In vivo clonal analysis for cell lineage in neurodevelopmental studies



Lineage tracing by clonal analysis

- Consists in quantitative analyses of clones originated from a single stem/progenitor cell
- Allows for **specific labeling** of single stem/progenitors
- Provides **spatial and/or temporal resolu**tion to investigate the lineage progression and fate specification of stem cells



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Genetic tools for prospective lineage tracing :

- ✓ Cre-Reporter lines
- ✓ Mosaic expression of multiple genes:
- Brainbow technology
- Confetti mice
- Star-Track plasmids
- Retroviral vector-mediated gene transfer
- RV carrying reporter genes (i.e. GFP)
- Barcoded Retroviral libraries (i.e. QmGFP-OL)

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Genetic tools for prospective lineage tracing :

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✓ Cre-Reporter lines





Cre-lines

Conditional - Inducible





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 Mosaic expression of multiple genes: based on Cre/lox recombination system

Site-specific recombination Excision No Α Inversion XFP lox2272 .--IoxP *loxN* ЧЧХ Yes No Yes No Yes B Excision: Е No recombination **IoxP** Brainbow 1.0 recombination (default) 2 1 lox2272 loxN Expression lox2272 recombination Cre-mediated recombination 2 F C Inversion: Brainbow 2.0 **IoxP** Expression Stochastic Cre-mediated Cre-mediated recombination recombination -D Excision and inversion: Brainbow 2.1, Brainbow AAV +•D> 2 IoxP 1 -----**IoxP** Expression G Multi-vector: 440 TIE-DYE, CLoNe Excision 2 Excision 1 Expression CHP Cre

Brainbow technology



Stochastically express two to four XFPs in a cellular population from a single promoter

When multiple copies of Brainbow are present in a cell, each copy recombines independently.

3 copies of Brainbow can generate 10 distinct colors and more copies will generate even greater color diversity.



Mosaic expression of multiple genes:

R26R Confetti Mice (Brainbow 2.1)



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✓ Mosaic expression of multiple genes:

Star Track method – a genetic tracing strategy for long term lineage of individual glial progenitors in vivo -



→ Following PBase-mediated transposition, fluorophores are randomly integrated into the cell genome

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Genetic tools for prospective lineage tracing :

- Retroviral vector-mediated gene transfer
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Retrovirus iv injection

- allows viruses to introduce recombinant DNA into the genome of a host cell (proliferating)
- the integrated exogenous DNA is then inherited by all the descendants of the infected cell

Multiple progenitor domains in the developing telencephalon





Multiple progenitor domains in the developing telencephalon





Lineage clonal analysis using Retroviral vector-mediated gene transfer to study MGE-derived inhibitory interneurons



Neuron Article

Wide Dispersion and Diversity of Clonally Related Inhibitory Interneurons

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http://dx.doi.org/10.1016/j.neuron.2015.07.030



Harwell et al., Neuron 2015

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Intraventricular retroviral injection



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RV-GFP intraventricular injection allows clonal lineage analysis BUT only if the progeny remains spatially confined

Need for an alternative method to follow the distribution of cells with broad migratory capacity in relation to their clonal siblings





RV-GFP intraventricular injection allows clonal lineage analysis BUT only if the progeny remains spatially confined

Need for an alternative method to follow the distribution of cells with broad migratory capacity in relation to their clonal siblings



Barcoded Retroviral libraries

EnvA-pseudotyped retrovirus library carrying oligonucleotide sequence tags or barcodes





Cells were labeled from each injection (n = 8 hemispheres, 4 brains) - between 14 and 234

The probability of neurons containing the same tag arising from two independent progenitor infections is low

→ Multiple cells sharing the same barcode are presumed to be clonally related siblings



Immunofluorescence for INs subtype markers PV and SOM

Spatial mapping of GFP/PV + and GFP/SOM + cells



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Brain sectioning

Immunofluorescence for INs subtype markers PV and SOM

Spatial mapping of GFP/PV + and GFP/SOM + cells

Laser capture microdissection of labelled cells

Viral tag amplification by PCR







Interneuron Clones Consist of Widely Dispersed Cells of Diverse Subtypes







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Wide Dispersion and Diversity of Clonally Related Interneurons



MGE interneurons derived from a common progenitor are widely dispersed across different regions of the brain

 $\rightarrow \mbox{the majority of clustered cells observed with sparse viral labeling are likely not clonally related$



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Barcoded Retroviral libraries (i.e. QmGFP-OL)

Advantages: allow determining lineage relationships in populations of cells regardless of their pattern of migration

→tags recovered from only 43% of GFP-positive cells

Limits: gene silencing \rightarrow can account for some reduction in clone size









courtesy of A. Puche





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Adult neurogenesis: a multi step process



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Maturation & selection of adult born OB interneurons

quantitative estimation of the absolute number of new neurons recruited into the bulb



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Embryonic Origin of Postnatal Neural Stem Cells

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http://dx.doi.org/10.1016/j.cell.2015.05.041



Retroviral barcode library: each retrovirus expresses mGFP and contains a unique 24 bp sequence



Retroviral barcode library injected intraventricularly at different embryonic stages



- individually labeled cells were identified as neurons, astrocytes, and oligodendrocytes:
- 1) based on their morphology, revealed by mGFP
- 2) by the expression of NeuN, GFAP, and Olig2

Their locations were mapped, and individual labeled cells were collected using laser capture





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Clonal Relationships between Postnatal OB Interneurons and Cx, Hp, St, and Sp Neurons



Inhibitory GABA-ergic Interneurons belong to different sub-types



GL, glomerular layer; GCL, granule cell layer





Mosaic Organization of Neural Stem Cells in the Adult Brain Florian T. Merkle, *et al. Science* **317**, 381 (2007); DOI: 10.1126/science.1144914

Model of SVZ stem cell potential





(Merkle et al., 2007)



label radial glia in a regionally specific manner



Specifically targeting stem cells and follow their progeny in vivo by Ad-Cre (Adenovirus expressing Cre recombinase) injection into GFP reporter mice.

Little diffusion \rightarrow localized injection (small volumes)

Ad-Cre infected radial glia and their progeny become permanently labeled with GFP



Targeting 15 different populations of radial glial cells at six different rostrocaudal levels (i to vi)



Specific target of spatially restricted radial glia by Ad-Cre and of adult SVZ stem cells they generate

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→different types of OB interneurons are derived from different locations in the SVZ

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A mosaic of early specified neural stem/progenitor cells contribute to the different OB interneurons subtypes



Cheuk Ka Tong and Arturo Alvarez-Buylla SnapShot Neuron 2014

NSC are regionally specified in both the neonatal and adult SVZ



Activity dependent regulation of Adult neurogenesis in the OB

Two basic experimental paradigms to study the effect of the **environment** on **neural plasticity in the OB**



Odor enrichment





Neural plasticity in the OB: focus on TH+ dopaminergic cells Dopaminergic interneurons





Odor deprivation by naris occlusion

Is reversible

decreased TH expression

Baker et al., J Neurosci 1983 Baker et al., Brain Res. 1993 Nasal plug removal



Cummings and Brunjes, J Neurosci. 1997



Odor enrichment increses TH+in the OB





42 Bonzano et al., EJN 2014

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