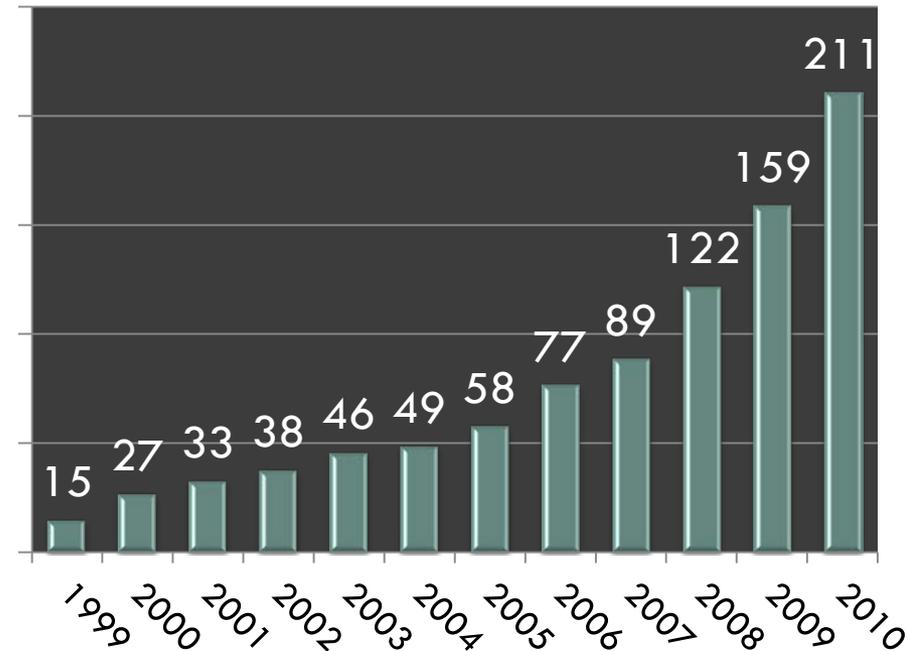
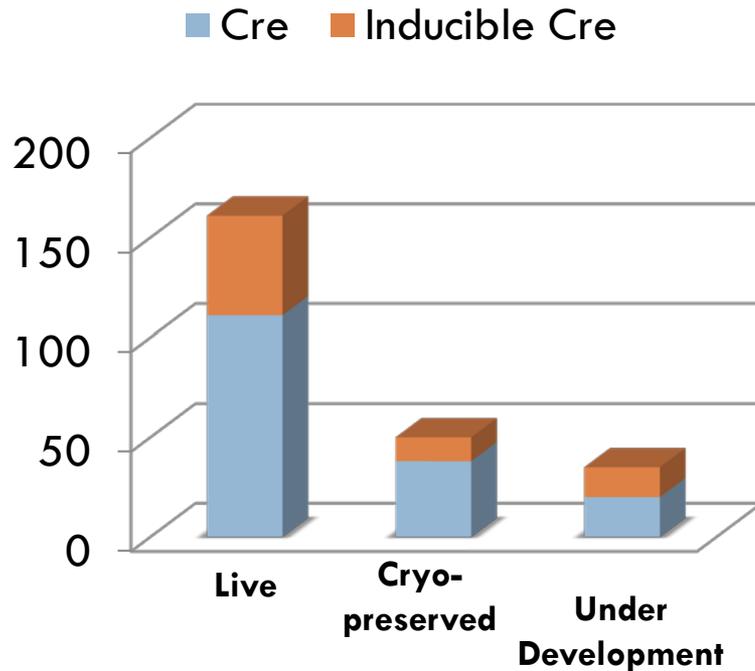


Leveraging conditional IKMC alleles: Cre driver strain resources

- Enabling access: critical role for Repositories
 - Live versus cryo
 - Quality control: transfer of responsibility
 - Directed versus “laissez faire” promotion strategies
- Detailed characterization
 - Add to and complement published data
 - Particularly important for Cres developed in large sets
 - Positive AND negative data
- Data dissemination
 - Centralized sources of Cre functional data
 - CREATE BioMART and CrePortal

Cre strain distribution

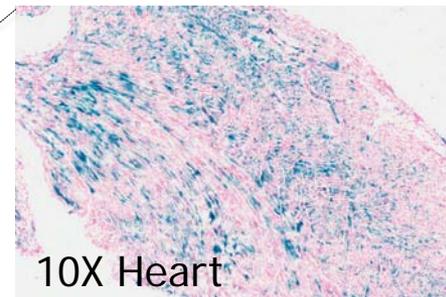
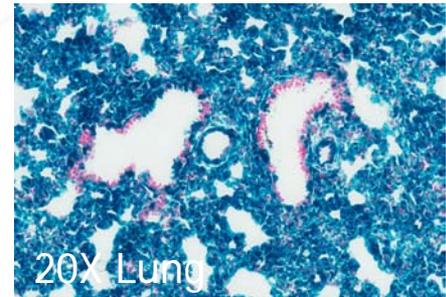
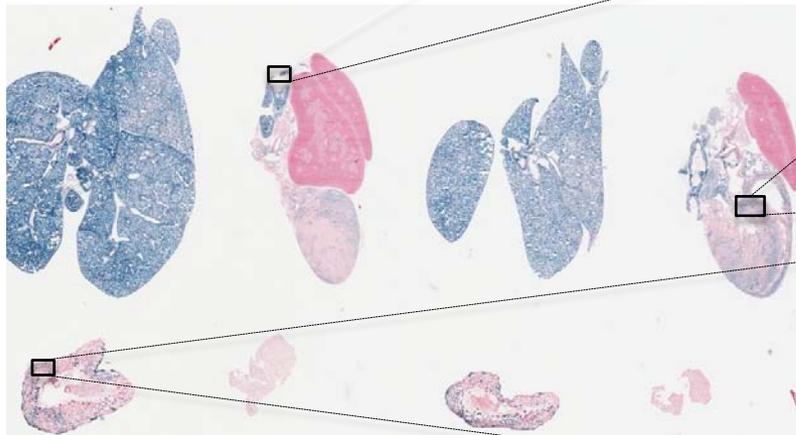


Growth increasingly driven by large sets (e.g. Neuroscience Blueprint) that have limited community use prior to distribution.

Open question: how do we best manage our Cre strain distribution to best serve the community?

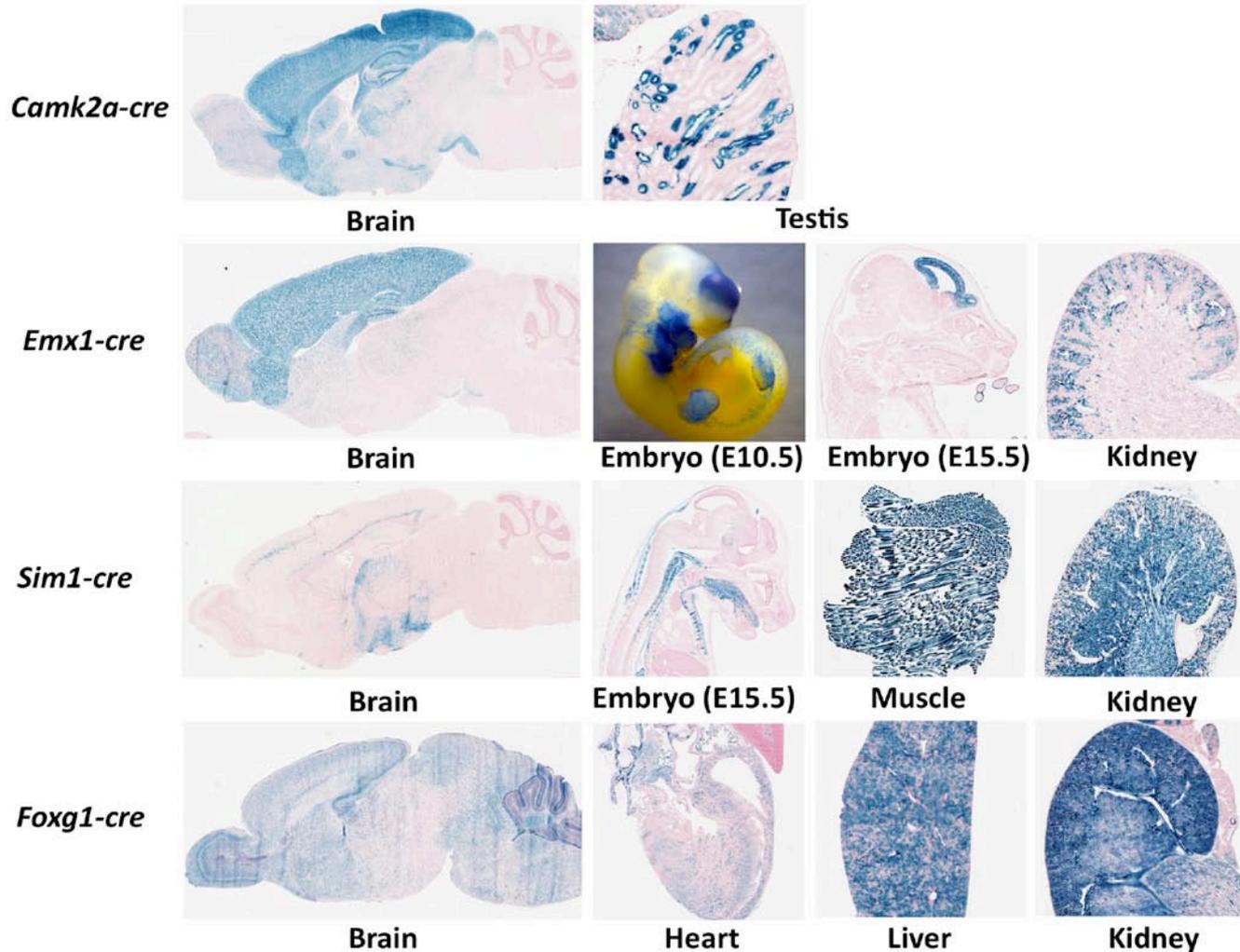
Comprehensive characterization of Cre driver pipeline (cre.jax.org)

- R26R-LacZ Reporter strain
- Four time points: E10.5, E15.5, P7 and P56 (adult)
- Whole-mount (E10.5) or frozen sections
- Full necropsy at P7 and P56 (11 organ systems, 30 organs/structures, 89 substructures)
- NanoZoomer slide scanner to capture data at 20X
- Data collected for 45 strains, 44 in progress
- Data delivered to creportal.org

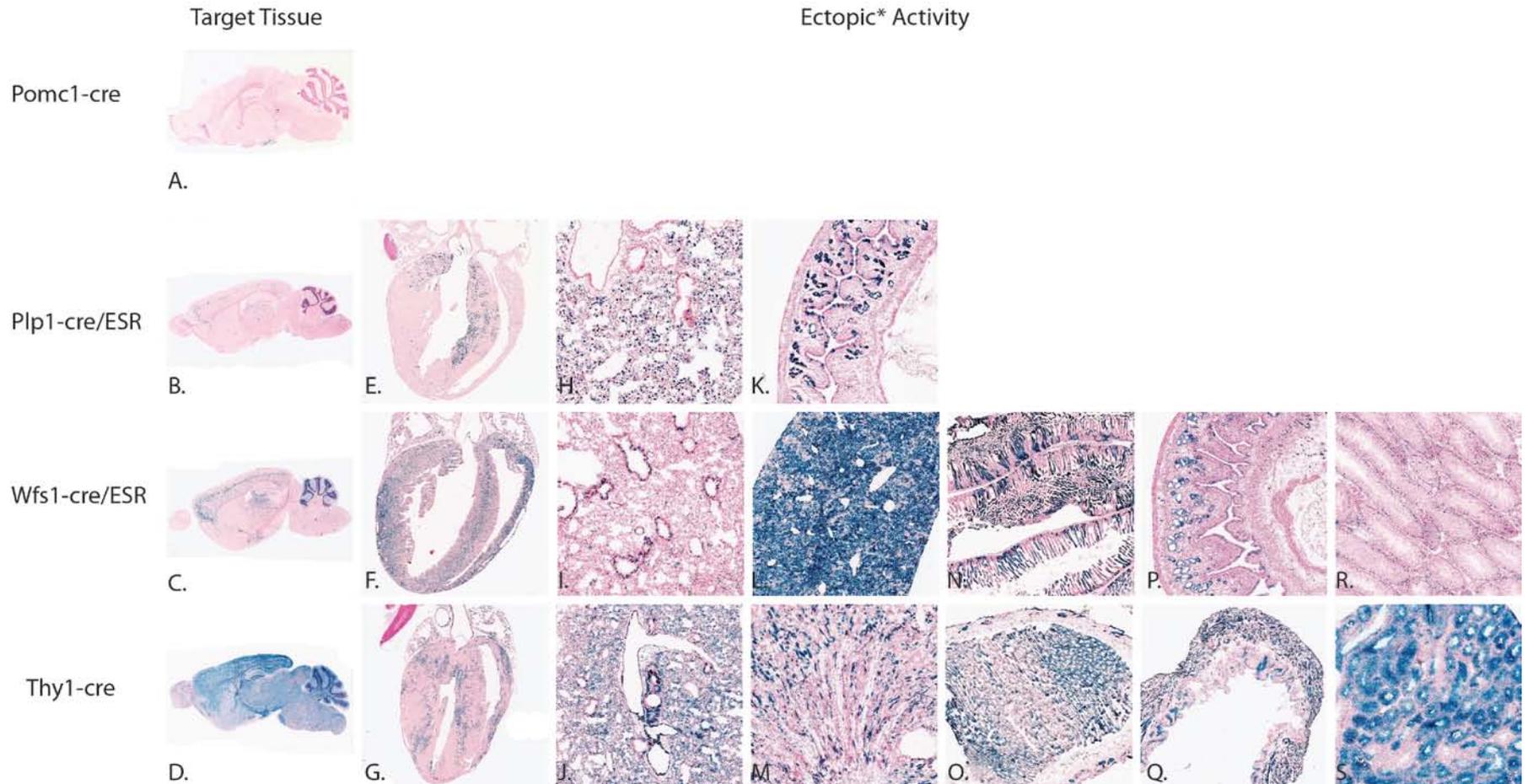


129(Cg)-Foxg1^{tm1(cre)Skh}/J Characterized with R26R LacZ Reporter

Off-target Cre activity: a surprisingly common problem



Ectopic* activity continued

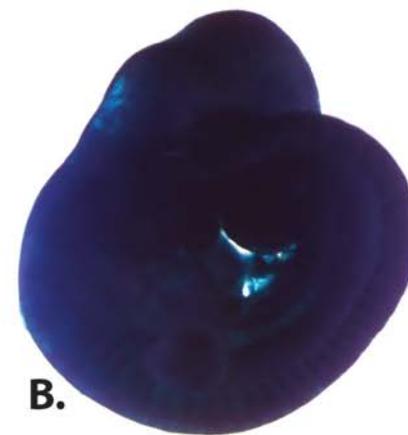
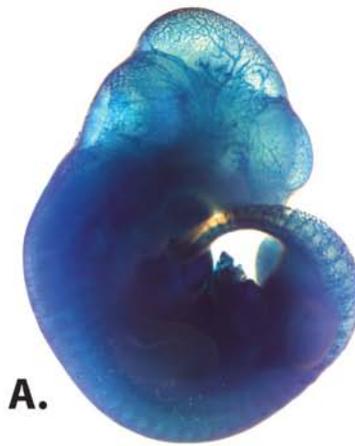


Inconsistent activity

Regionally Restricted

Widespread

B6.Cg-Tg(Tek-cre)12Flv/J



B6.Cg-Tg(Vav1-cre)A2Kio/J



Recombinase (cre) Home (www.creportal.org)

Recombinase (cre) Specificity

MGI collects and annotates expression and activity data for recombinase-containing transgenes and knock-in alleles.

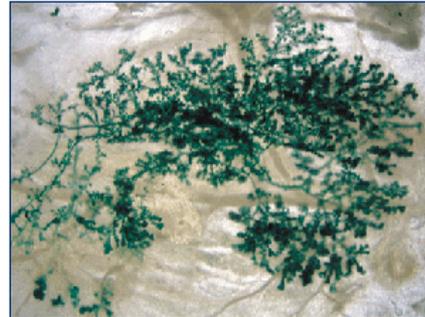


Image reproduced with permission of the Journal of Cell Science. Rijnkels M and Rosen, JM, J Cell Sci 2001, 114(17): 3147-3153.

Access Data

FIND RECOMBINASE-CARRYING ALLELES

Search for alleles assayed for specificity/expression in a particular anatomical system.

Recombinase specificity in: (choose one)

Search for alleles by promoter/driver specificity.

Recombinase driven by: (choose one)

RETRIEVE ALL ALLELES

Retrieve a list of all recombinase-containing transgenes and knock-in alleles.

[MGI Recombinase Alleles Report](#)

[Tab-delimited version](#)

RELATIONS

Search for Alleles

Search for Alleles

1,516 total Cre driver strains

~50% fully curated for Cre activity

FAQs

How do I...

.. find existing recombinase-expressing transgenes and knock-ins that have a given promoter (driver)? [FAQ](#)

ie? [FAQ](#)

12 Mar 2010

248 Drivers in recombinase knock-in alleles

2,165 Alleles in recombinase specificity assays

[More...](#)

October 29, 2009

curated from the scientific literature and integrated from projects underway at collaborating organizations. (See the **Collaborators** tab, below).
Alleles) consortium.

Search for cre activity by anatomical system or by driver/promoter

Get a report of all recombinase knock-ins and transgenes



Tg(AMH-cre)1Flor - Reproductive System

Recombinase Specificity Detail

[Allele Information](#) | [Tissue Information](#) | [Images](#) | [Recombinase Specificity](#) | [References](#)

| | | |
|---------------------------|---|---|
| Allele Information | Allele: Tg(AMH-cre)1Flor transgene insertion 1, Florian Guillou | Driver: AMH Type: Transgenic (Cre/Flp) |
| | Synonym: AMH-cre | |
| | Molecular description: The transgene shows Cre activity in AMH-expressing cells of the testis. | Gene, and genomic sequences from MT1 to male mice and granulosa cells of 16 day old mice are available. |
| | Find mice (IMSR): Mice with this allele are available from the IMSR. | |
| | Additional Tissues: Tg(AMH-cre)1Flor is also active in the following tissues: Alimentary | |

- Molecular description of the transgene or knock-in
- Links to mice in IMSR
- Other tissues with cre activity

| | | |
|---------------------------|----------------------------|---|
| Tissue Information | Reproductive System | Other tissues: Alimentary Brain Cardiovascular Endocrine Eye Gastrointestinal Hematopoietic Immune Integument Musculoskeletal Nervous Respiratory Skin Skeletal Stomach Urogenital Whole Body |
|---------------------------|----------------------------|---|

Other recombinase knock-ins/Tgs with activity in reproductive system

Images *Drag images to compare to others*

J:81650 Fig. 3

Gallery of images for this cre in this tissue/organ

Recombinase Specificity *Click heading to resort table.*

| Structure | Assay | Assay Note |
|---------------------|------------------|--|
| testis | embryonic 15 | |
| seminiferous tubule | postnatal week 9 | Present Regionally restricted J:157337 Fig. S1C right panel Figure not shown |

| Assays | Genotypic Background | Result Notes | |
|----------------------|----------------------|------------------|------------|
| Assay Type | Reporter Gene | Detection Method | Assay Note |
| Recombinase reporter | lacZ | Direct Detection | |
| Recombinase reporter | lacZ | Direct Detection | |

Recombinase activity data

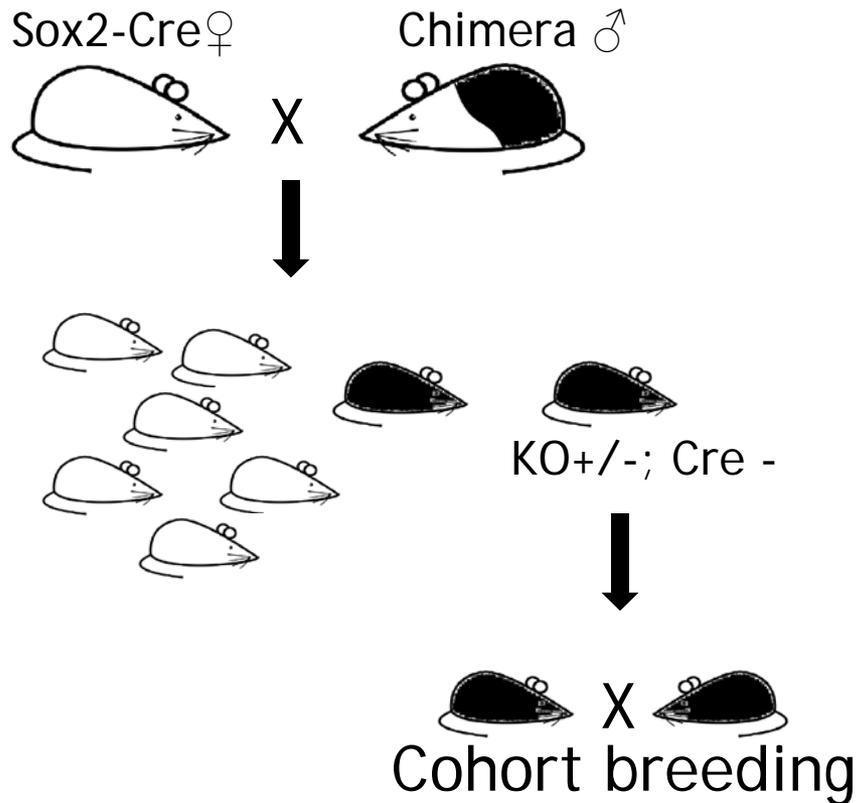
| | |
|-------------------|---|
| References | All for this allele: 34 reference(s) |
|-------------------|---|

[Link to References](#)

Cre strains for KOMP2, IMPC

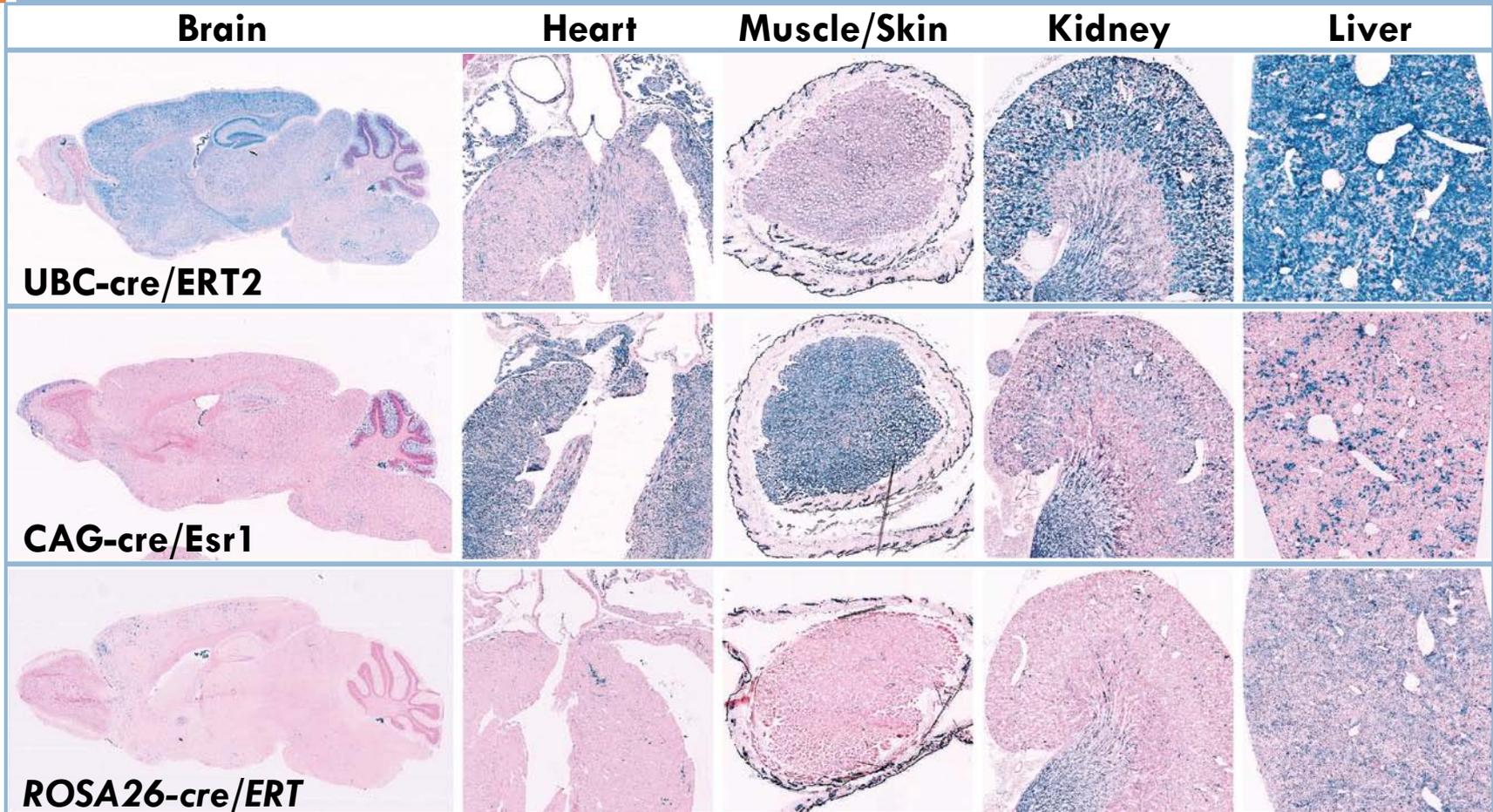
- Cre deleter
 - C57BL/6N background
 - Highly efficient
 - Maternal deletion 'feature' (Sox2-Cre; ICS Cre-GFP)
- FLPe germline deleter strains on B6N needed to convert to conditional alleles
- Inducible Cre (ERT2) and tissue specific Cre strains for use with lethals
- Need to move well-characterized, well-published Cres to B6N

JAX Cre deleter choice: Sox2-Cre



- Deletion activity independent of Cre allele segregation
- ▣ Speed congenic to C57BL/6N using new N/J SNP panel

Comparison of ubiquitous CreER lines



Thanks to...

Cre Characterization Pipeline

Caleb Heffner

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Cat Lutz

Steve Rockwood

Mike Sasner

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RR032656

CrePortal

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Peter Frost

Martin Ringwald

James Kadin

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