

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



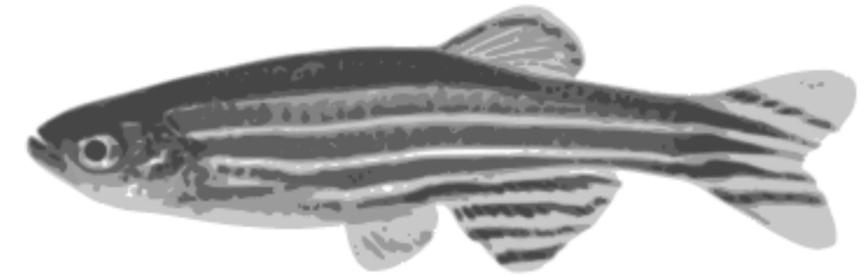
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Adult mouse



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

Adult zebrafish



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

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Adult mouse



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- Mathematical model of NSC dynamics

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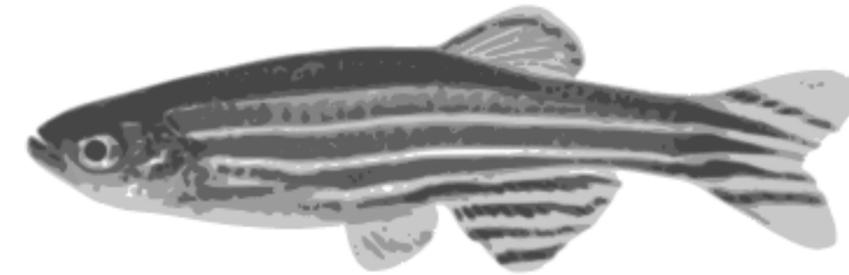
Unraveling regulatory feedback mechanisms in adult neurogenesis through mathematical modelling

[Diana-Patricia Danciu](#) , [Filip Z. Klawe](#), [Alexey Kazarnikov](#), [Laura Femmer](#), [Ekaterina Kostina](#), [Ana Martin-Villalba](#) & [Anna Marciniak-Czochra](#) 

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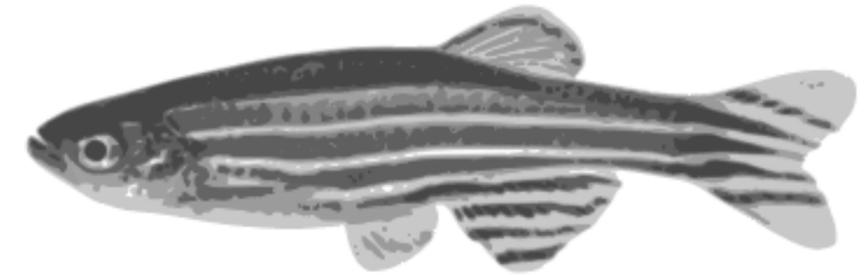
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Adult zebrafish



- 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

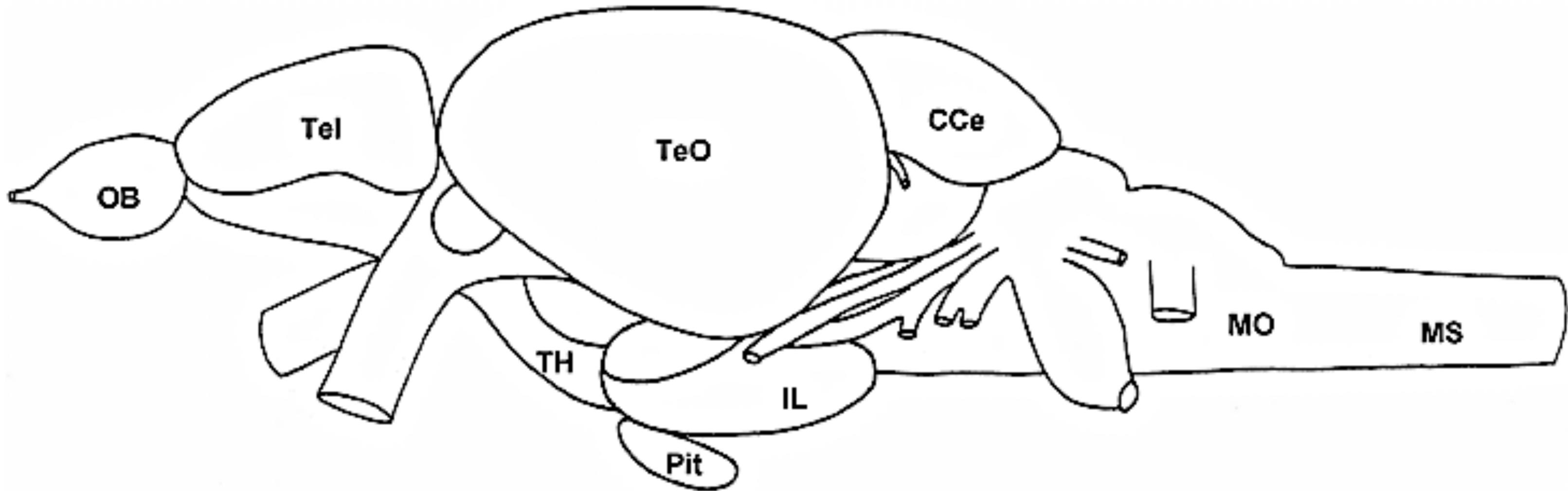


Figure: Lateral view of the adult zebrafish brain (*credit to Daniel R. Dietrich*)

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

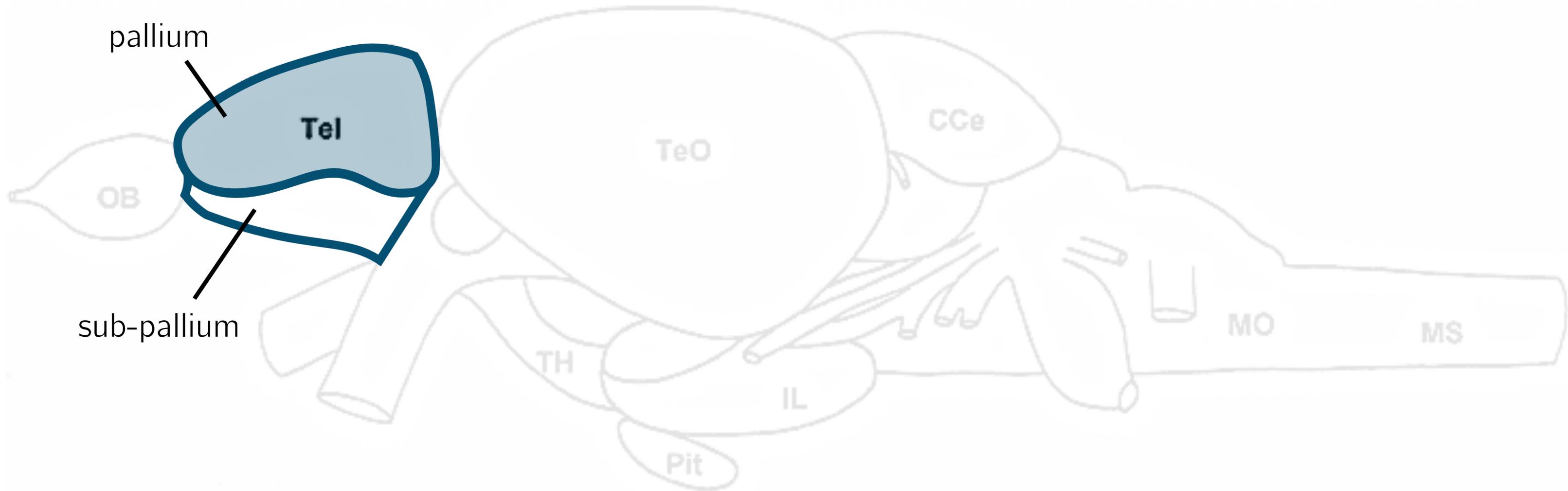


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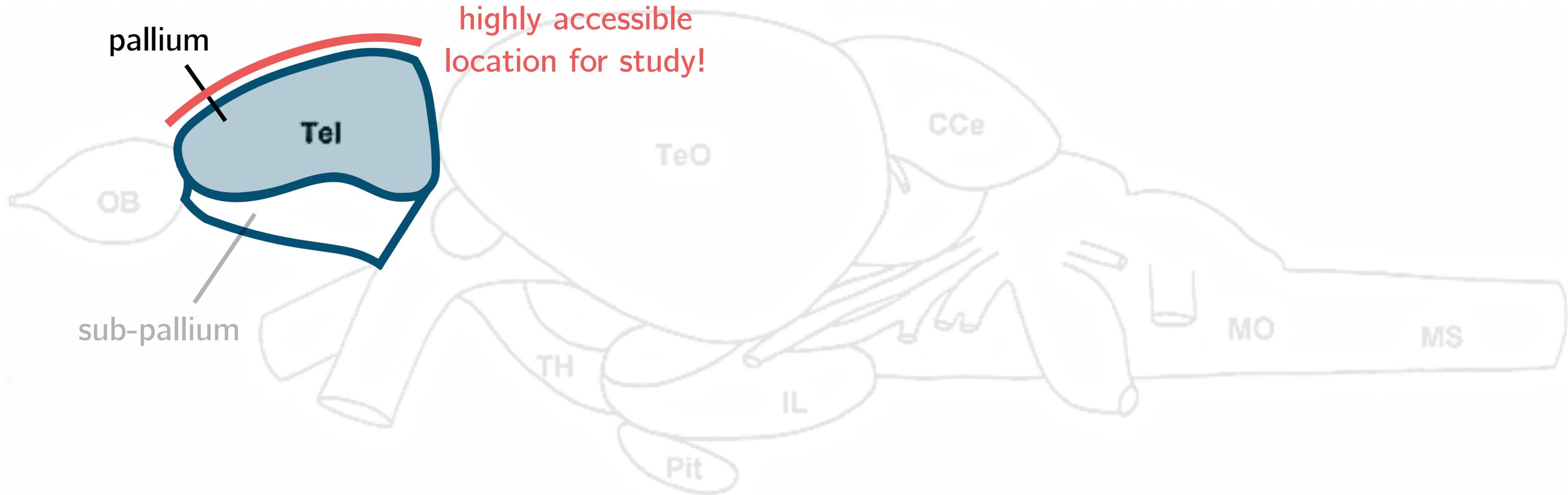


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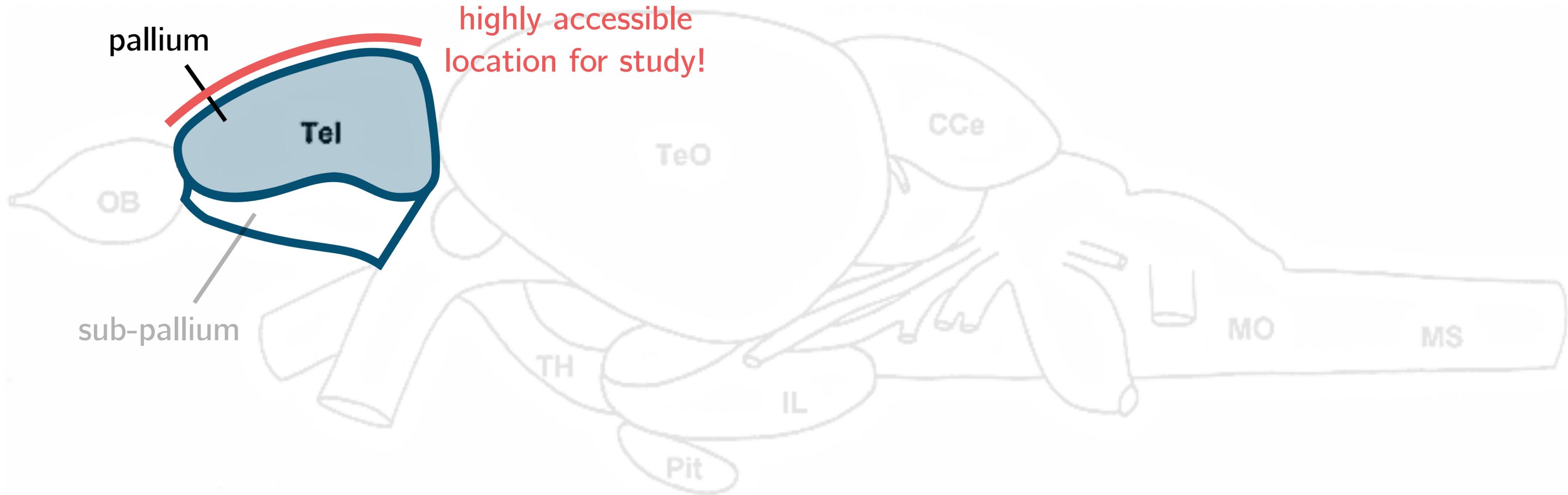


Figure: Lateral view of the adult zebrafish brain (*credit to Daniel R. Dietrich*)

Radial glia as NSCs:
GFAP/S100 β /GS
Her4 (=Hes5)
Sox2



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

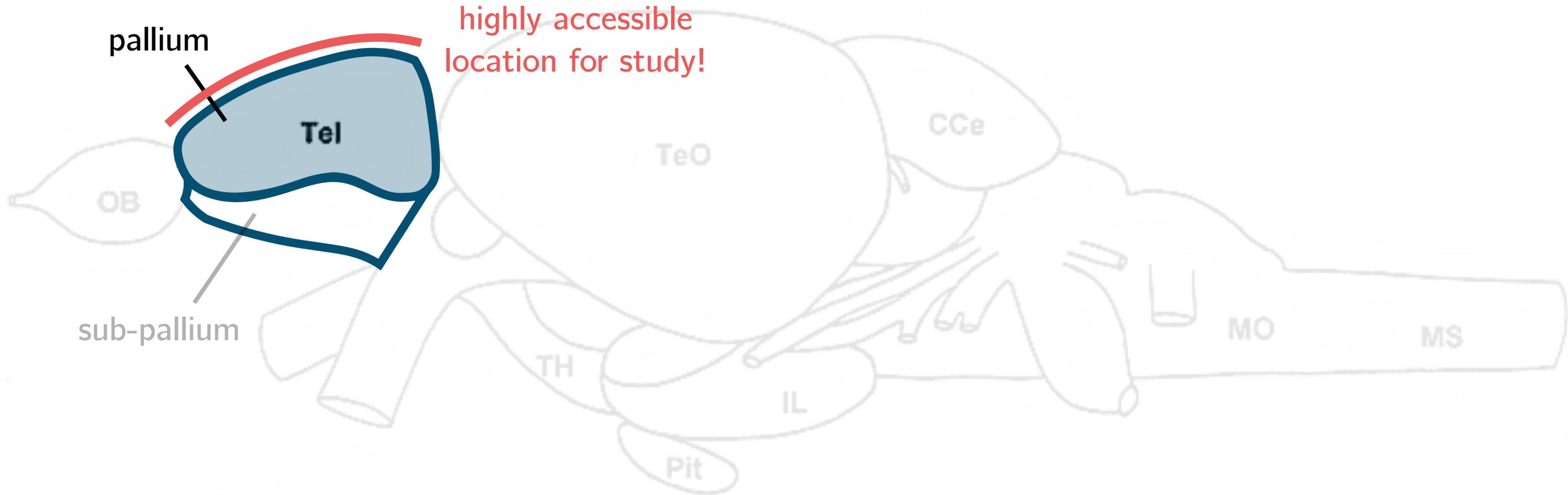


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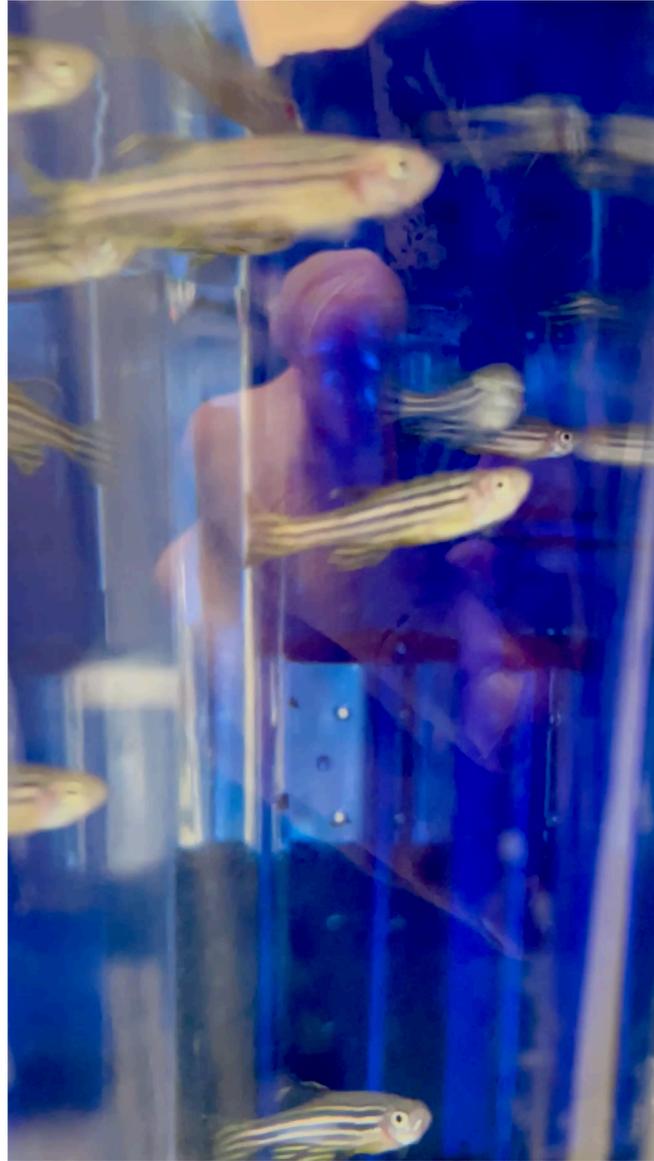
Radial glia as NSCs:
GFAP/S100 β /GS
Her4 (=Hes5)
Sox2



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

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

Wild Type



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

Wild Type



Mutant

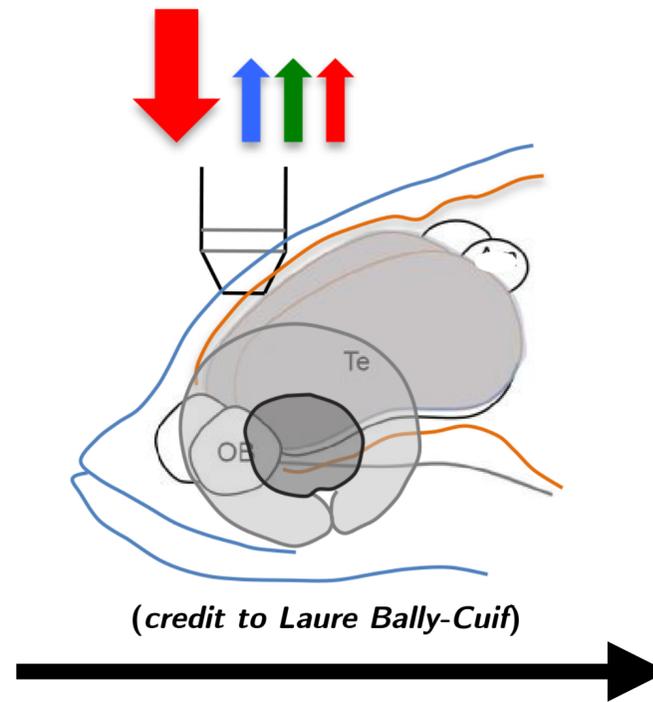


Biological Context: NSCs anatomy and lineage progression in the fish brain

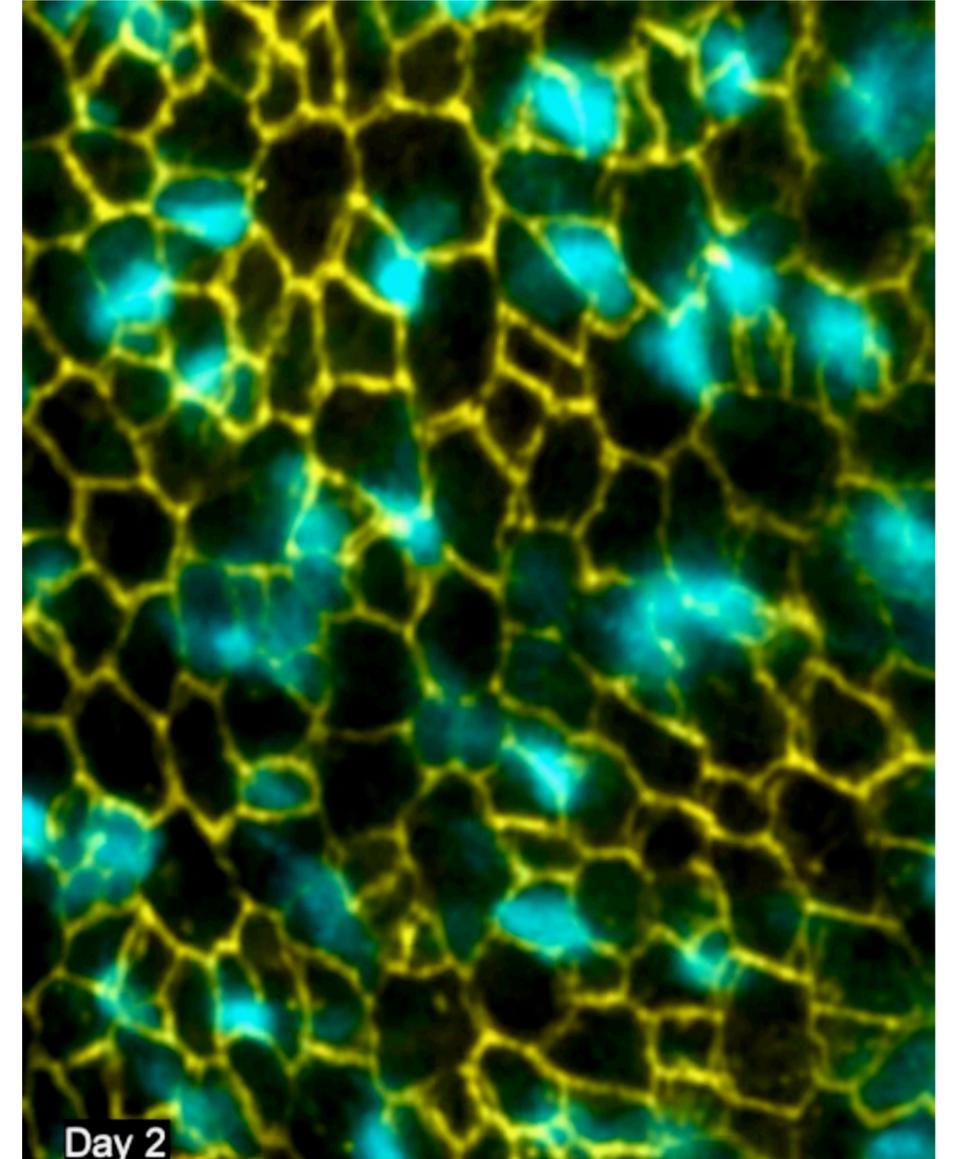
Wild Type



Mutant

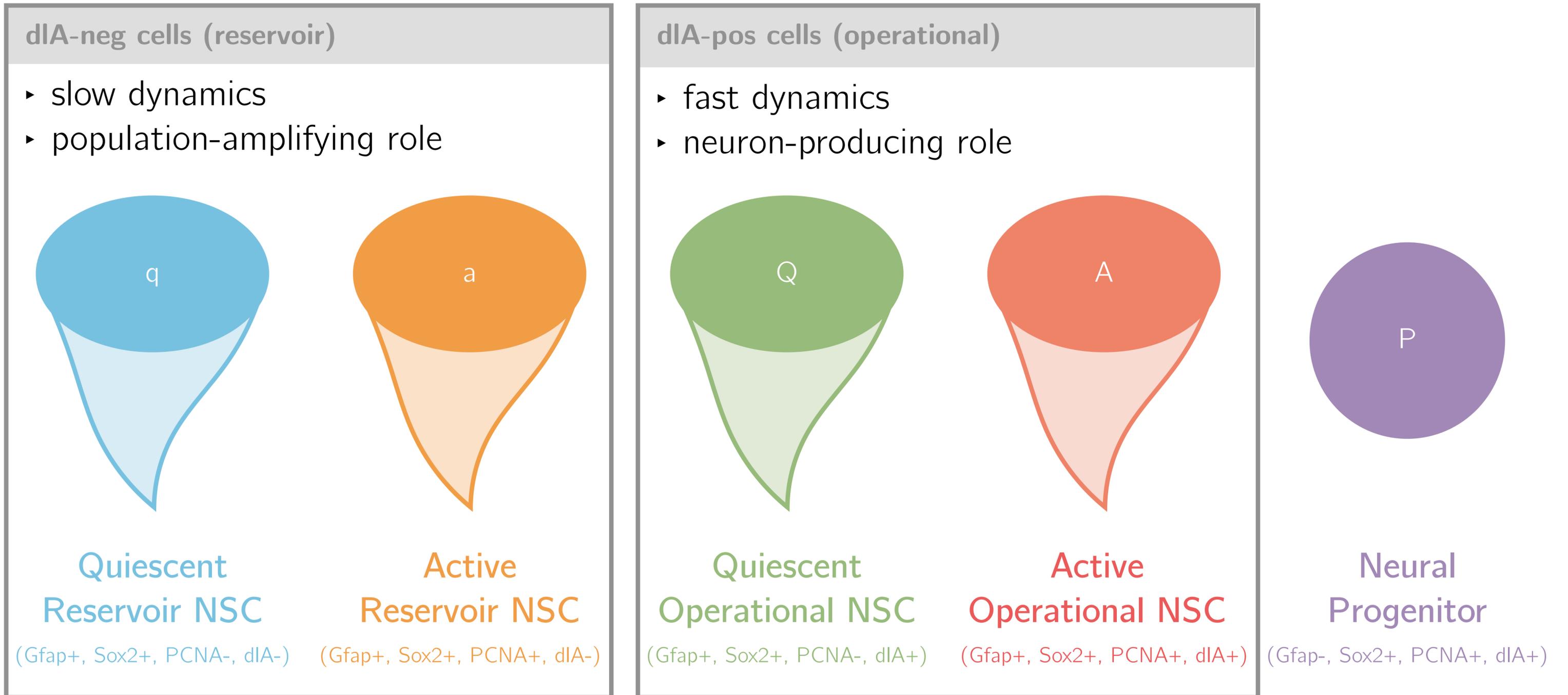


Pallial NSC population

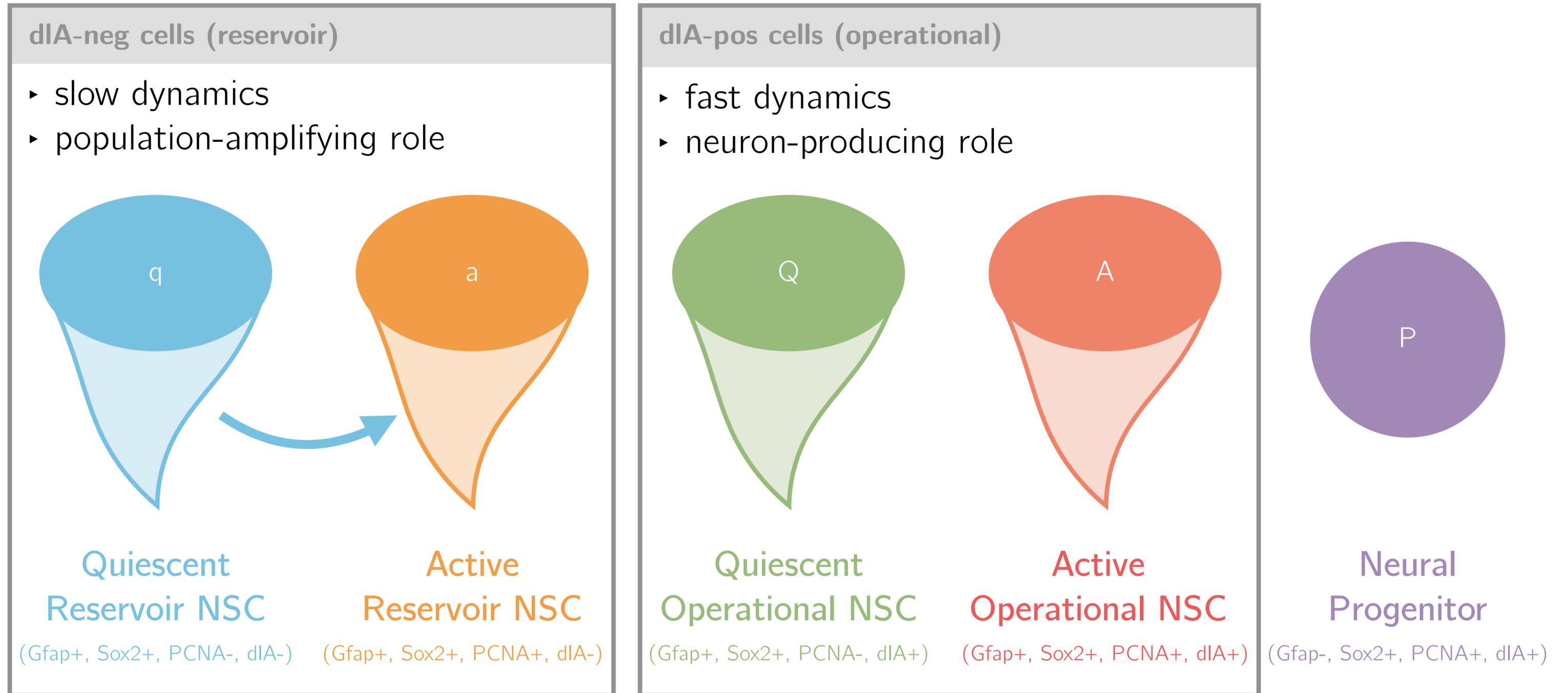


source: <https://doi.org/10.1126/sciadv.adg7519>

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

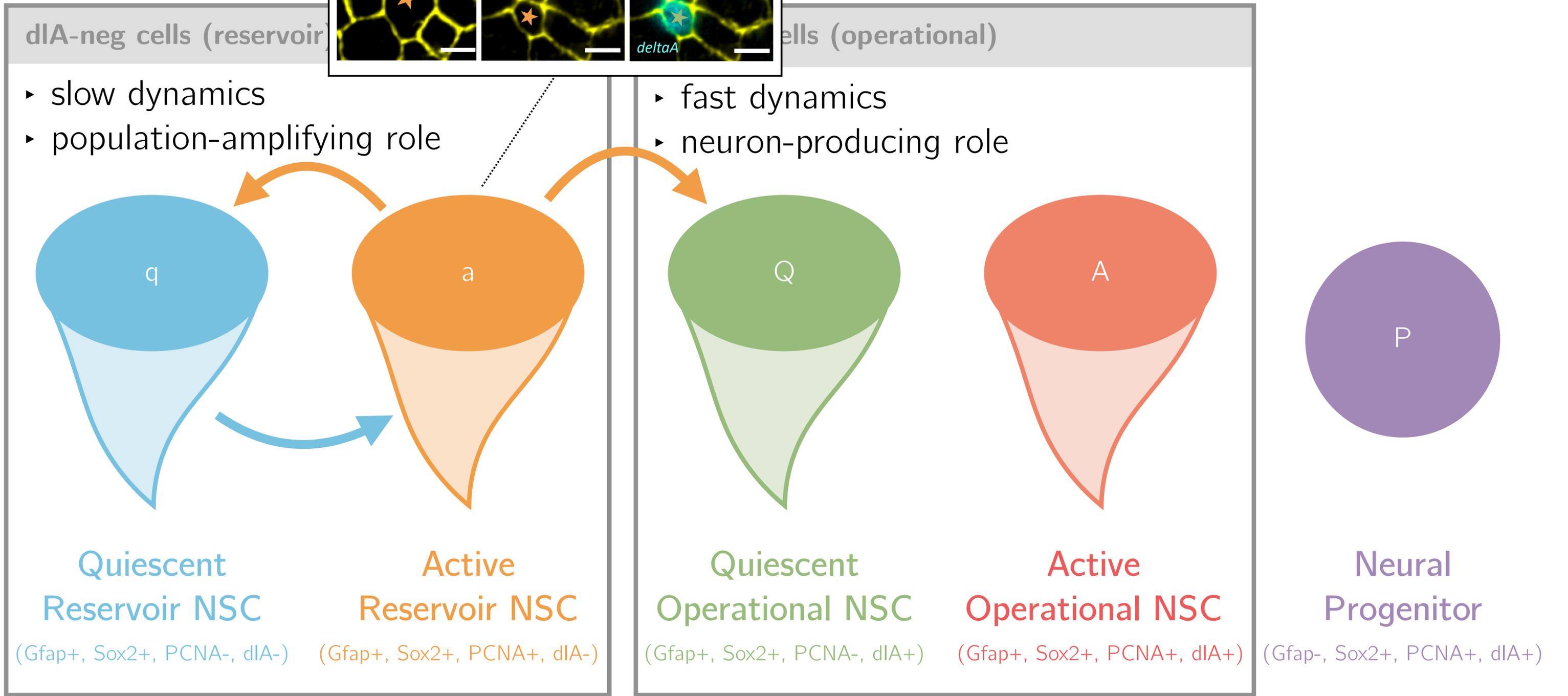
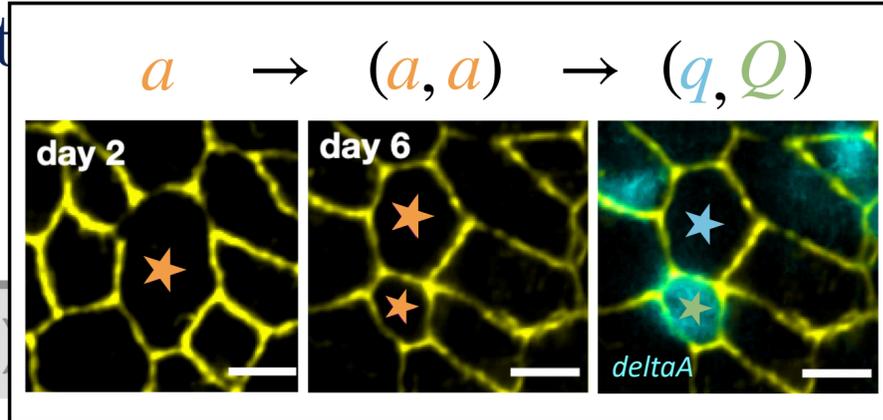


Biological Context: NSCs anatomy and lineage progression in the fish brain

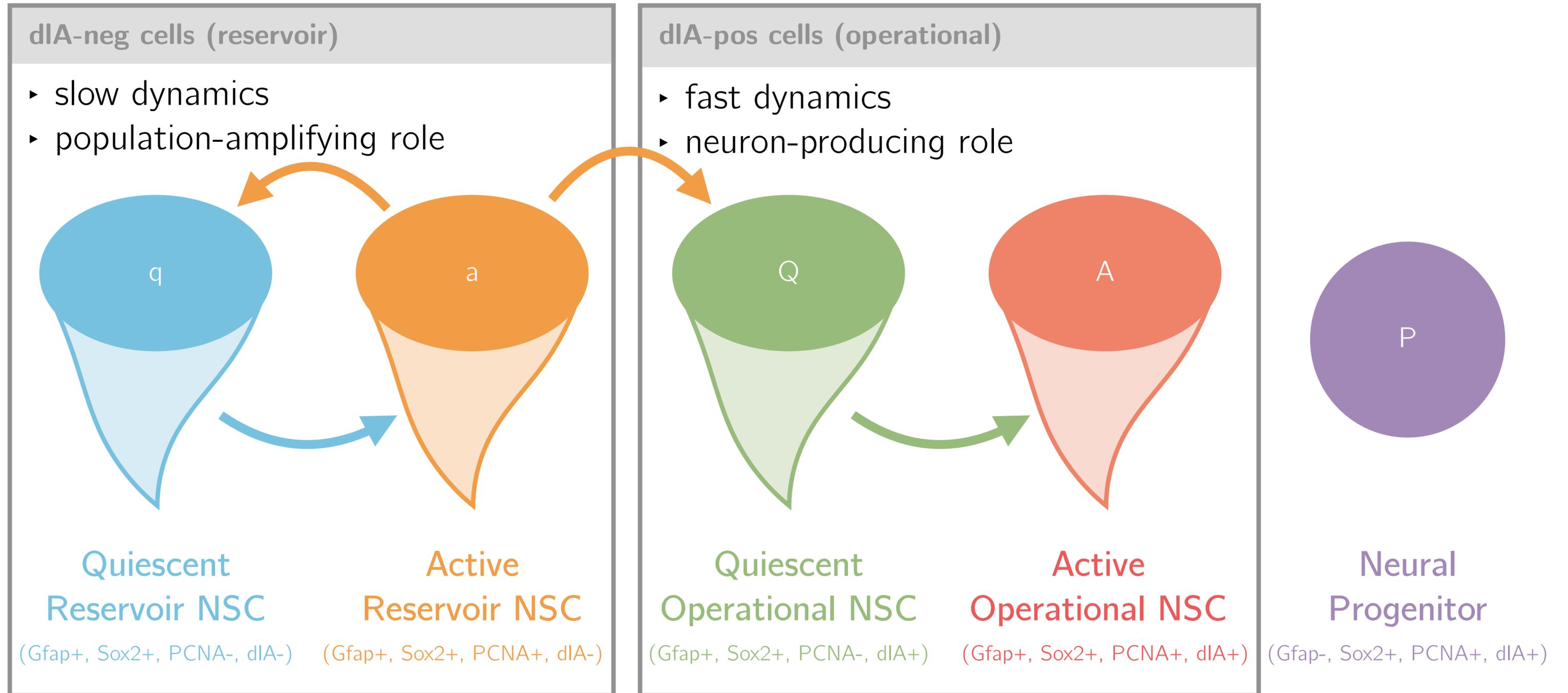


Biological Context

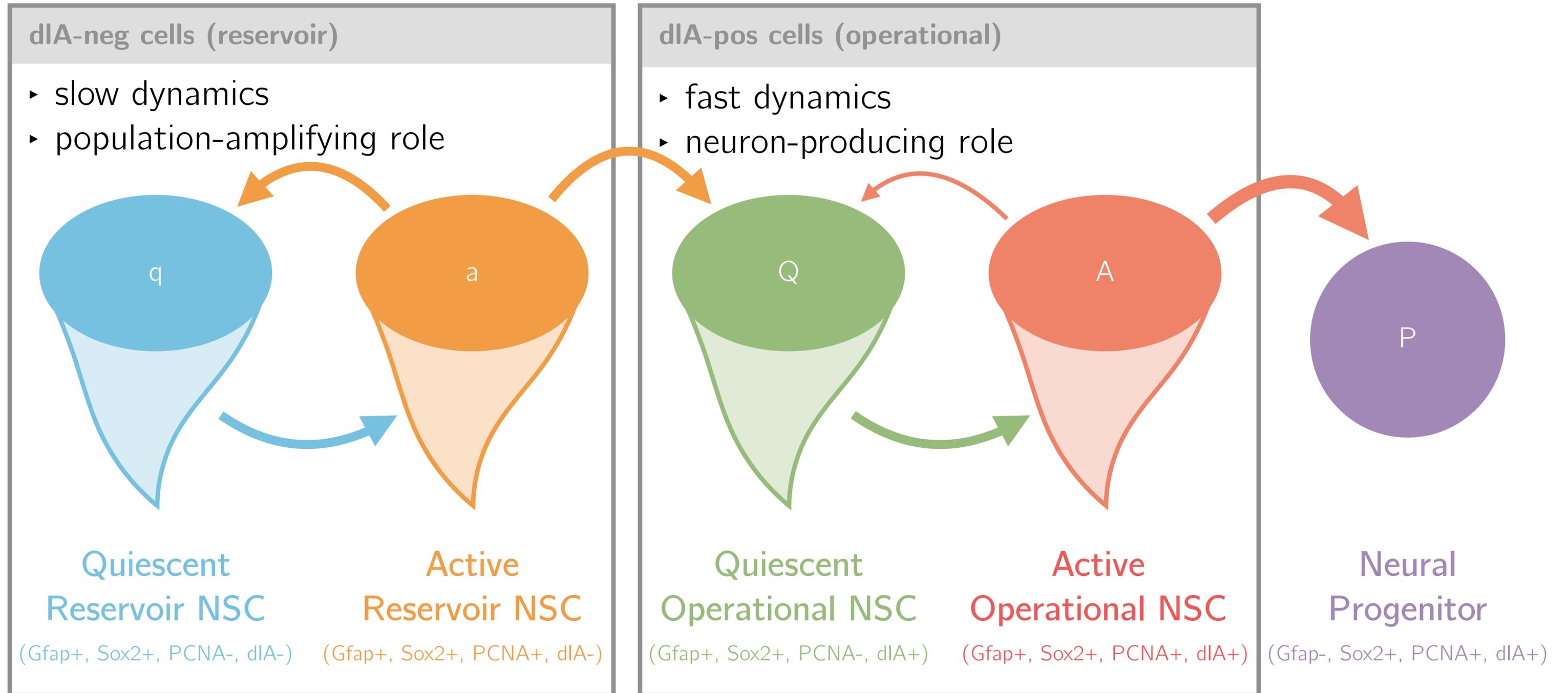
Age progression in the fish brain



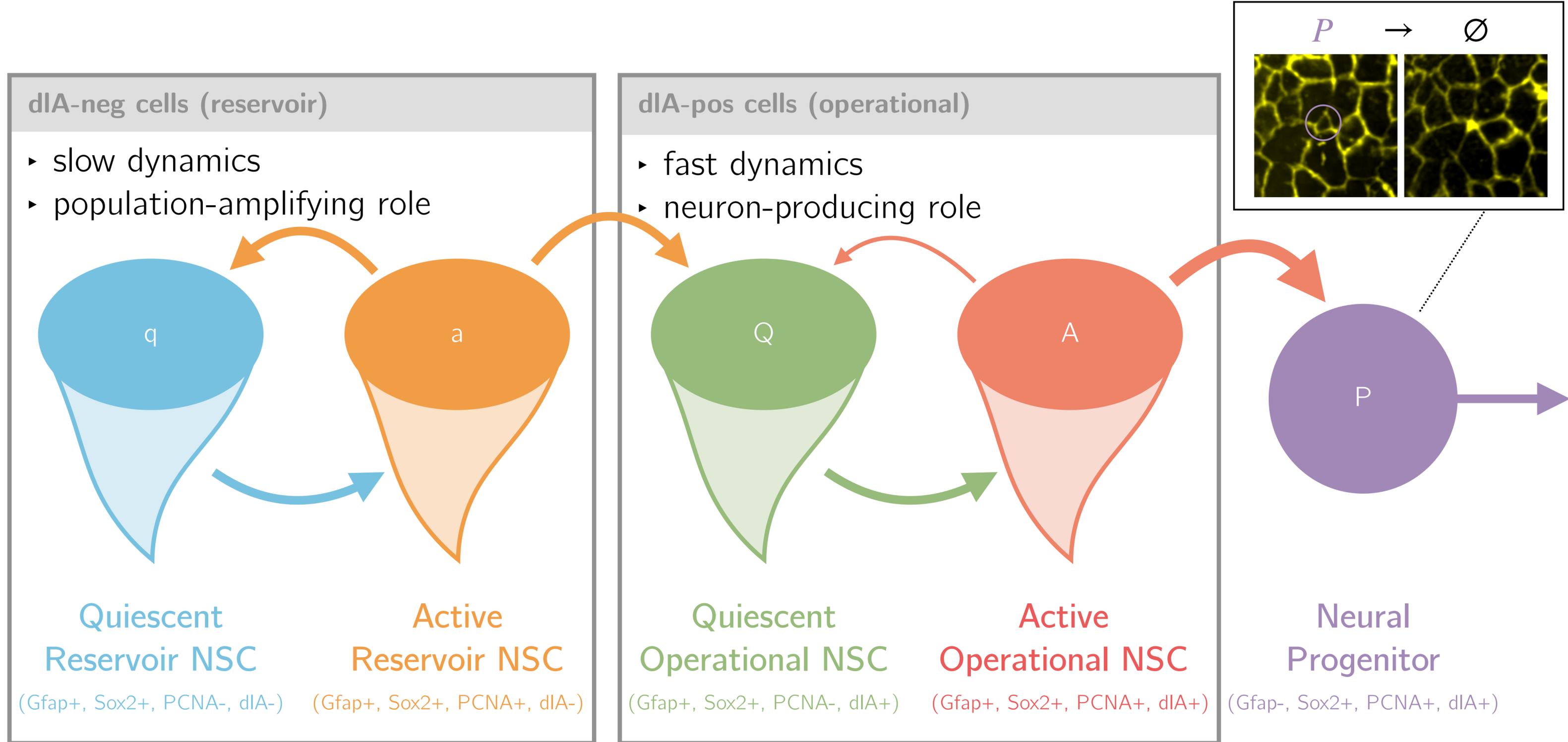
Biological Context: NSCs anatomy and lineage progression in the fish brain



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

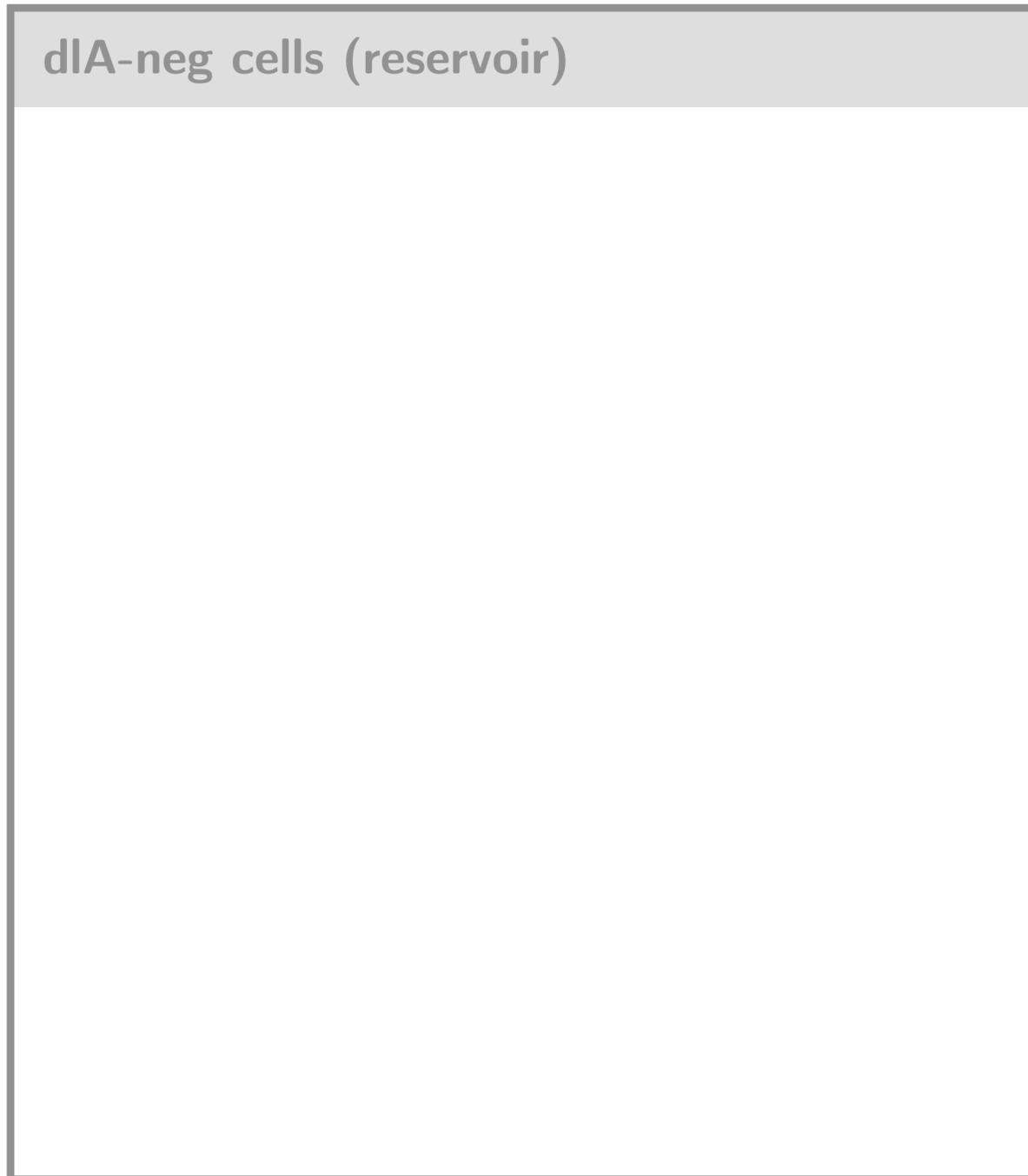


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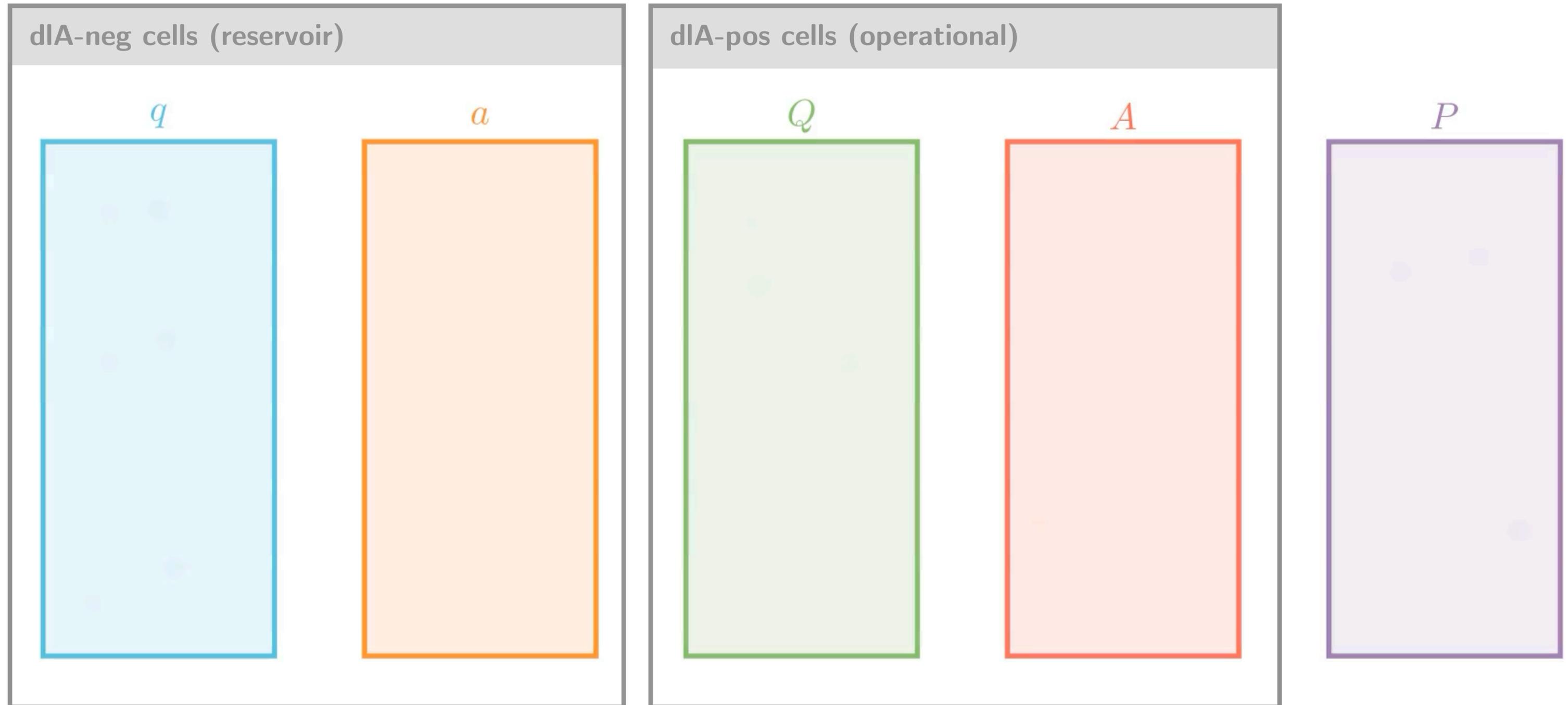
Model Visualization: Compartmental evolution

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



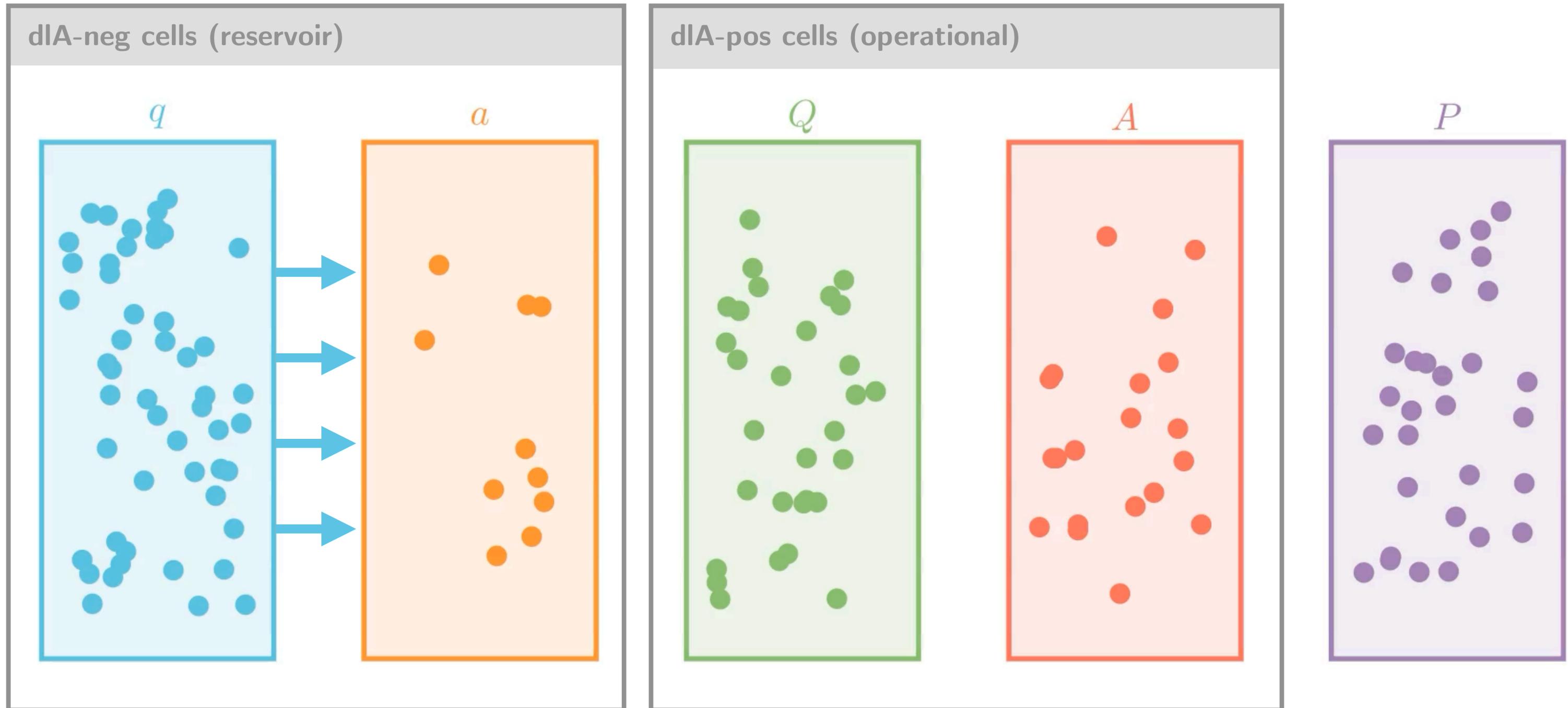
Model Visualization: Compartmental evolution

Step 2: Populate each compartment (initial condition)



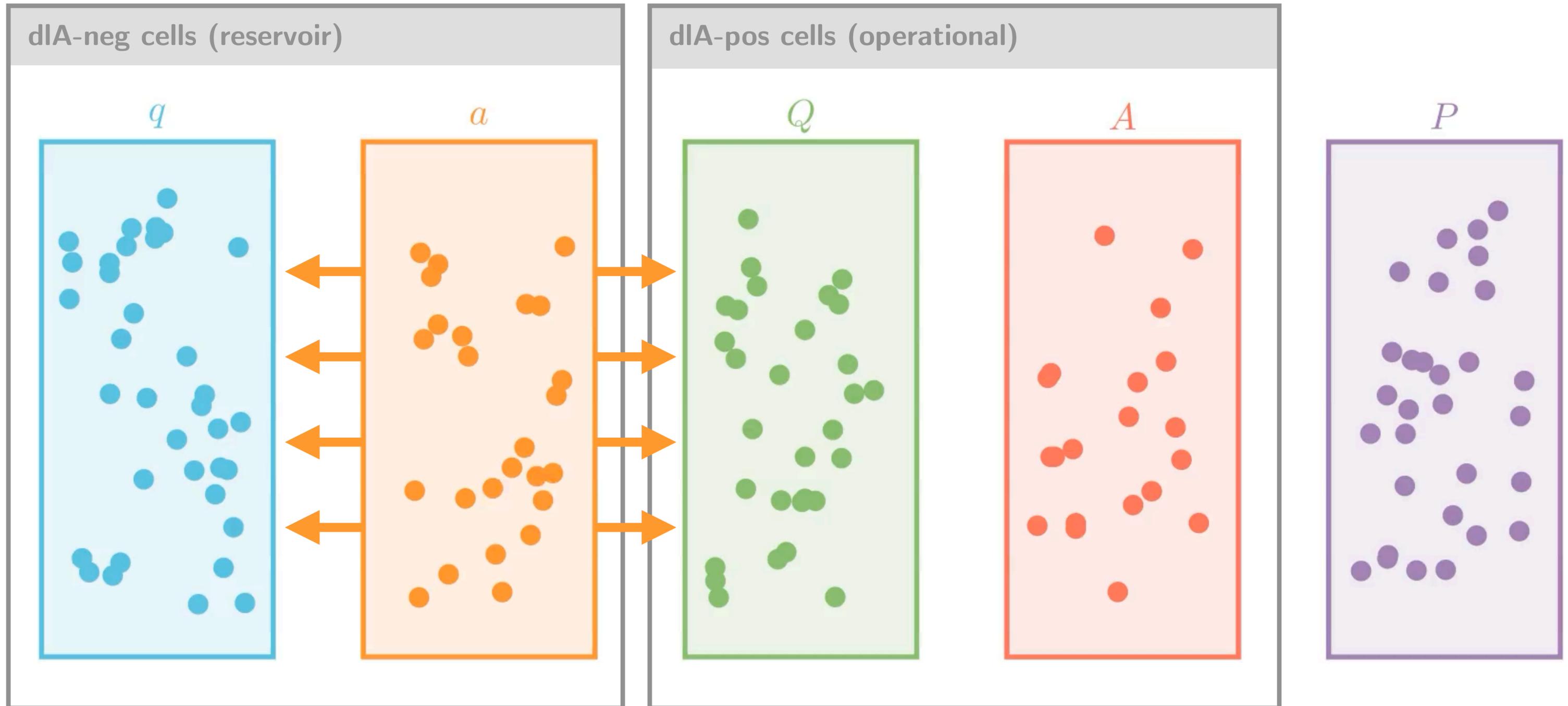
Model Visualization: Compartmental evolution

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



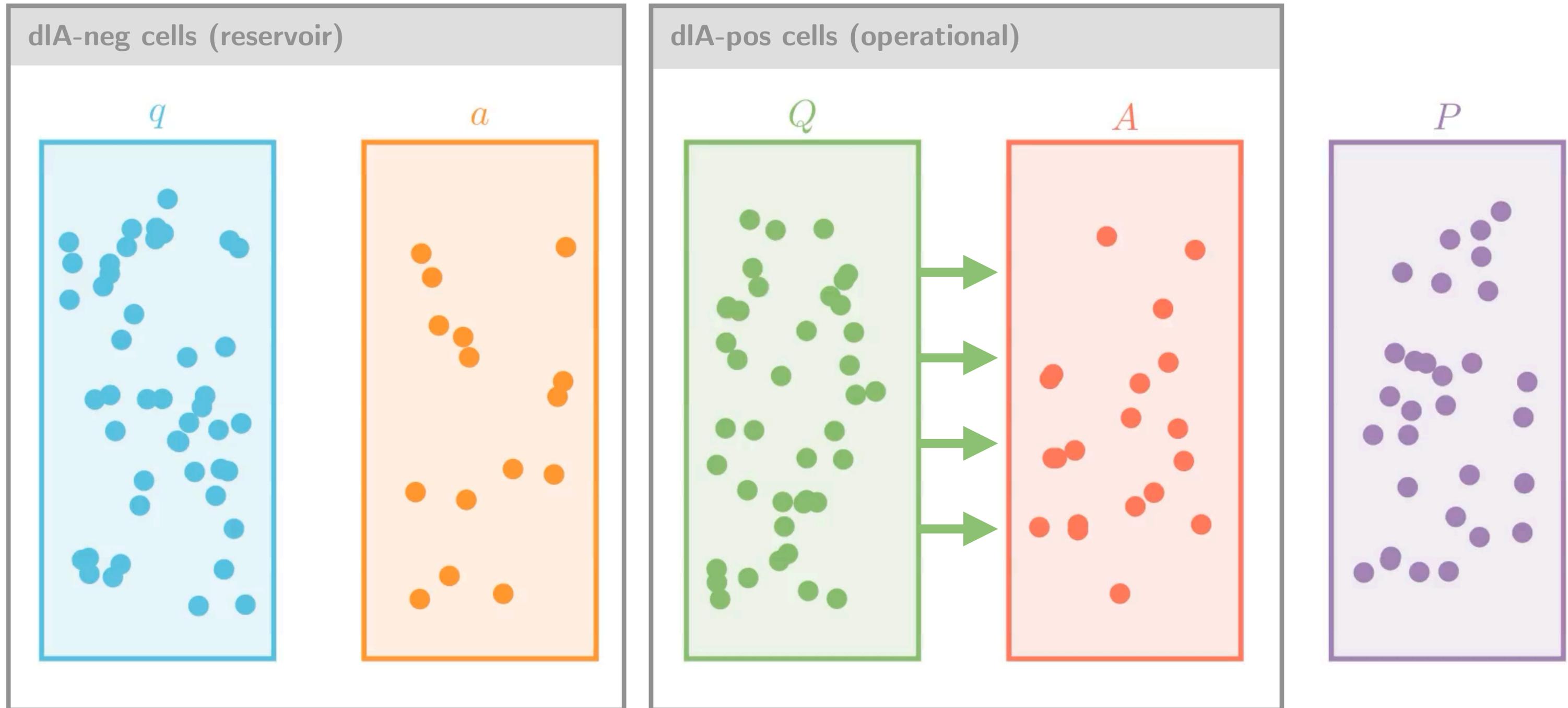
Model Visualization: Compartmental evolution

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



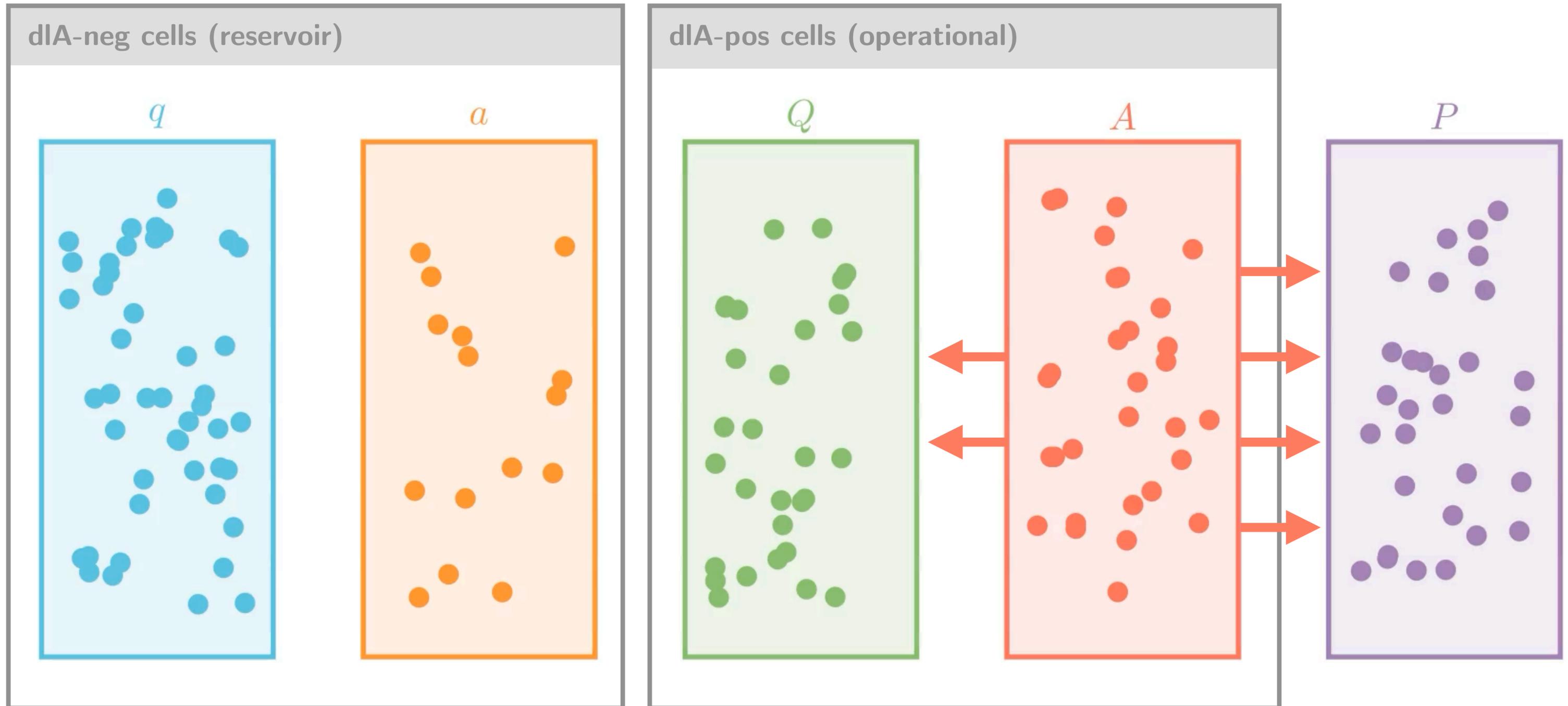
Model Visualization: Compartmental evolution

Step 3.c: operational qNSC activate ($Q'(t) = -r_1Q(t)$, and $A(t) = +r_1Q(t)$)



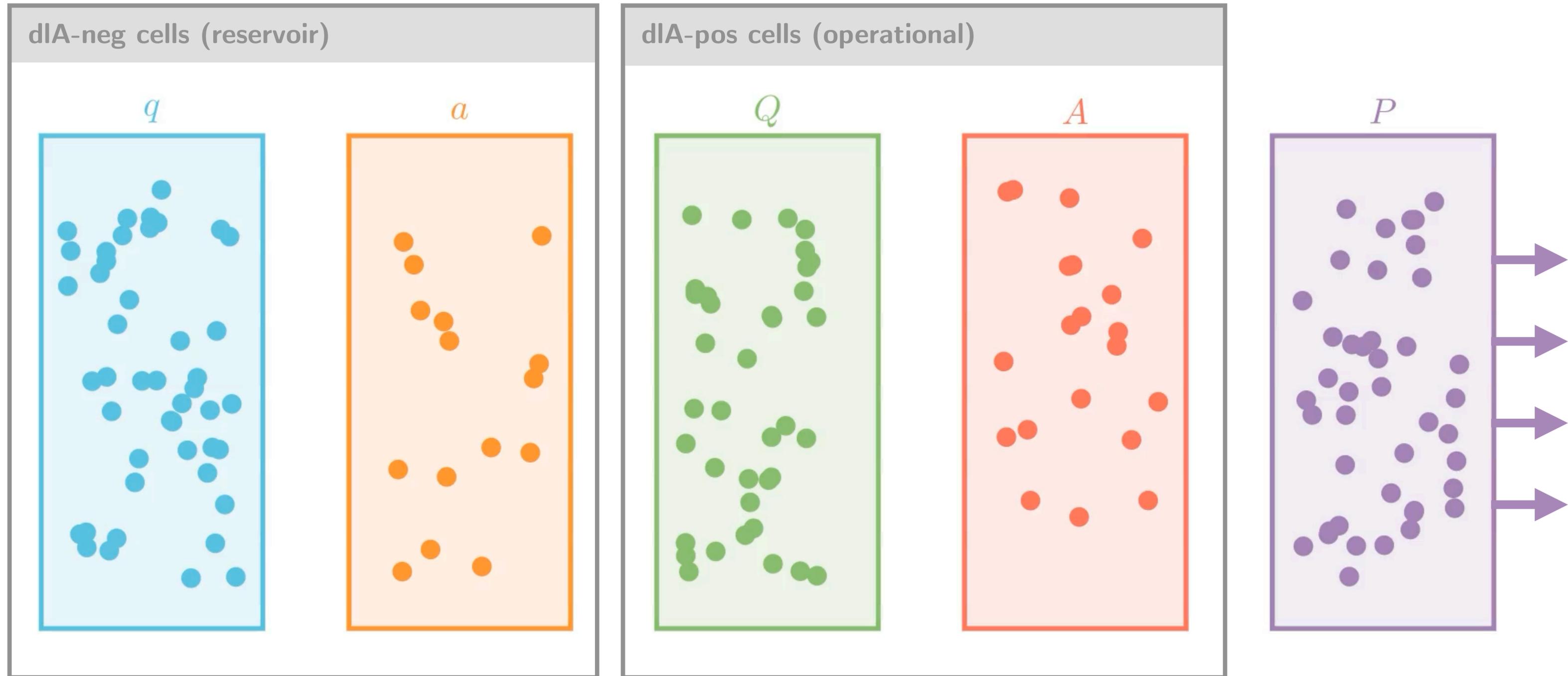
Model Visualization: Compartmental evolution

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

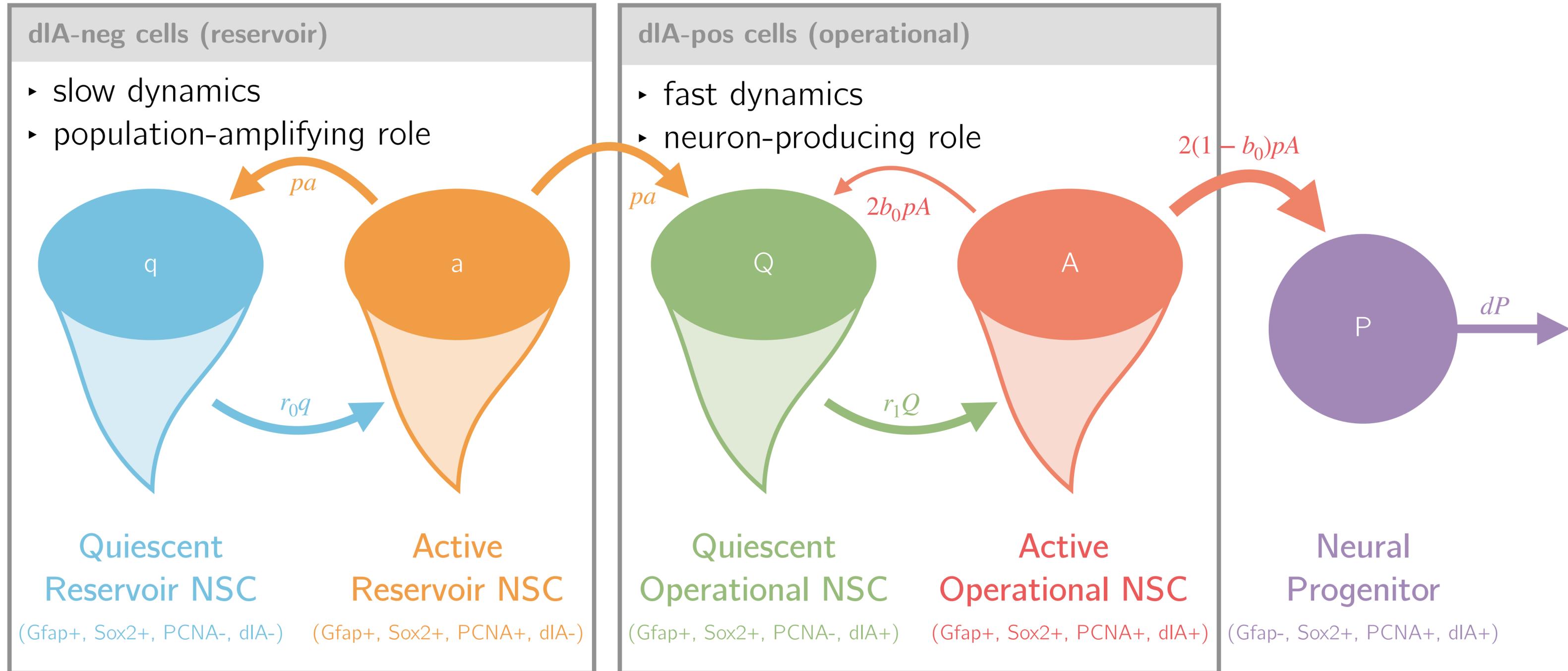


Model Visualization: Compartmental evolution

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



Model Equations (Summary)



Model Equations

dIA-neg cells (reservoir)



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



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

dIA-pos cells (operational)



$$Q'(t) = -r_1 Q(t) + 2b_0 pA(t) + pa(t),$$



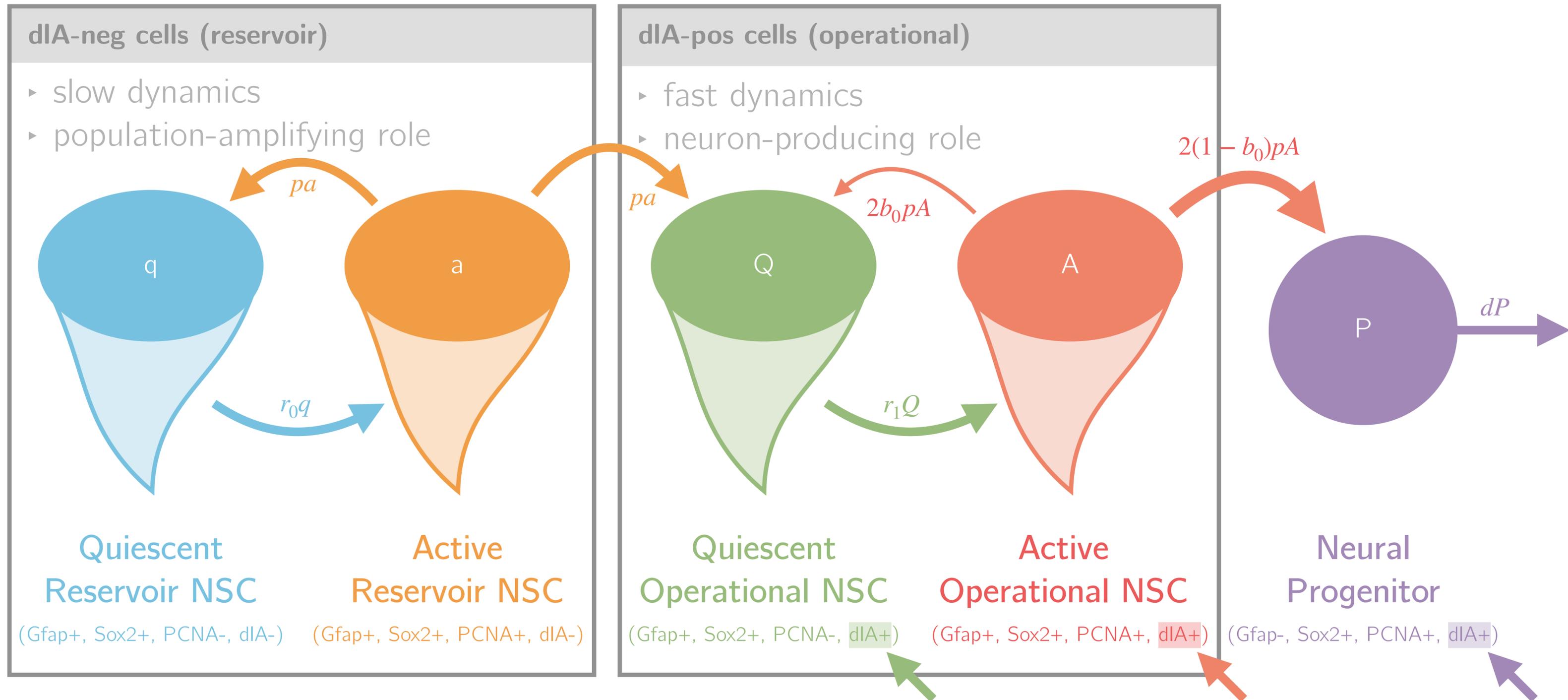
$$A'(t) = r_1 Q(t) - pA(t),$$



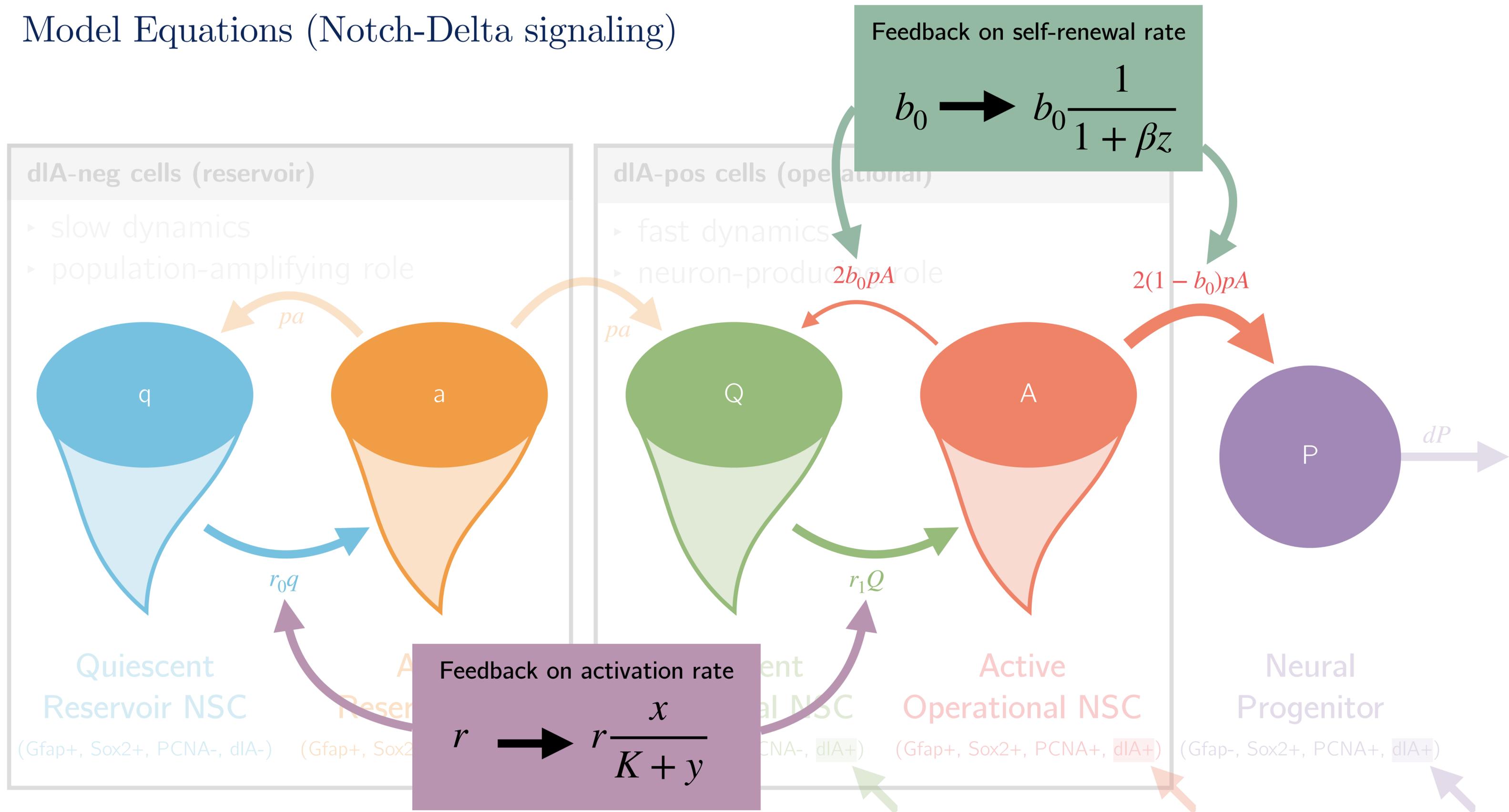
$$P'(t) = 2(1 - b_0)pA(t) - dP(t)$$

Modeling Notch-delta Signaling

Model Equations (Notch-Delta signaling)



Model Equations (Notch-Delta signaling)



dIA-neg cells (reservoir)



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



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

dIA-pos cells (operational)



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



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



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

dIA-neg cells (reservoir)



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



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

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



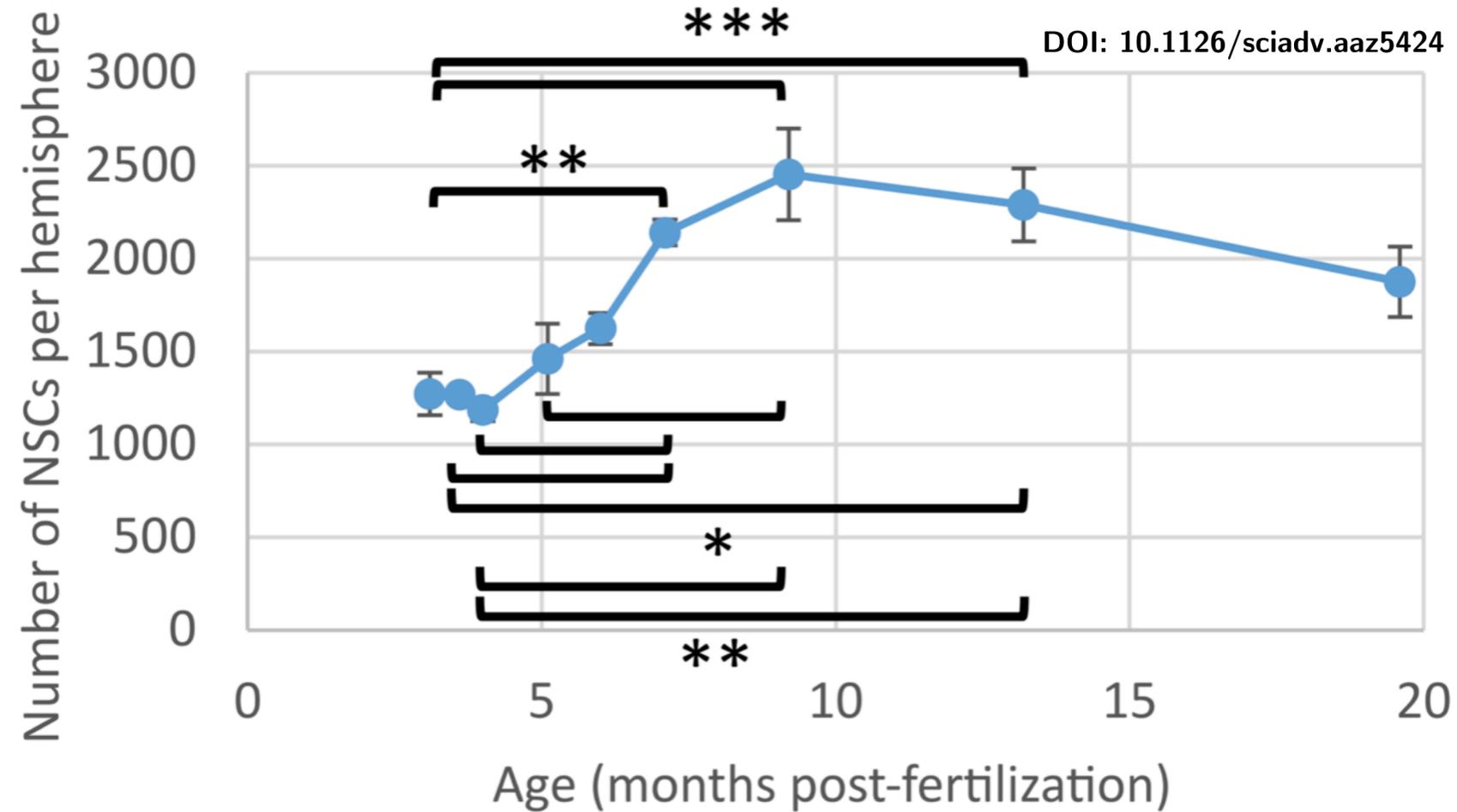
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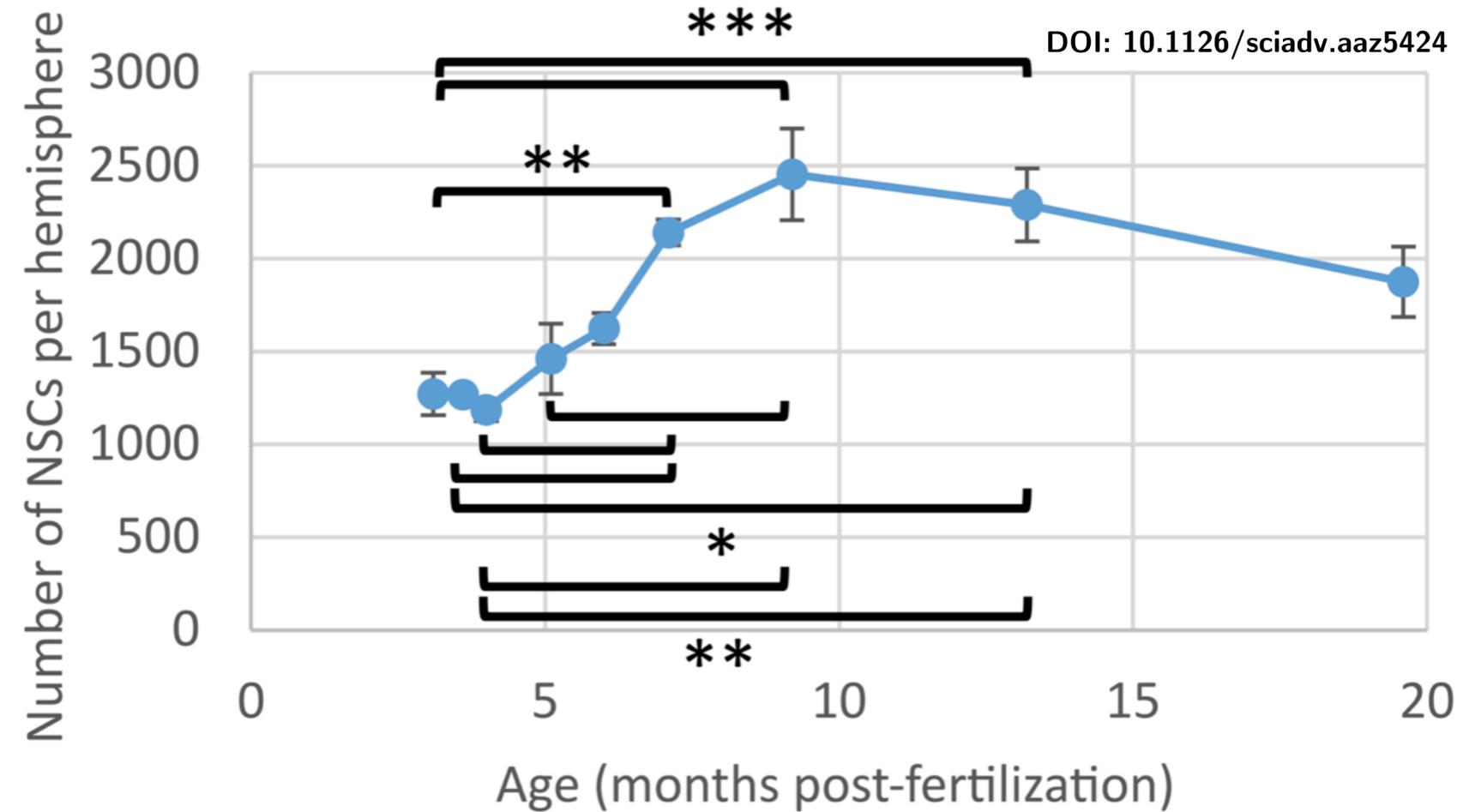
**Mouse
System**

Calibration: Investigating the data



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

Calibration: Investigating the data

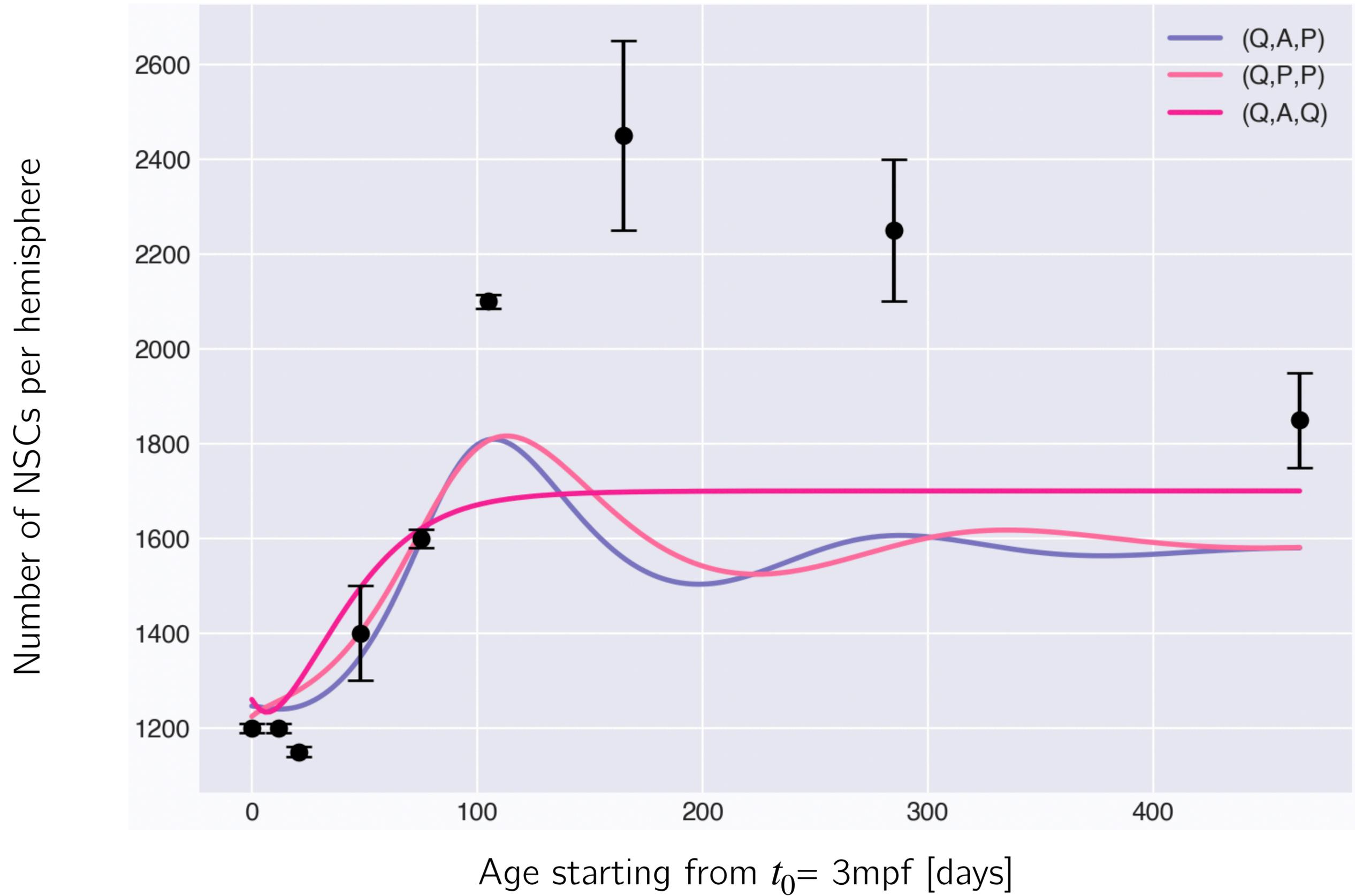


- a. Average number of cells per hemisphere (n=7 brains), reported by Sox2 expression (NSCs + NPs) over 500 days

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

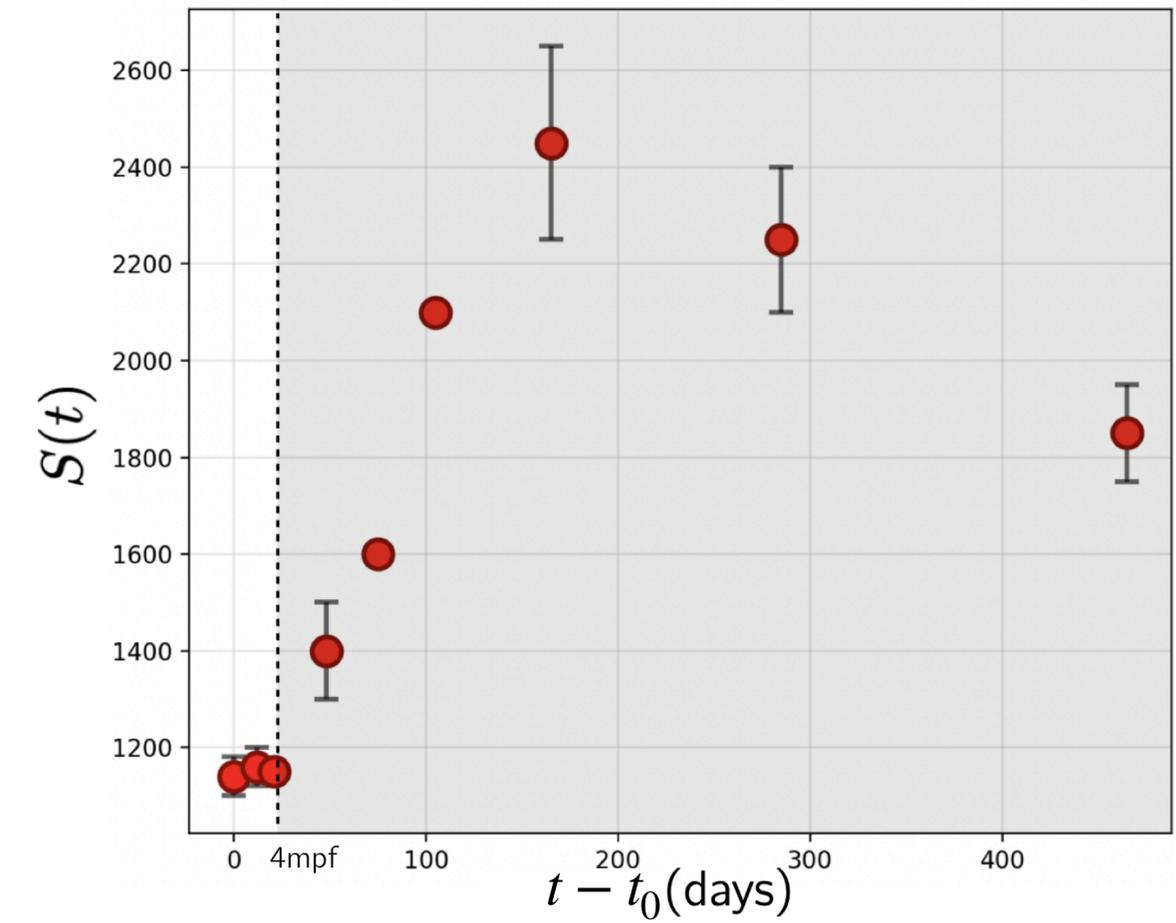


Modeling the source

Investigation: Modeling the source

Total population

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

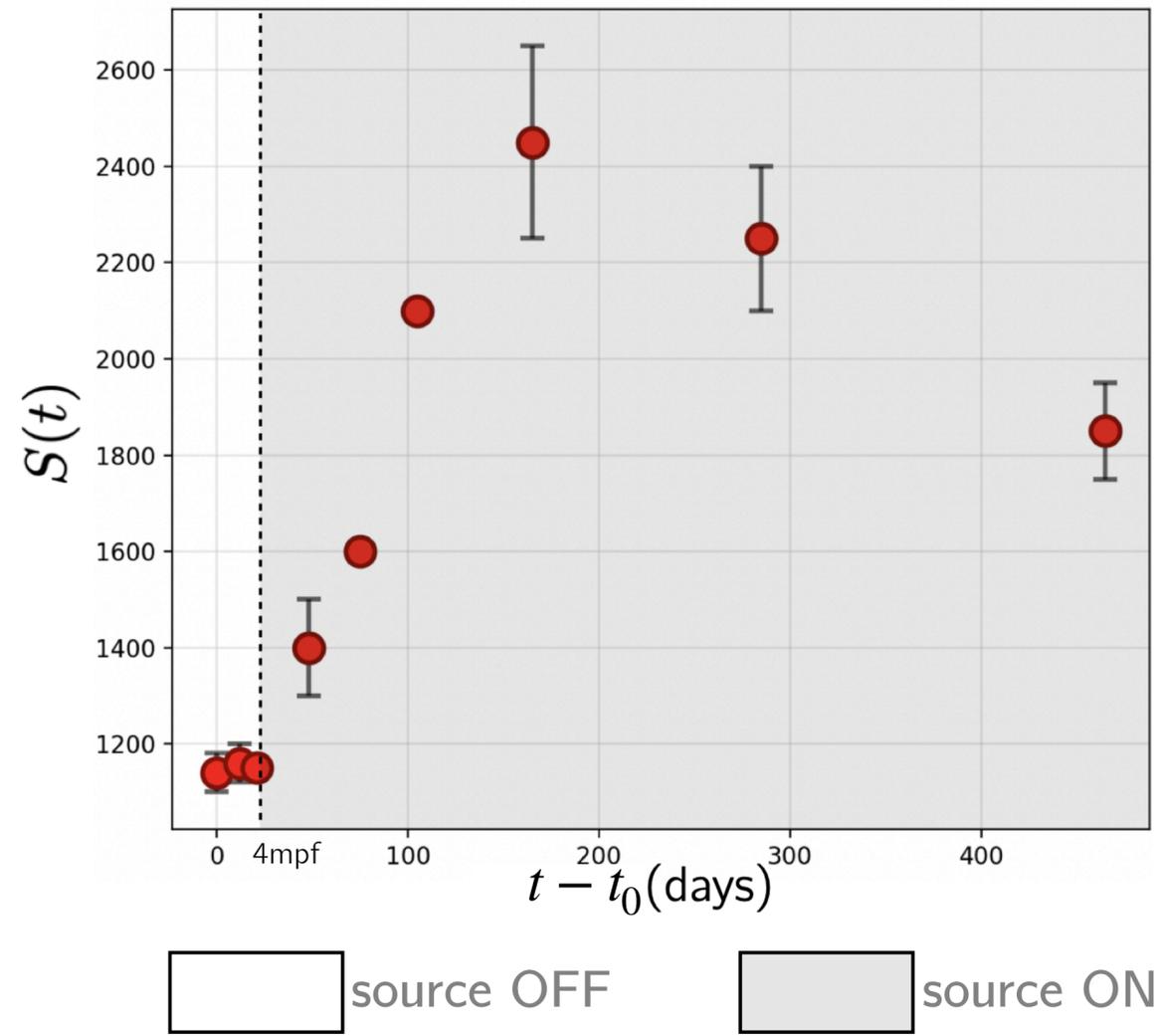


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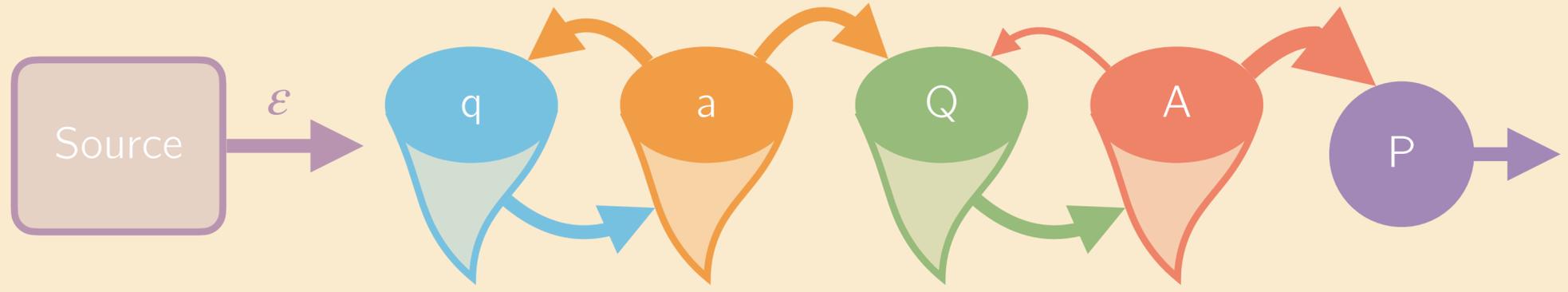
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Investigation: Modeling the source

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

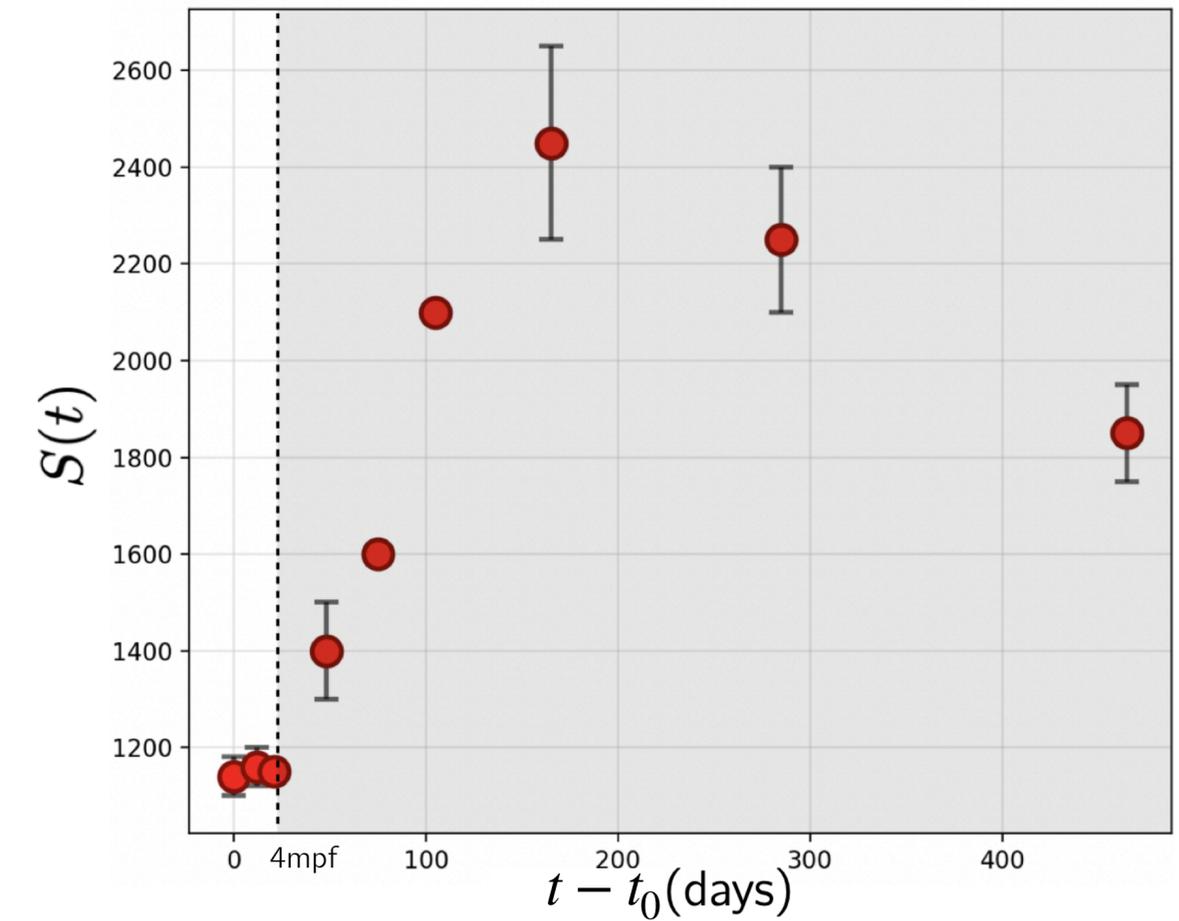


a. Introduce cells in the population, at the top of the hierarchy, and at a small constant rate $\epsilon > 0$.



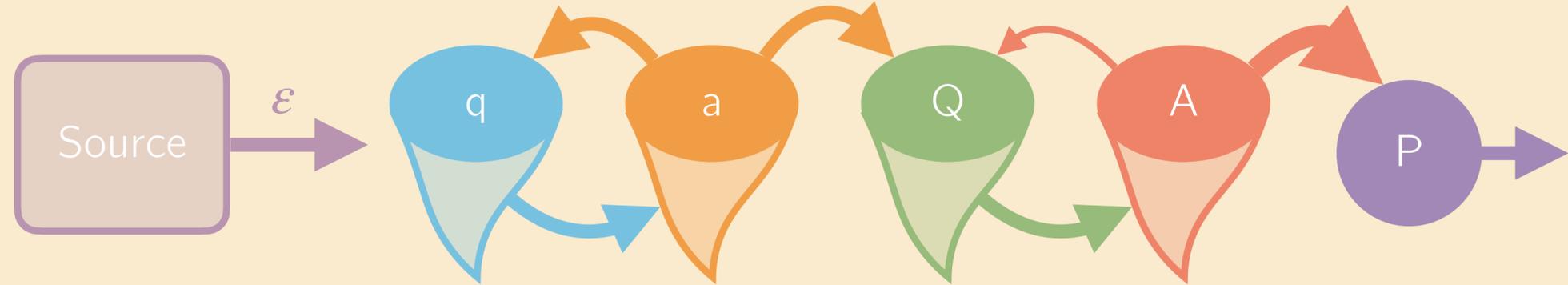
Investigation: Modeling the source

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

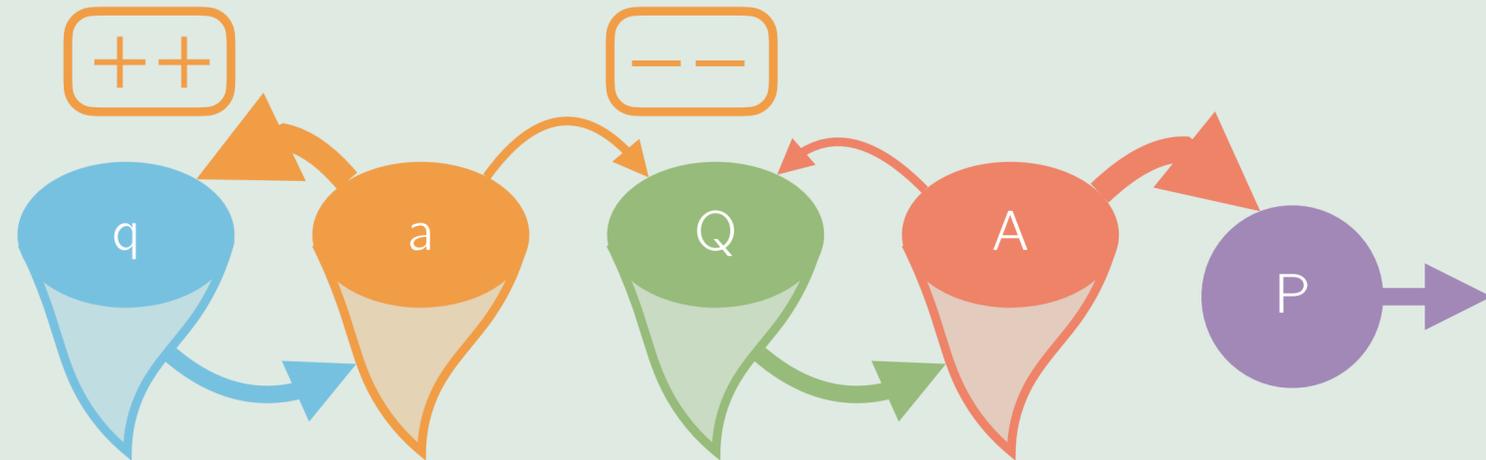


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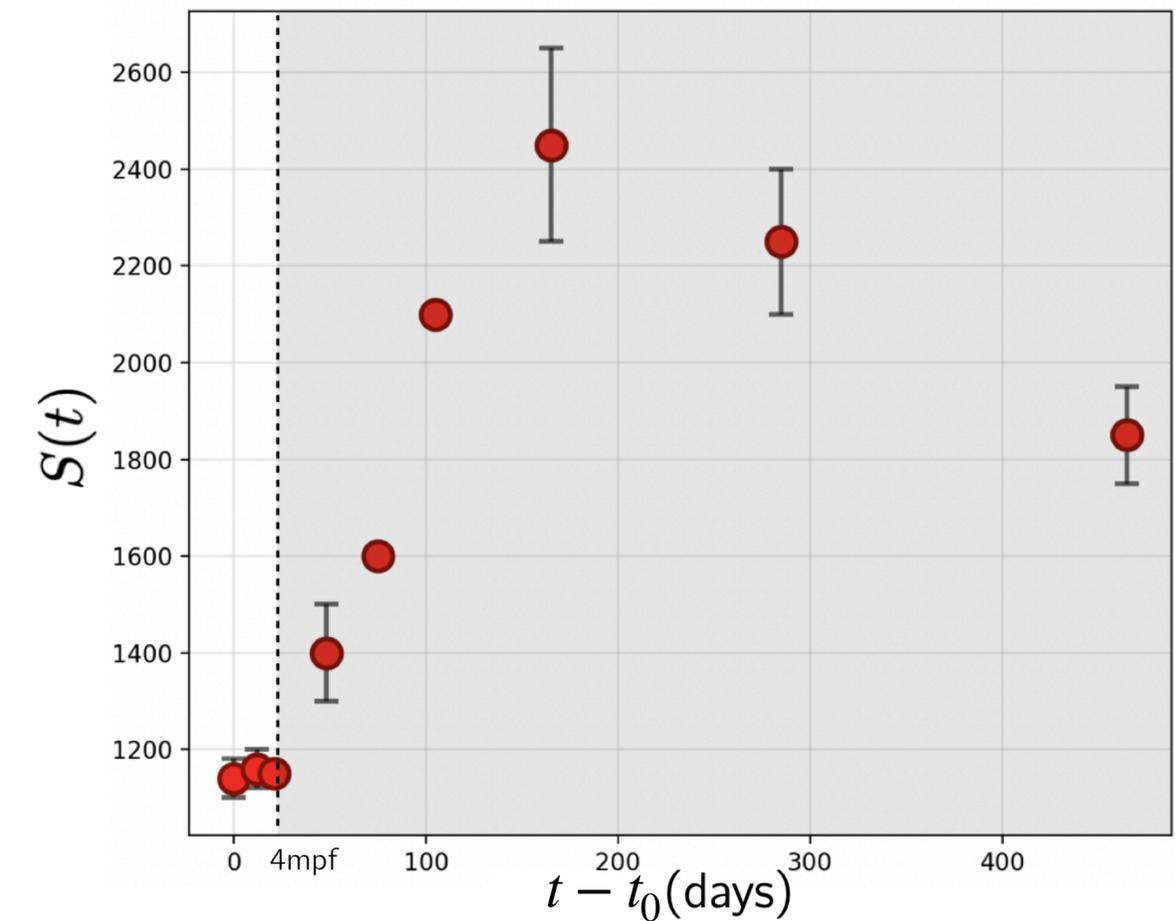


b. Slightly increase the self-renewal rate of reservoir aNSCs by $\epsilon > 0$.



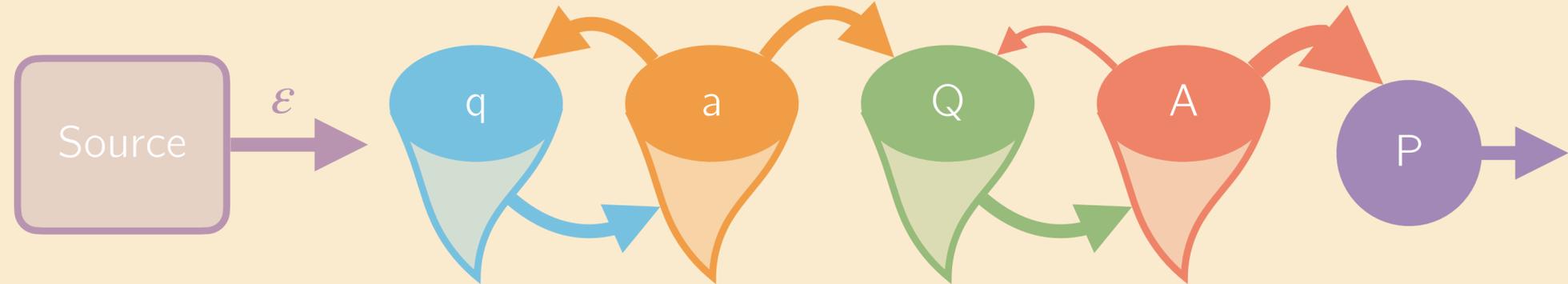
Investigation: Modeling the source

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

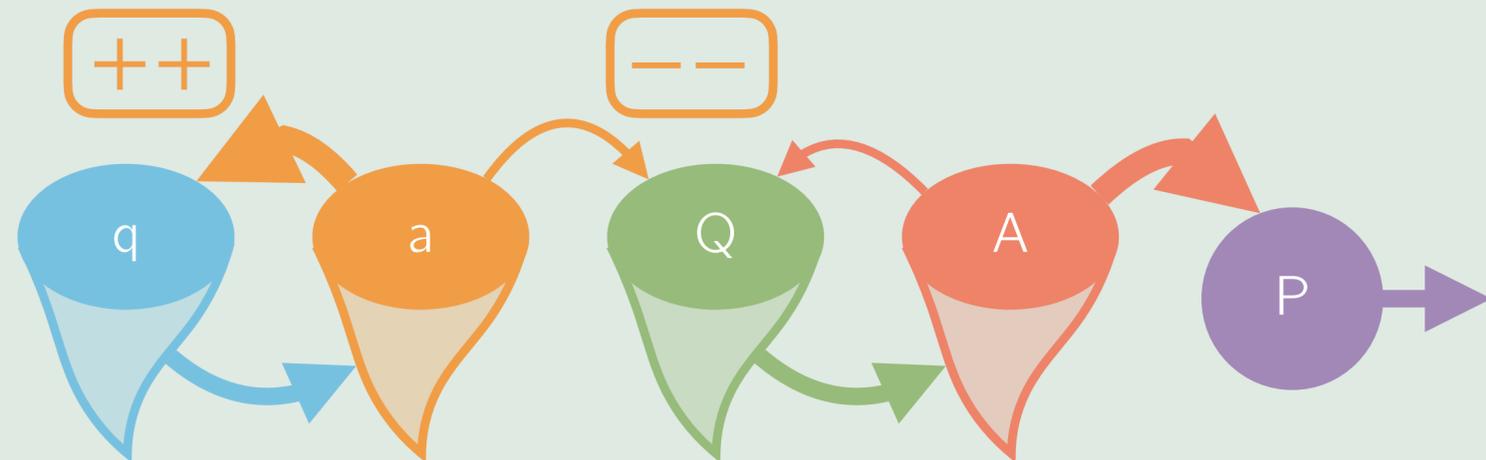


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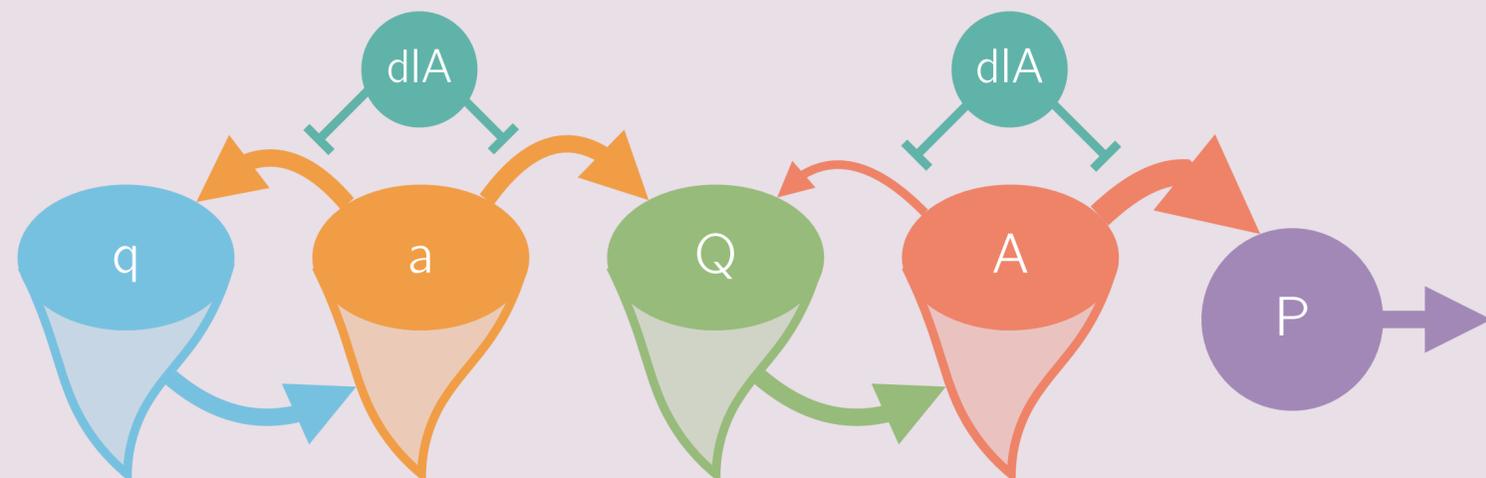
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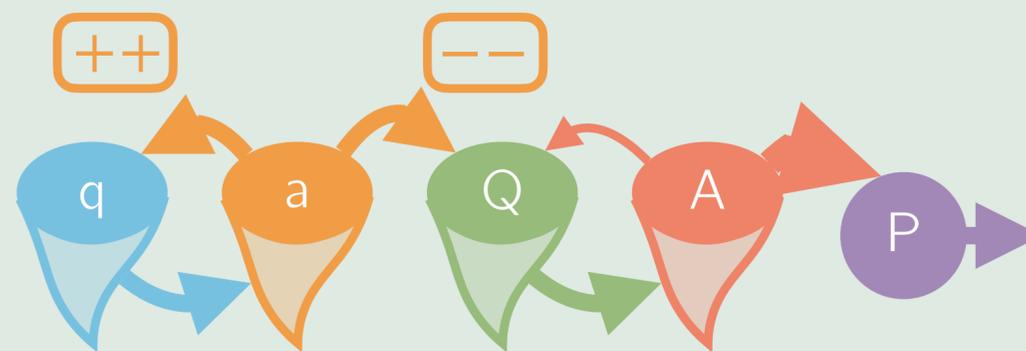
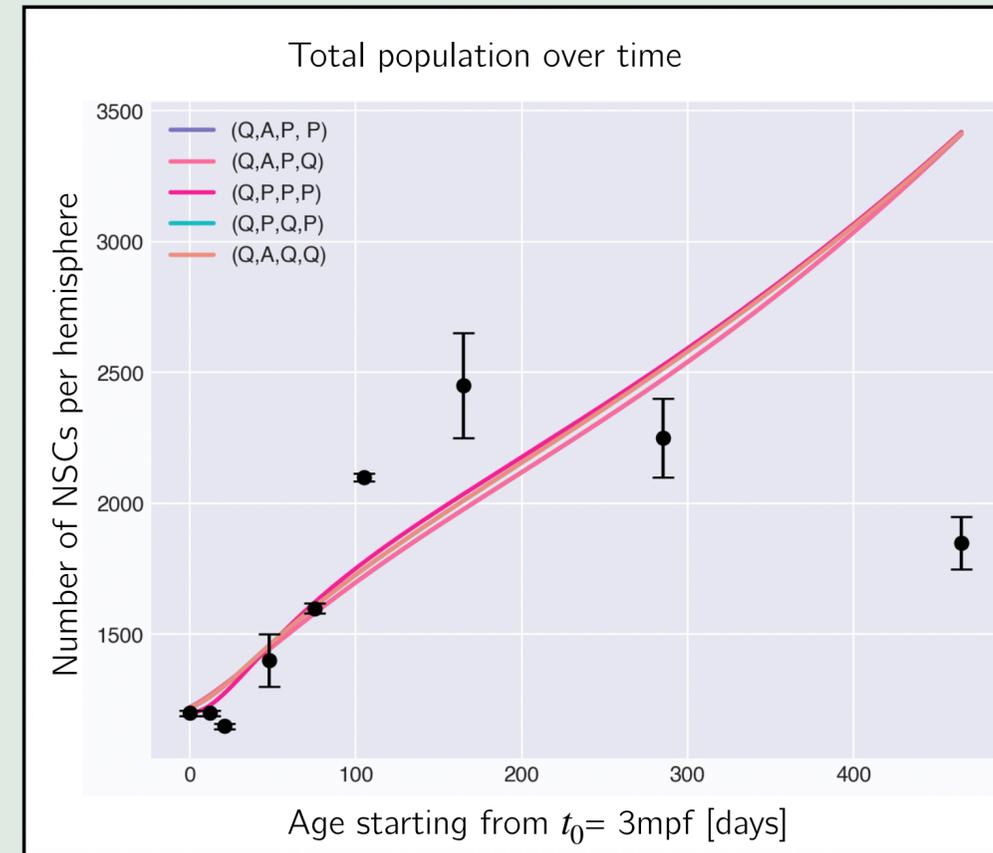
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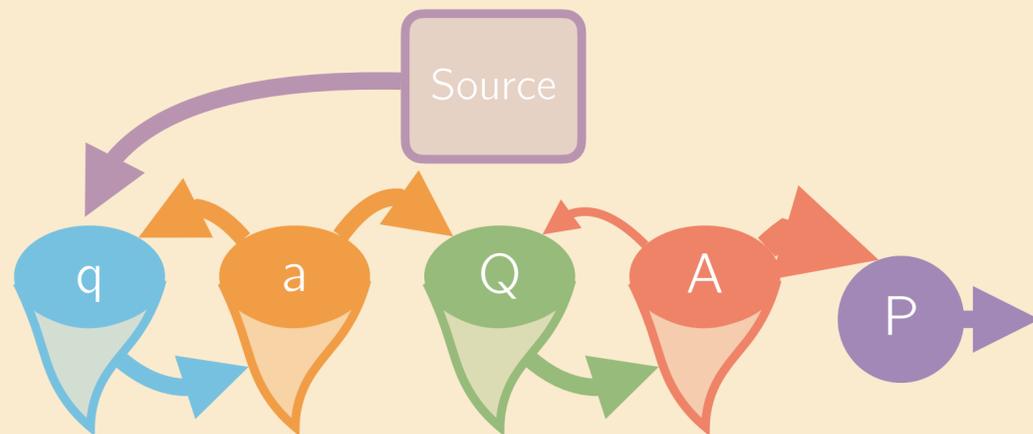
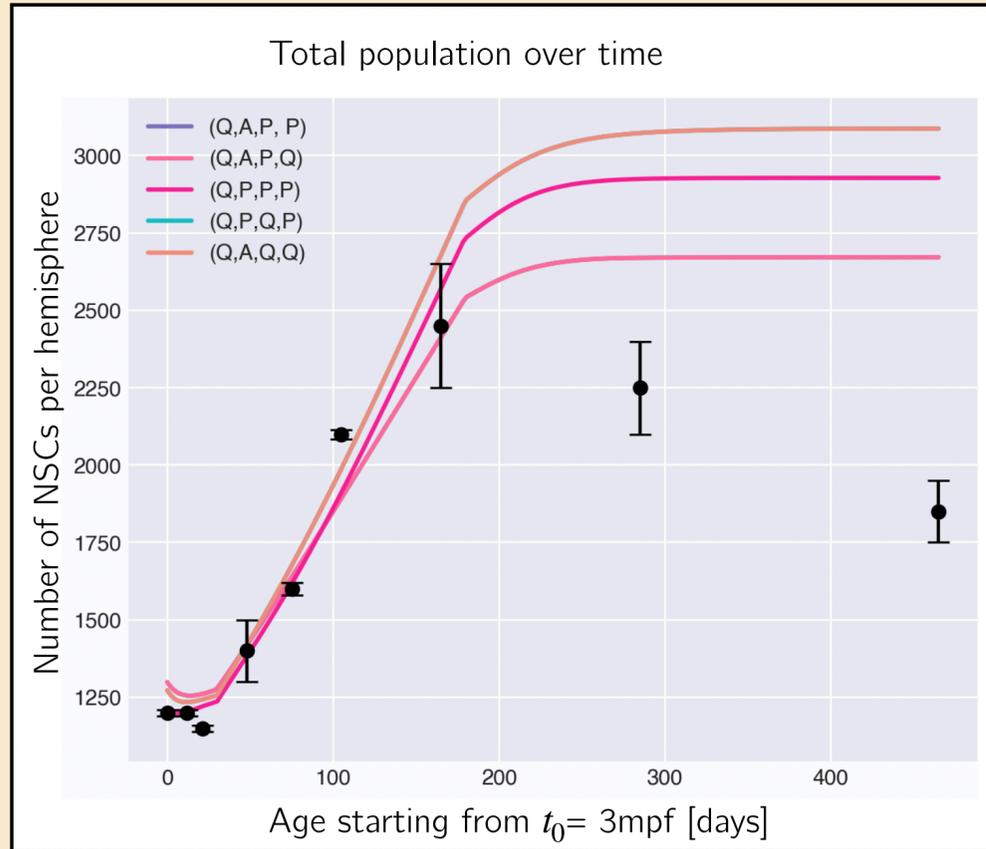
c. Notch-Delta signaling controls the self-renewal rate of aNSCs



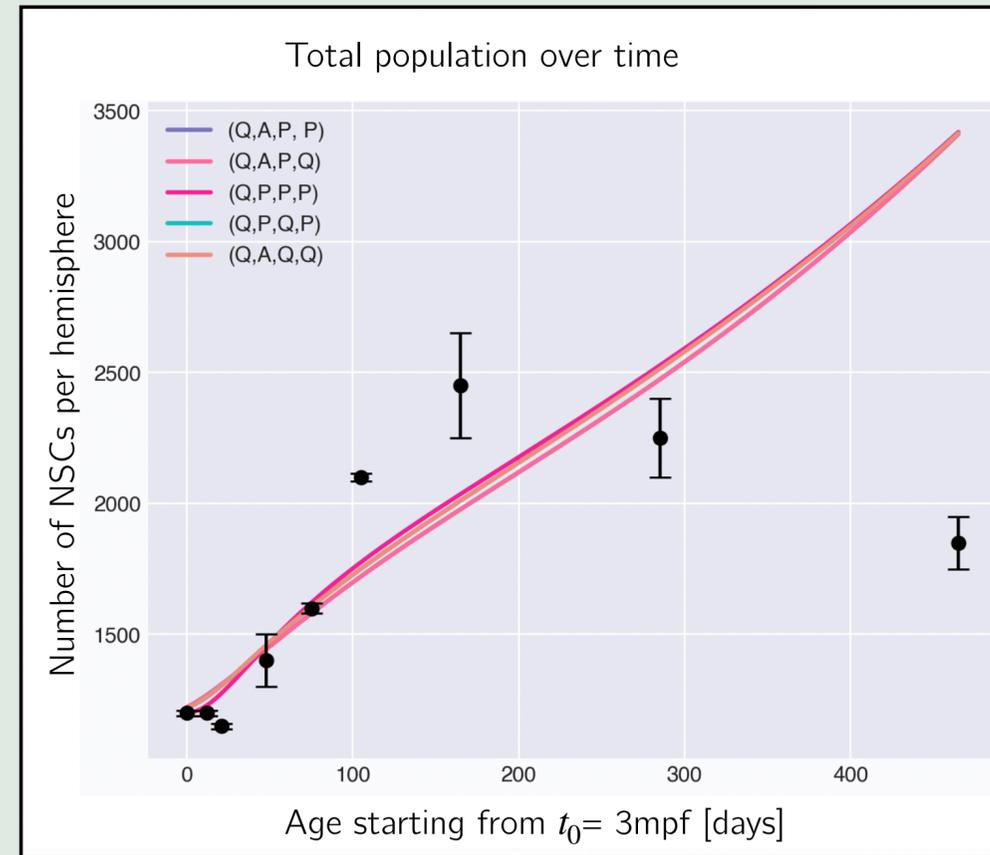
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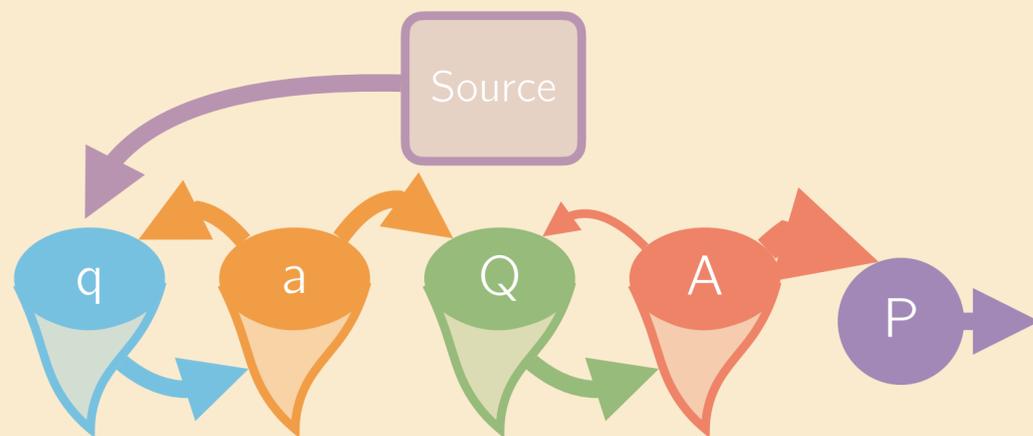
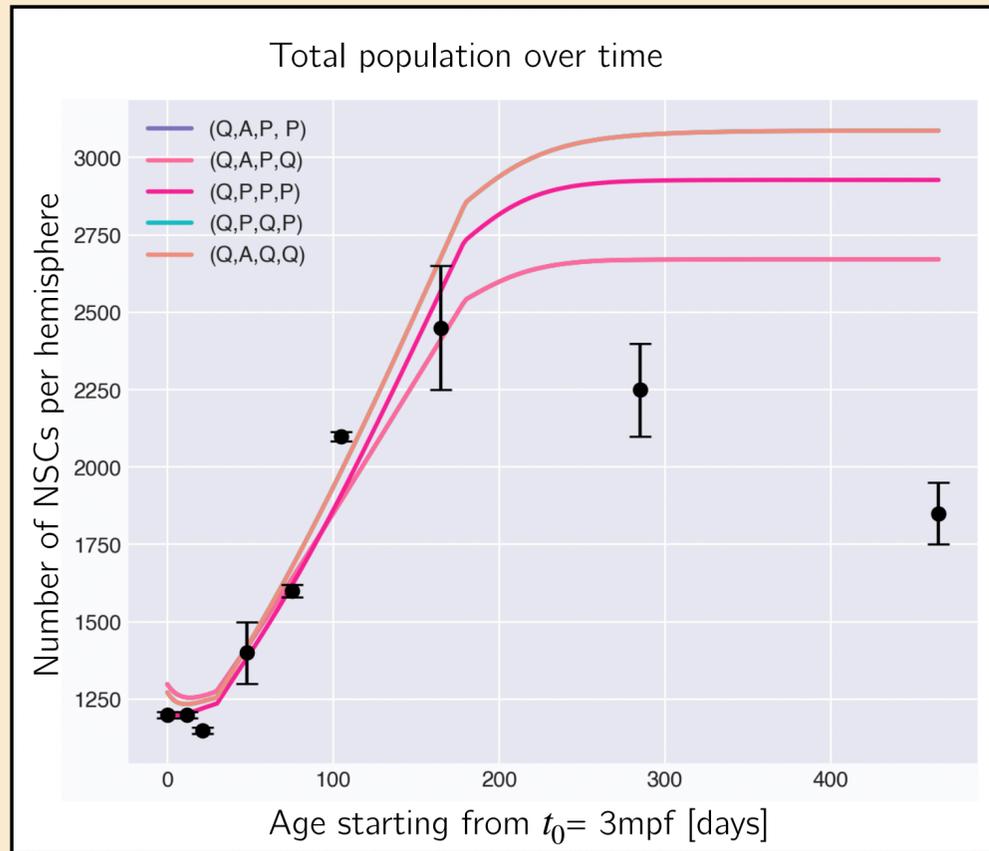
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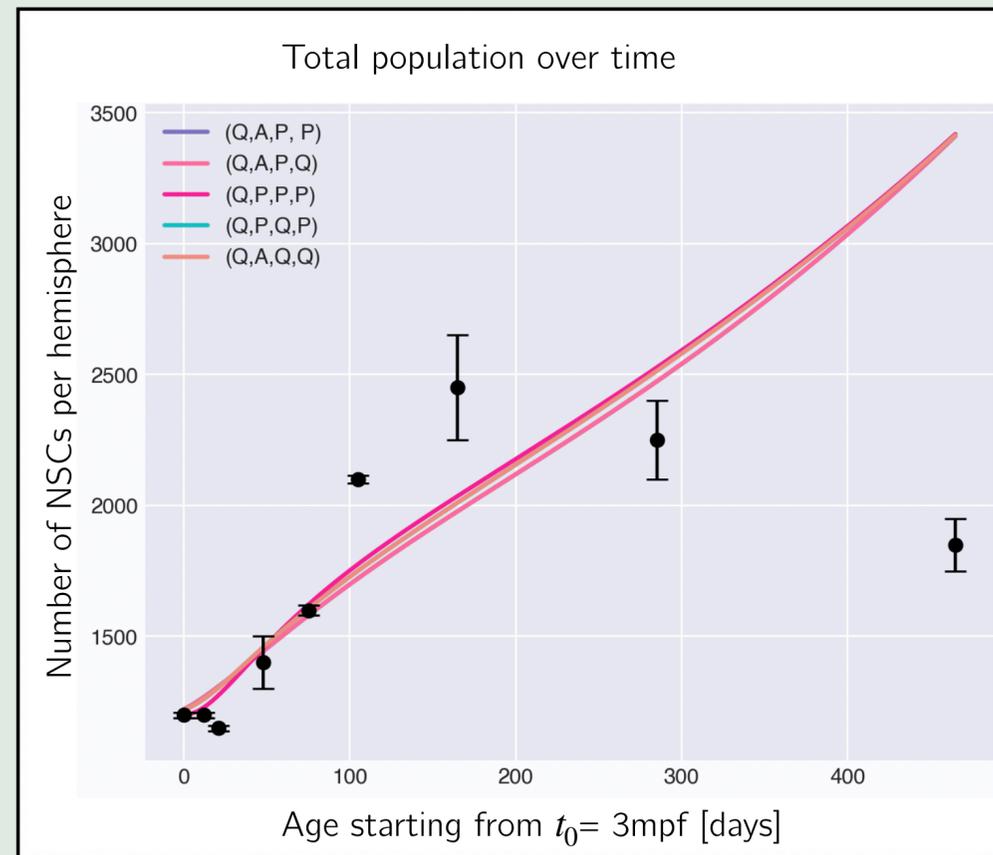
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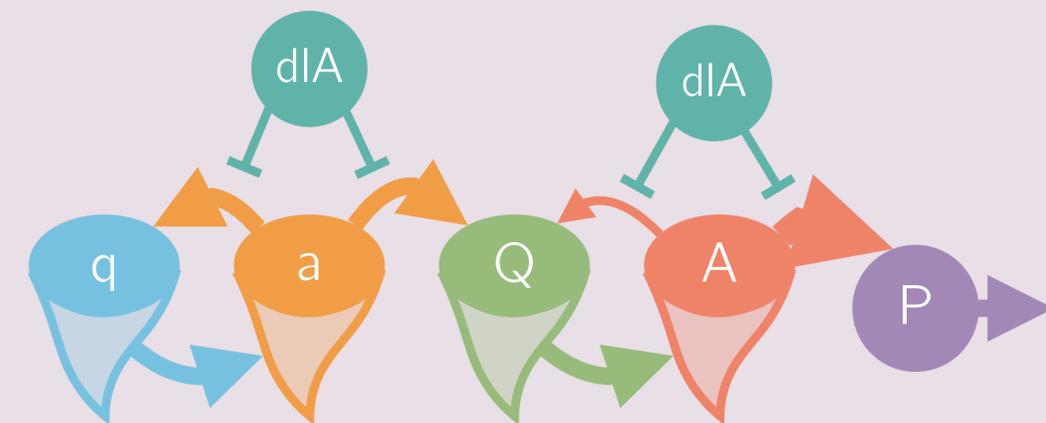
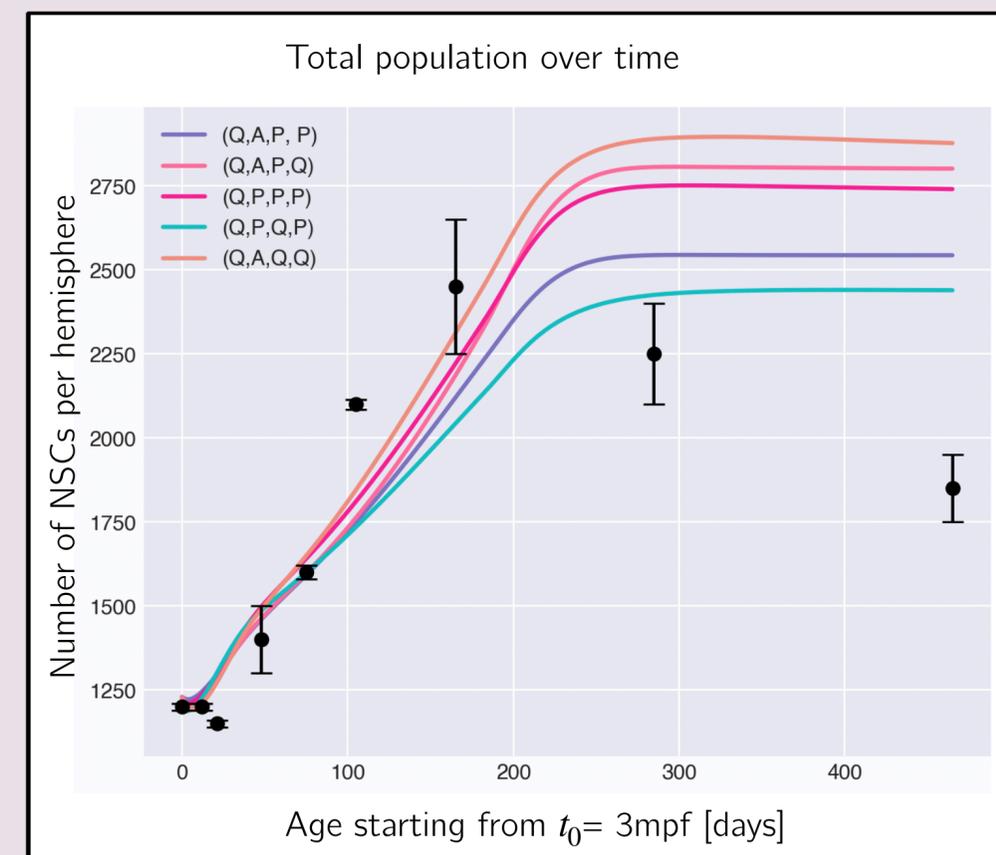
a. Introduce cells in the population, at a small constant rate $\epsilon > 0$.



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



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

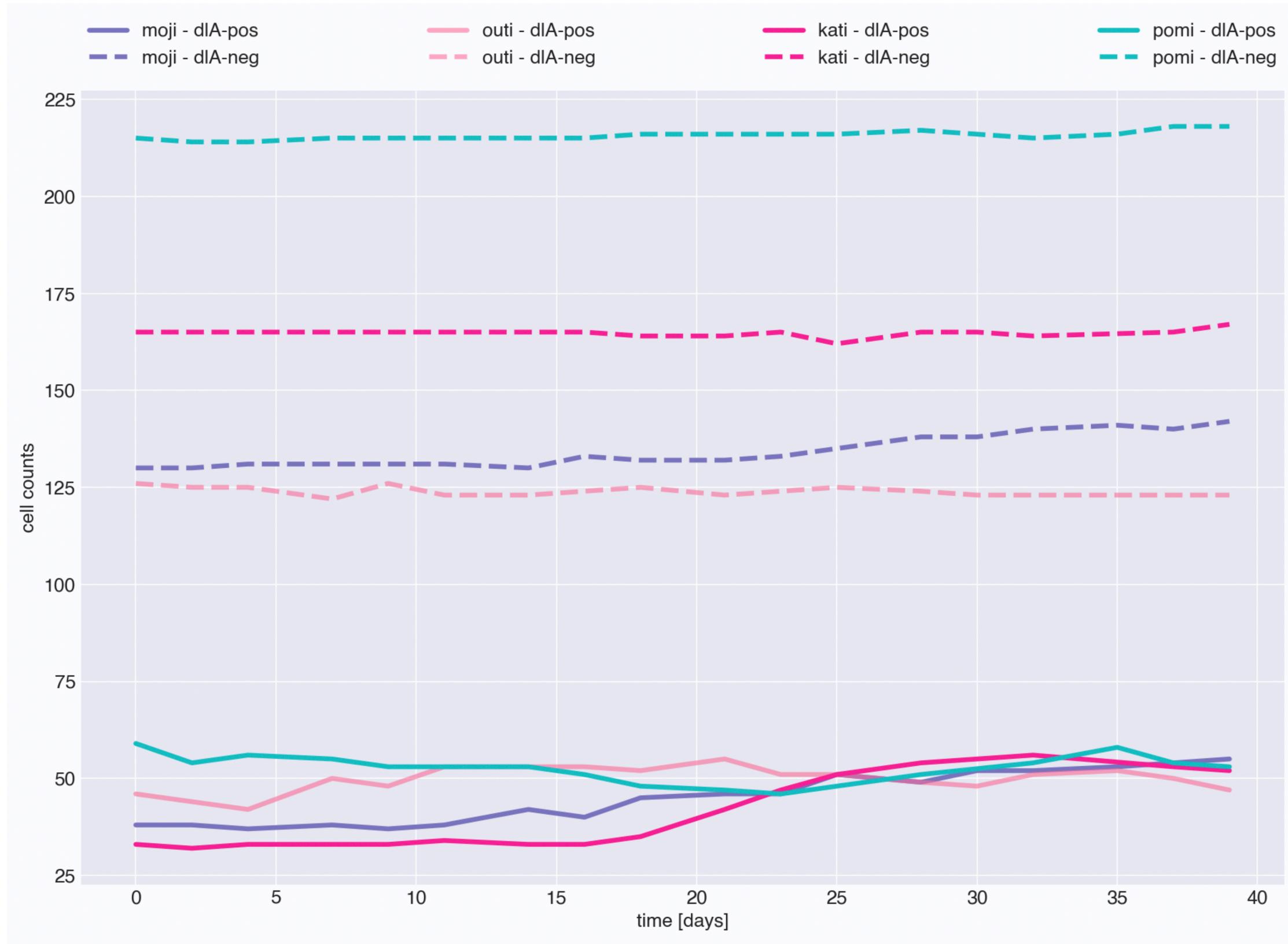
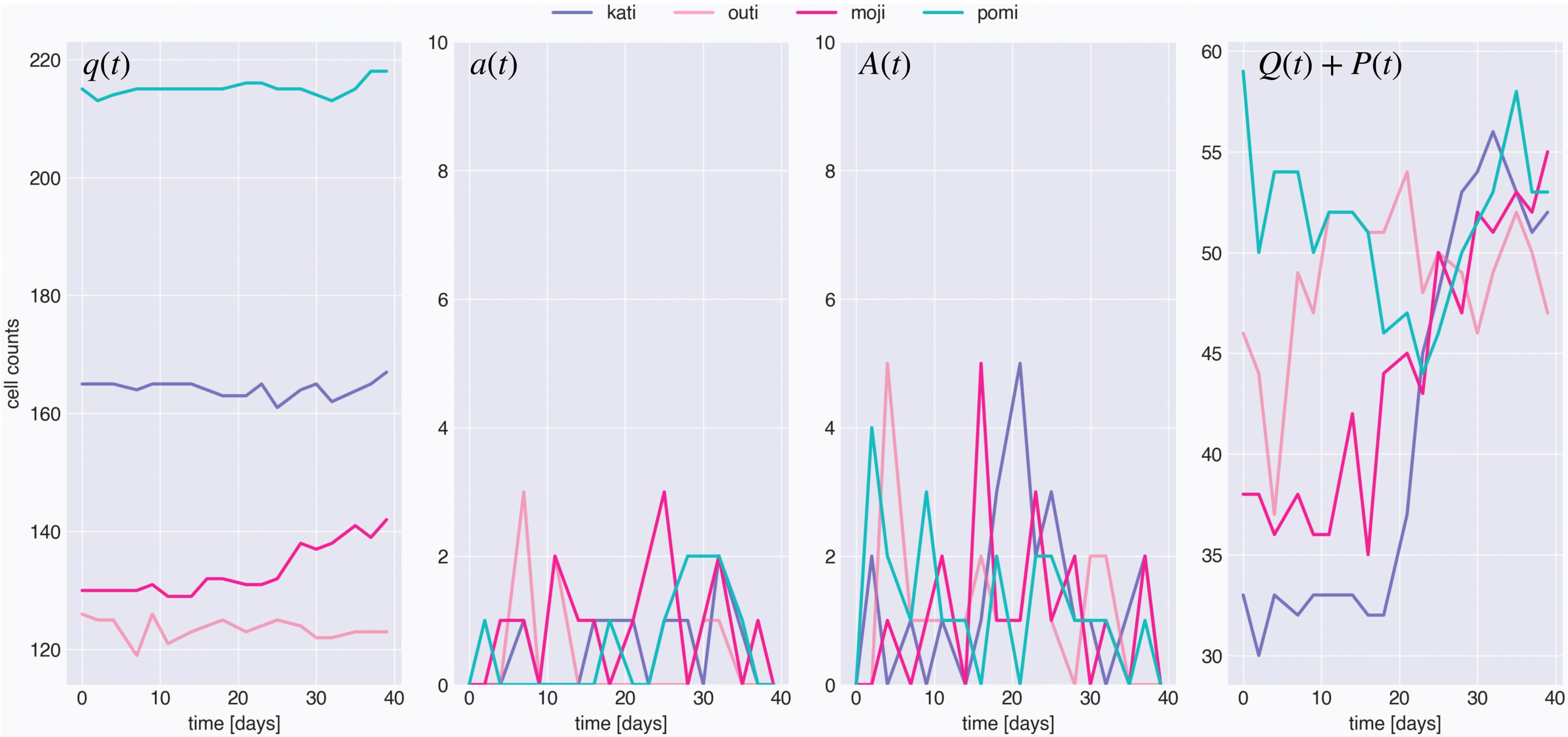


Figure (temp): Number of cells of each type over time per fish



Thank you for your attention!