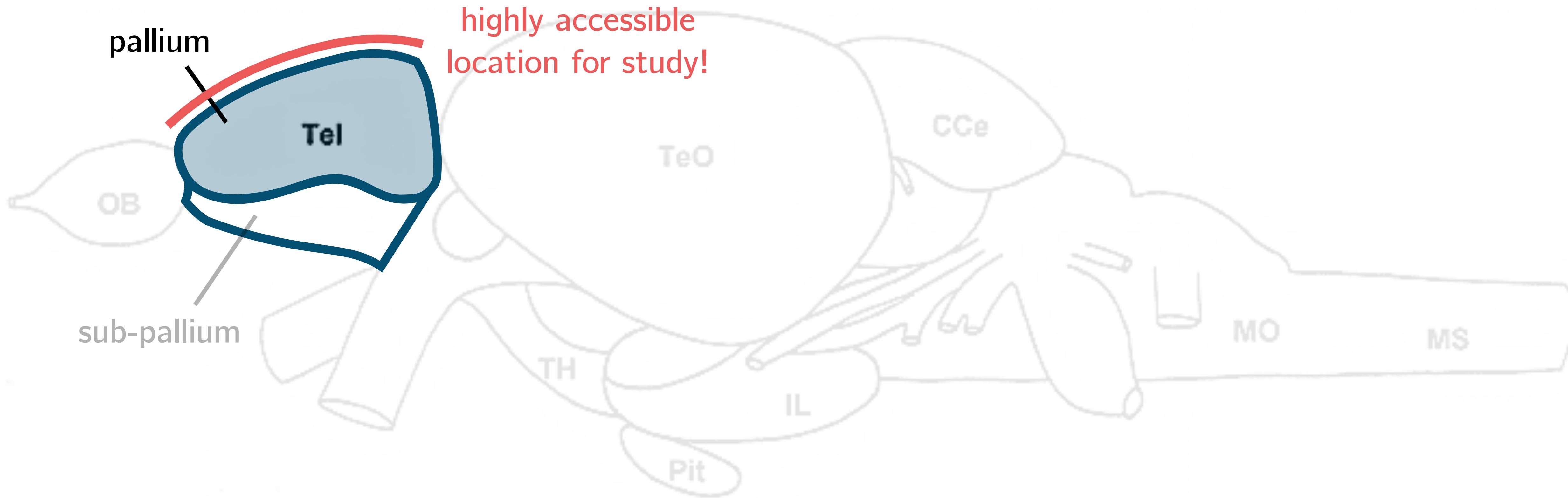
**Figure: Lateral view of the adult zebrafish brain (*credit to Daniel R. Dietrich*)**

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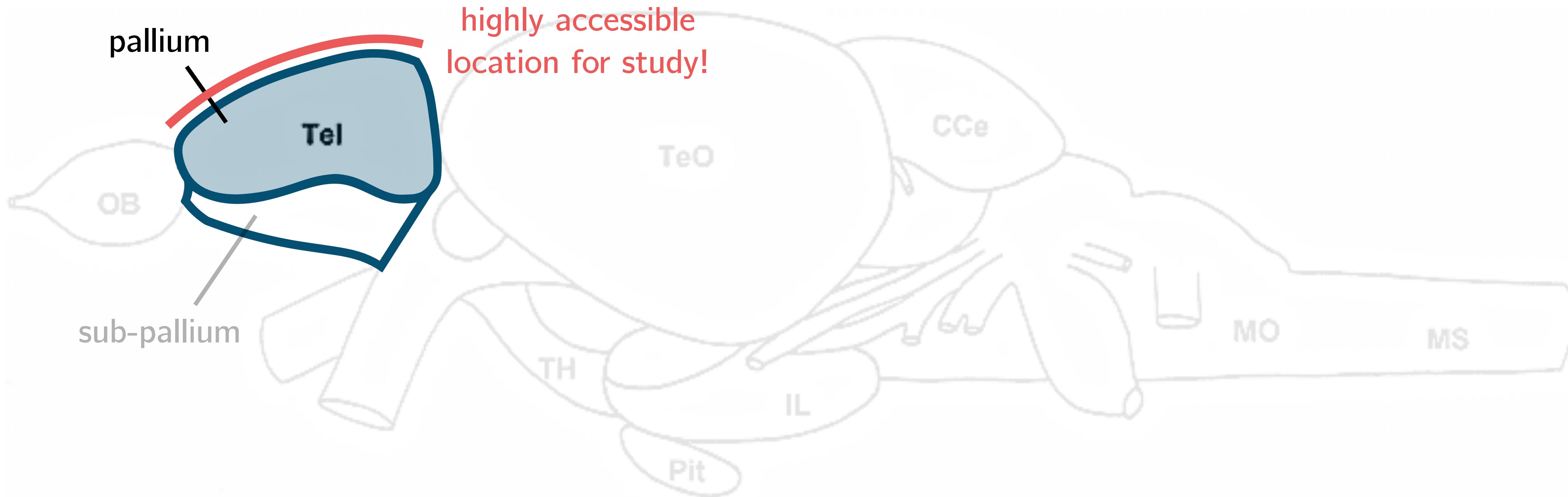
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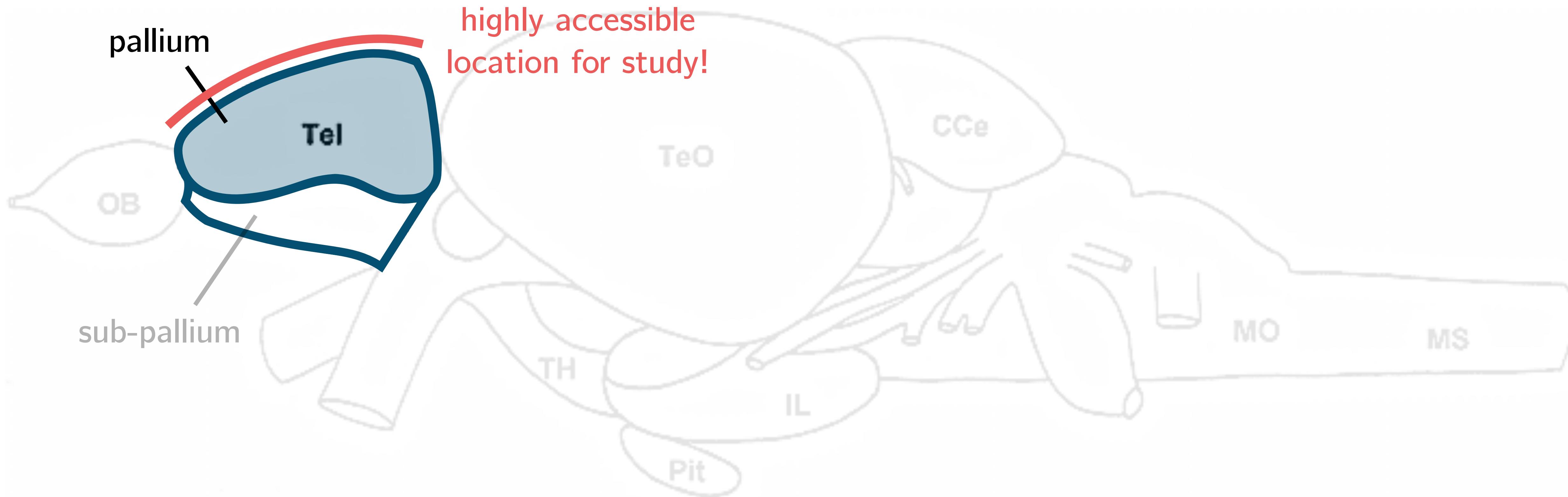


**Figure: Lateral view of the adult zebrafish brain (credit to Daniel R. Dietrich)**

Radial glia as NSCs:  
GFAP/S100 $\beta$ /GS  
Her4 (=Hes5)  
Sox2

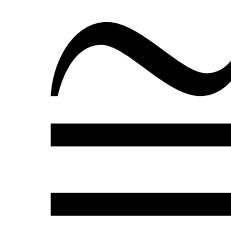


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**Figure: Lateral view of the adult zebrafish brain (*credit to Daniel R. Dietrich*)**

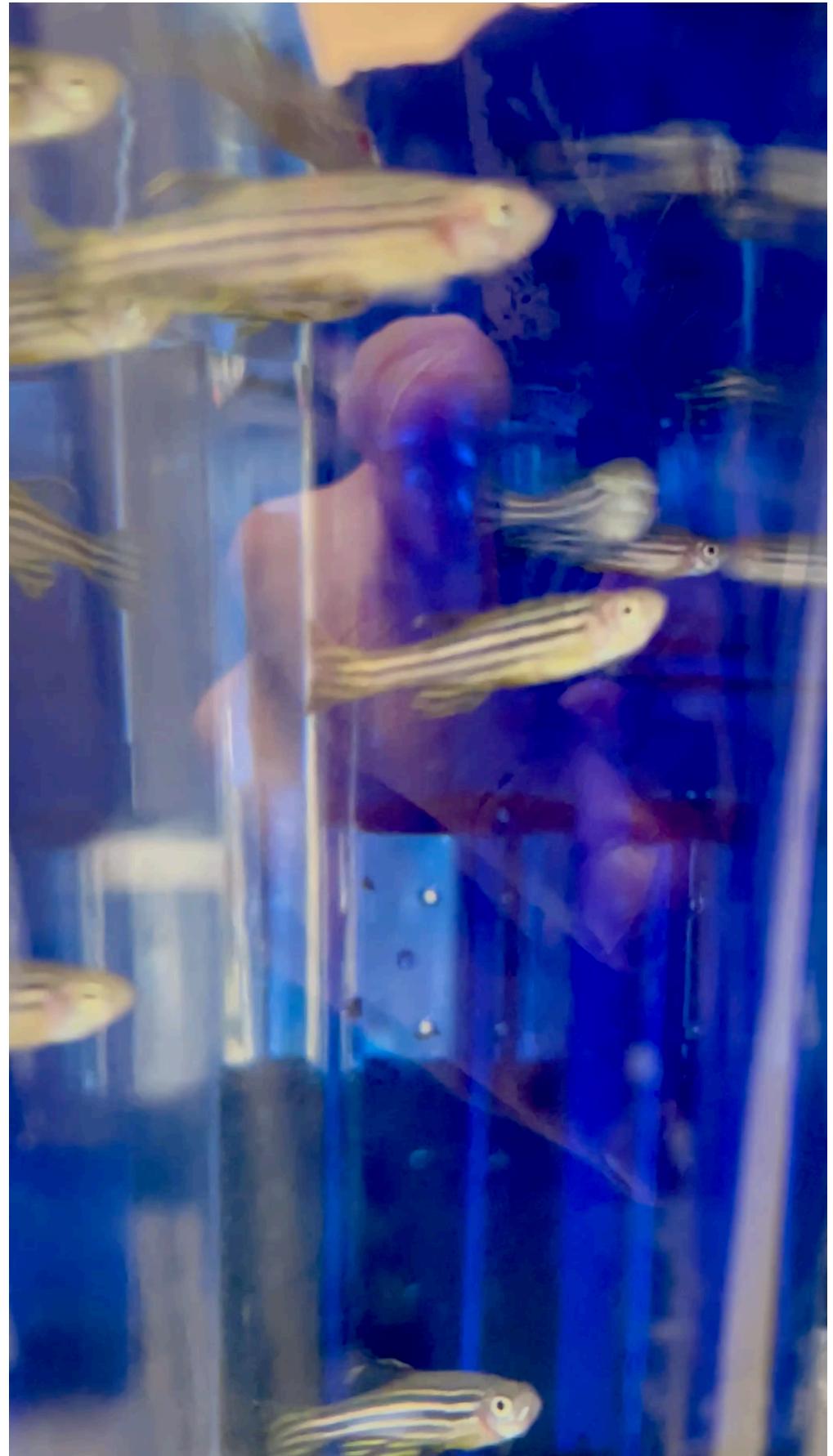
Radial glia as NSCs:  
GFAP/S100 $\beta$ /GS  
Her4 (=Hes5)  
Sox2



SVZ/SEZ of the lateral ventricle  
and  
DG of the hippocampus

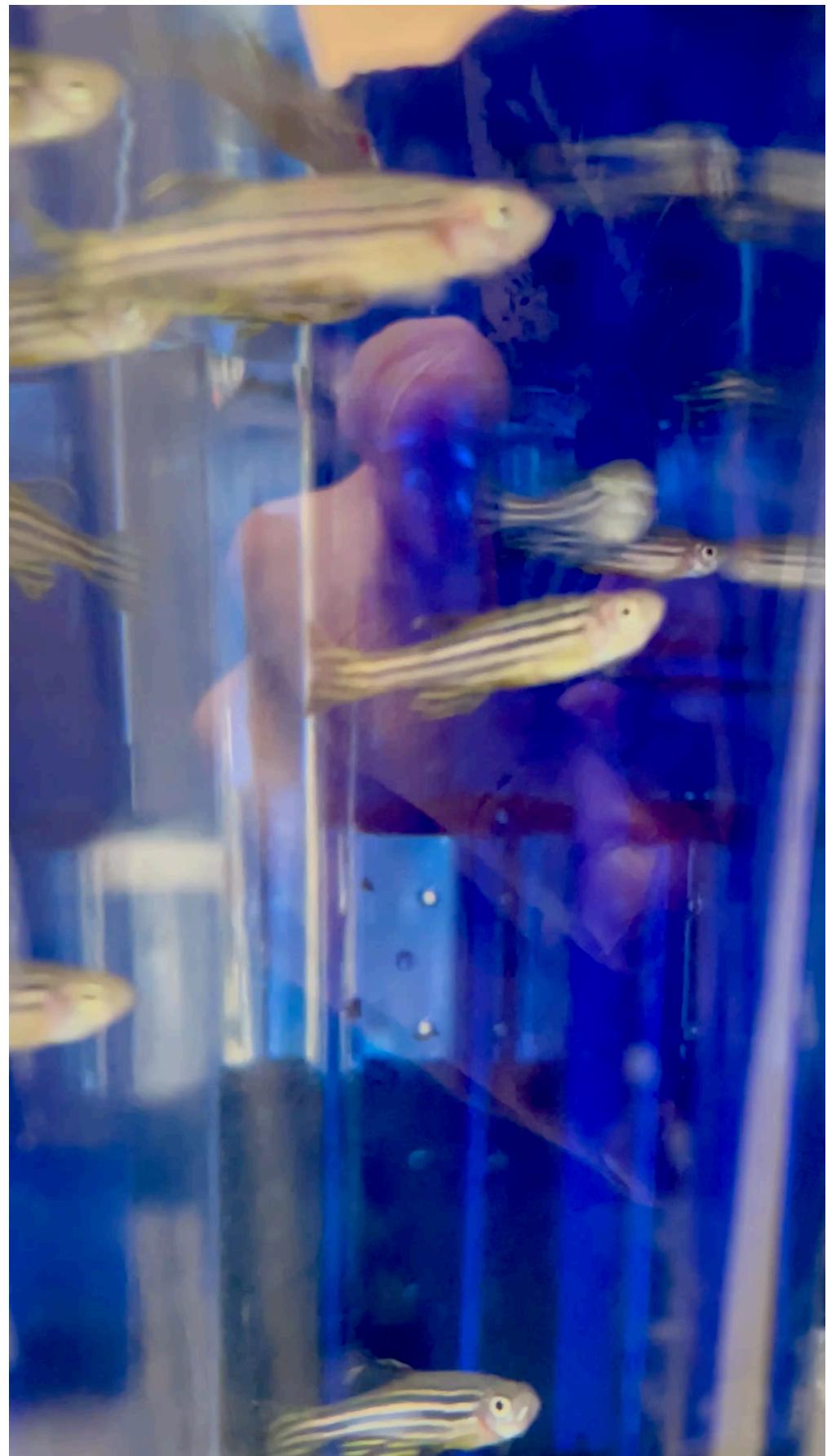
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## Wild Type



# Biological Context: NSCs anatomy and lineage progression in the fish brain

**Wild Type**



**Mutant**



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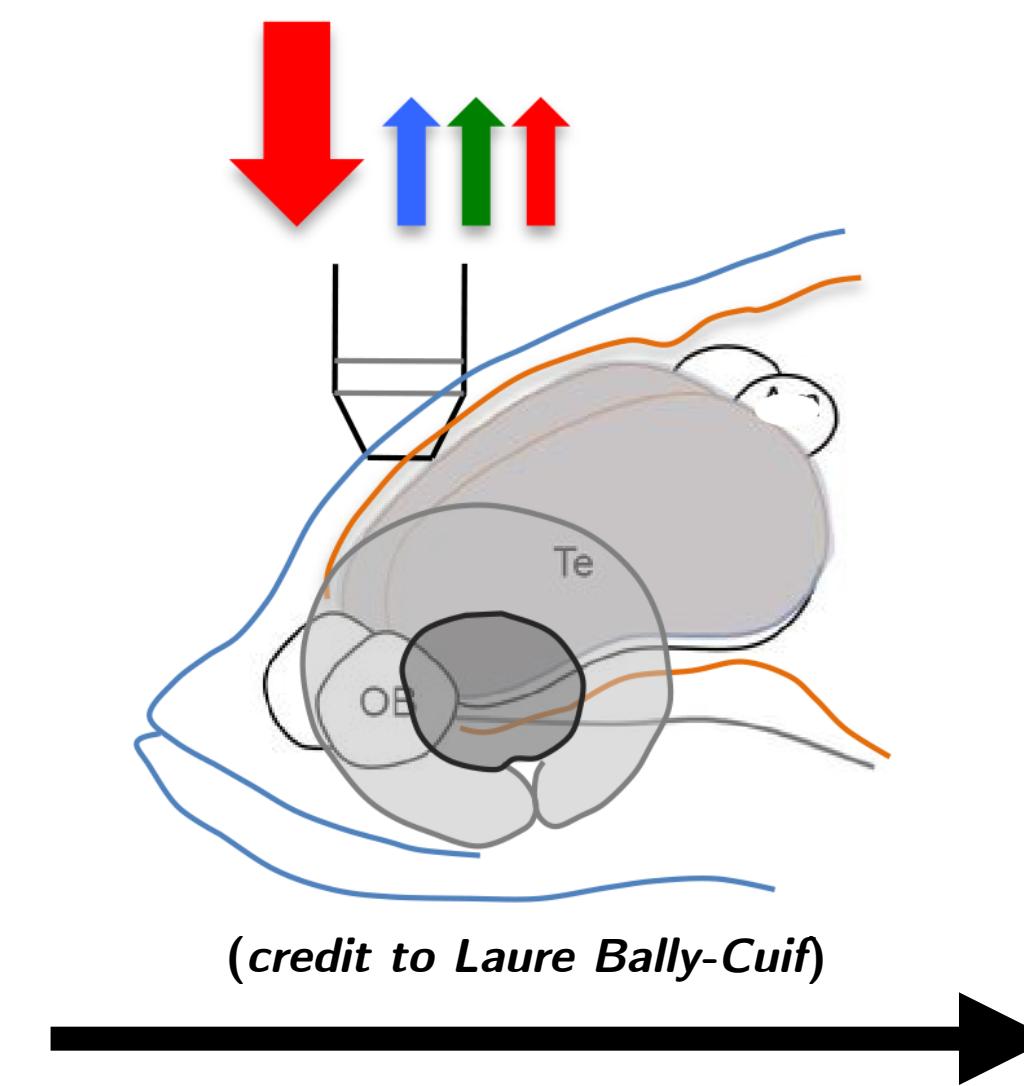
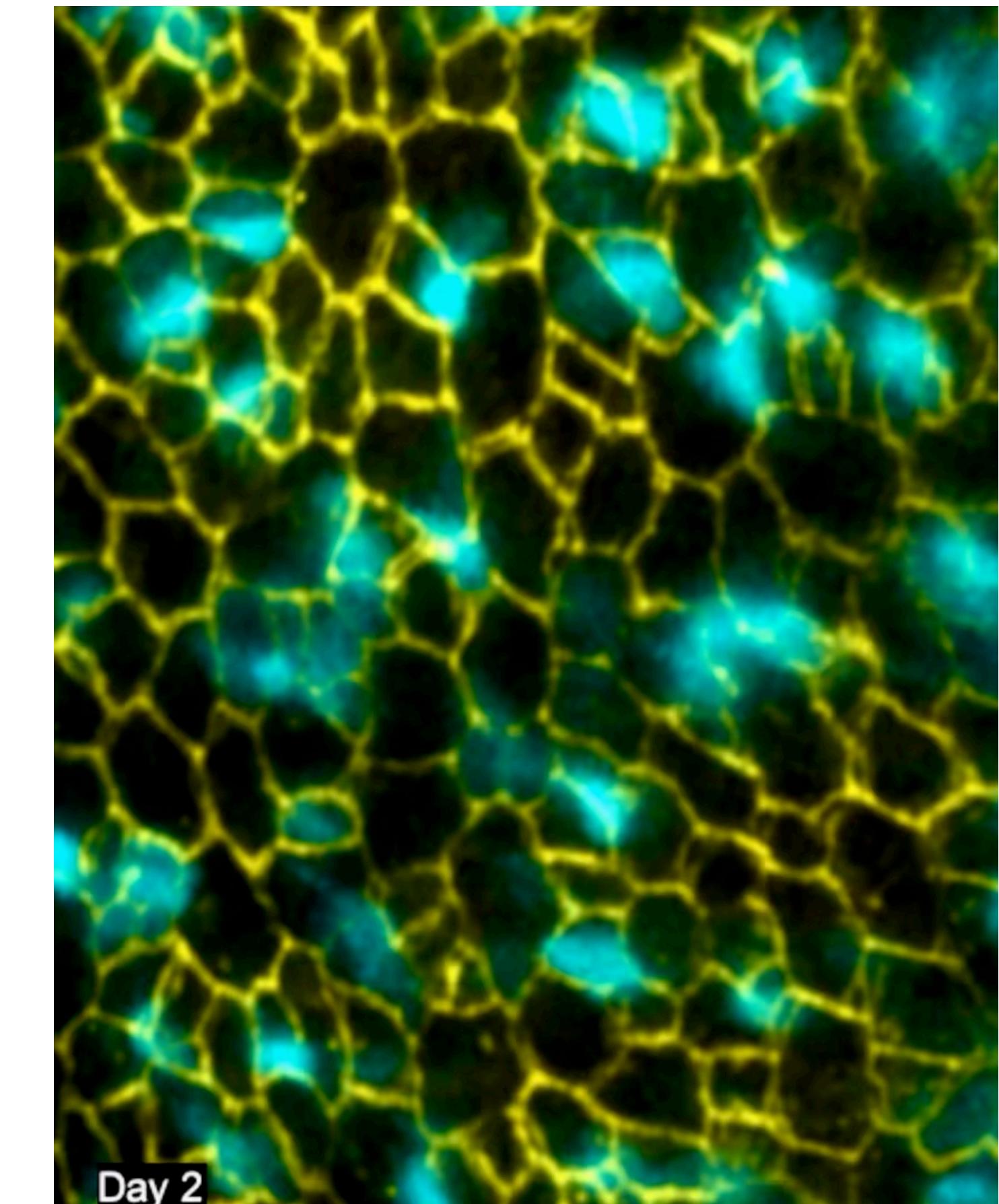
**Wild Type**



**Mutant**

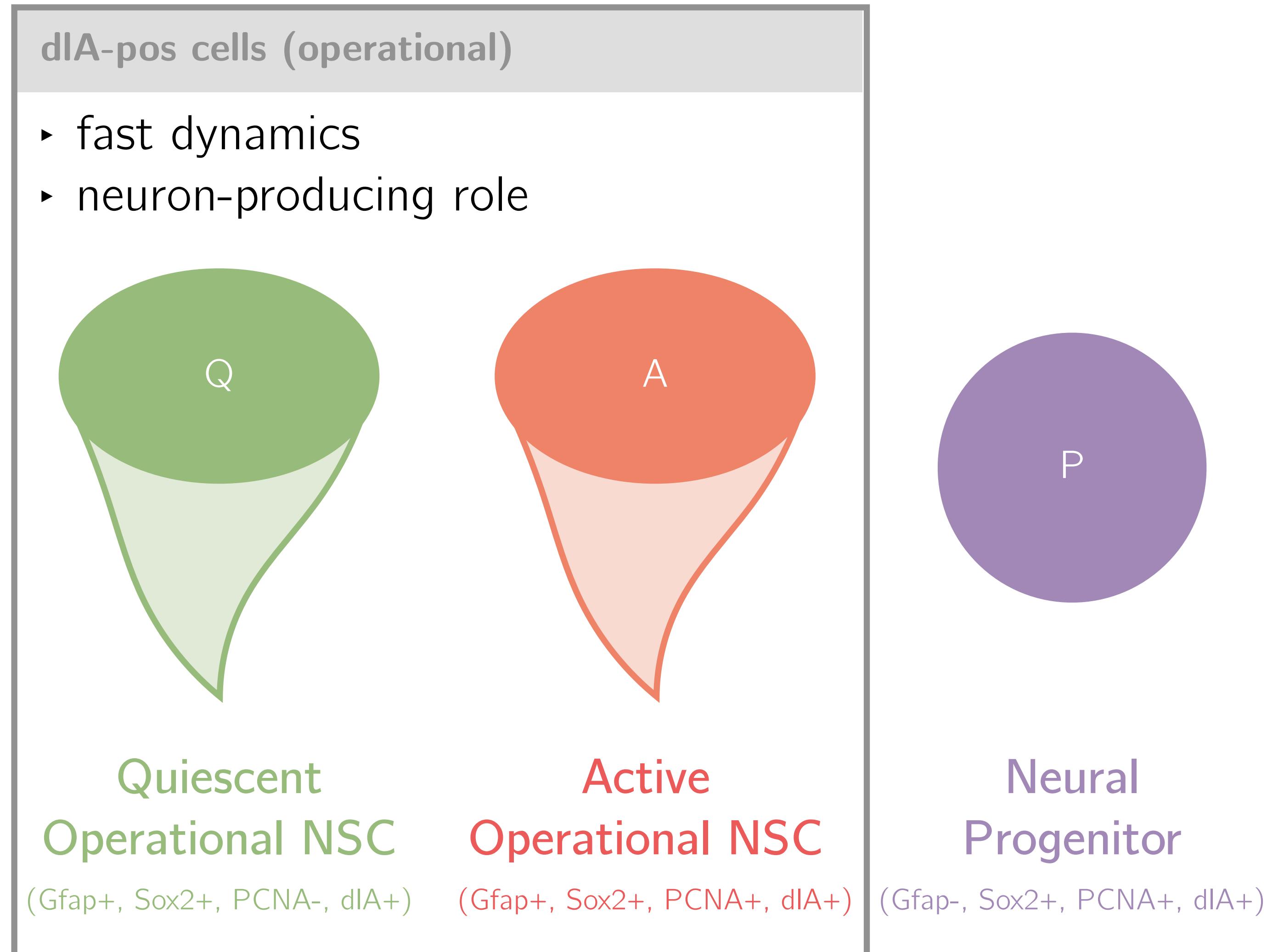
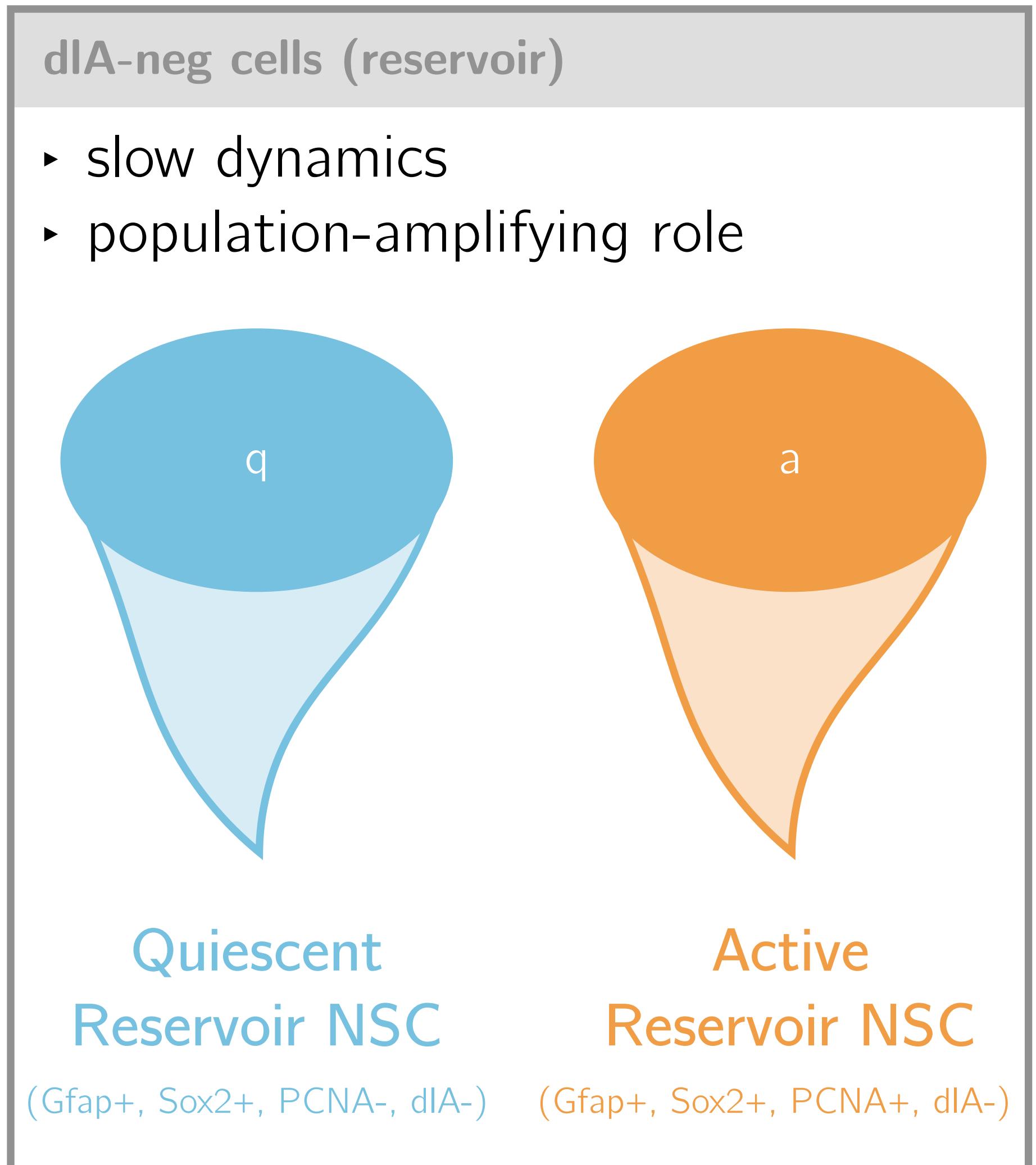


**Pallial NSC population**

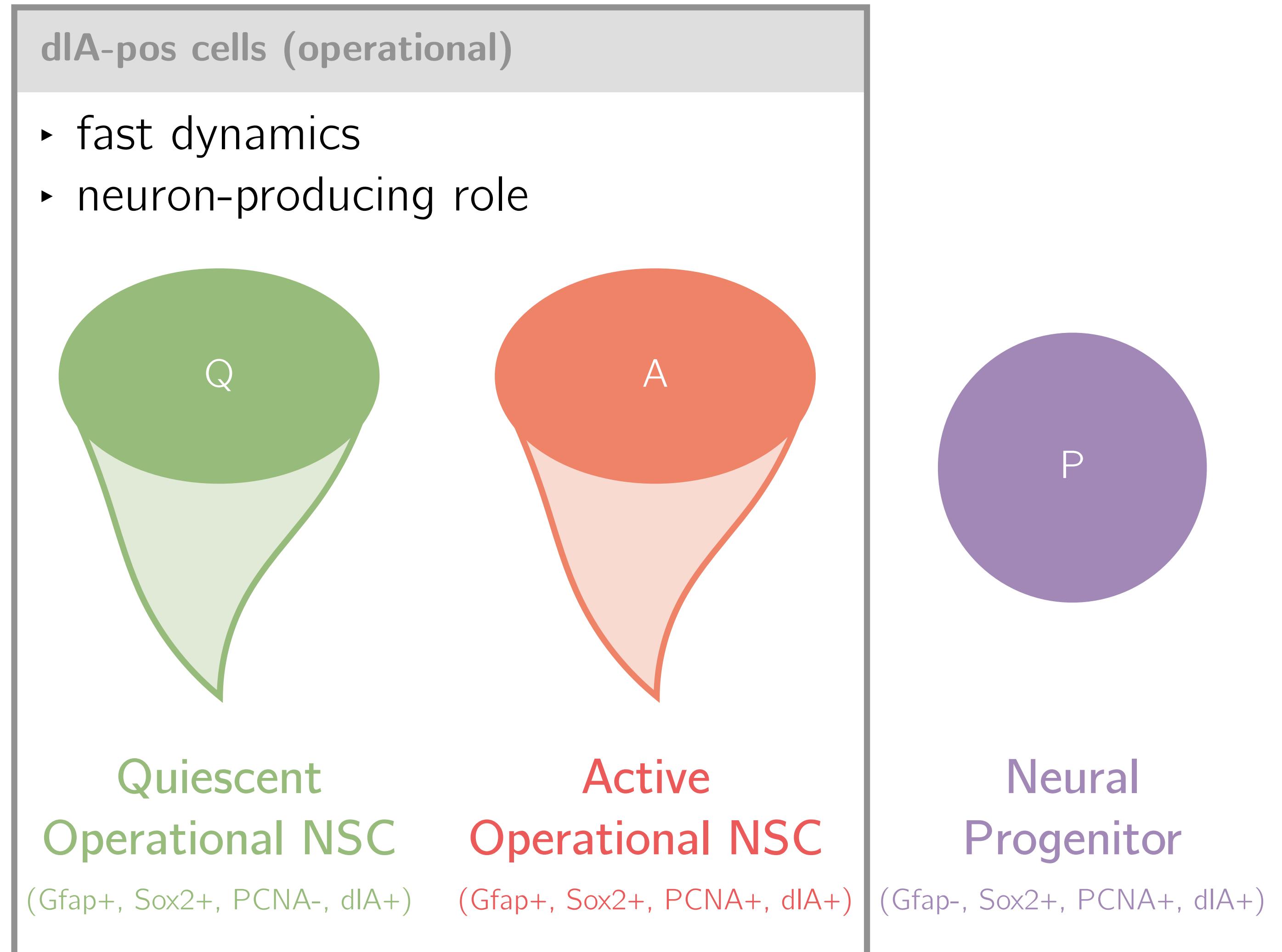
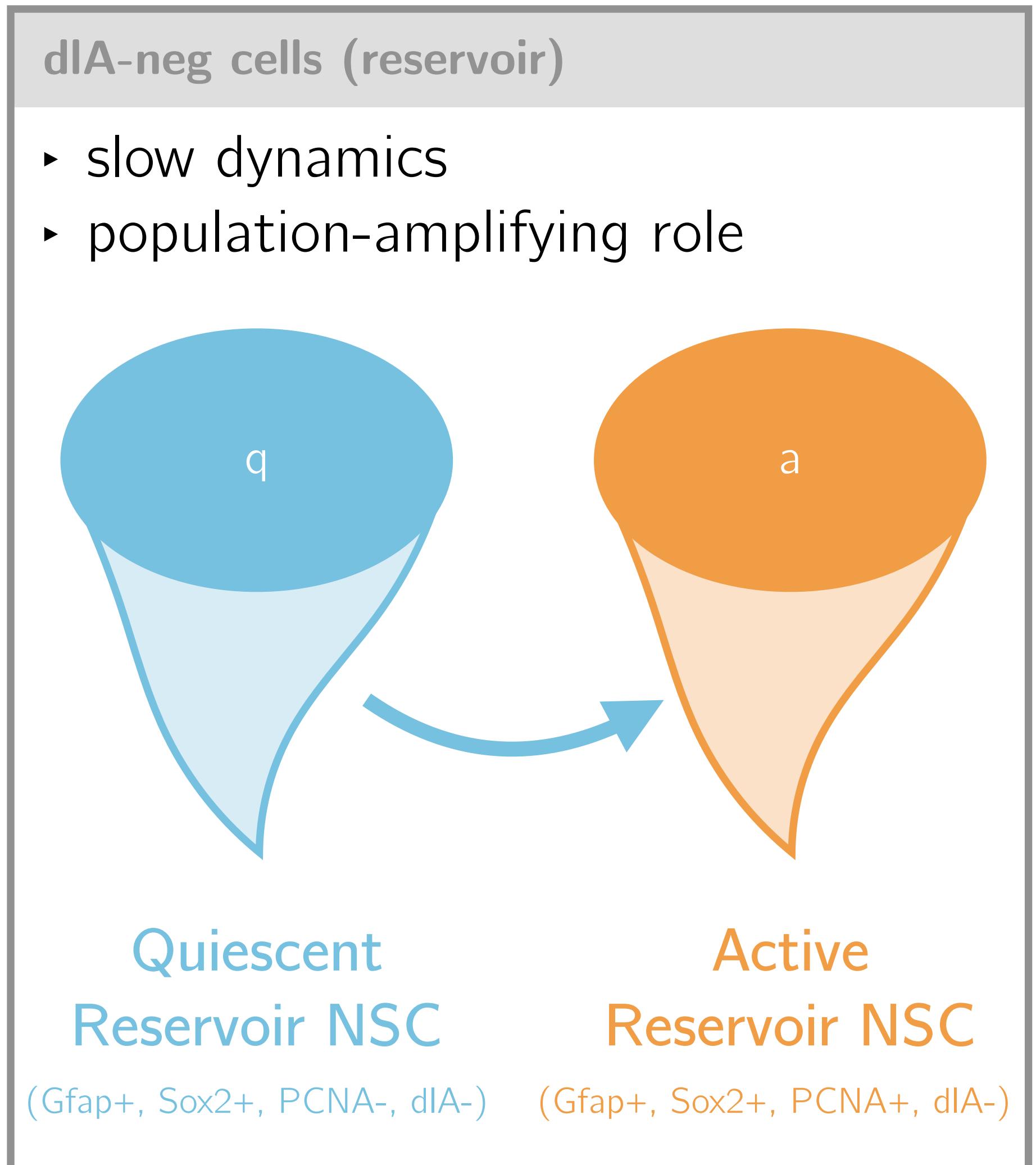


source: <https://doi.org/10.1126/sciadv.adg7519>

# Biological Context: NSCs anatomy and lineage progression in the fish brain

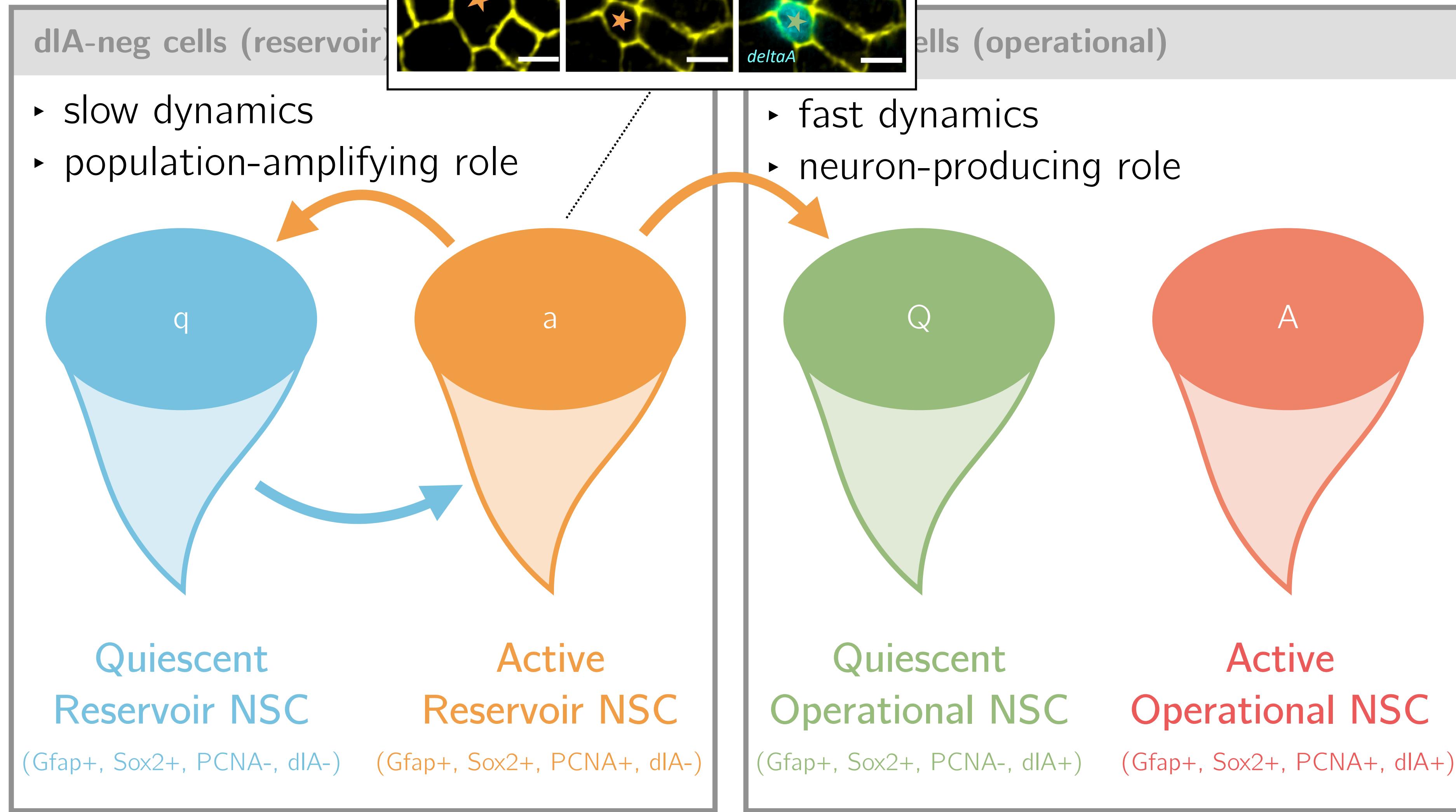


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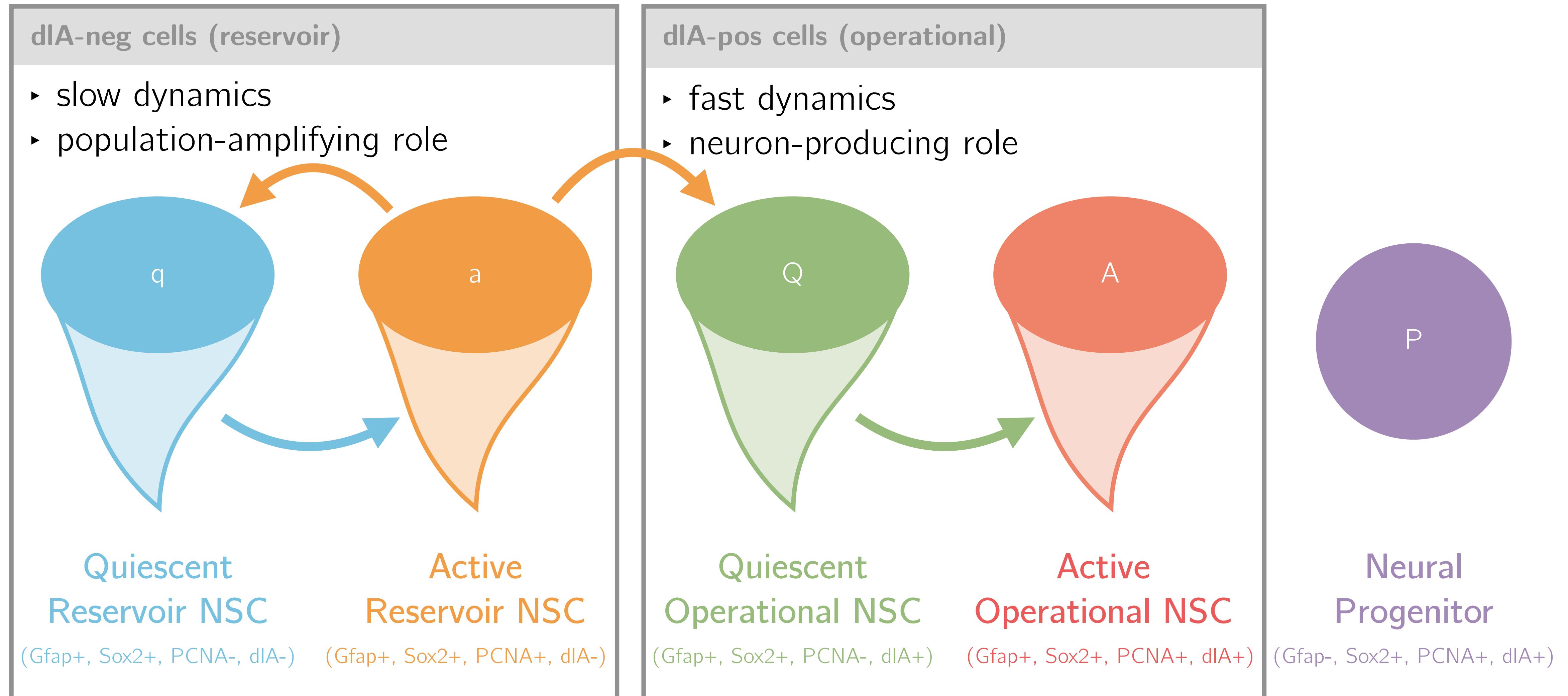


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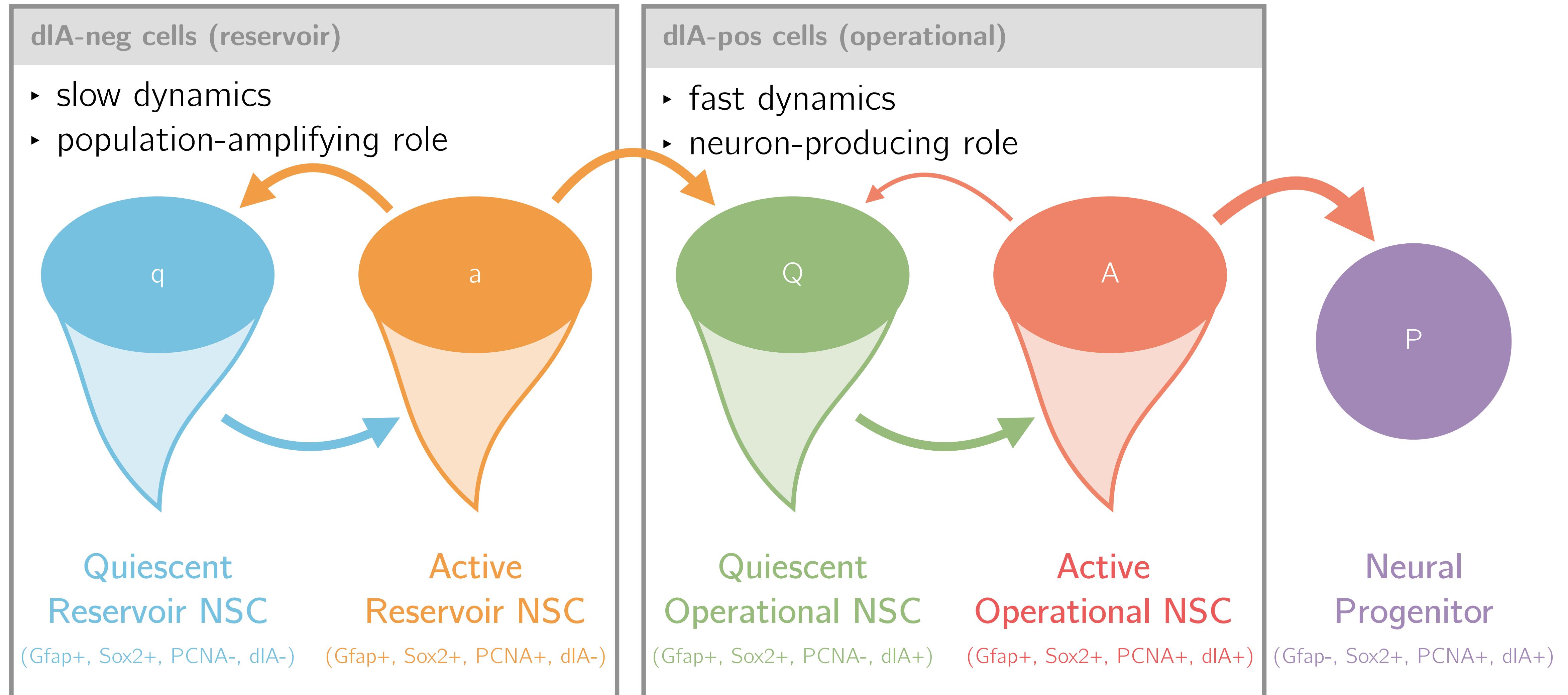
# Stage progression in the fish brain



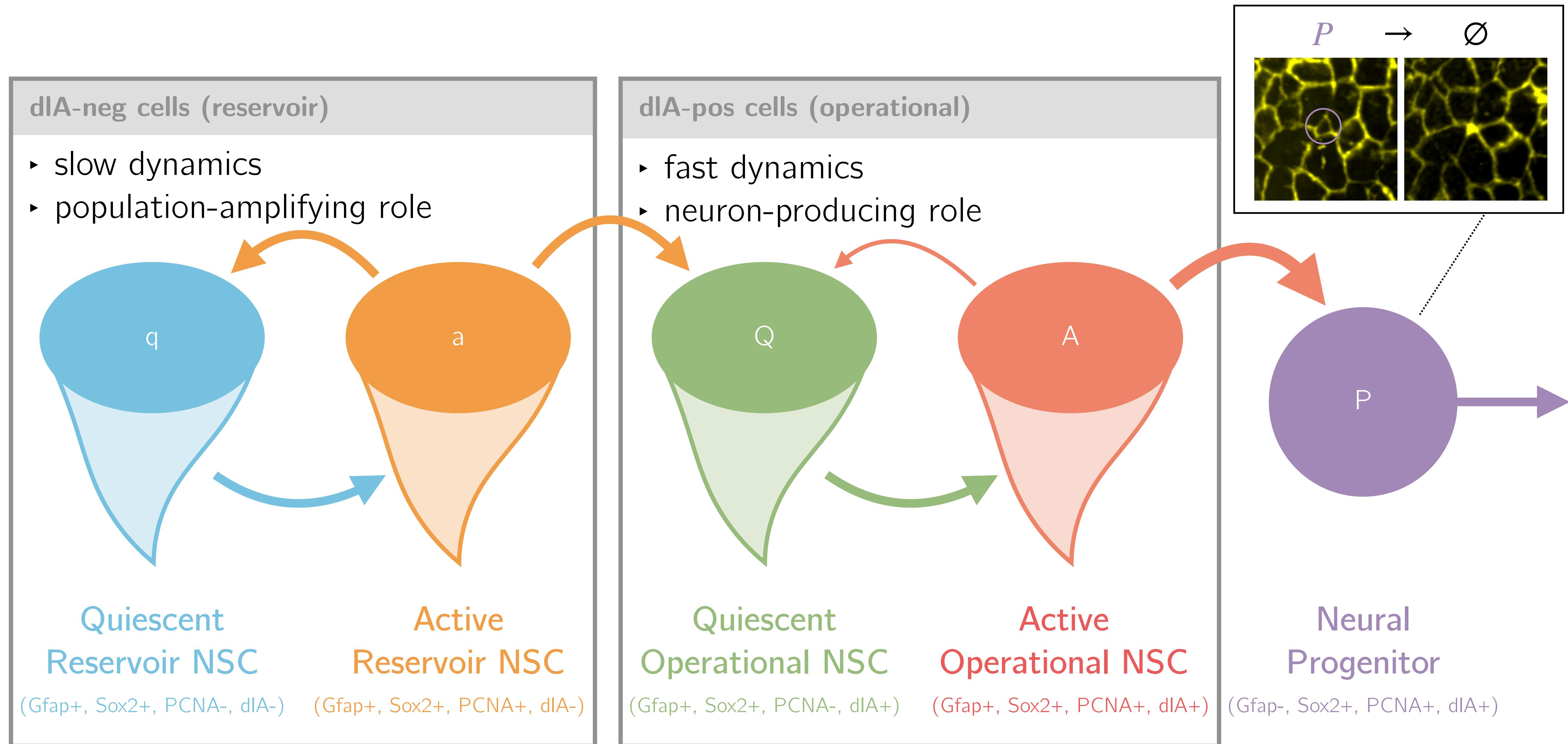
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# Biological Context: NSCs anatomy and lineage progression in the fish brain

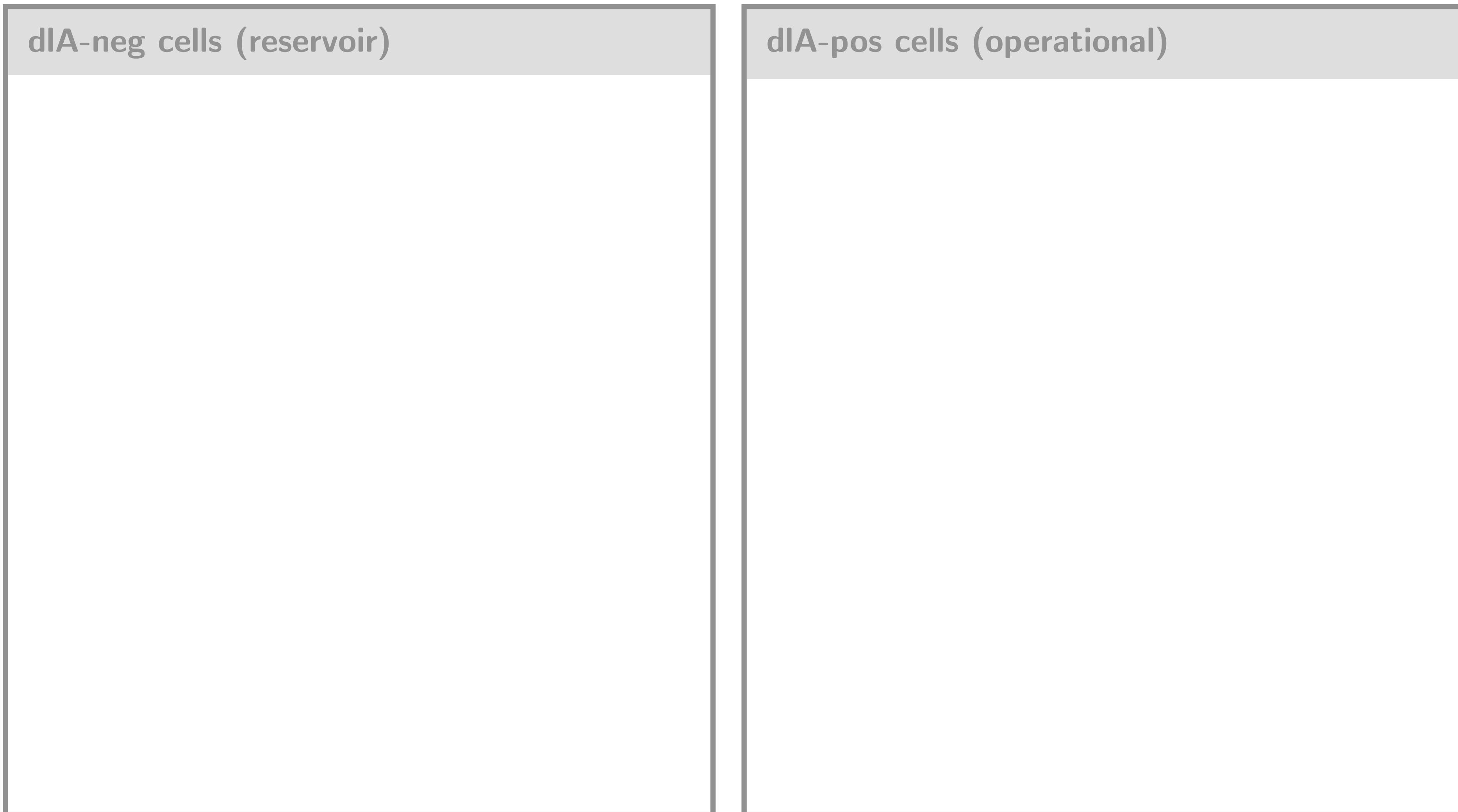


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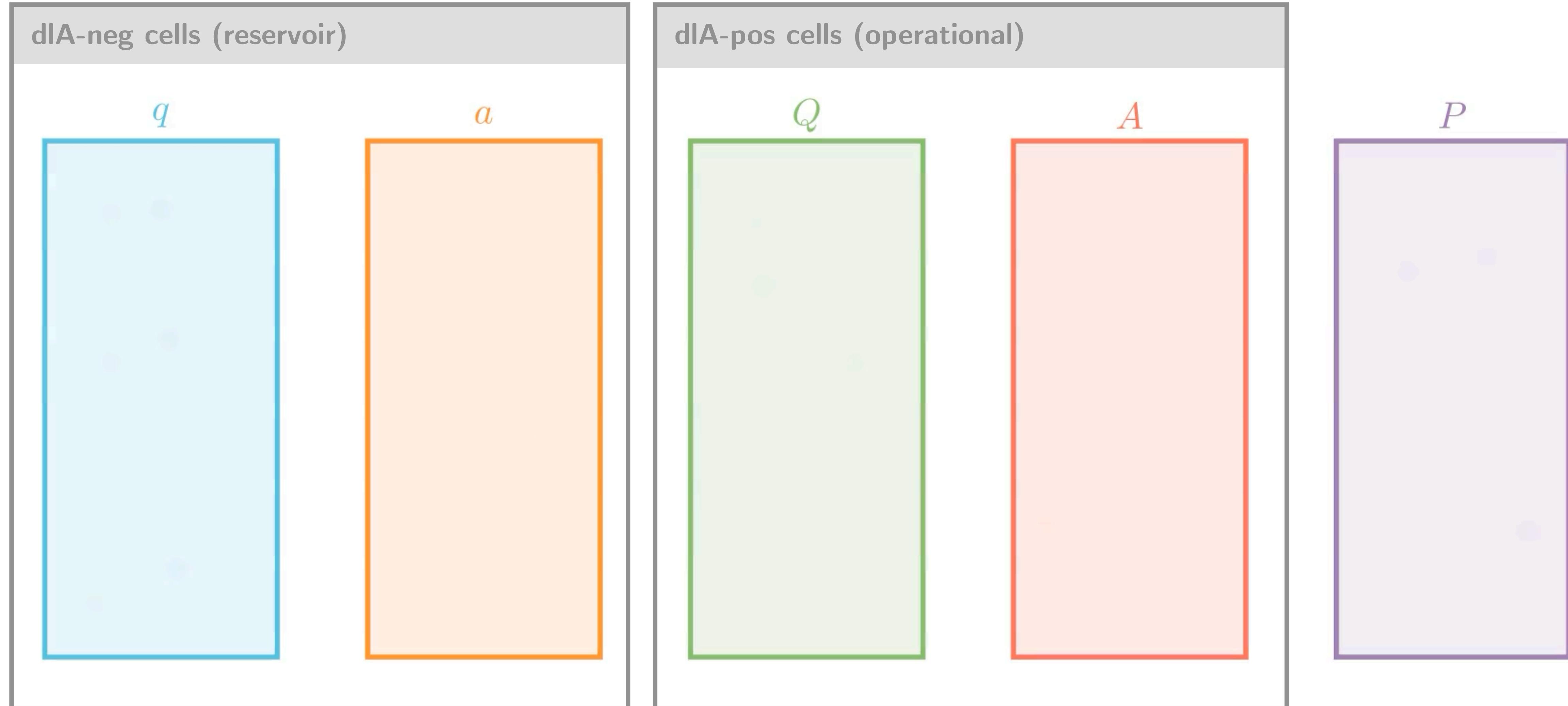
# Model Visualization: Compartmental evolution

Step 1: Define state variables (who's who)



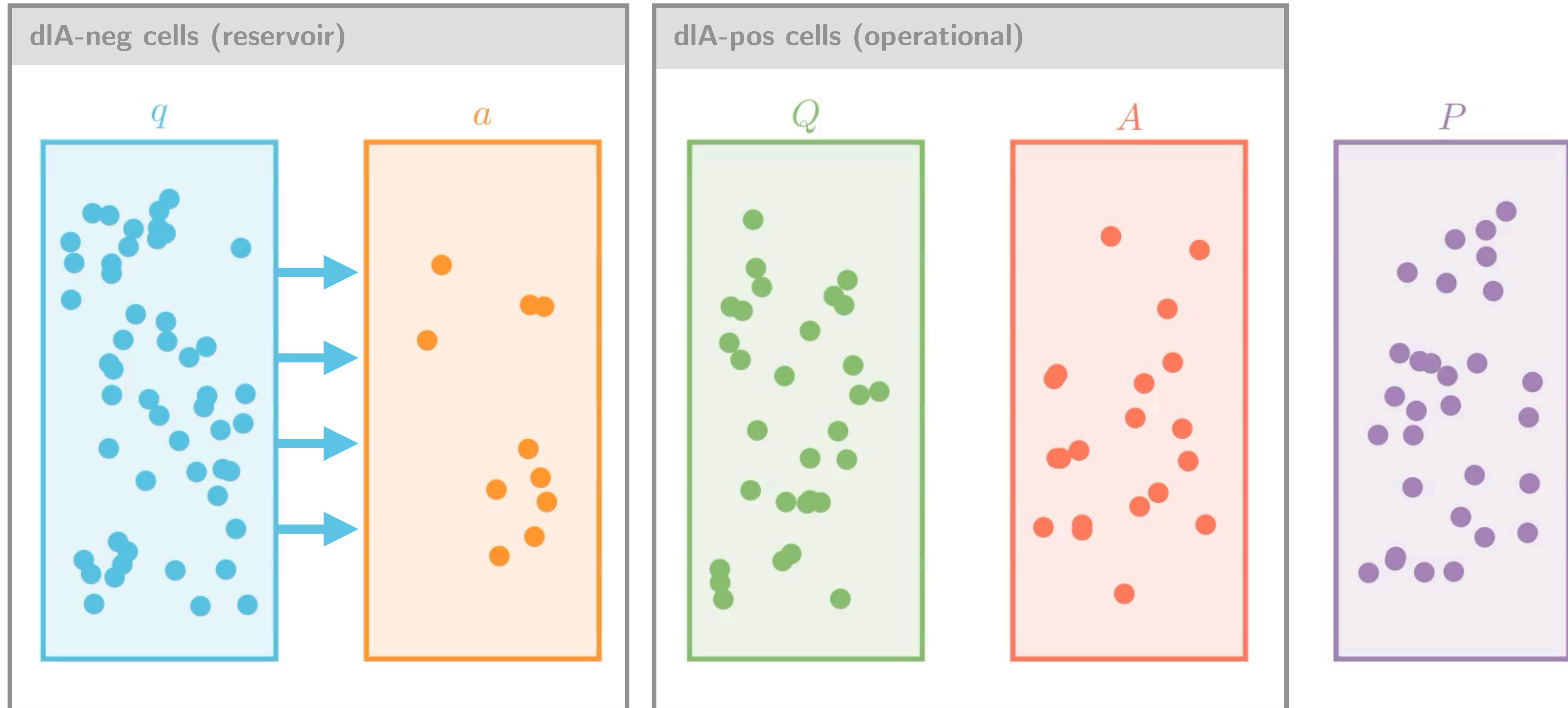
# Model Visualization: Compartmental evolution

Step 2: Populate each compartment (initial condition)



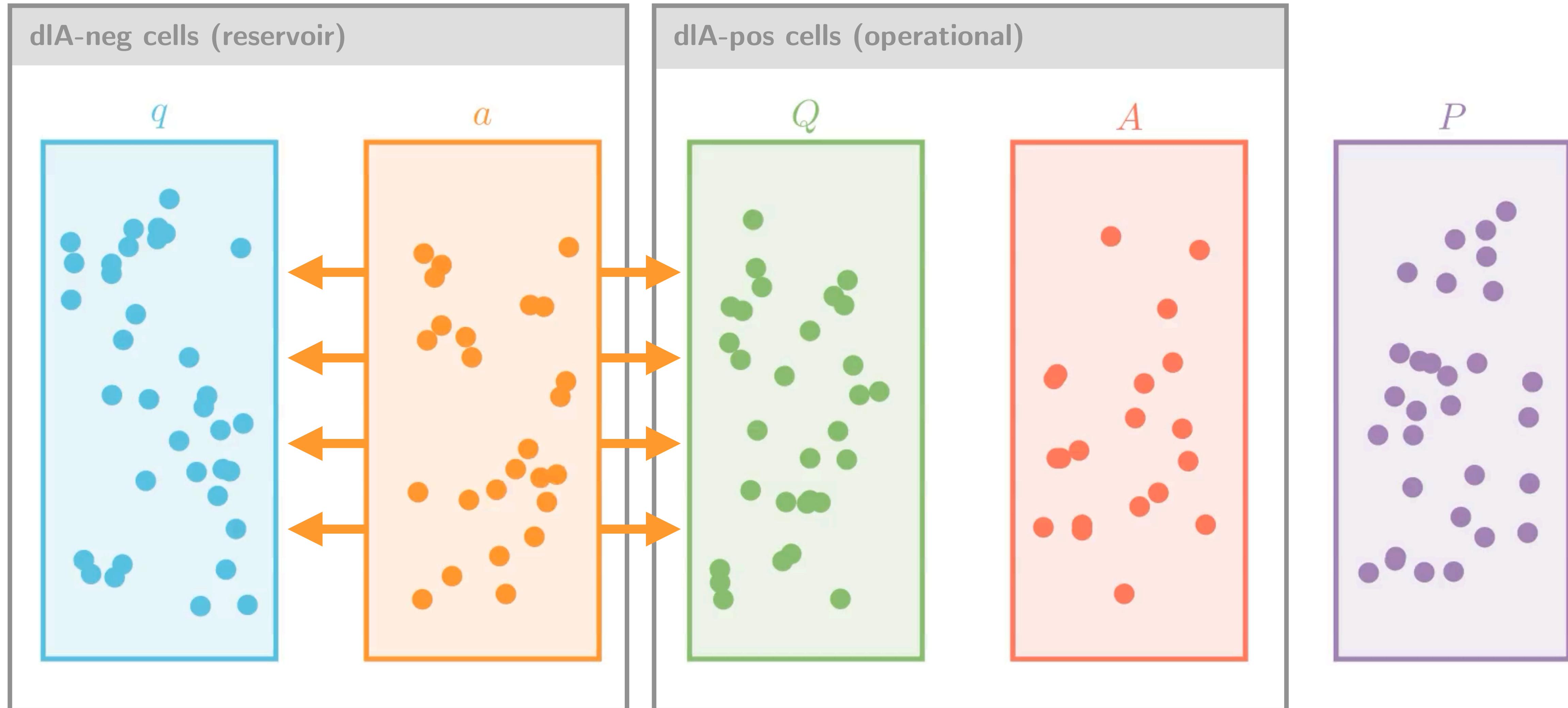
# Model Visualization: Compartmental evolution

Step 3.a: reservoir qNSC activate ( $q'(t) = -r_0q(t)$  and  $a'(t) = +r_0q(t)$ )



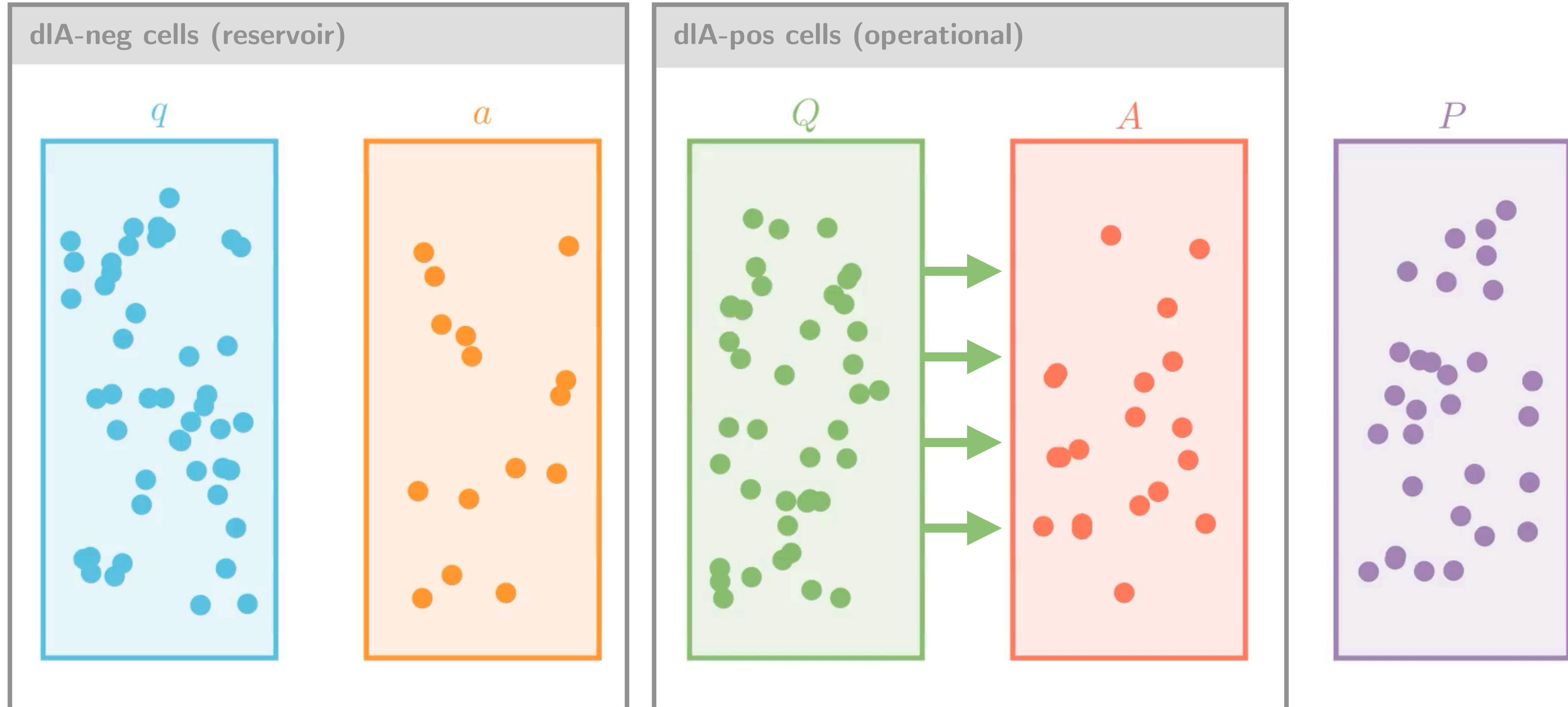
# Model Visualization: Compartmental evolution

Step 3.b: reservoir aNSC divide asymmetrically ( $a'(t) = -pa(t)$ ,  $q'(t) = +pa(t)$ , and  $Q'(t) = +pa(t)$ )



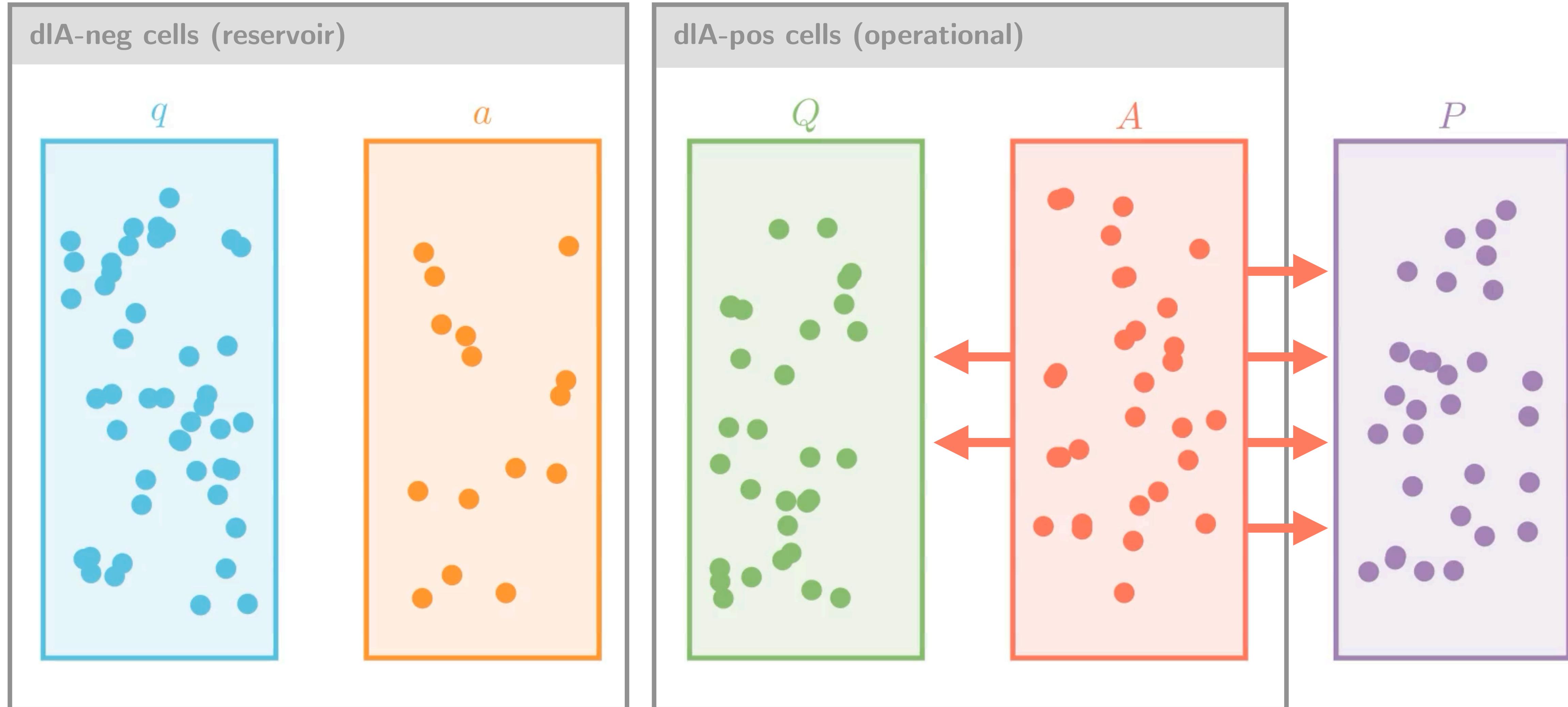
# Model Visualization: Compartmental evolution

Step 3.c: operational qNSC activate ( $Q'(t) = -r_1 Q(t)$ , and  $A(t) = +r_1 Q(t)$ )



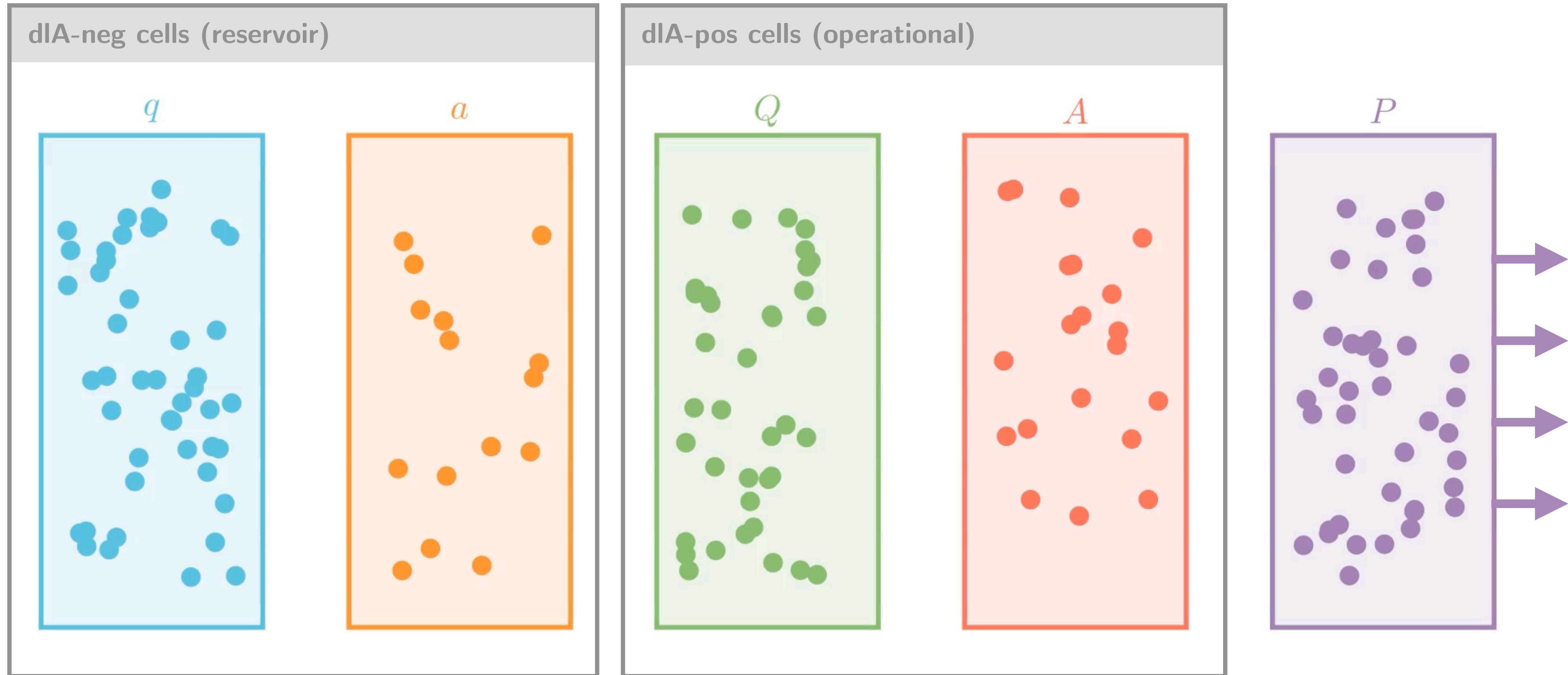
# Model Visualization: Compartmental evolution

Step 3.d: operational aNSC divide ( $A'(t) = - pA(t)$ ,  $Q'(t) = + 2b_0pA(t)$ , and  $P'(t) = + 2(1 - b_0)pA(t)$ )

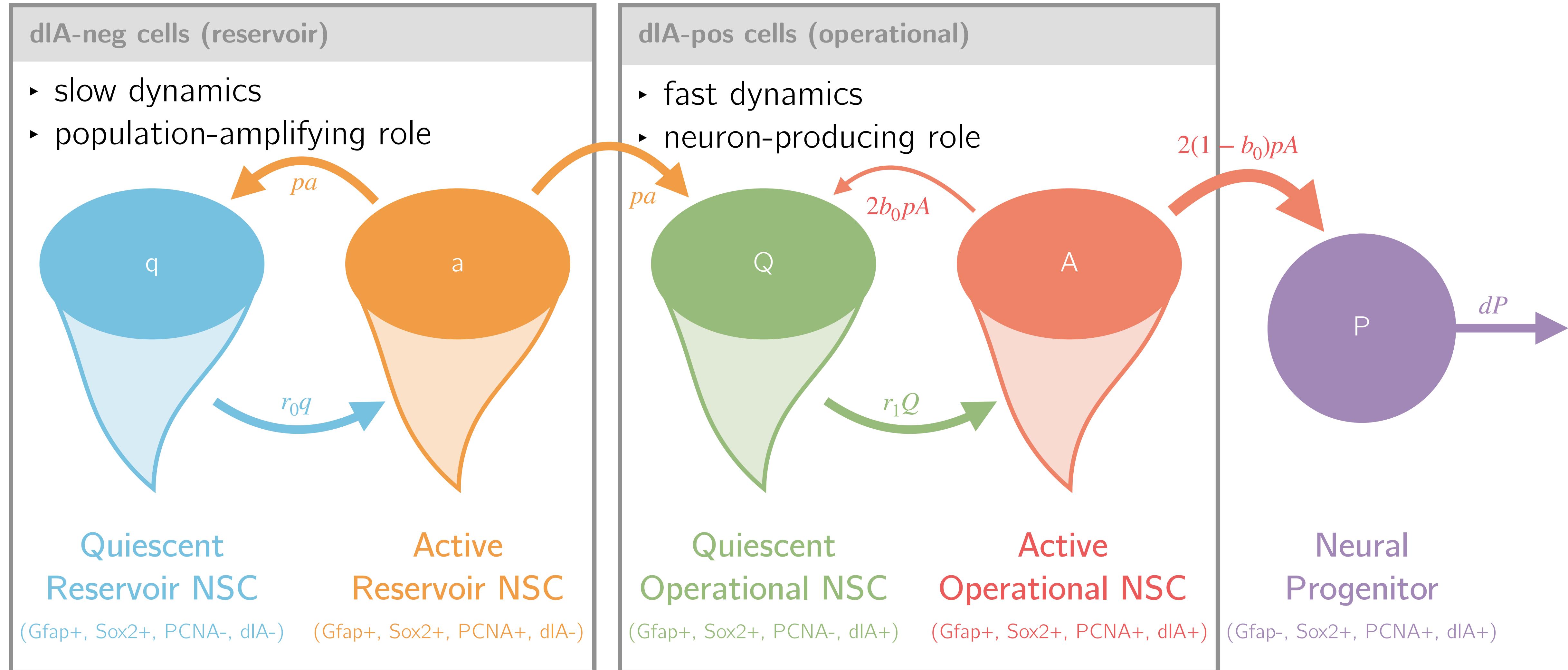


# Model Visualization: Compartmental evolution

Step 3.e: neural Progenitors delaminate ( $P'(t) = - dP(t)$ )



# Model Equations (Summary)



# Model Equations

dIA-neg cells (reservoir)



$$q'(t) = -r_0 q(t) + p_a(t),$$



$$a'(t) = r_0 q(t) - p_a(t),$$

dIA-pos cells (operational)



$$Q'(t) = -r_1 Q(t) + 2b_0 p_A(t) + p_a(t),$$



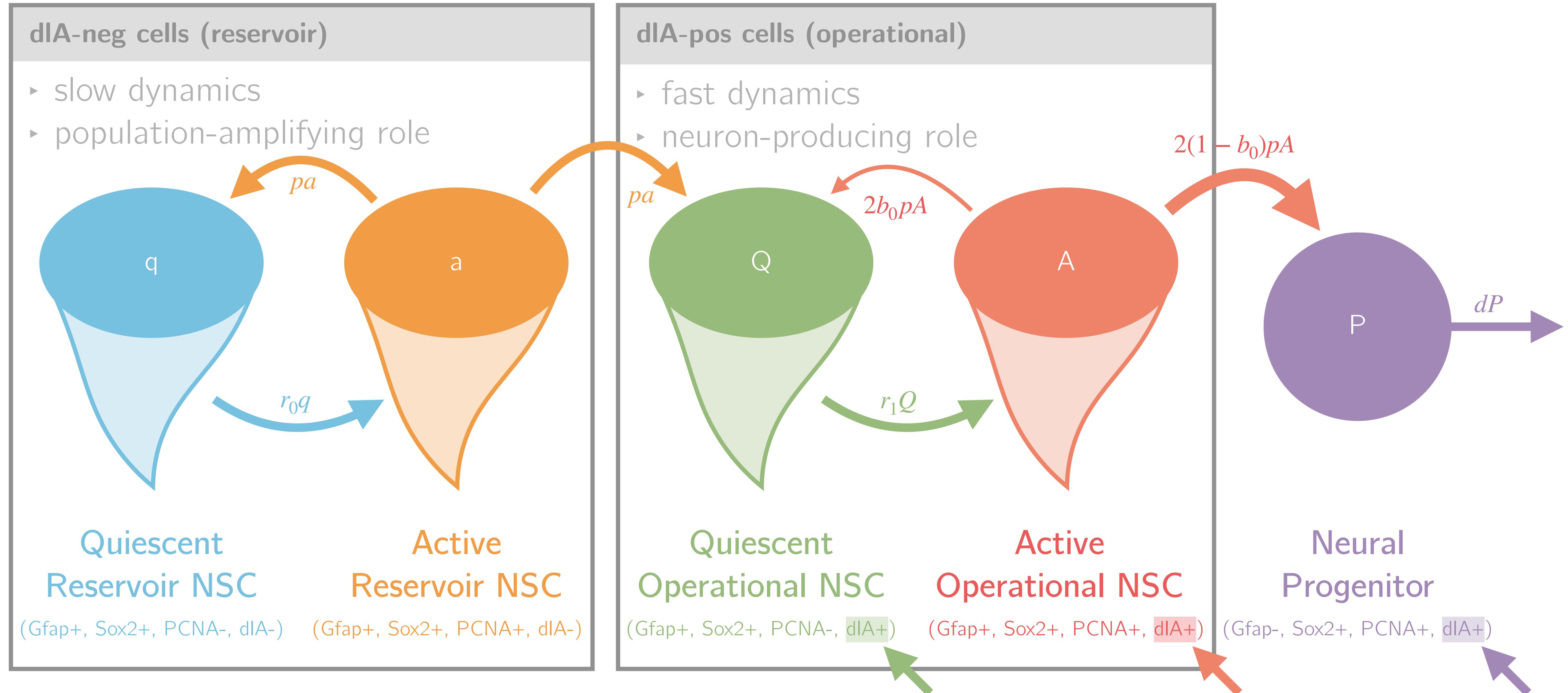
$$A'(t) = r_1 Q(t) - p_A(t),$$



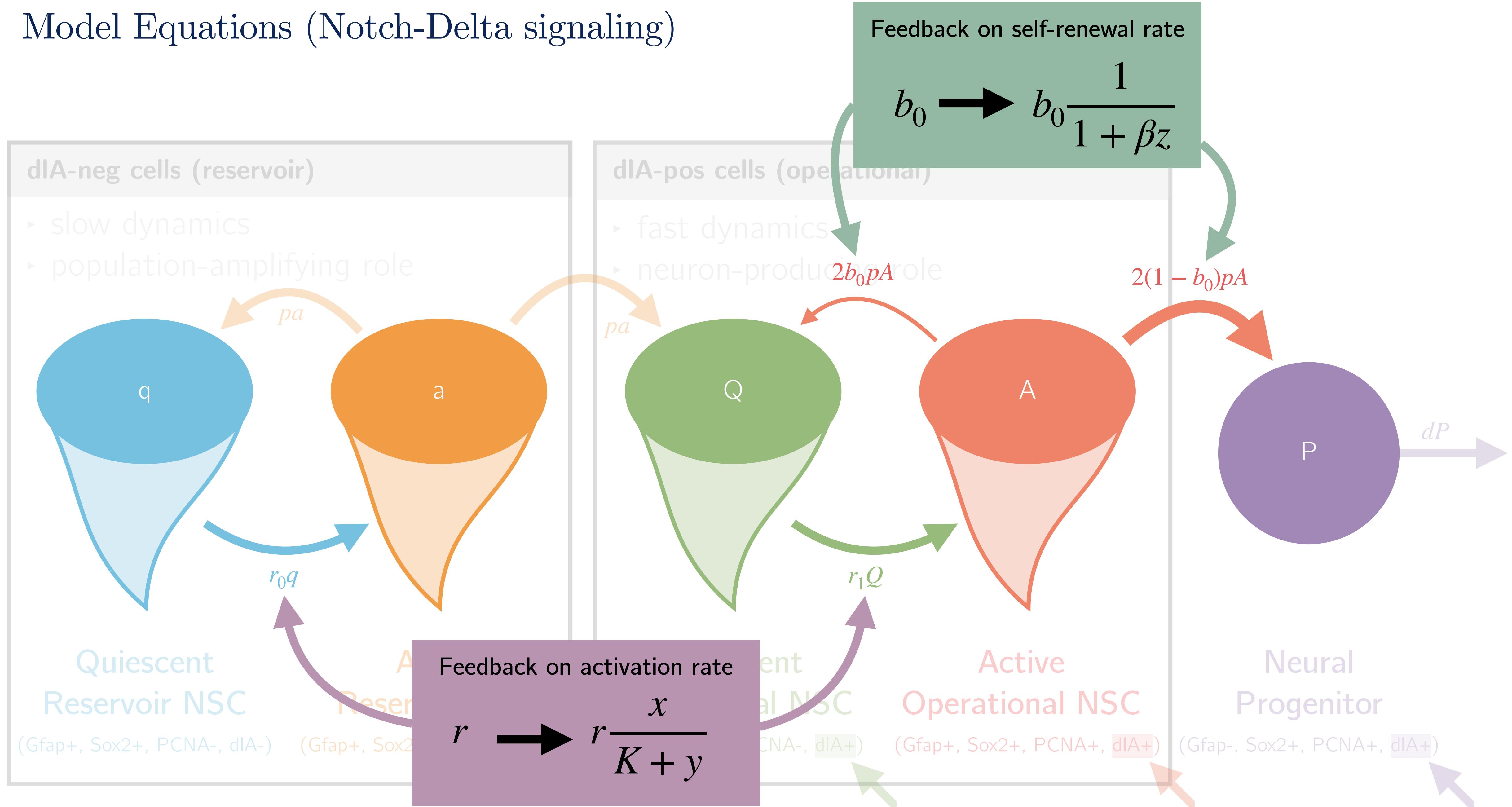
$$P'(t) = 2(1 - b_0)p_A(t) - dP(t)$$

# Modeling Notch-delta Signaling

# Model Equations (Notch-Delta signaling)



# Model Equations (Notch-Delta signaling)



dIA-neg cells (reservoir)



$$q'(t) = -r_0 \frac{x(t)}{K + y(t)} q(t) + p_a(t),$$



$$a'(t) = r_0 \frac{x(t)}{K + y(t)} q(t) - p_a(t),$$

dIA-pos cells (operational)



$$Q'(t) = -r_1 \frac{x(t)}{K + y(t)} Q(t) + 2b_0 \frac{1}{1 + \beta z(t)} p_A(t) + p_a(t),$$



$$A'(t) = r_1 \frac{x(t)}{K + y(t)} Q(t) - p_A(t),$$



$$P'(t) = 2 \left( 1 - b_0 \frac{1}{1 + \beta z(t)} \right) p_A(t) - d_P(t)$$

dIA-neg cells (reservoir)



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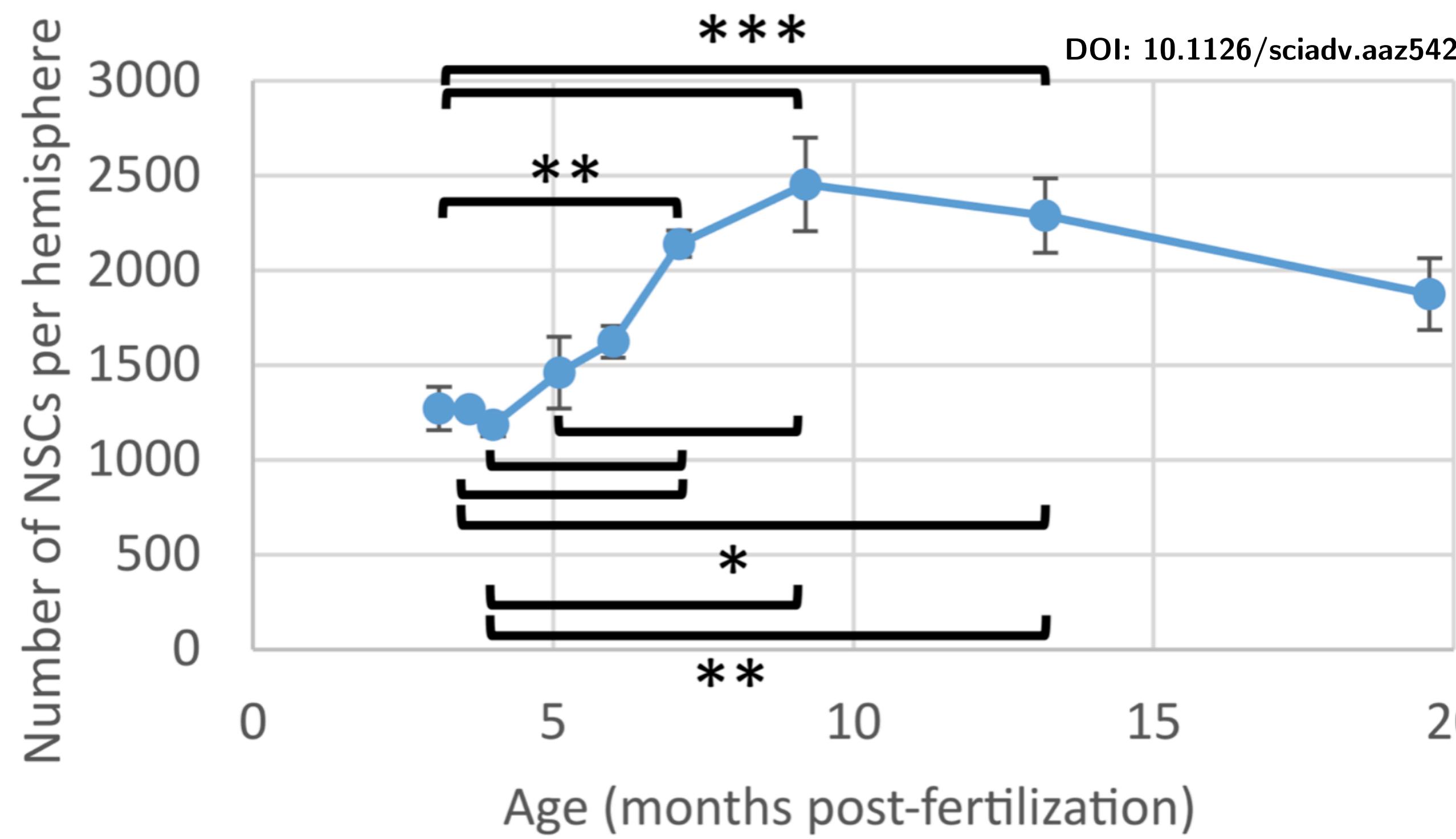
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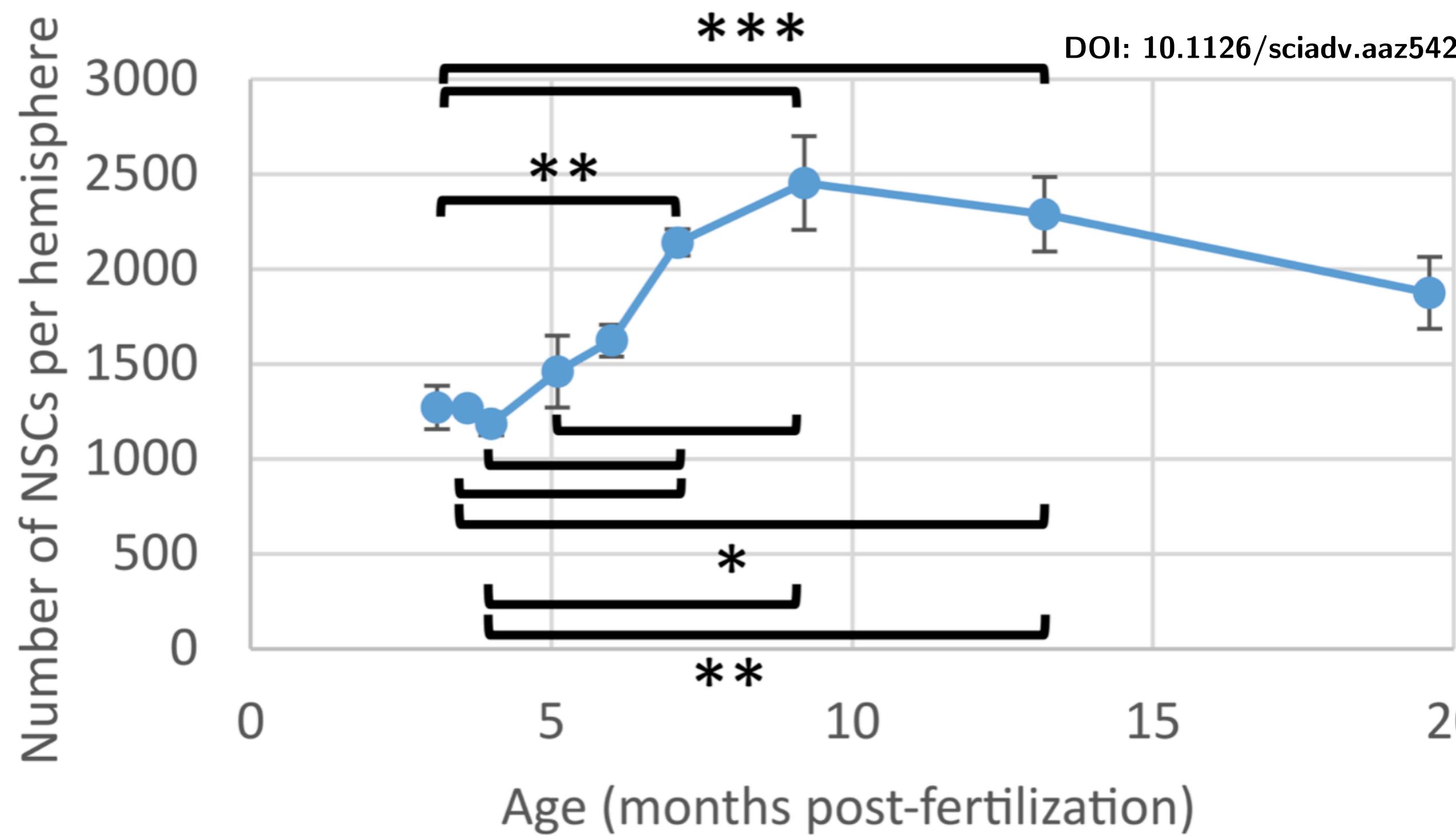
**Mouse  
System**

## Calibration: Investigating the data



- a. Average number of cells per hemisphere ( $n=7$  brains), reported by Sox2 expression (NSCs + NPs) over 500 days

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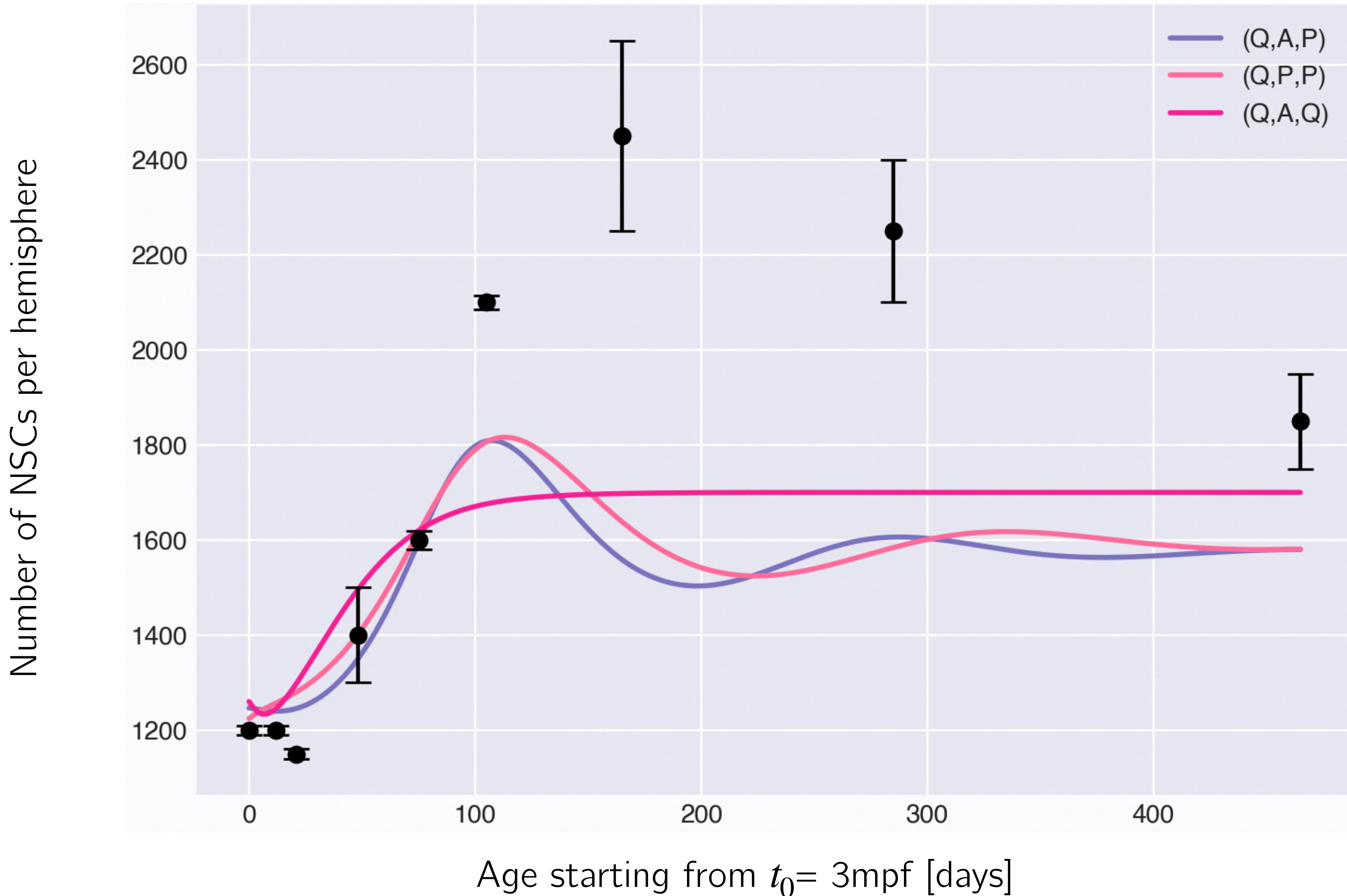


- a. Average number of cells per hemisphere ( $n=7$  brains), reported by Sox2 expression (NSCs + NPs) over 500 days

Provides a measure of

$$S(t) := q(t) + a(t) + Q(t) + A(t) + P(t)$$

# Comparison of multiple plausible scenarios for Notch-DeltaA signaling

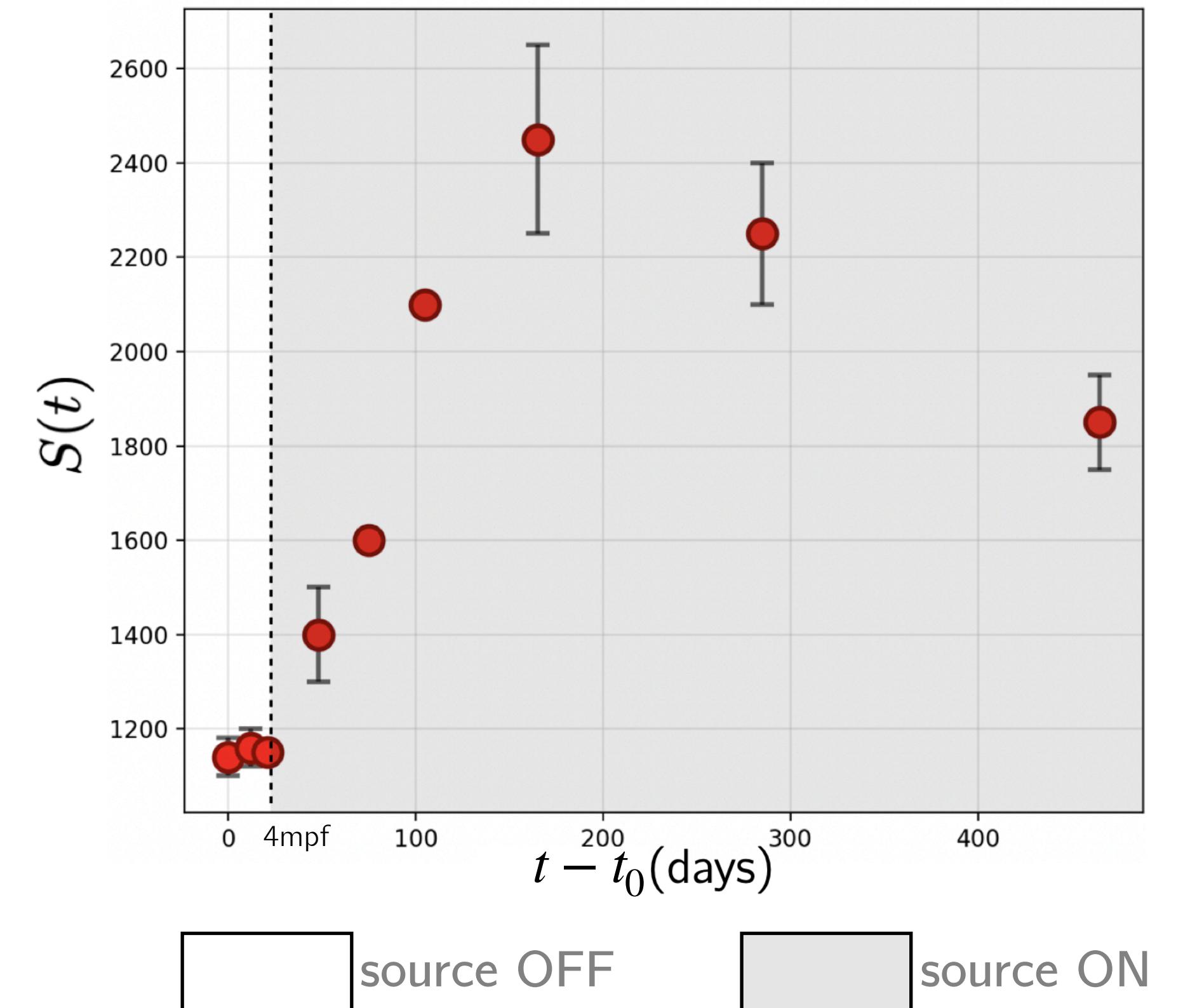


# Modeling the source

## Investigation: Modeling the source

Total population

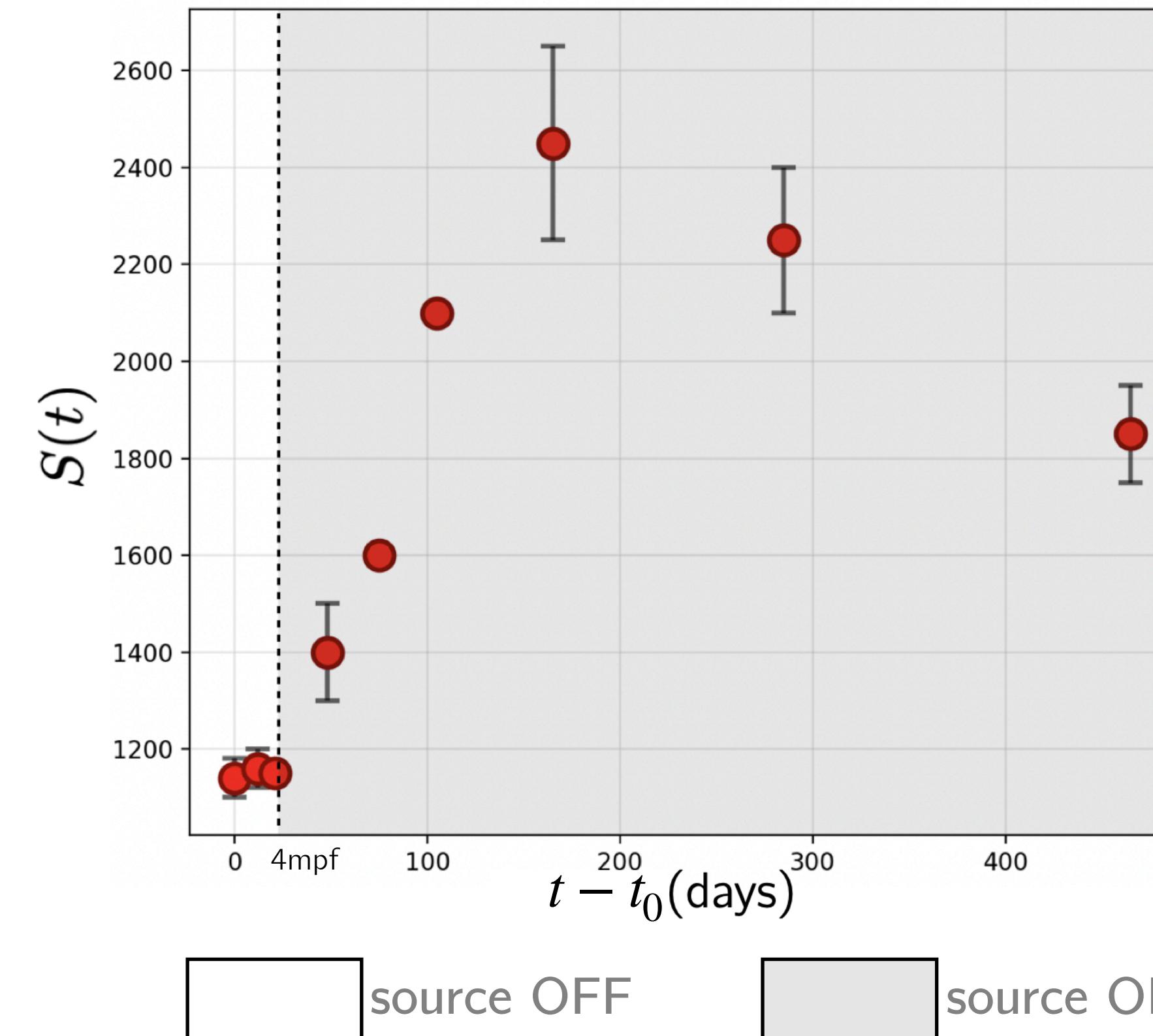
$$S(t) = (q + a + Q + A + P)(t)$$



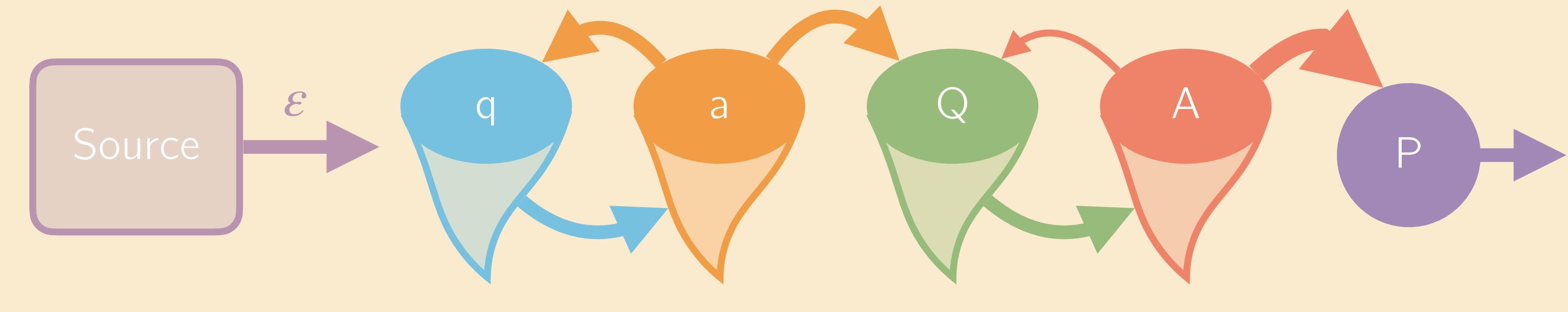
## Investigation: Modeling the source

Total population

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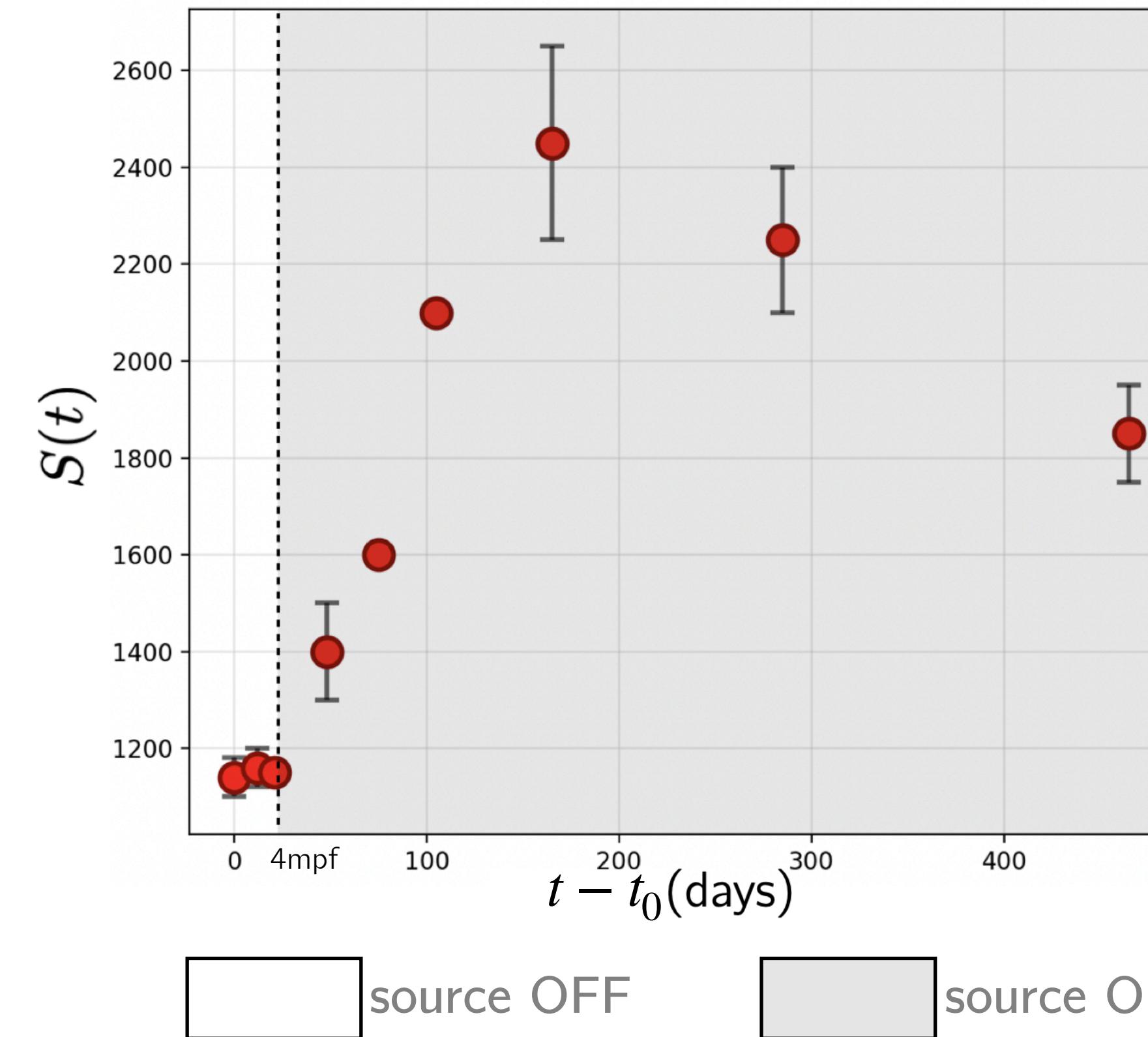
- a. Introduce cells in the population, at the top of the hierarchy, and at a small constant rate  $\varepsilon > 0$ .



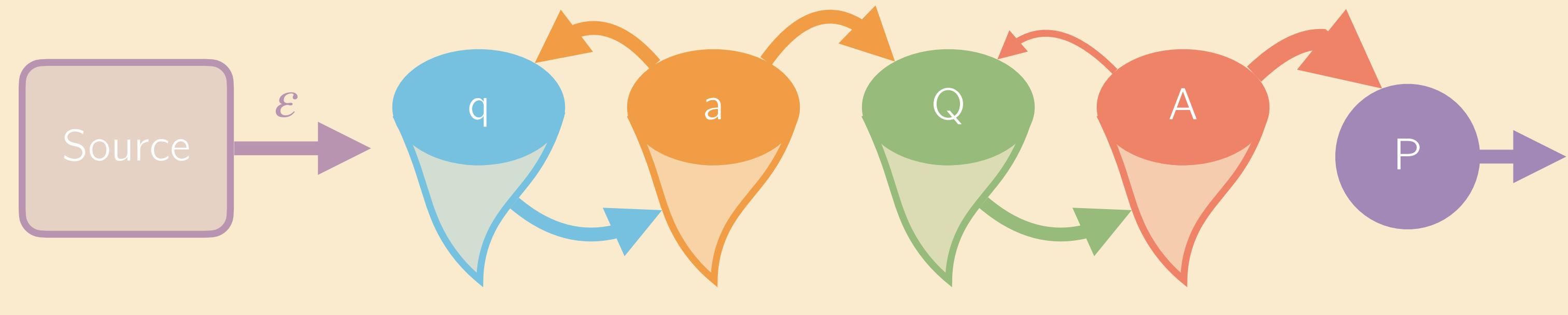
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Total population

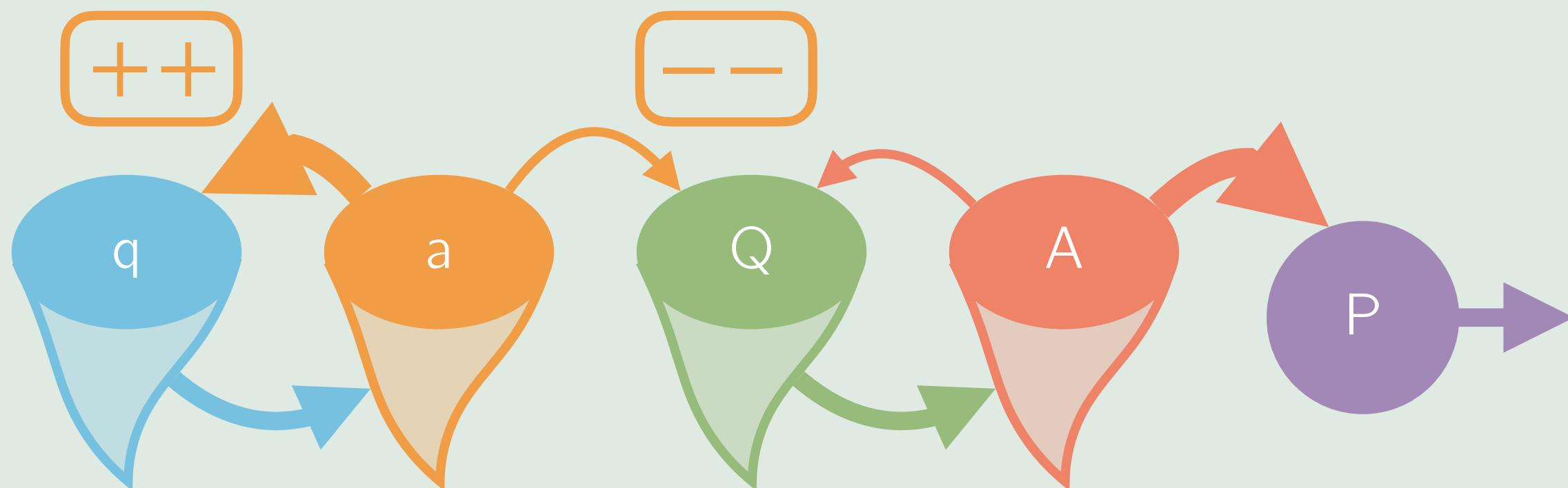
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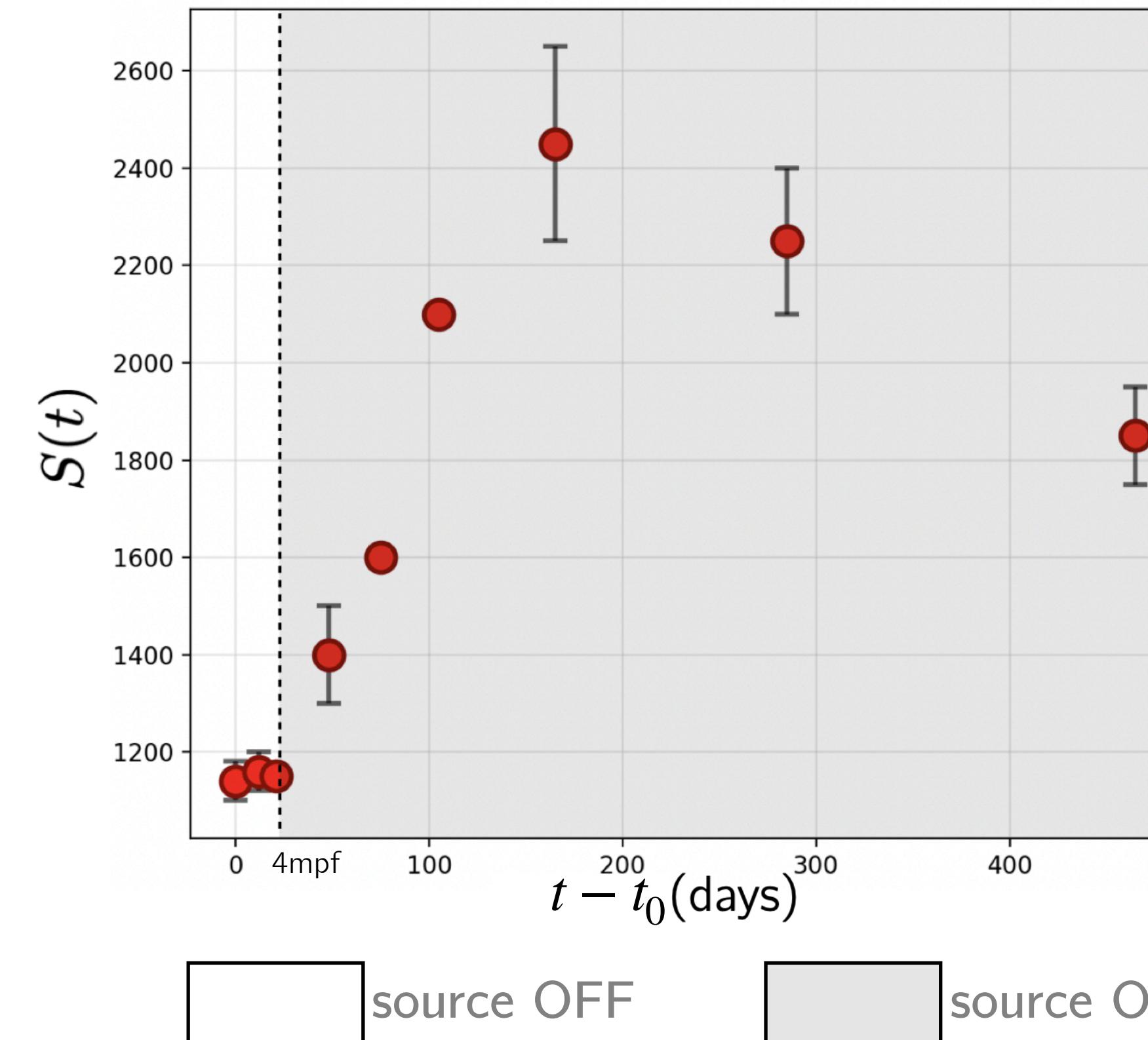


- b. Slightly increase the self-renewal rate of reservoir aNSCs by  $\varepsilon > 0$ .

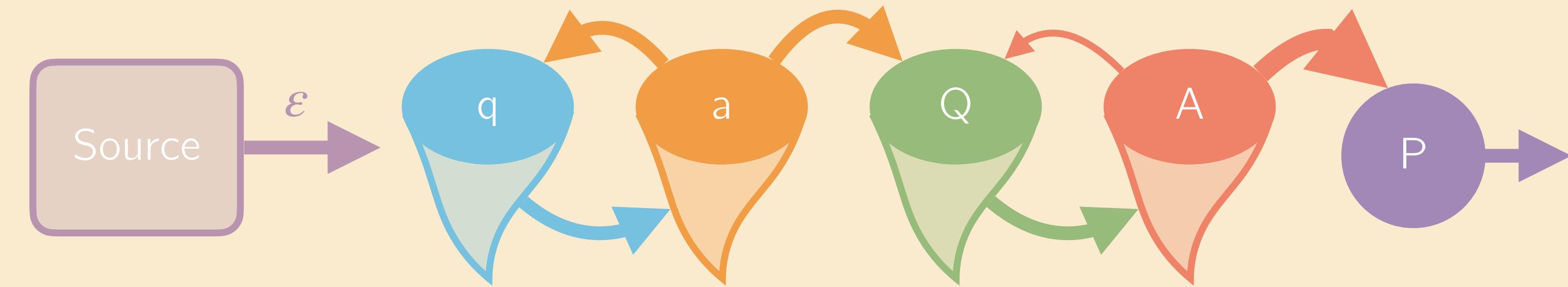


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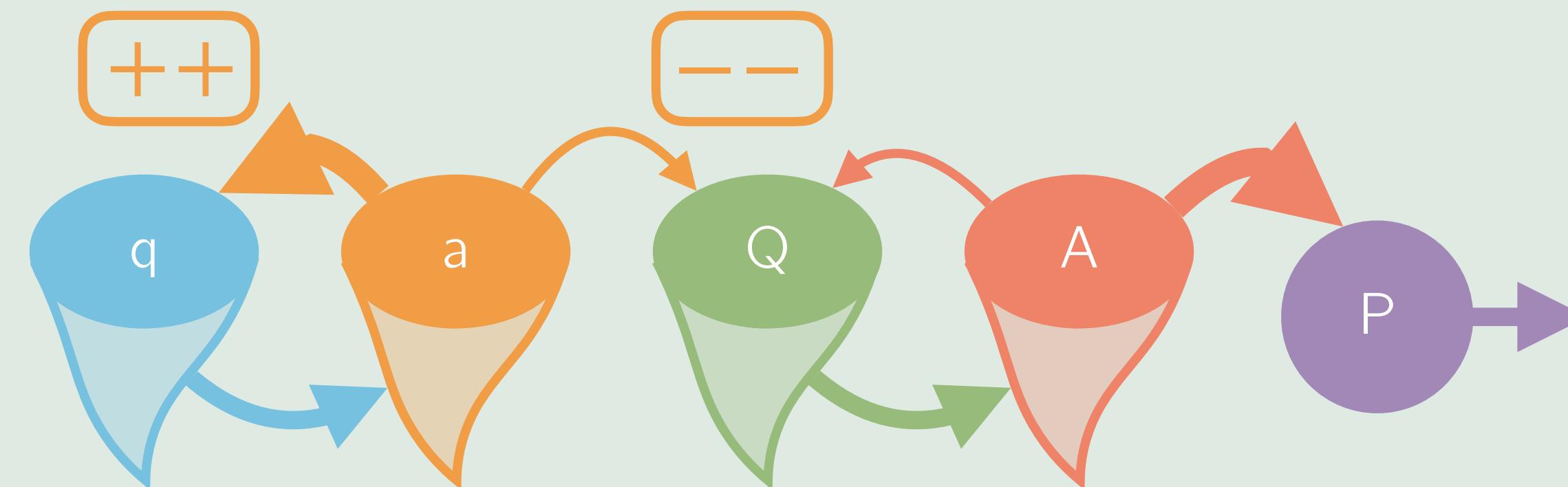
Total population  
 $S(t) = (q + a + Q + A + P)(t)$



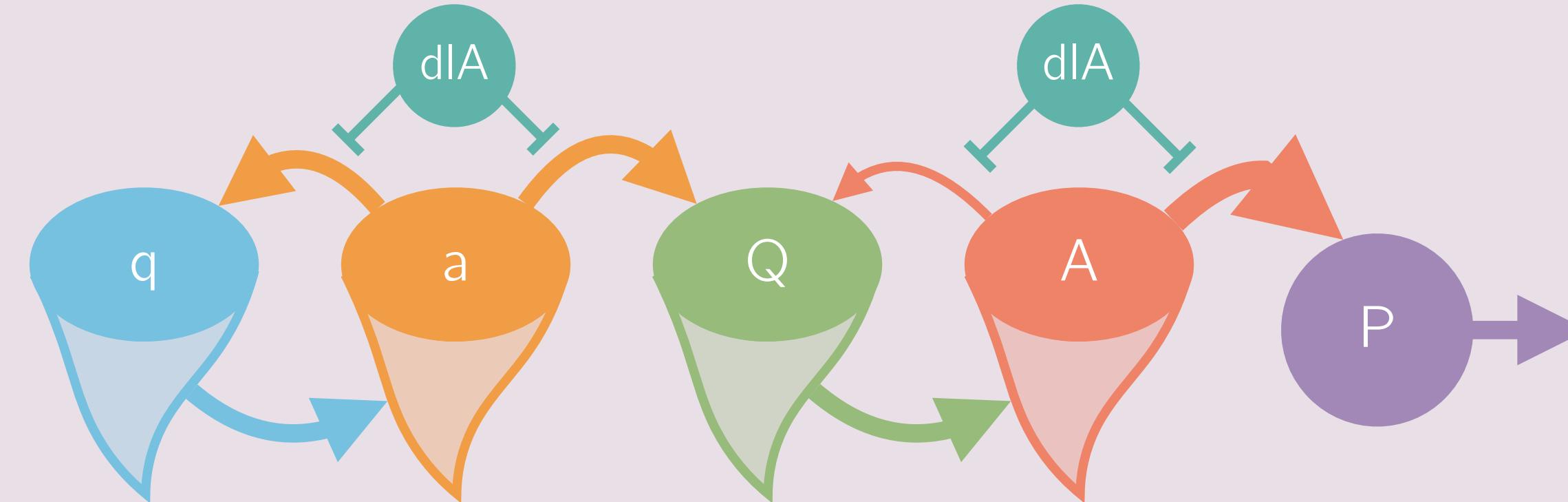
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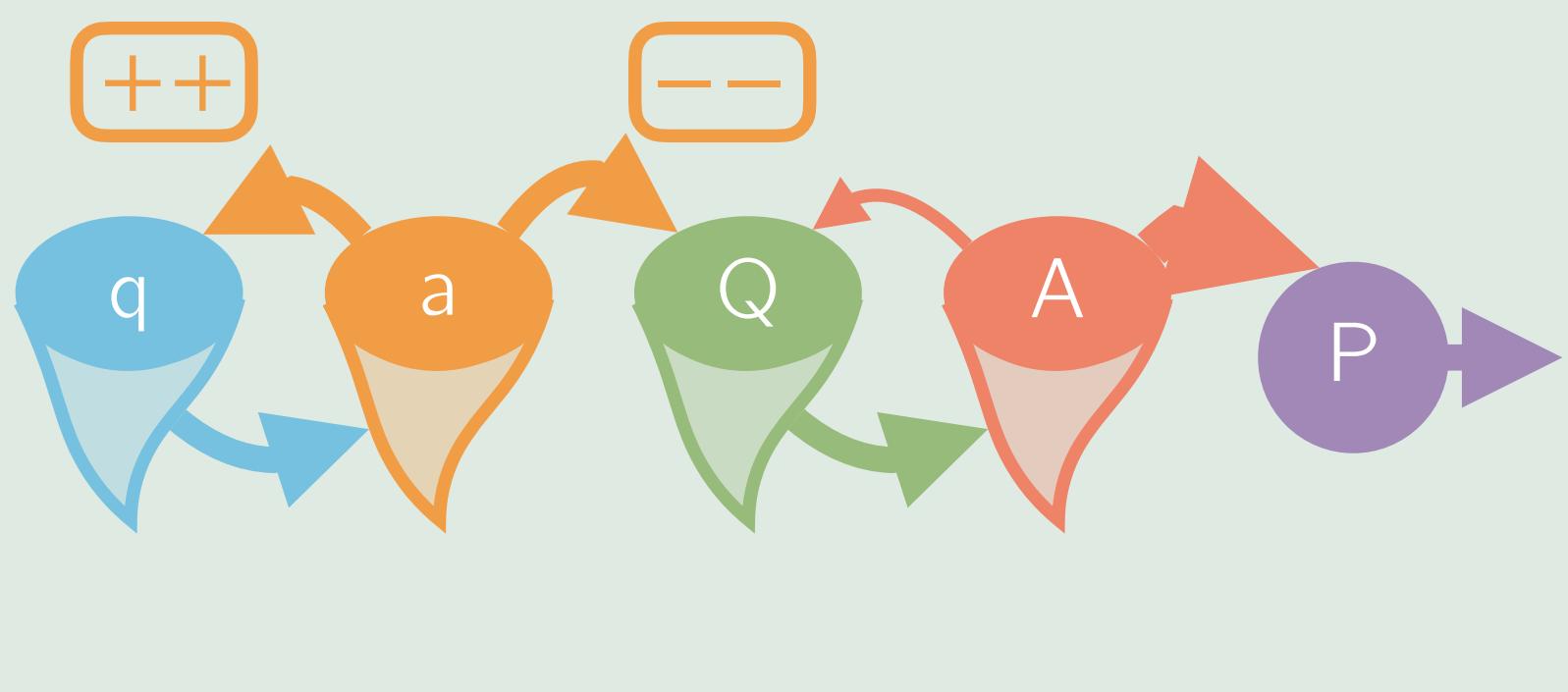
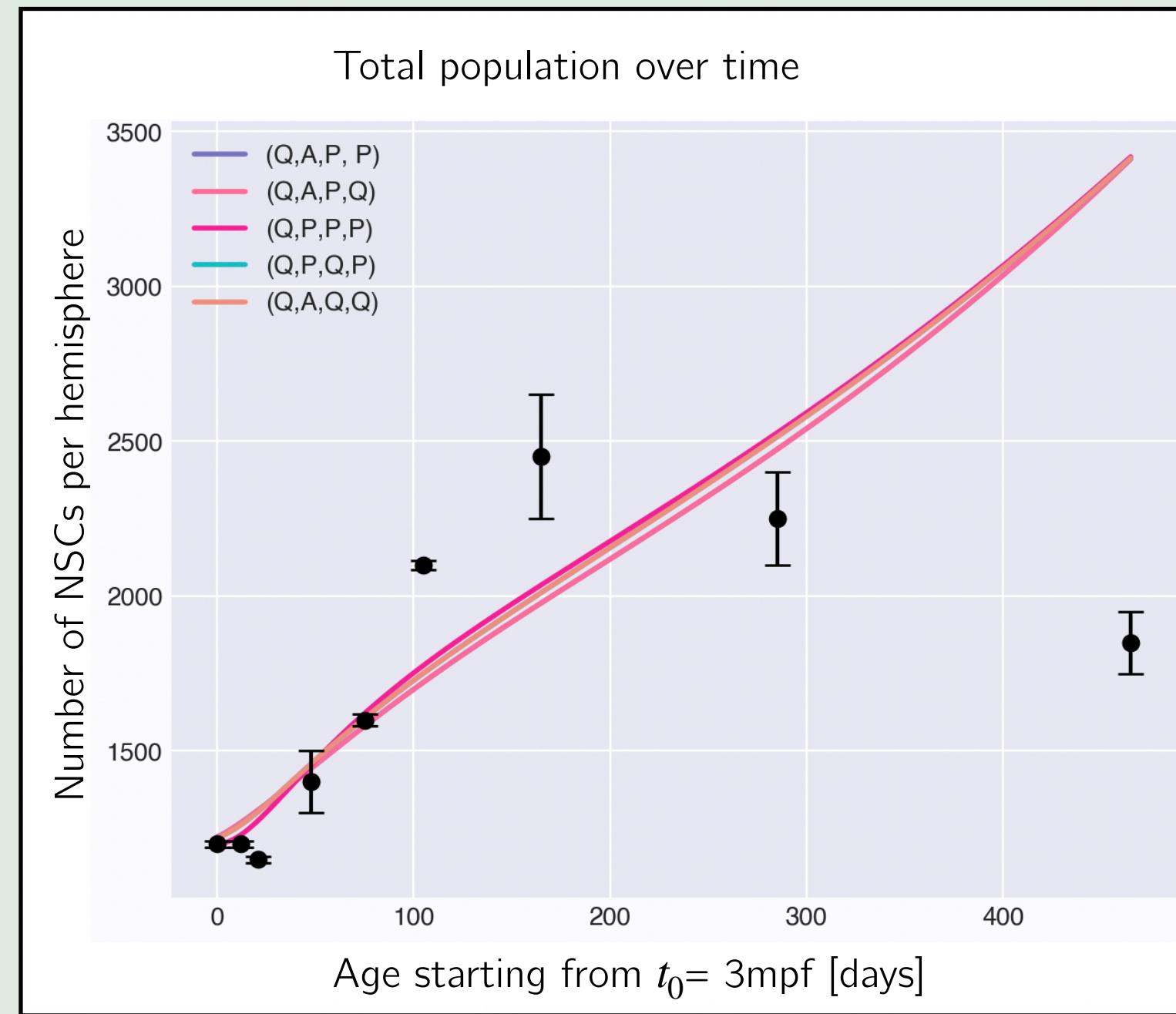
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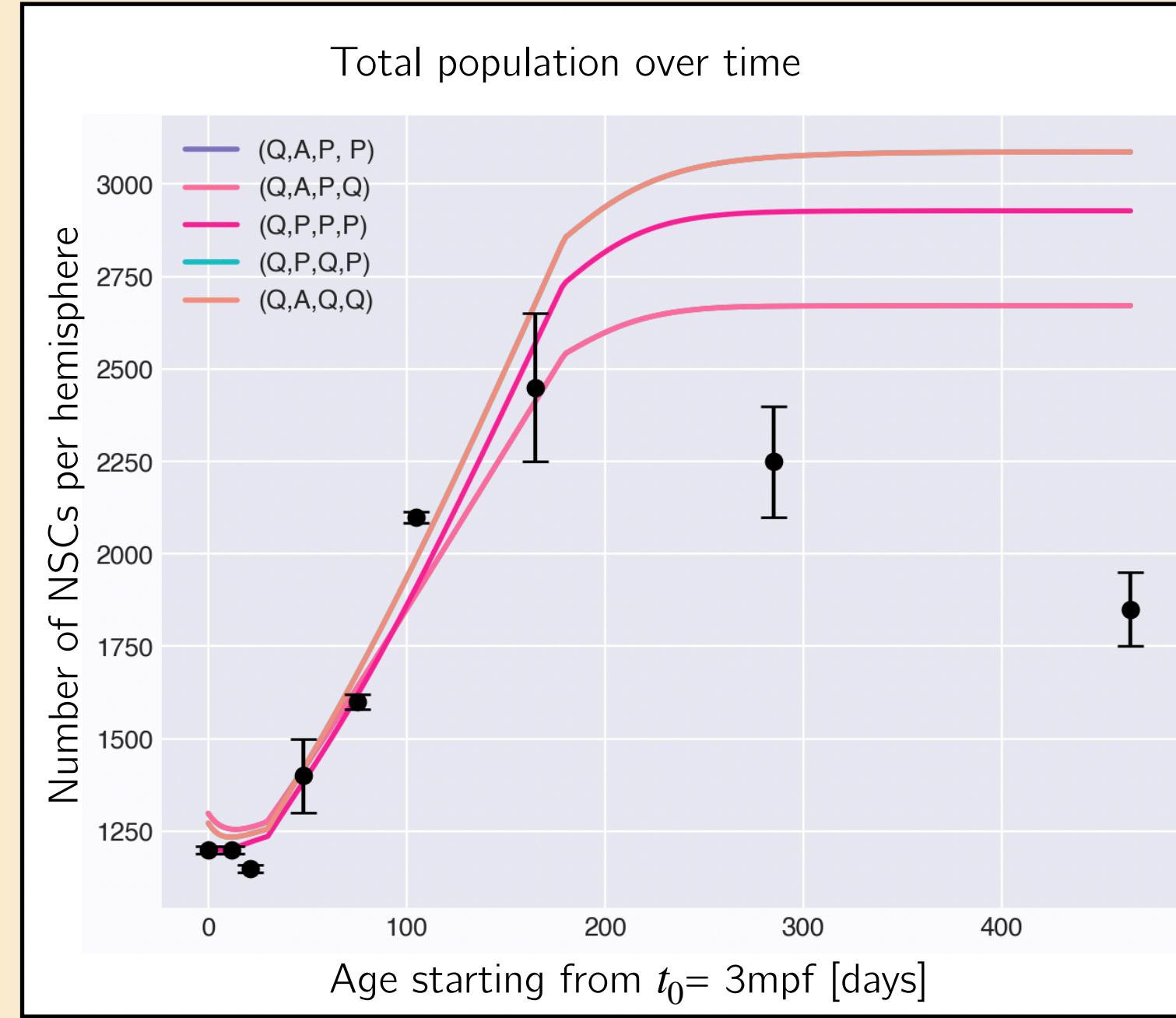
c. Notch-Delta signaling controls the self-renewal rate of aNSCs



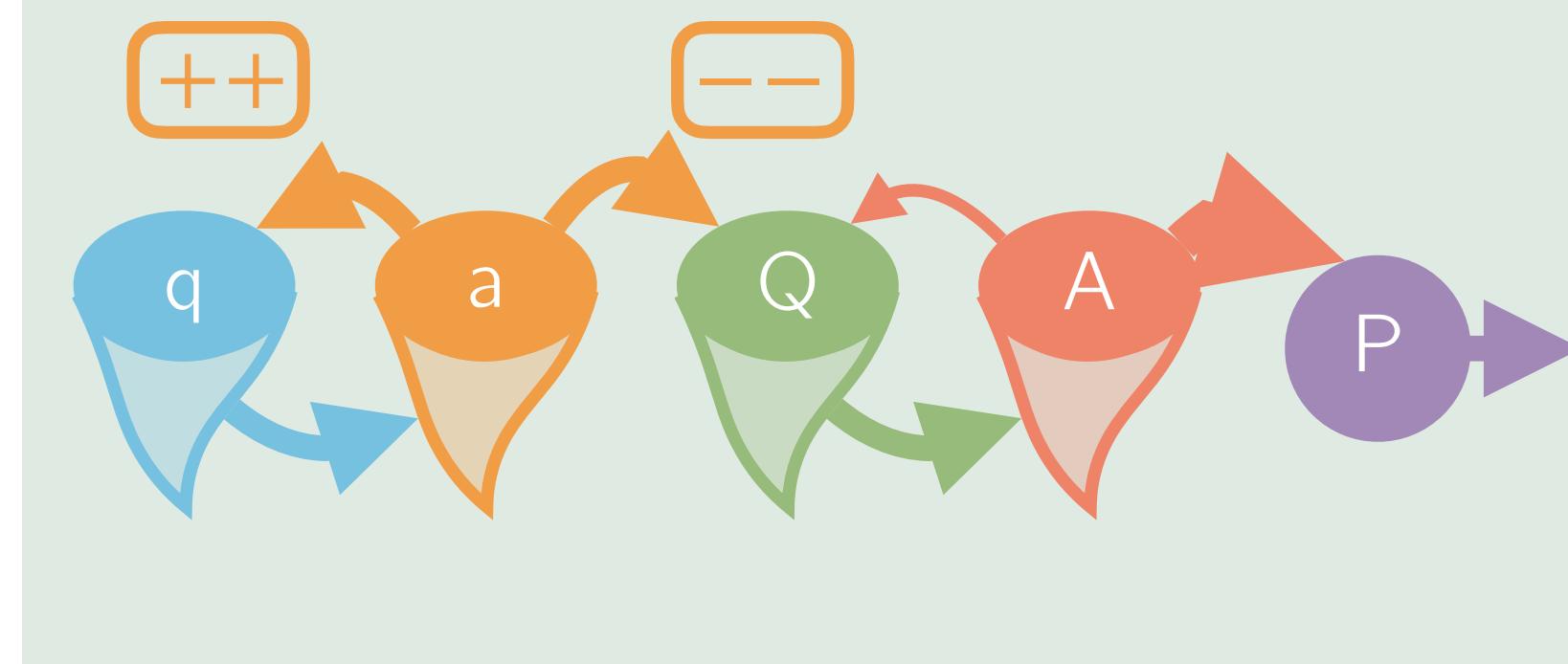
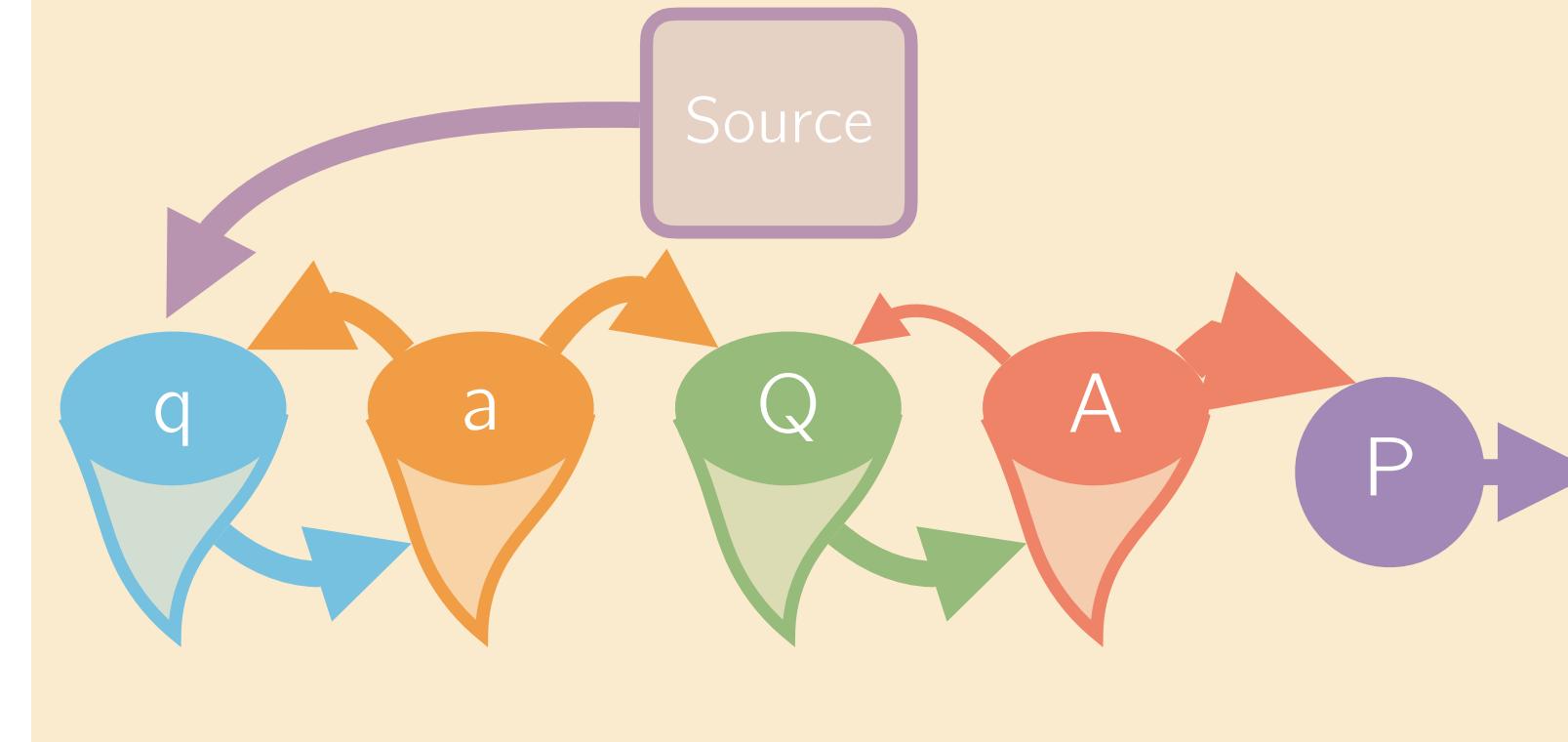
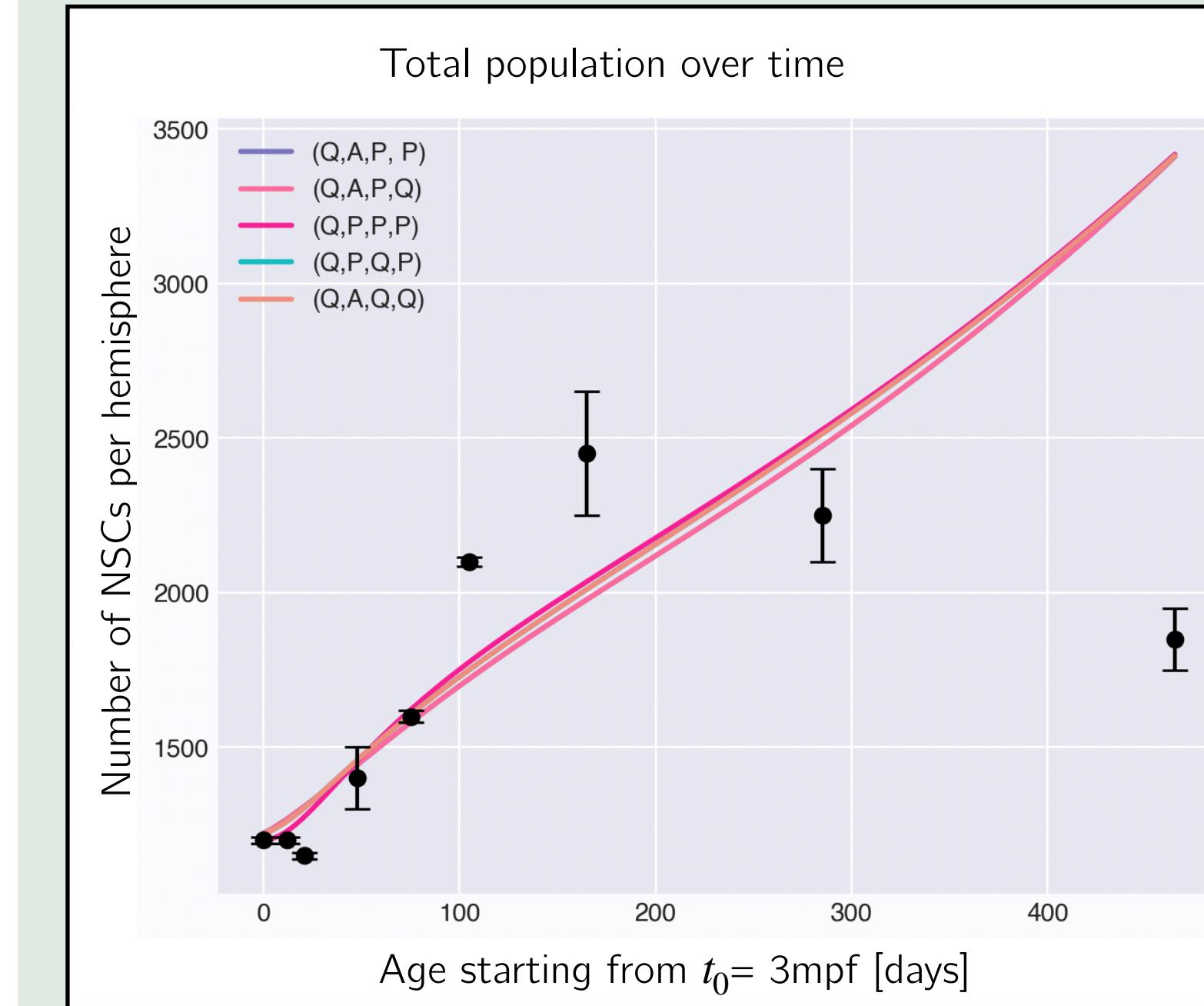
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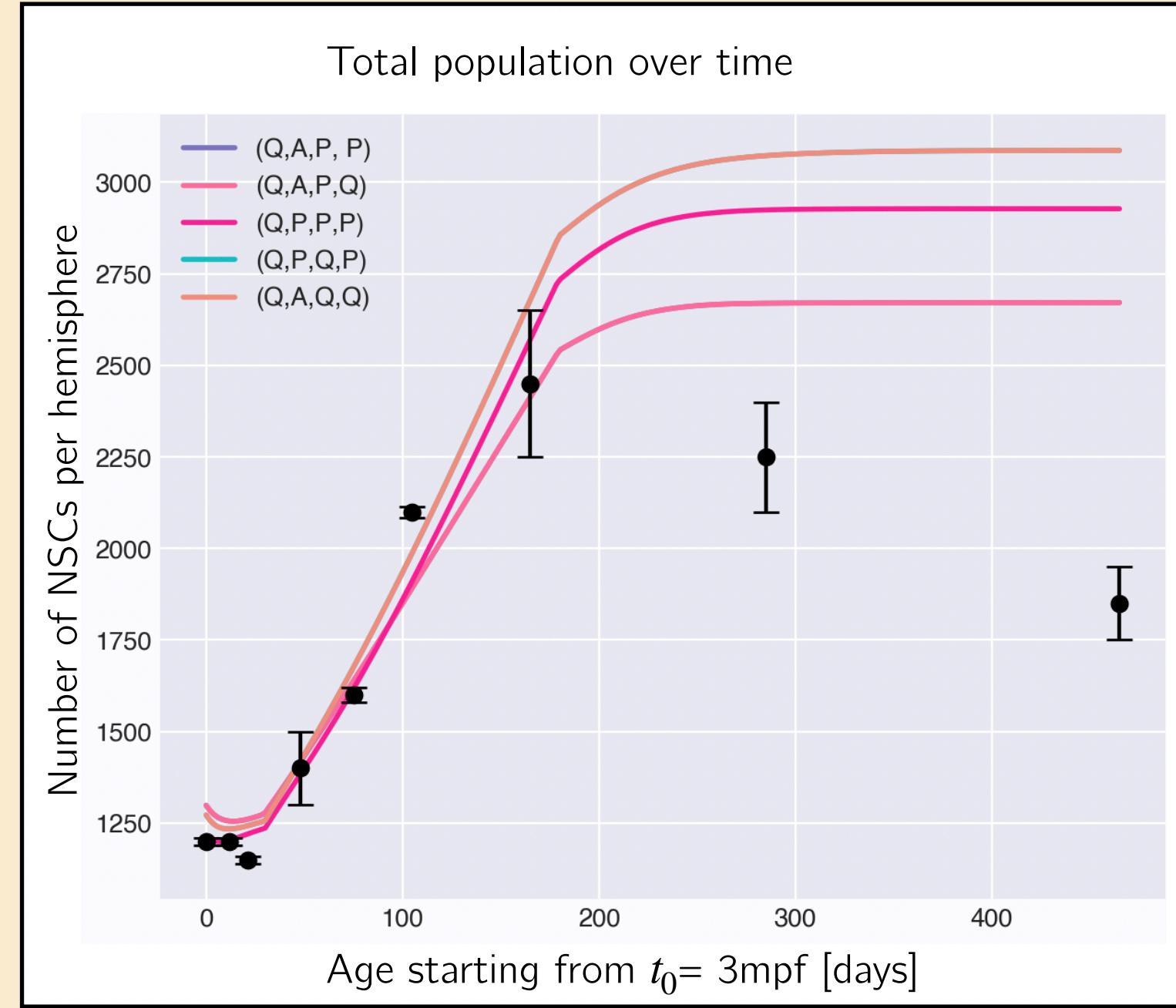
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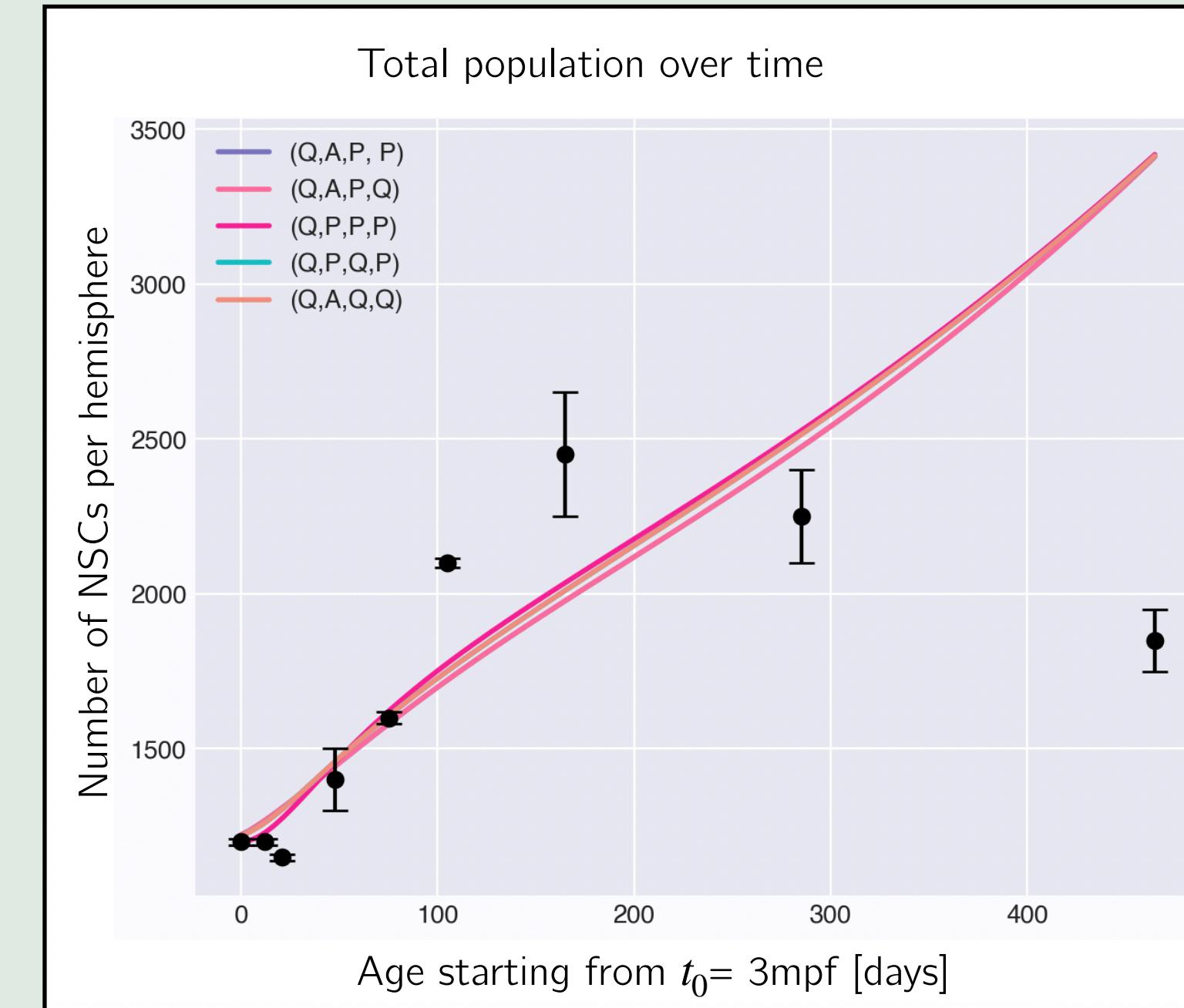
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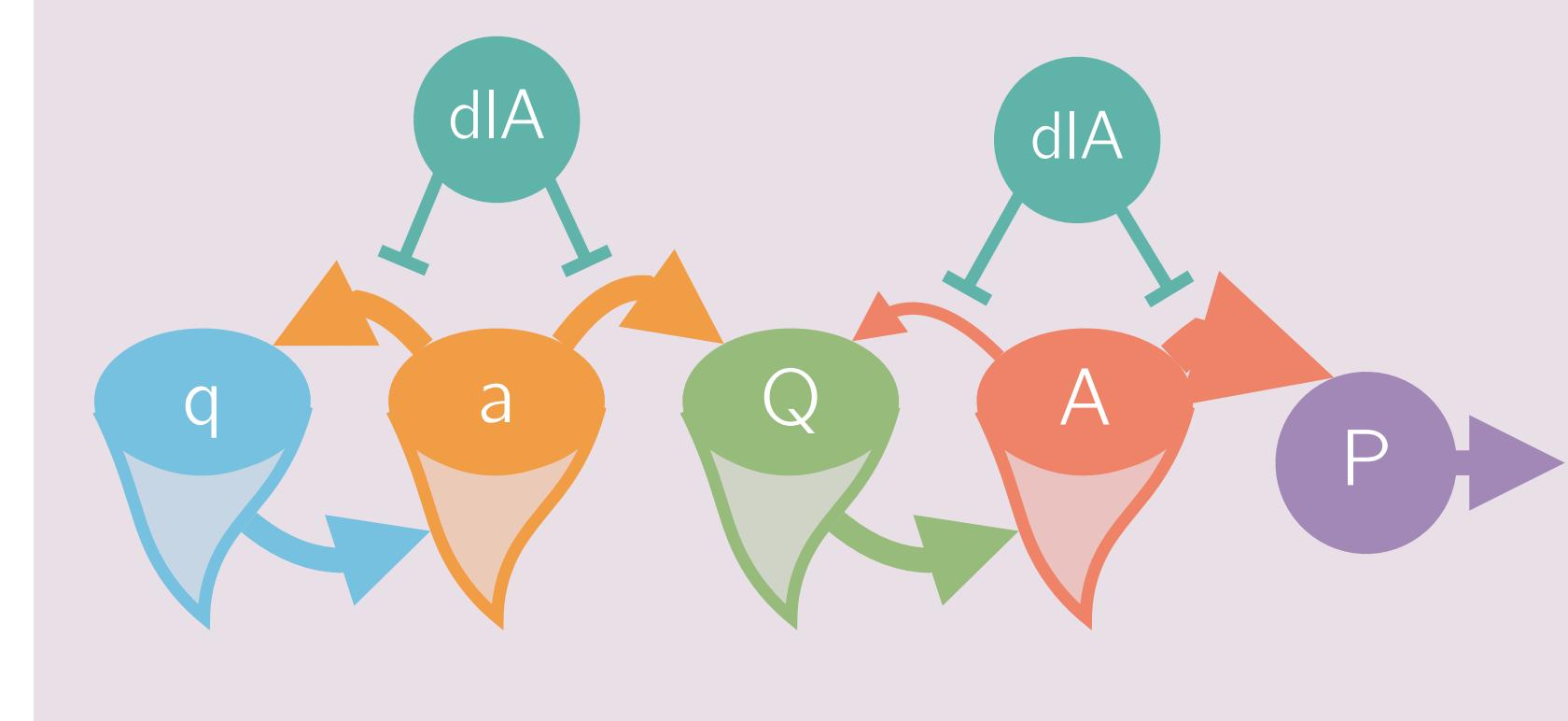
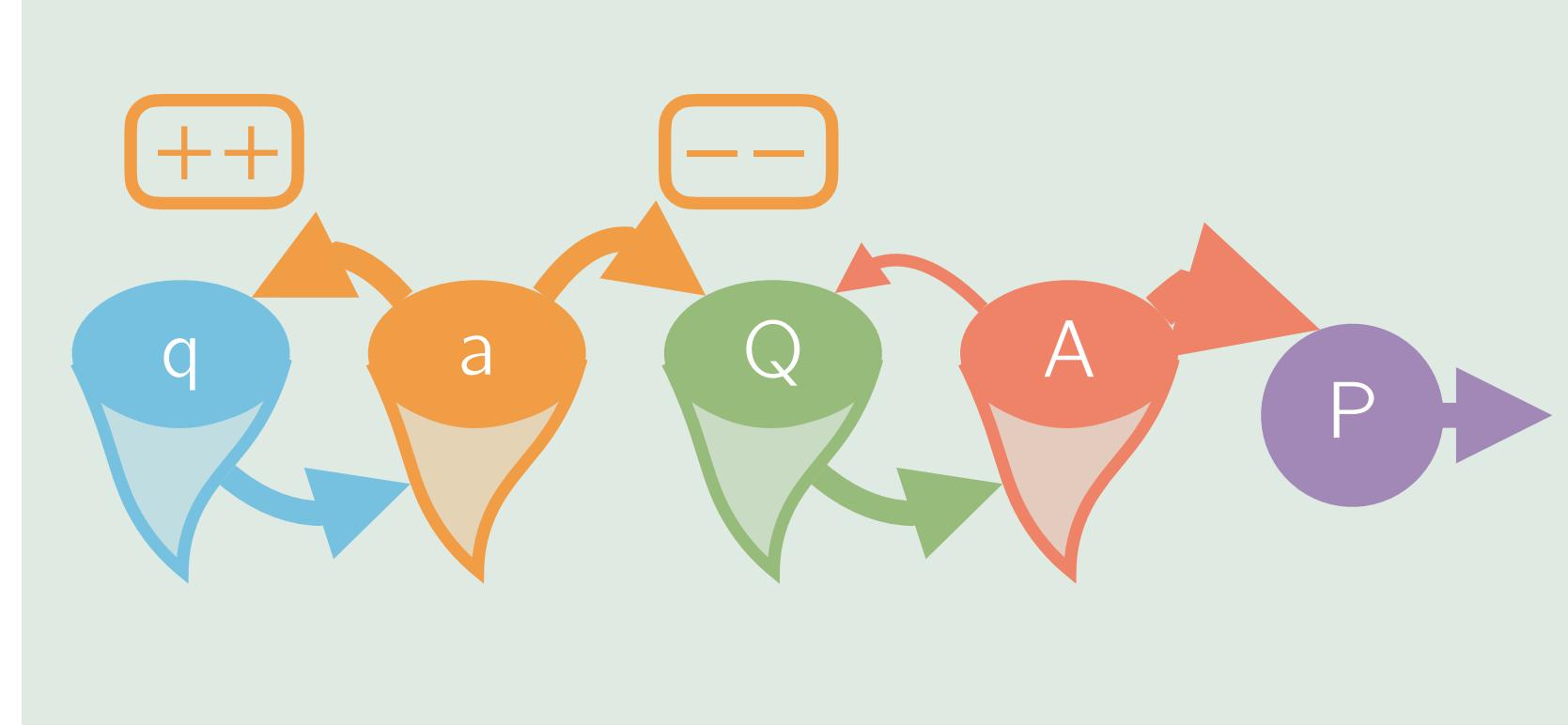
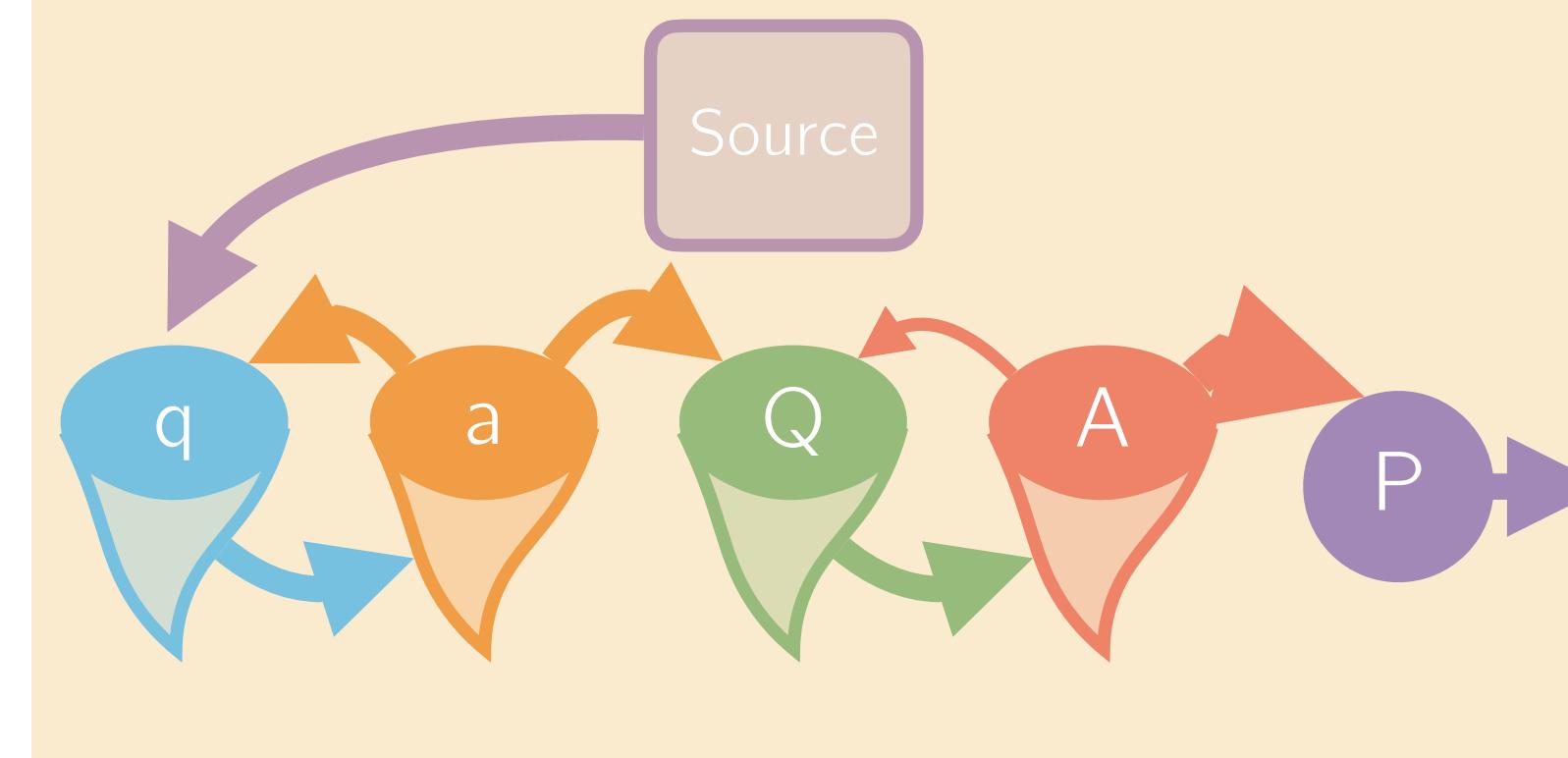
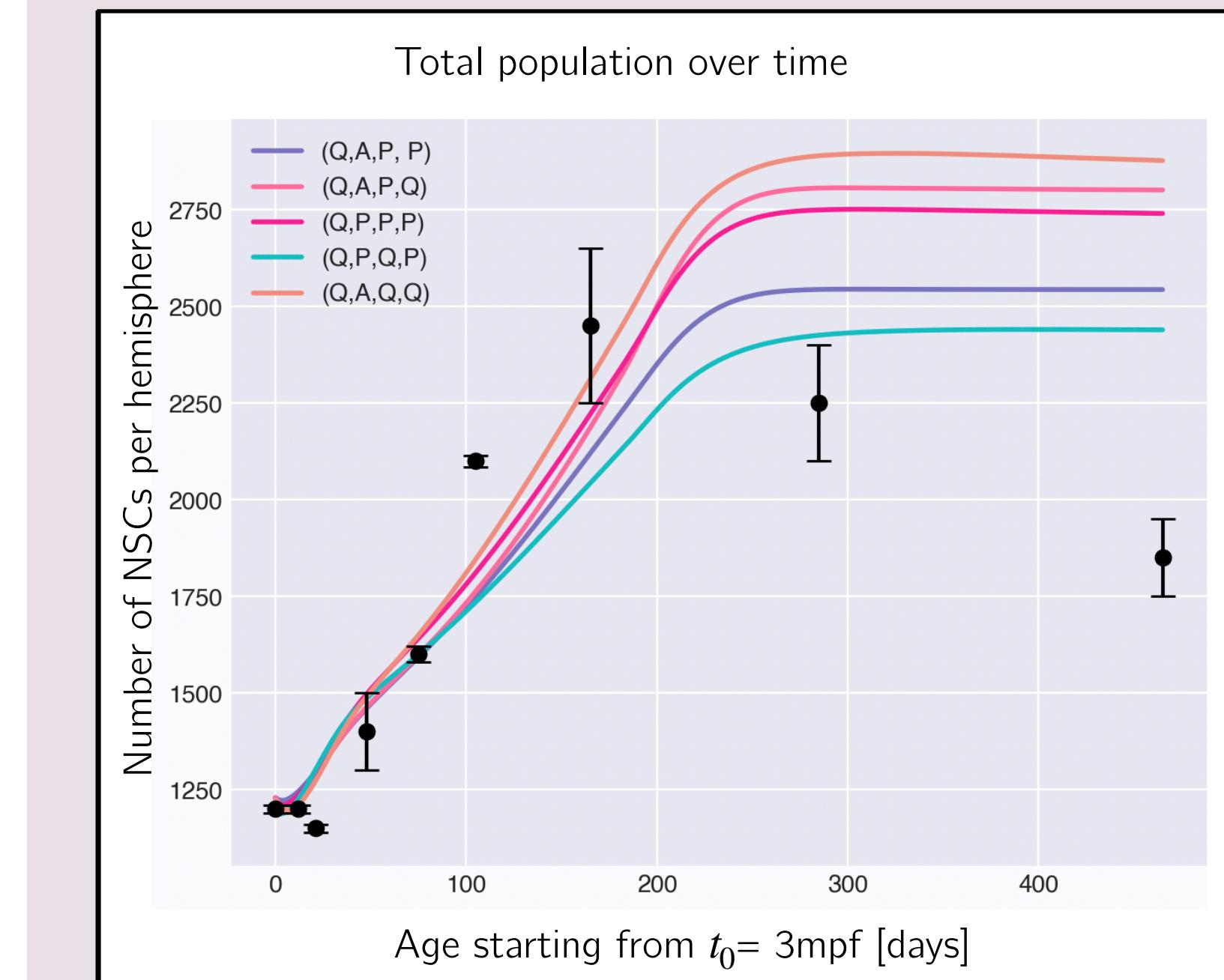
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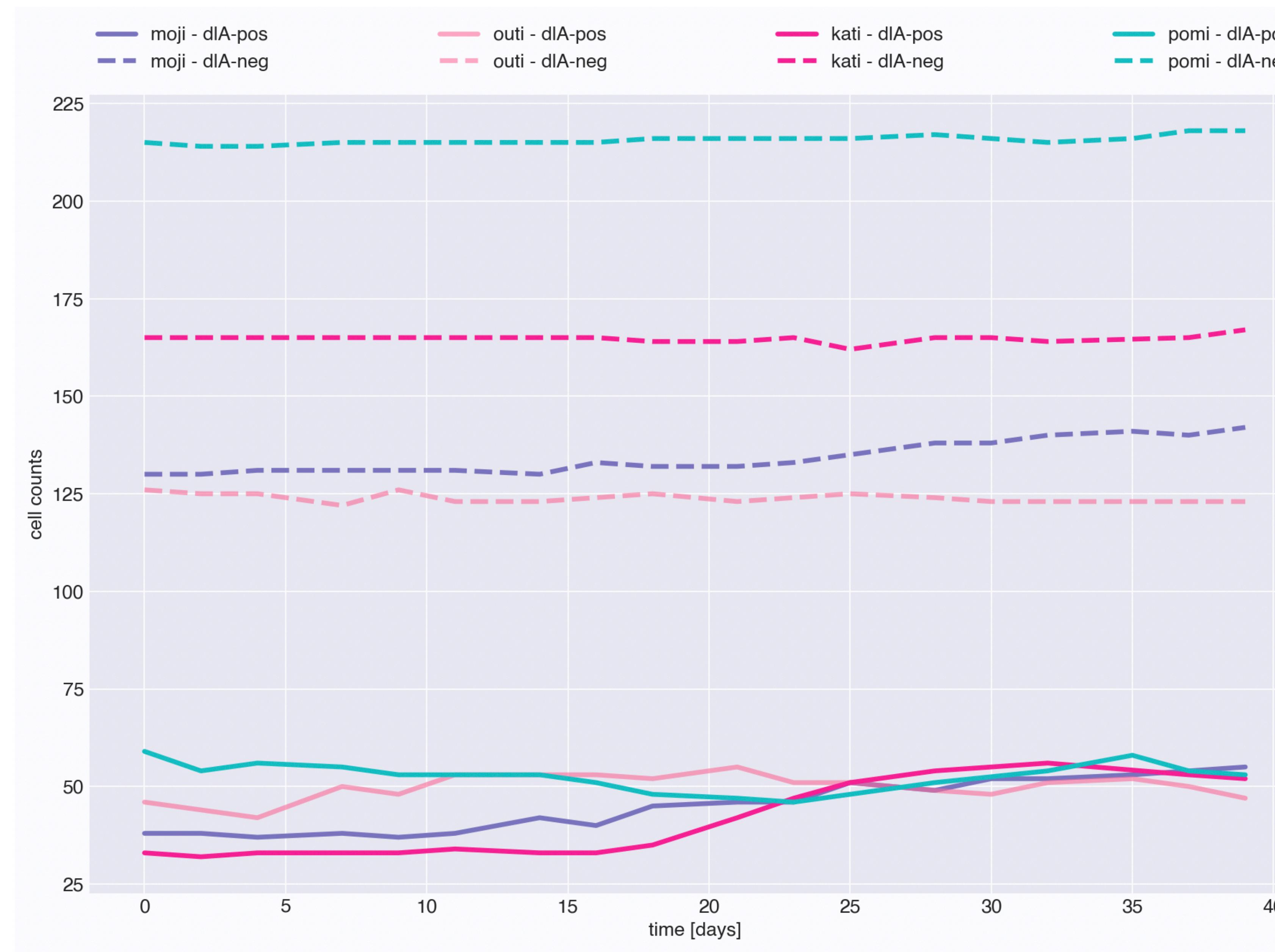


c. Notch-Delta signaling controls the self-renewal rate of aNSCs too.

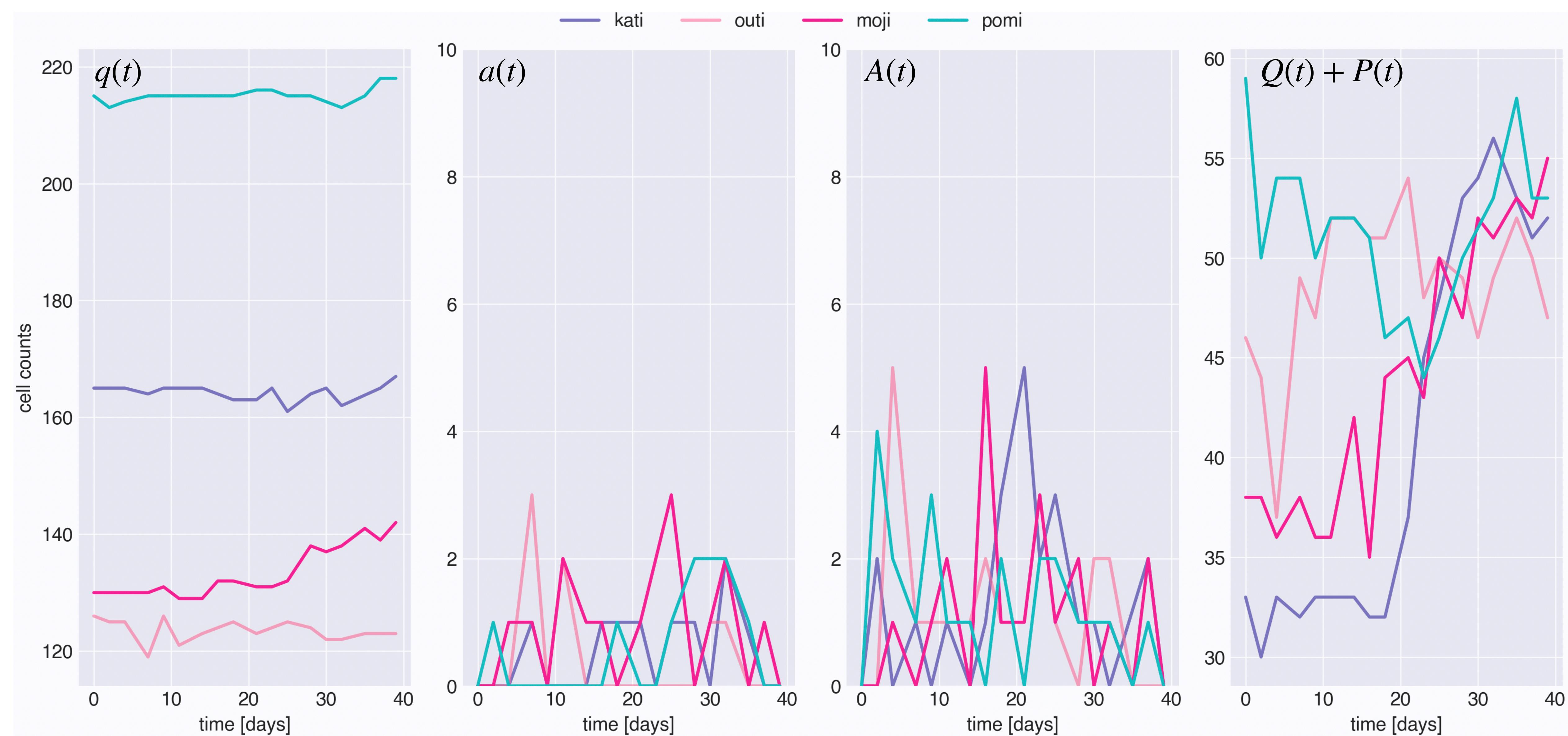


*In vivo* data now available!

# Comparison of deltaA-neg (Reservoir) and deltaA-pos (Operational) cell numbers over time per fish



**Figure (temp): Number of cells of each type over time per fish**



**Thank you for your attention!**