

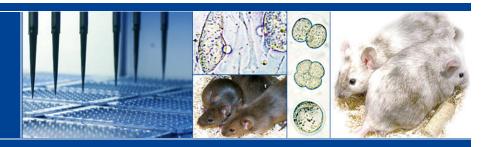
Cre-drivers EUCOMMTOOLS

European Conditional Mouse Mutagenesis Program: TOOLS for Functional Annotation of the Mouse Genome

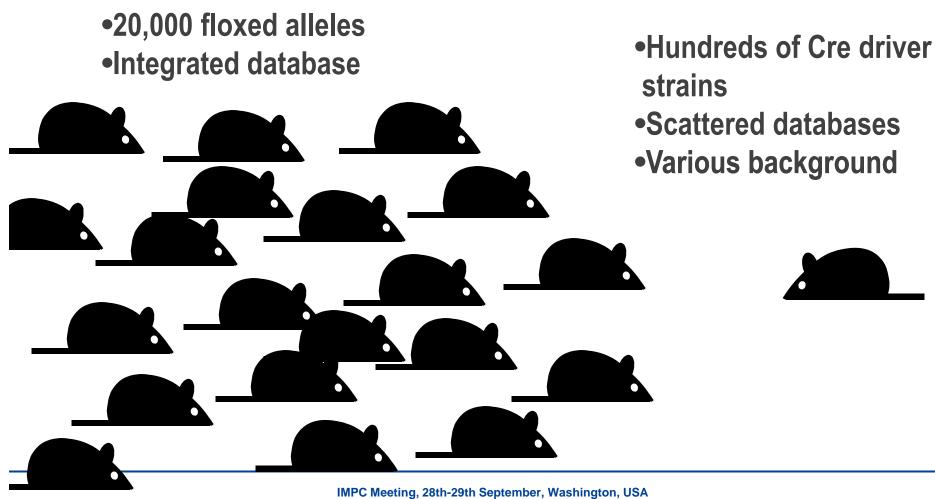
> *IMPC meeting* 28th-29th September, 2011 Washington, USA

Wolfgang Wurst Helmholtz Zentrum Munich

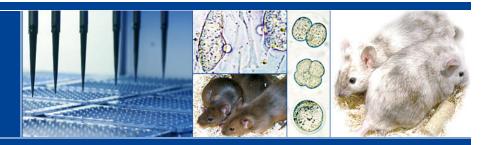




The Challenge



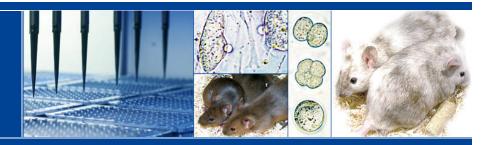


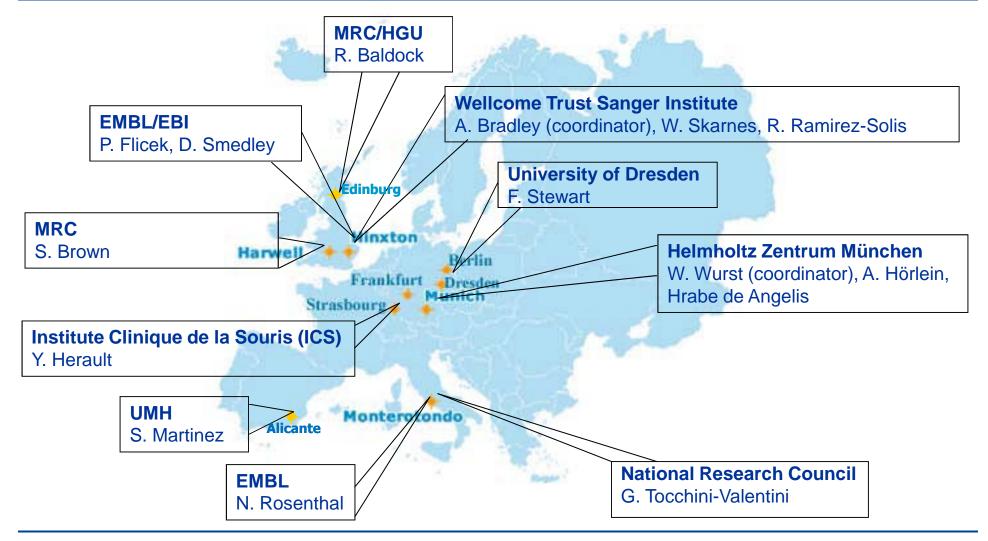


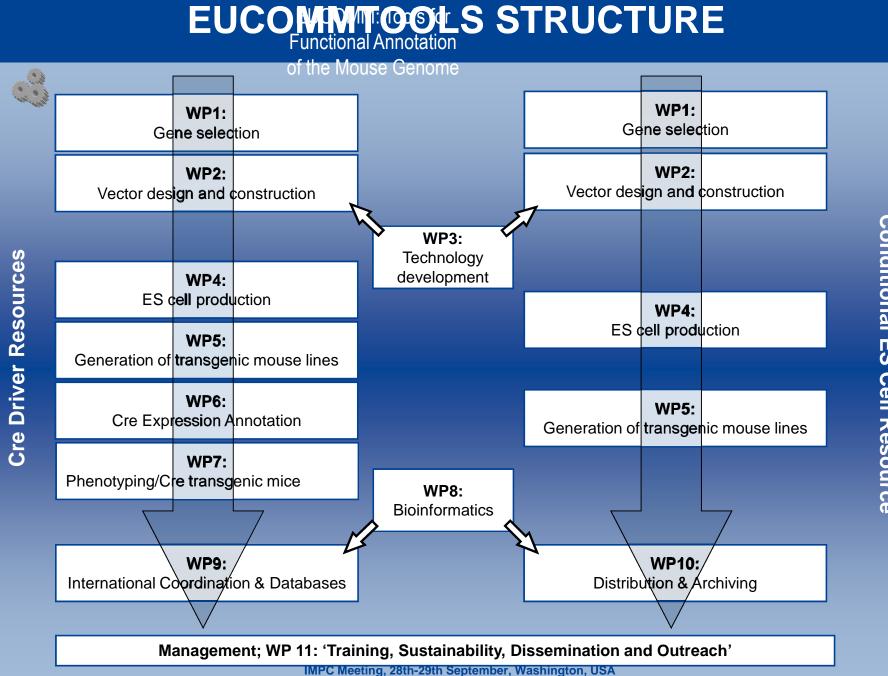
Objectives:

- » Identification of 500 genes and their promoters suitable as Cre drivers covering all organs and major cell types
- » Establishment of **500** BAC-CreERT2T2A-eGFP/BAC-Cre-T2A-eGFP (in the Rosa26 or Hprt locus) constructs and recombinant ES cells Cre driver transgenic mouse lines
- » Establishment of 250 BAC-CreERT2T2A-eGFP/BAC-Cre-T2A-eGFP driver transgenic mouse lines
- » Archiving and distribution by EuMMCR/EMMA





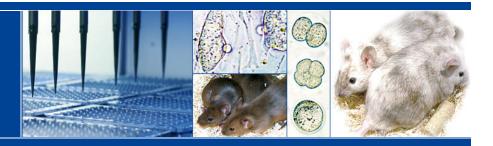




Conditional ES Cell Resource



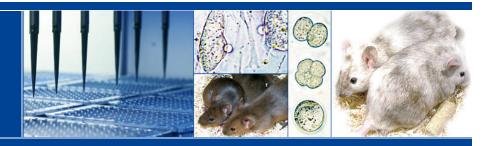
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WP1 Gene/Promoter Selection: (Damian Smedley, Nadia Rosenthal)

- » Create categorize existing Cre lines
- » Create survey collecting requests from the scientific community
- » **Coordinate** production with e.g. NorCOMM2, CanEuCre, Pleiades, NIH Blueprint, Allen Brain Institute, Riken, Jax
- » **Experts** opinions and suggestions
- » Identify specific expression Eurexpress (embryo), GXD (adult), Sanger LacZ
- » Remake most popular cre lines not on C57BL6/N avaibale
- » More than 250 genes have been selected

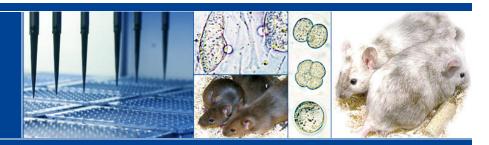




Overall summary of Cre driver selection

ANALYSIS	DRIVERS
Suggested genes from survey	17
Suggested tissues from survey	18
Specific adult expression identified from Eurexpress	64
Specific adult expression identified from GXD	94
Specific adult expression identified from BioGPS	71
Sanger LacZ expression	11
Popular Cre lines to remake as inducible B6N	41
Tissues not covered by current Cre lines	7
Expert suggestions	26
TOTAL	349



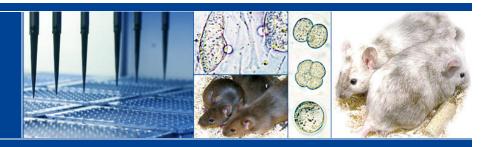


Selection status:

- » Curated to a non-redundant list of 275 genes with official MGI symbols and IDs
- » Excluded those with a known Cre line at <u>www.creline.org</u>
- » Excluded those with a Cre line in the GENSAT, CDN or ICS projects
- » Identify specific expression in normal adult samples at GXD, ArrayExpress and BioGPS (GNF microarray datasets) with possible backup from Eurexpress embryonic expression patterns



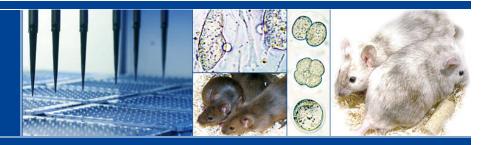
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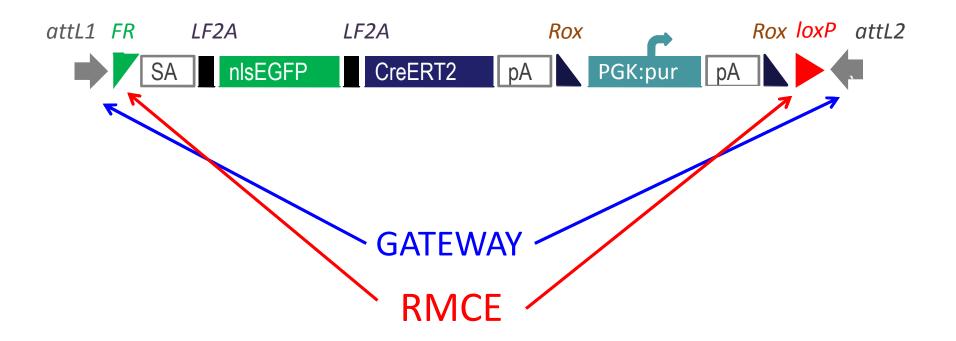
Strategy: Cre Knock-ins

- » Exploit EUCOMM/KOMP Modular Vector Resources to rapidly build a set of Cre Knock-ins
 - » Mixture of re-targeting and RMCE
- » Using Cre and inducible CreERT2
- » CreERT2 Knock-in vector design
 - » Vector includes nlsEGFP reporter
 - **»** T2A or LF2A for signal sequence containing genes
- » Comparison of Targeted Knock-ins versus BAC RMCE
 - » Coordination with Haydn Prosser from Bradley laboratory





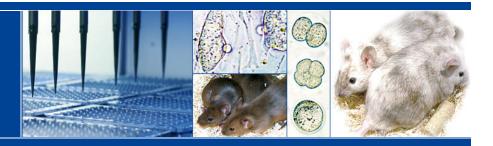
» CreERT2 Gateway/RMCE Cassette:



Barry Rosen, Bill Skarnes



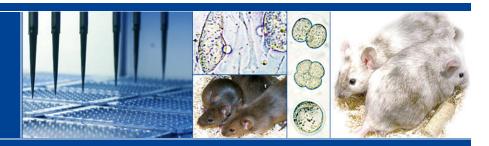
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Considerations of EUCOMMTools Cre Knock-ins

- » Simple to make with existing technology and resources
- » Pro's of Cre-Knock-ins
 - Re-utilization of previously validated rate
 - No new technology needed, only CreERT2 vector cassette
 - Knocked-in Cre should faithfully recapitulate endogenous expression
- » Con's of Cre Knock-ins
 - Difficult to breed animals to homozygosity in most cases
 - Potential for haplo-insufficient phenotypes





Alternative Cre Alleles

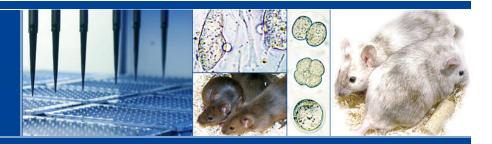


<u>3'UTR IRES Cre Insertion</u> Leaves endogenous protein intact

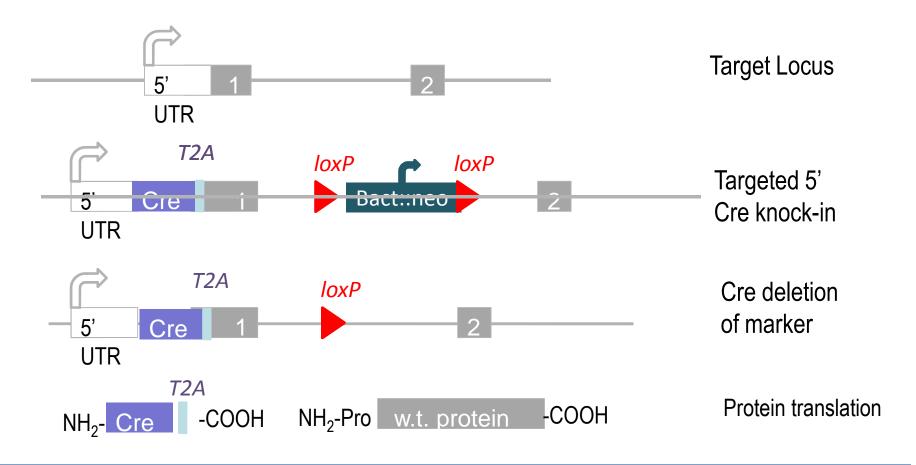


<u>2A Mediated Cre 3'</u> <u>Insertion</u> N-terminal fusion, adds COOH extension to endogenous protein

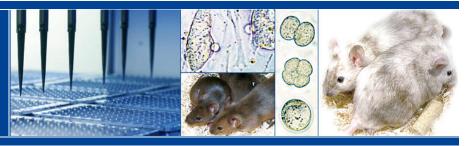




Alternative Cre alleles: 5' Cre 2A-mediated insertion







Pros and Cons of Different Alternative Cre Alleles

Pro

Con

Universal LF2ACre Knockin

<u>3'UTR IRES Cre Insertion</u>

-Applicable to all genes -Still deletes an allele -Exploits existing IKMC resources

-Applicable to all genes -Endogenous protein intact -Validated in literature -Reduced Cre from IRES

-Adds 3' aa's to endogenous protein -Additional aa's with LF2A -new vector, no RMCE

5' Cre 2A-mediated insertion -Minimal perturbation of locus -Promoter specific applications

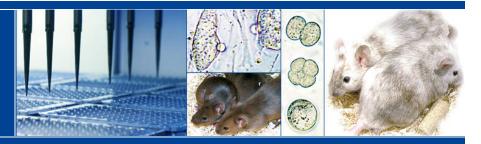
-HT vector construction

2A Mediated Cre 3' InsertionShould express high levels

of Cre

-Initiating ATG must be defined carefully -new vector, no RMCE

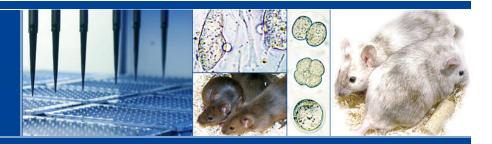




CreERT2 Knock-ins: Progress Re-Targeting

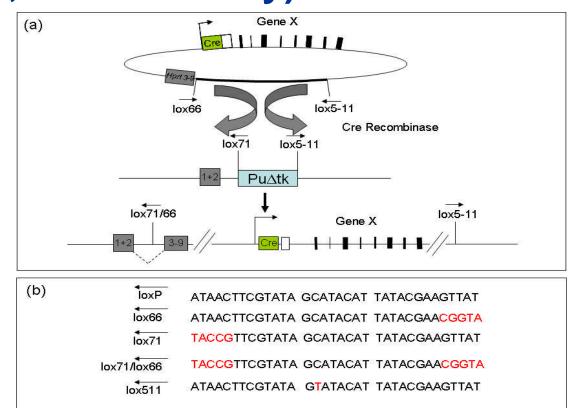
- » 49 Intermediate Vectors Identified
- » Criteria: Targeted ES Cells Status in EUCOMM/KOMP
- » Avoid signal sequence/secreted for moment(50% of genes)





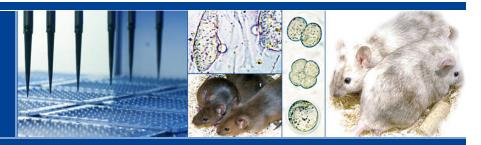
EUCOMMTOOLS – BAC knock-in Technology (Hayden Prosser, Allan Bradley)

RMCE strategy for introducing BAC eGFPT2ACreERT2/Cre-T2A-eGFP transgenes into the ES cell genome





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Cre-BAC transgenesis by site specific integration

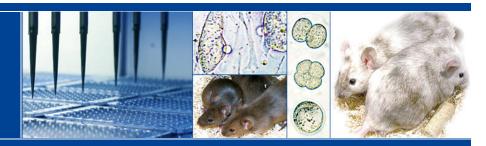
Pros:

- » Can use Human or Mouse BACs
- » Avoids disruption of endogenous alleles as would occur in targeted Cre knock-in to "Driver" gene
- » Avoids variation that might occur with random integration
- » Cassette exchange removes extraneous prokaryotic vector DNA

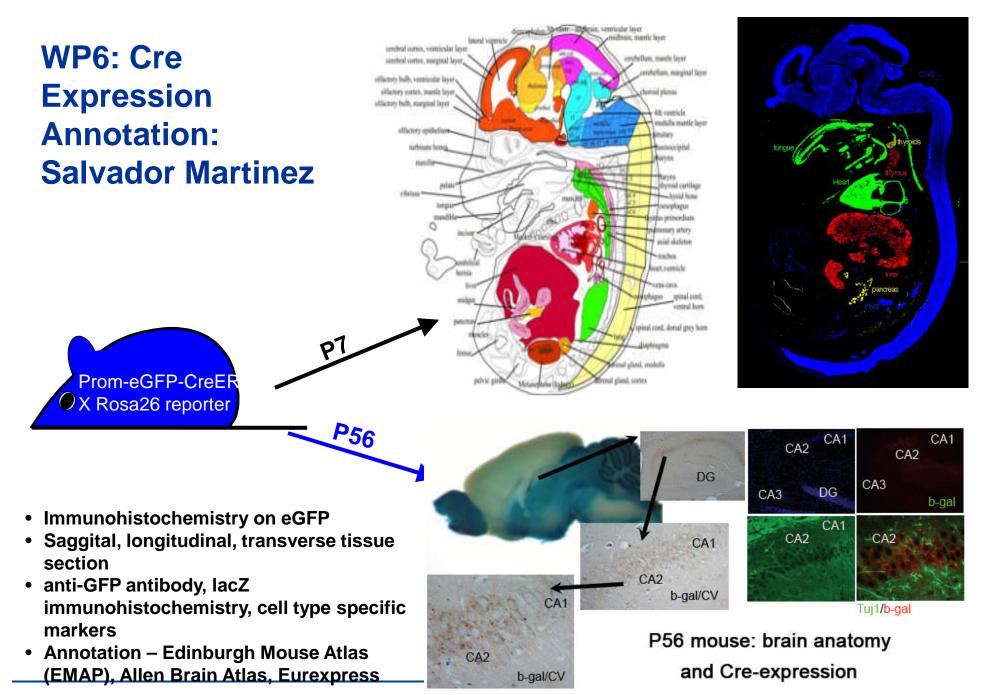
Cons:

- Integrated BACs do exhibit deletions in a proportion of cases
- » "Passenger" genes may be associated with Cre driver transgene
- » Gene size limits

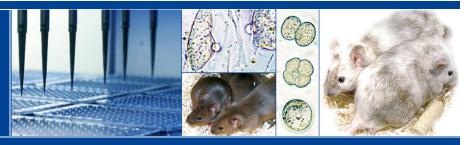




- » Currently:
- » Direct comparison of Knock-in versus Bac-knock-in strategy!

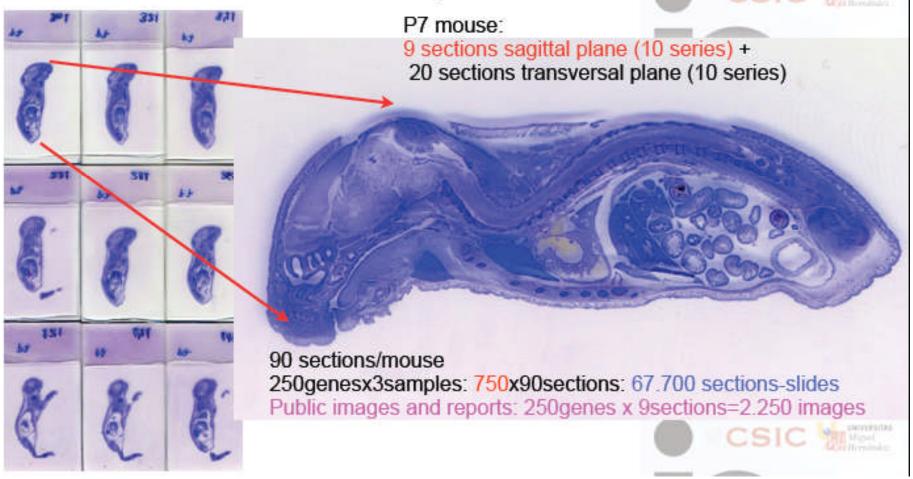




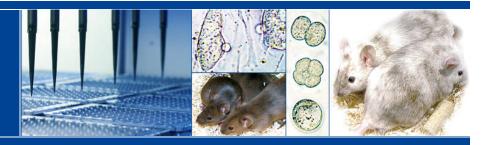


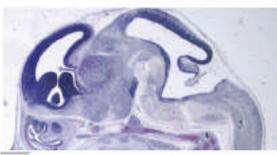
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WP6: Cre Expression Annotation



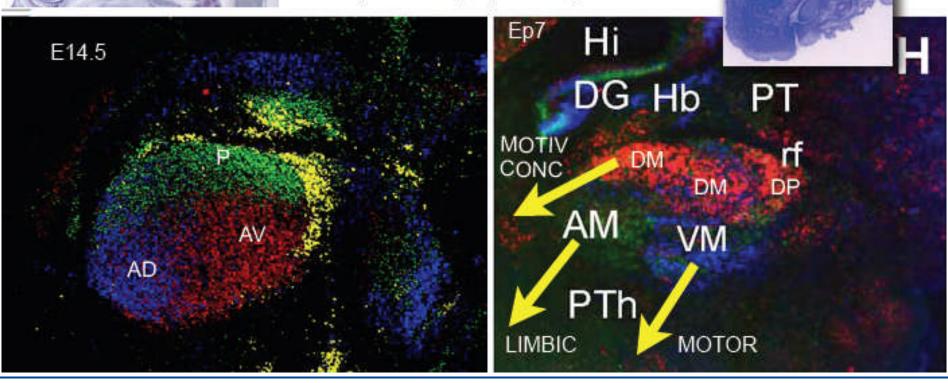




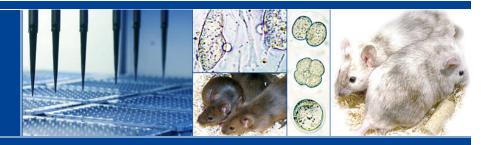


DESCIBE GENE EPRESSION IN RELATION TO ANATOMY (CELLS IN THE SPACE)

Cells (neurons+glia) in the space: BRAIN

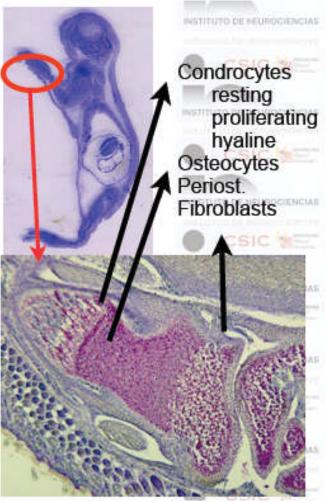




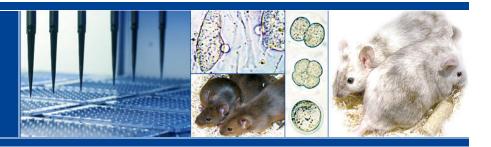


P7 mouse: 15 sections sagittal plane NEW BODY ORGANS ONTOLOGY

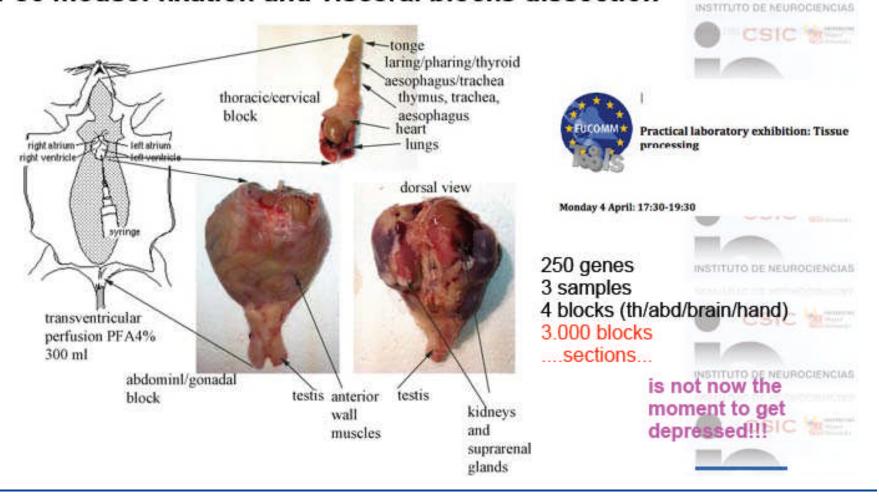
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WE NEED AN ON	1000 C		left ventricle	
VVLIL			right ventricle	



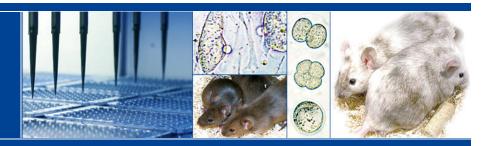




P56 mouse: fixation and visceral blocks dissection

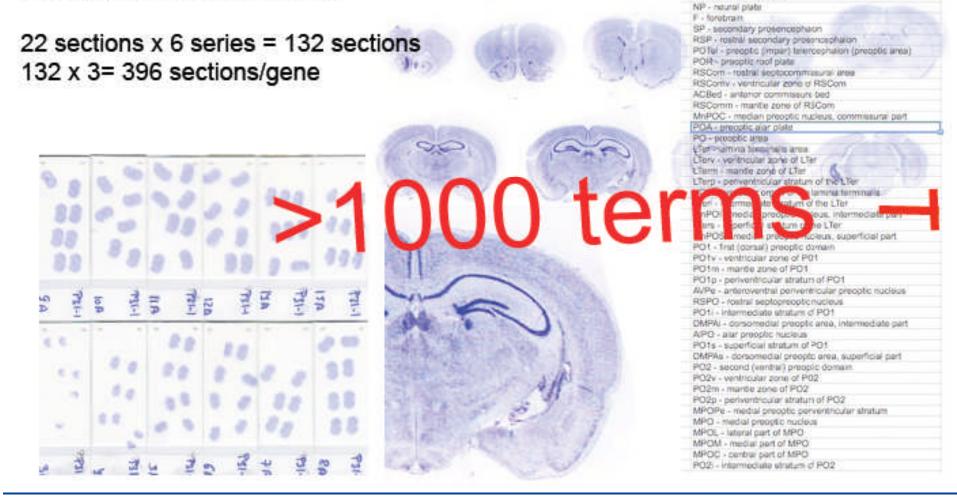




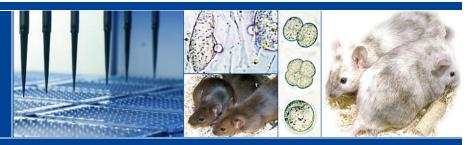


mouse - Mus musculus

BRAINS, transversal sections

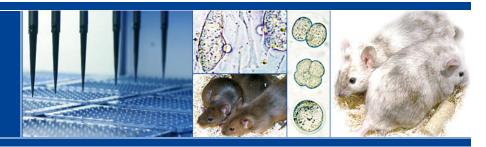


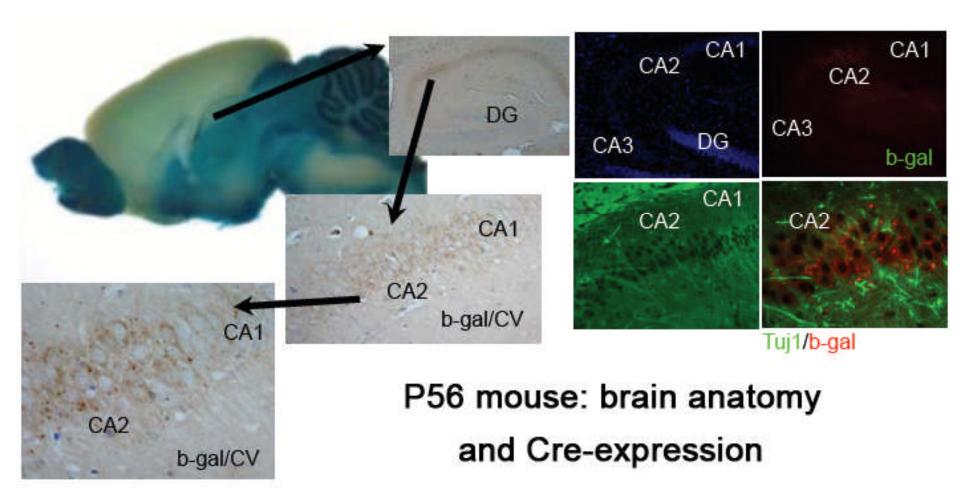




THORACIC/CERVICAL REGION	522 sections/ gene
ABDOMINAL REGION	3600 sections/gene
BRAINS, transversal sections	396 sections/gene
BRAINS, sagittal sections	255 sections/gene
TOTAL	4783 sections/gene WORK TO DO
In practical terms: 29 sections THX 200 sections ABD 22 (Brain T) 10 (Brain S) 261 SECTIONS/GENE WORK TO SHO 200 sections THX 200 sections ABD 21 (Brain T) 22 (Brain T) 22 (Brain T) 22 (Brain T) 20 (Brain S) 20 (





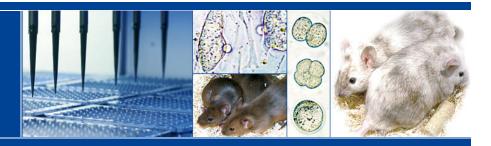












Distribution of EUCOMMTOOLS Material

Dissemination of • mutant ES cell lines http//:www.eummcr.org

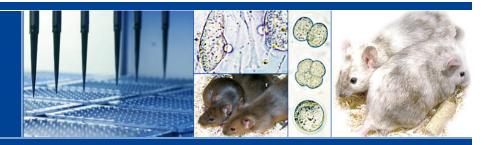


 mouse lines http//:www.emma.org



Dissemination asap as QC confirmed





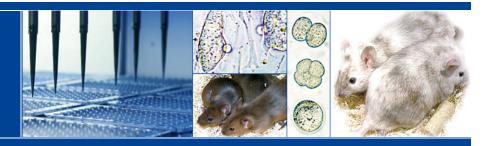
EUCOMMTOOLS integration in IKMC portal (www.knockoutmouse.org)



Long term **SUSTAINABILITY**



EUCOMM: Tools for Functional Annotation of the Mouse Genome



- » Outlook:
- » Technology and reagent development
 - » IKMC toolkit extension
 - » New ligand/SSR-LBD pairs
 - » New conditional switches
 - » Additional optogenetic tools, markers, cell survival



