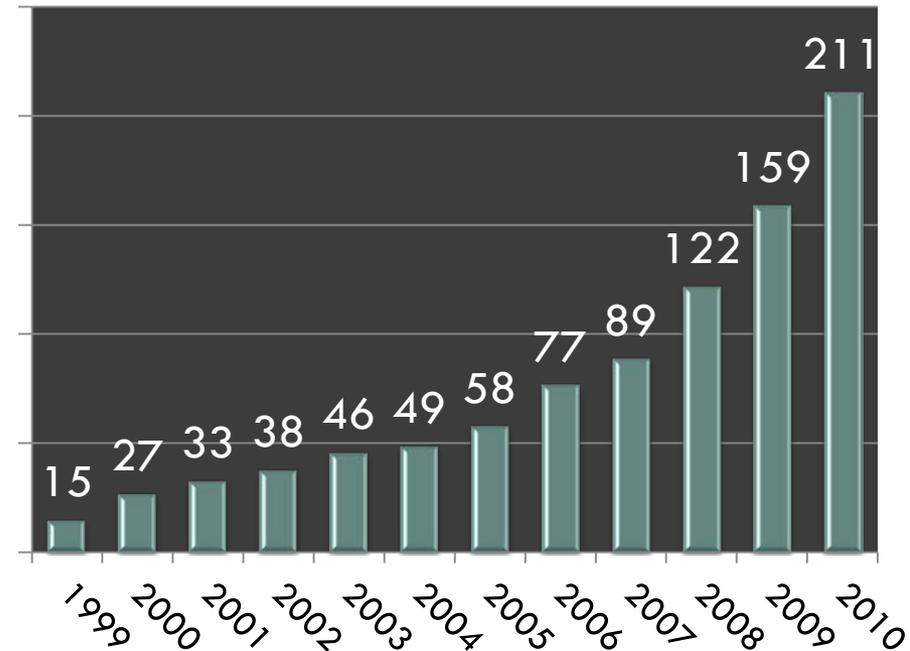
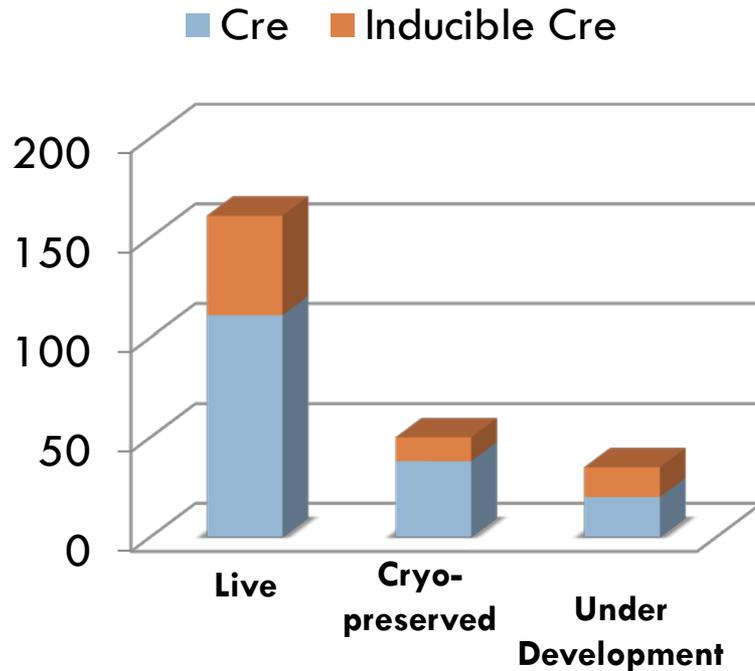


Leveraging conditional IKMC alleles: Cre driver strain resources

- 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

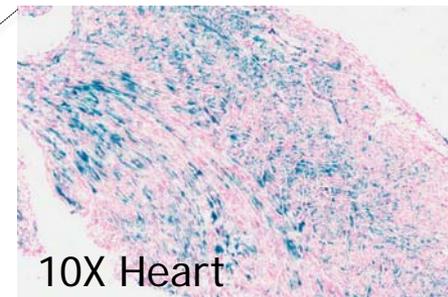
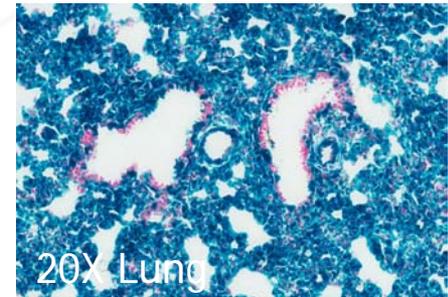
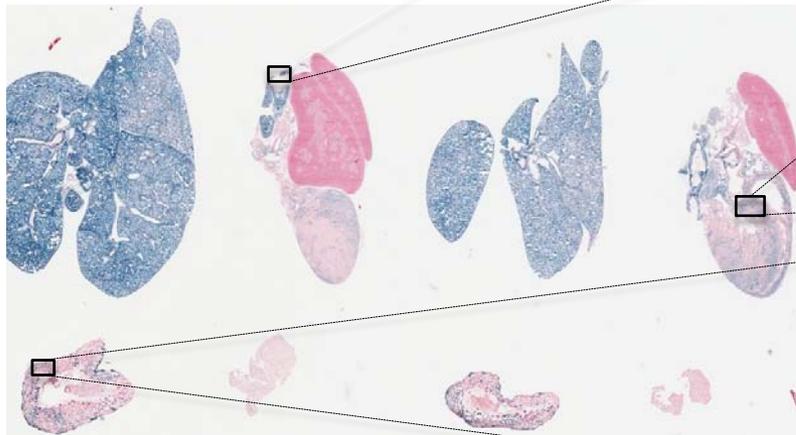


Growth increasingly driven by large sets (e.g. Neuroscience Blueprint) that have limited community use prior to distribution.

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

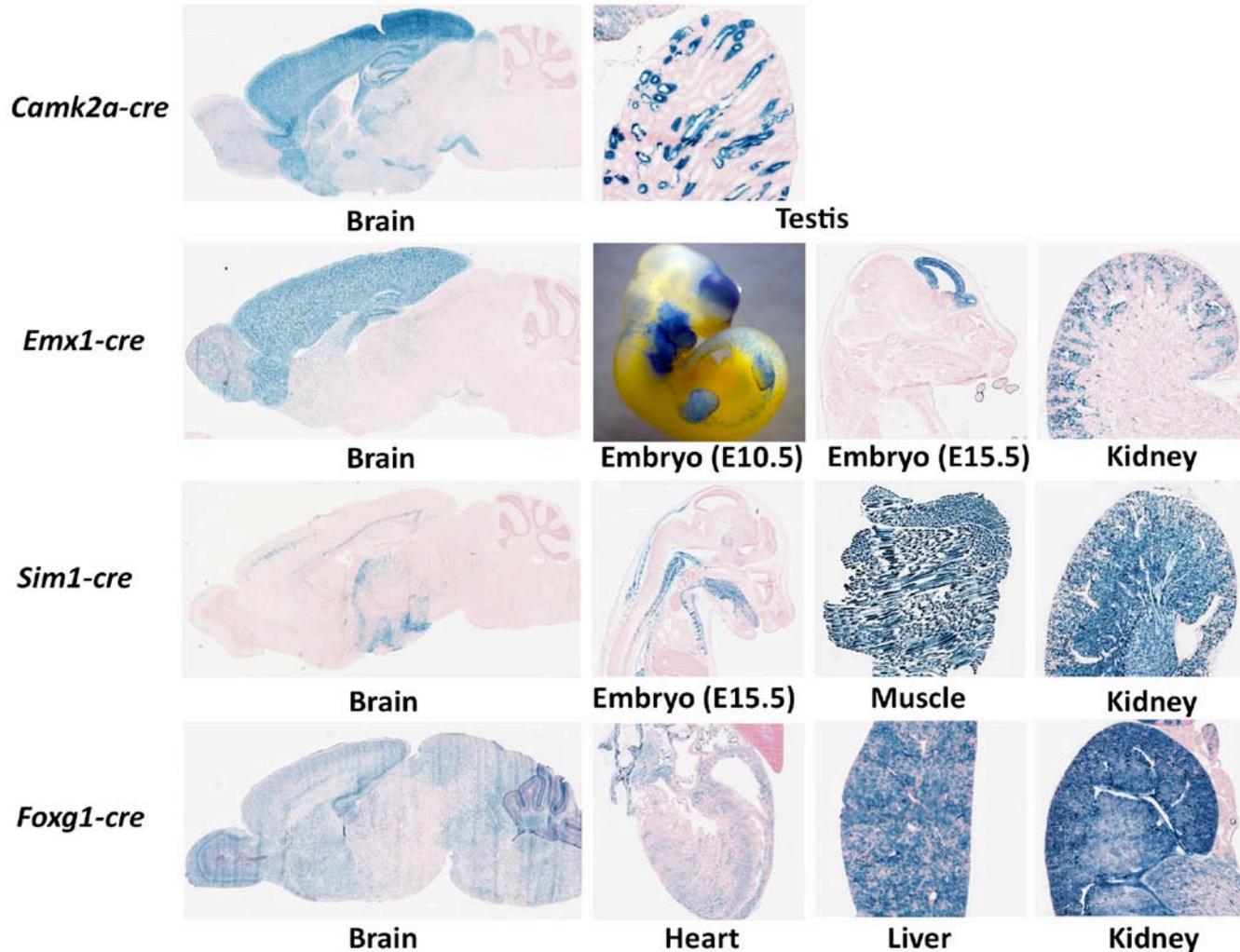
Comprehensive characterization of Cre driver pipeline (cre.jax.org)

- 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

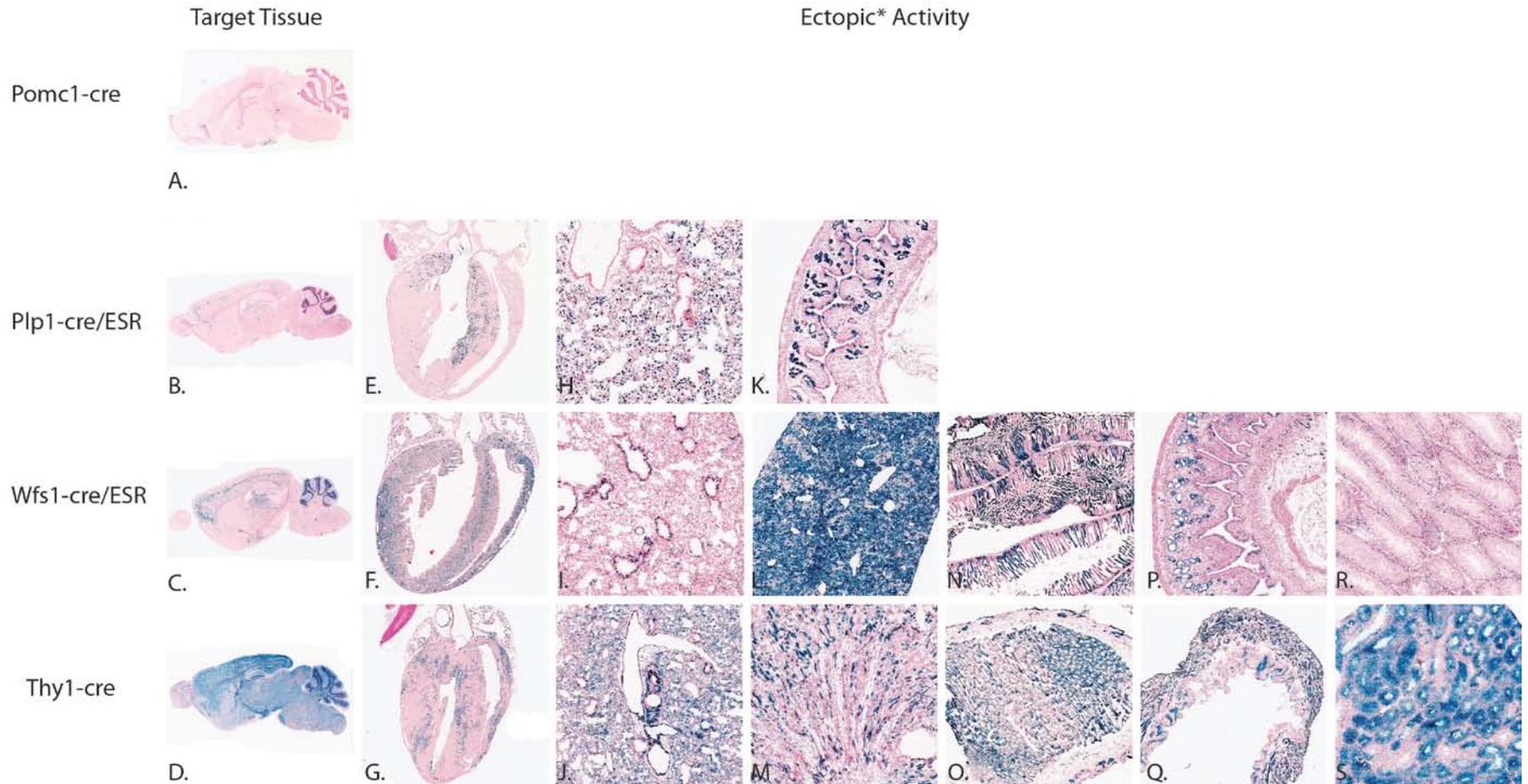


129(Cg)-Foxg1^{tm1(cre)Skh}/J Characterized with R26R LacZ Reporter

Off-target Cre activity: a surprisingly common problem



Ectopic* activity continued

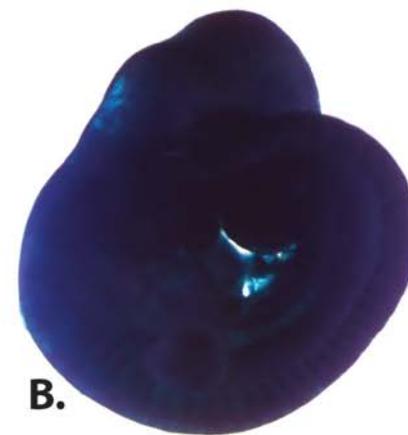
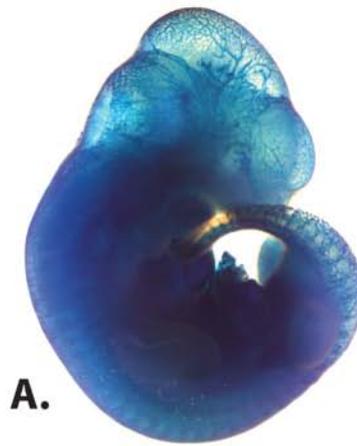


Inconsistent activity

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.

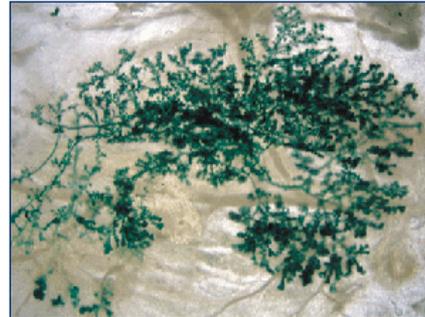


Image reproduced with permission of the Journal of Cell Science.
Rijnkels M and Rosen, JM, J Cell Sci 2001, 114(17): 3147-3153.

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

FAQs

How do I...

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

1,516 total Cre driver strains
~50% fully curated for Cre activity

Search for Cre activity by anatomical system or by driver/promoter
Get a report of all recombinase knock-ins and transgenes



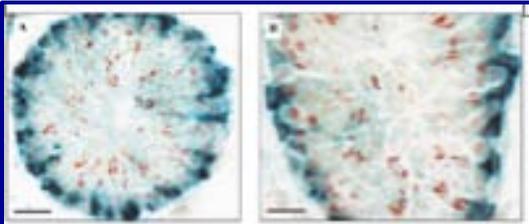
Tg(AMH-cre)1Flor - Reproductive System

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	
	Molecular description: The t show	ene, and genomic sequences from MT1 to male mice and granulosa cells of 16 day o
	Find mice (IMSR): Mous	able
	Additional Tissues: Tg(A alim	

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



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

Tissue Information	Reproductive System	Other ▶ Alp
---------------------------	----------------------------	----------------

em tissues:
m1(cre)Lowl ... (more)
e detail. [Reset Images](#)

Images

Drag images to compare to others o

J:81650 Fig. 3

Gallery of images for this cre in this tissue/organ

Recombinase Specificity

Click heading to resort table. [i](#)

Structure	Assay	Assay Note
testis	embryon 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

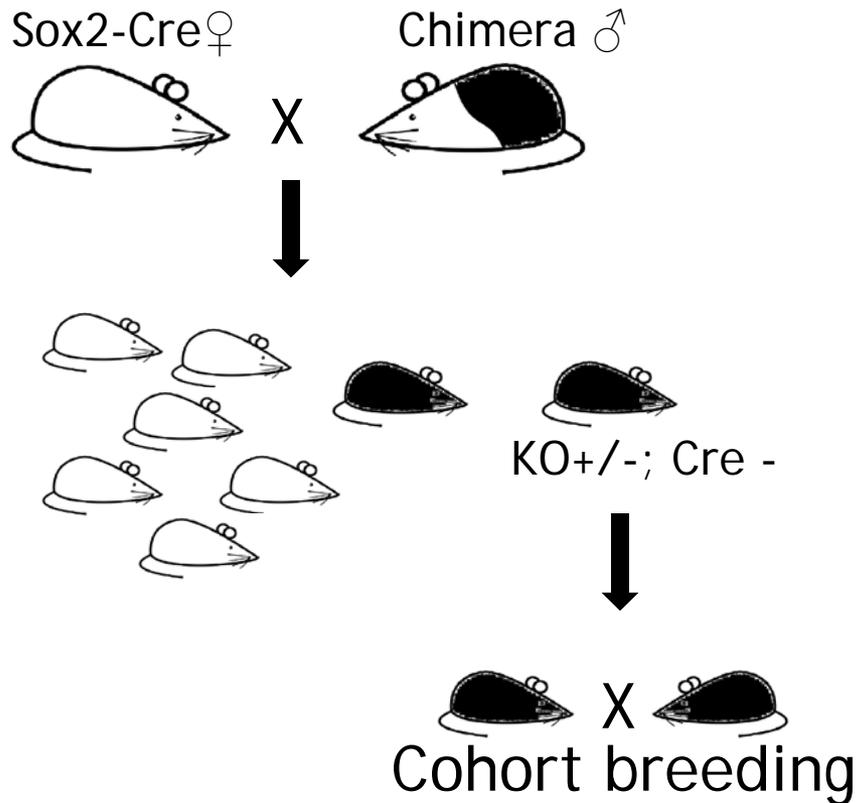
[Link to References](#)

References	All for this allele: 34 reference(s)
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Cre strains for KOMP2, IMPC

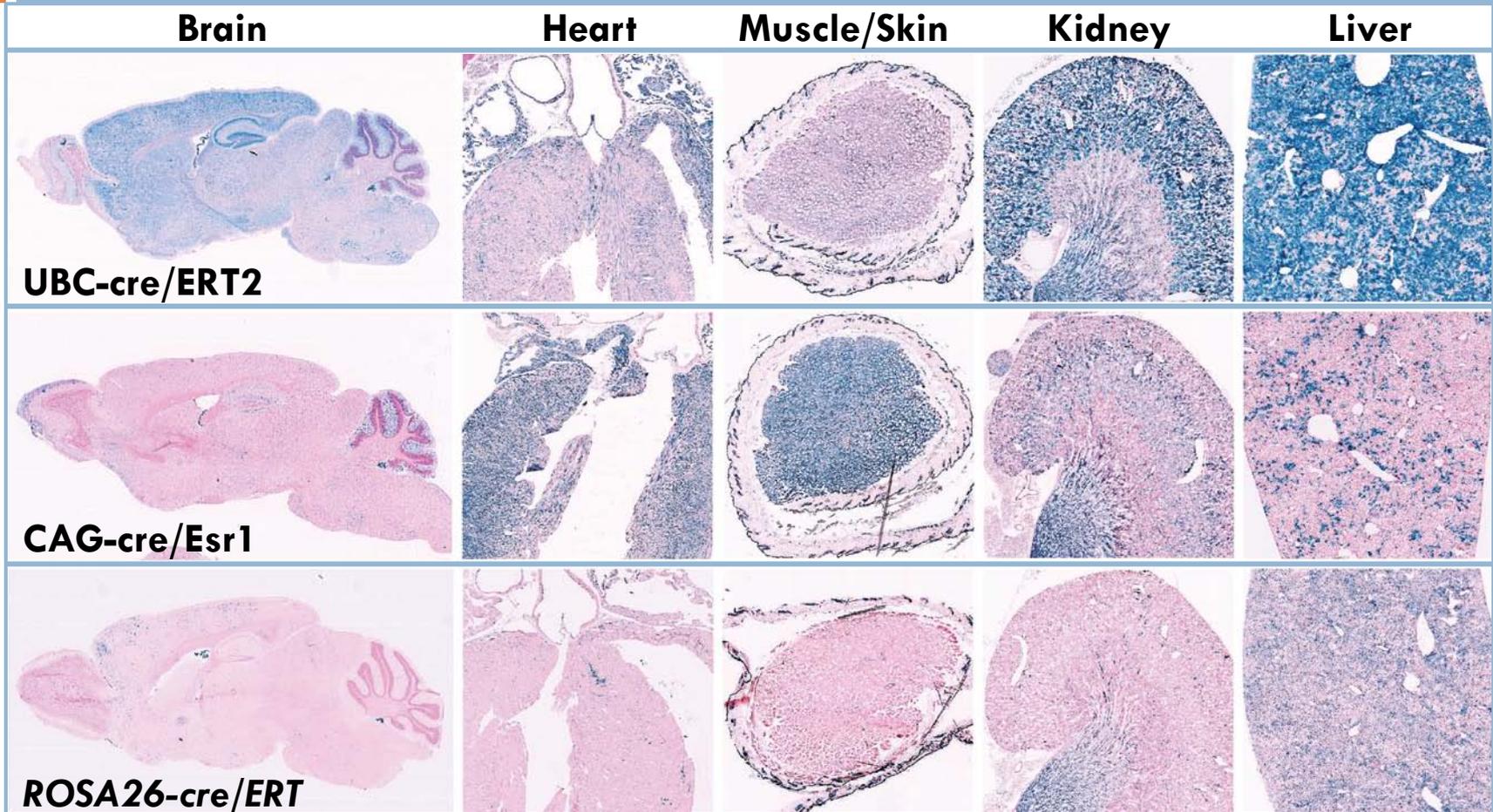
- Cre deleter
 - C57BL/6N background
 - Highly efficient
 - Maternal deletion 'feature' (Sox2-Cre; ICS Cre-GFP)
- FLPe germline deleter strains on B6N needed to convert to conditional alleles
- Inducible Cre (ERT2) and tissue specific Cre strains for use with lethals
- Need to move well-characterized,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

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Cat Lutz

Steve Rockwood

Mike Sasner

Supported by ARRA RR026117

RR032656

CrePortal

Randal Babiuk

Kim Forthofer

Peter Frost

Martin Ringwald

James Kadin

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