



**Lydia Teboul Head of Transgenics**The Mary Lyon Centre







#### **Targeting the genome**

- Genome sequence + Gene targeting technology
- Publicly available allele collections
- Quality control of the EUCOMM library

#### Large scale phenotyping

- EUMODIC: Large scale phenotyping proof of principle
- IMPC

#### Aging screen

- ENU mutagenesis
- Screens







#### **Targeting the genome**

- Genome sequence + Gene targeting technology
- Publicly available allele collections
- Quality control of the EUCOMM library

#### Large scale phenotyping

- EUMODIC: Large scale phenotyping proof of principle
- IMPC

#### Aging screen

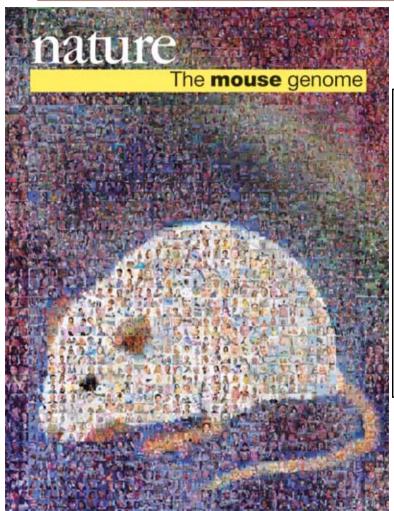
- ENU mutagenesis
- Screens



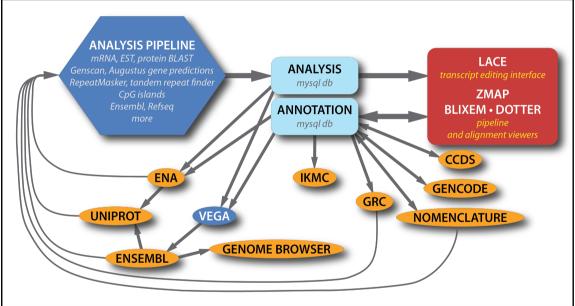


# The mouse genome





Sequence data



VEGA annotation by HAVANA group (WTSI) Manual functional annotation of the genome: where are the protein coding genes?



# **2007 Nobel prize in Physiology** and Medicine

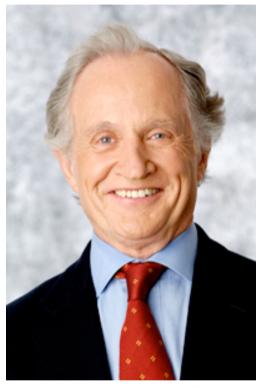




**Oliver Smithies** 



Martin J. Evans



Mario R. Capecchi

"principles for introducing specific gene modifications in mice by the use of embryonic stem cells"







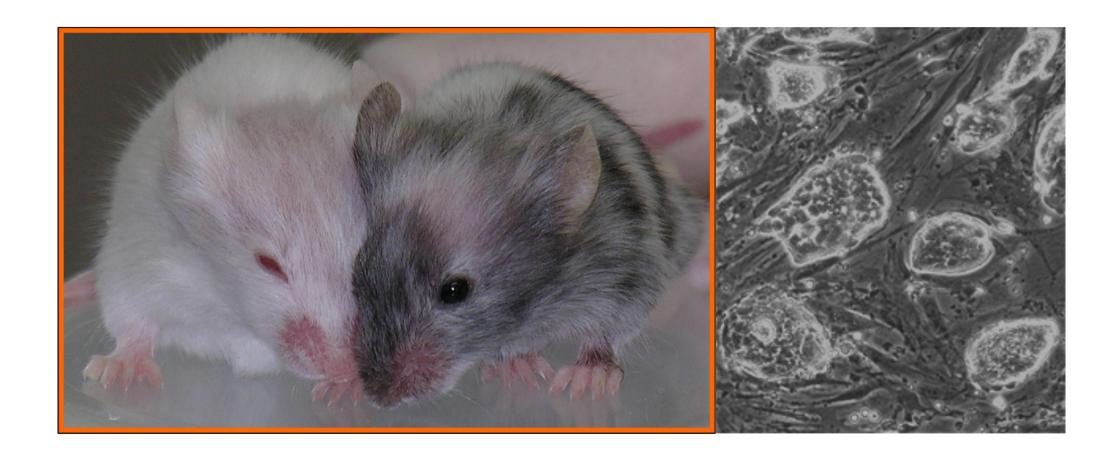














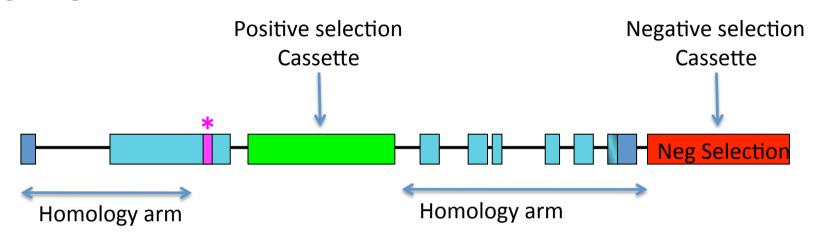


- Homologous recombination in ES cells
- Blastocyst injection
- Germ line transmission





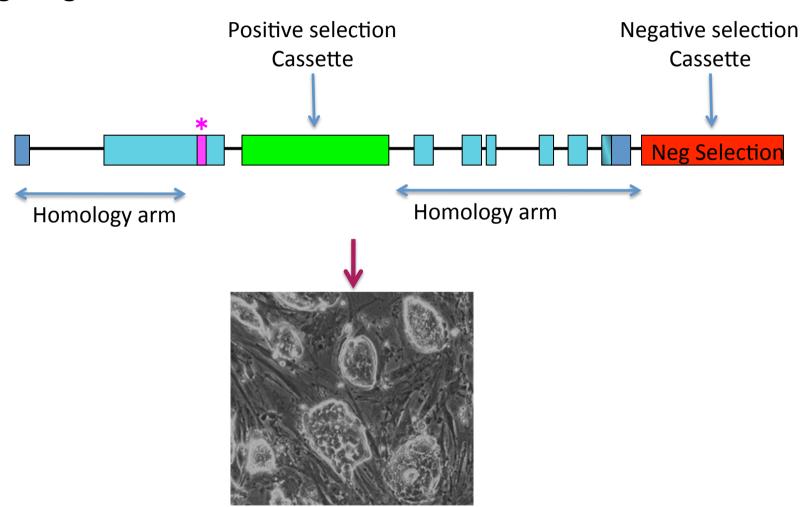
## Targeting construct





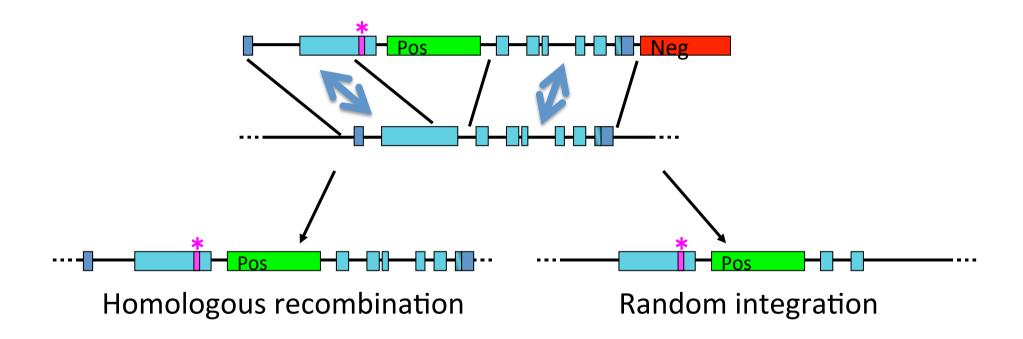


## Targeting construct









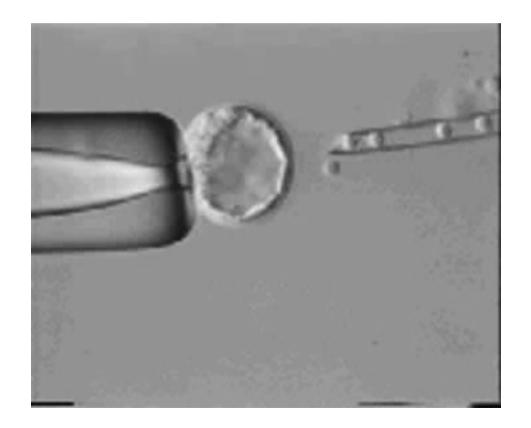
ES cells screen: finding homologous integration event

- Southern blotting
- Long Range PCR
- Loss of allele copy counting





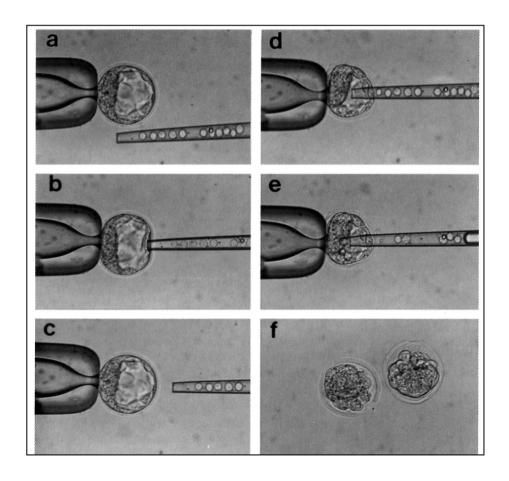
# Blastocyst injection







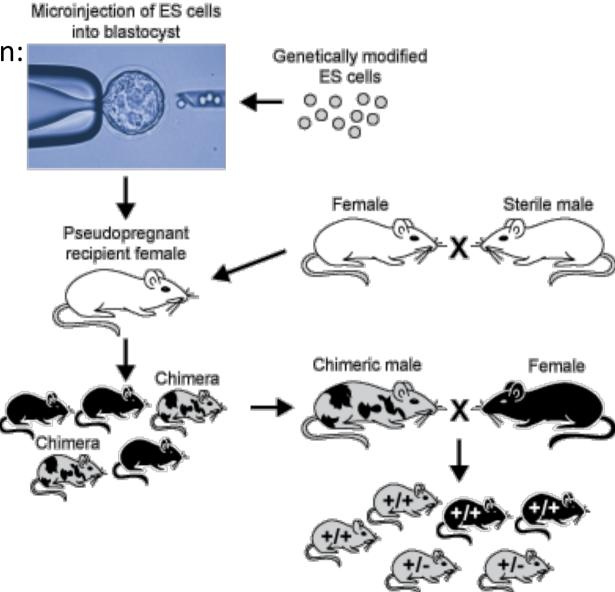
# Blastocyst injection







Germ Line Transmission:





# Targeting the genome











**Bill Skarnes** 

Francis Stewart

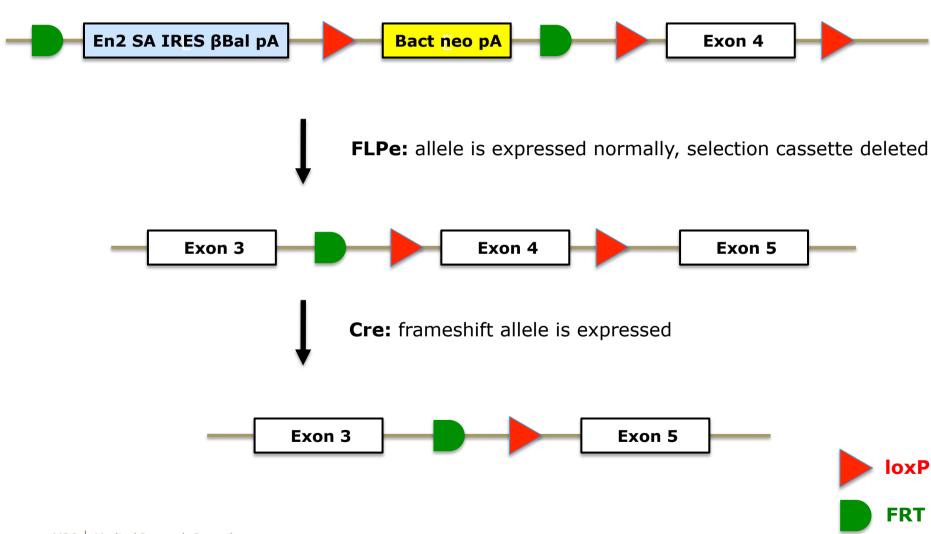
**Wolfgang Wurst** 

-> EUCOMM and KOMP programmes



# **EUCOMM** and **KOMP** Gene targeting: KO first







## **Gene targeting collections: IKMC**



IKMC: International Knockout Consortium: Libraries of targeted ES cells

Targeted deletions: Velocigene

• Conditional targeting: EUCOMM, KOMP, NorCOMM

Number of protein coding genes with mutant ES cell lines in the IKMC resource:

<b>Total Genes</b>	KOMP CSD	KOMP Regeneron	EUCOMM	NorCOMM
Vectors available	6590	4733	8837	839
ES cells available	5252	3959	7165	569
Mutant mice available	450	382	687	4

Figures on 2<sup>nd</sup> May 2012



## **Quality control of the EUCOMM library**

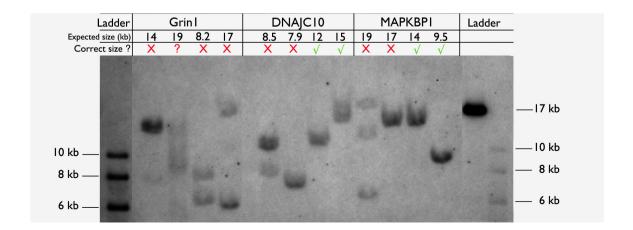


#### ES to mouse conversion:

- 568 conditional mouse lines (536 genes) produced for distribution through EMMA
- Germ Line Transmission rate (generally 1 injection session/clone): about 50%

#### Allele integrity:

- Early data suggested 1/5 clone was incorrect
- Clone screening strategy revised, extra check are now performed before cell distribution
- Advise to users/mouse producers: Check clones by Southern blot before injection





# Distribution of targeting vectors, ES cells and mouse mutants



# 2 gene targeting constructs and ES cell distribution centre:

**EuMMCR:** in Munich, distributes European collections

MMRRC: in Davies, CA, distributes US collections



### **European Mouse Mutant Archive:**

Node In Harwell

- Archiving of GA mice lines
- ~1500 mouse lines archived
- Rederivation
- Assisted reproduction







#### Targeting the genome

- Gene targeting technology + Genome sequence
- Publicly available allele collections
- Quality control of the EUCOMM library

#### Large scale phenotyping

- EUMODIC: Large scale phenotyping proof of principle
- IMPC

#### Aging screen

- ENU mutagenesis
- Screens

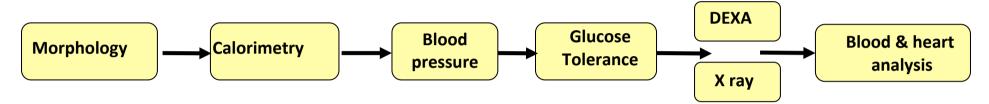




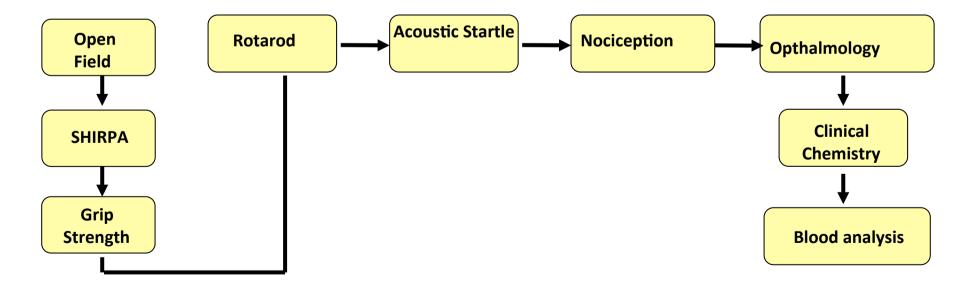
# **European Mouse Disease Clinic Multi-system phenotyping**



#### **Pipeline 1- Metabolic Profiling**



#### **Pipeline 2- Neurosensory Profiling**

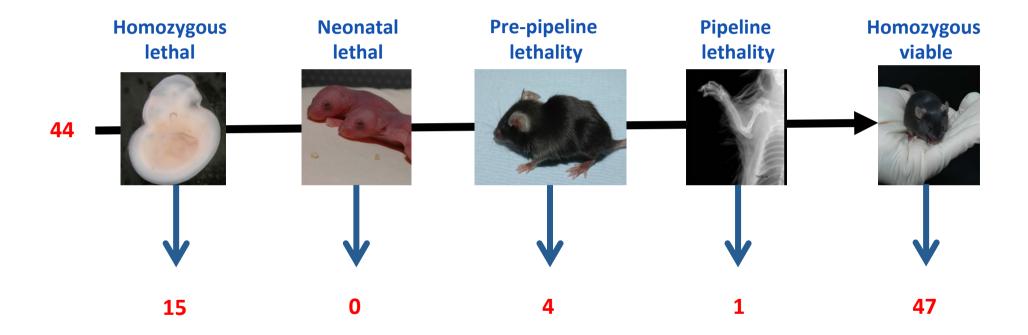






# **Viability**







# **Homozygous Lethal**



#### Homozygous lethal



**15** 

Secisbp2 Selenocysteine insertion sequence-binding protein 2 (dies before dpc9.5)



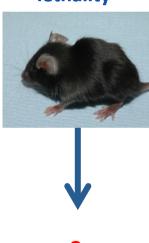
Lack of neural tube closure and turning defect



# **Pre-pipeline Lethality**



**Pre-pipeline** lethality



Bag3

Is involved in anti-stress and antiapoptotic pathways.

**Back limb paralysis** (d22-30)

Npc1 Niemann Pick type C1

Tremors, uncoordinated gait (d19-28)



## **Pipeline Lethality**



**Pipeline** lethality



#### Cisd2

Previously implicated in human longevity, mouse knockouts of this gene show premature-ageing phenotypes.

Piloerect coat Hunched **Rapid Breathing Tremors Abnormal gait** 

**Onset of disease** 

(average 132 days, earliest 96 days)

**Disease Progression** (2-32 days after onset)

No histological abnormalities



### **Phenotypes**





#### The message is:

That there are known knowns,

There are things we know that we know,

There are known unknowns,

That is to say there are things that we now know, we don't know

But there are also unknown unknowns,

There are things we do not know we don't know

And each year we discover a few more

Of those unknown unknowns.

-Feb. 12, 2002, Department of Defense news briefing

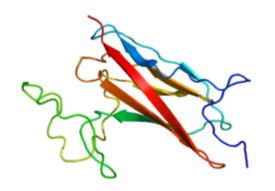
- **Known Knowns-** Confirming phenotypes in existing mutants
- **Unknown Knowns-** New phenotypes in existing mutants
- **Known Unknowns-** Confirming phenotypes seen in other mutations of the same gene
- **Unknown Unknowns-** New phenotypes in new mutants



### **EUCOMM Lines**



Mary Lyon Centre



#### MYBPC3

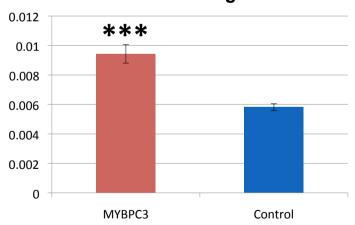
Myosin binding protein C.

Hypertrophic cardiomyopathy



## **Heart weight** 0.25 0.20 0.15 0.10 0.05 0.00 MYBPC3 Control

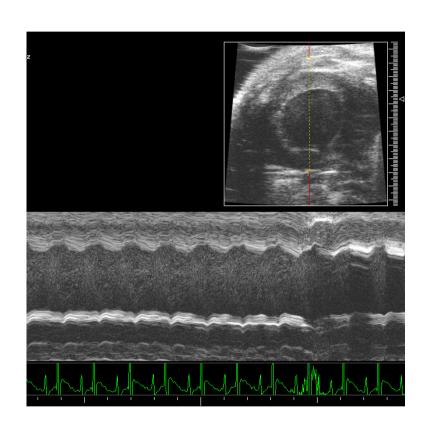
#### **Heart weight normalised with** Tibia length



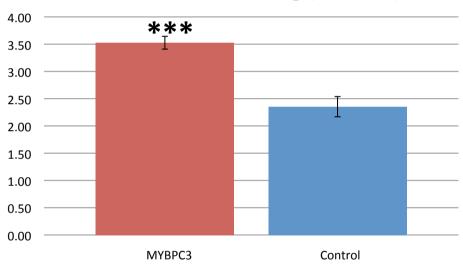


### **MYBPC3**





### Left Vent. Ant. Wall Avg (diastole)

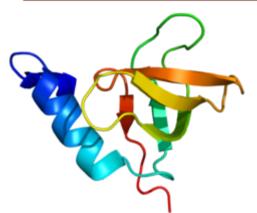


Left Ventricle Hypertrophy

### Echocardiogramme

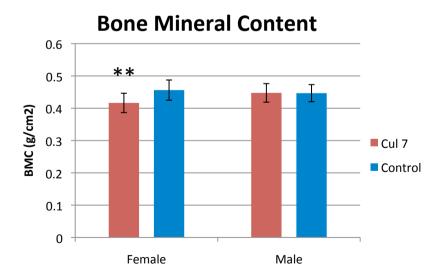
## Cul-7

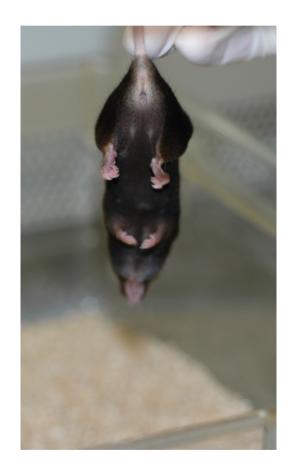




#### Cul-7 Component of an E3 ubiquitinprotein ligase complex.

Defects have been associated with 3M syndrome type 1.



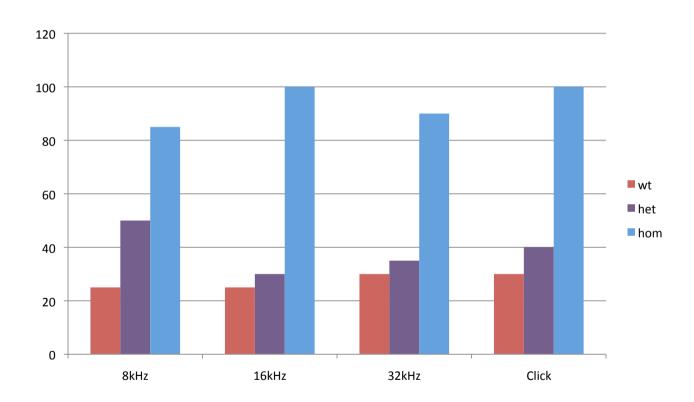




# Elmod-1



### ABR Thresholds (note n=1)

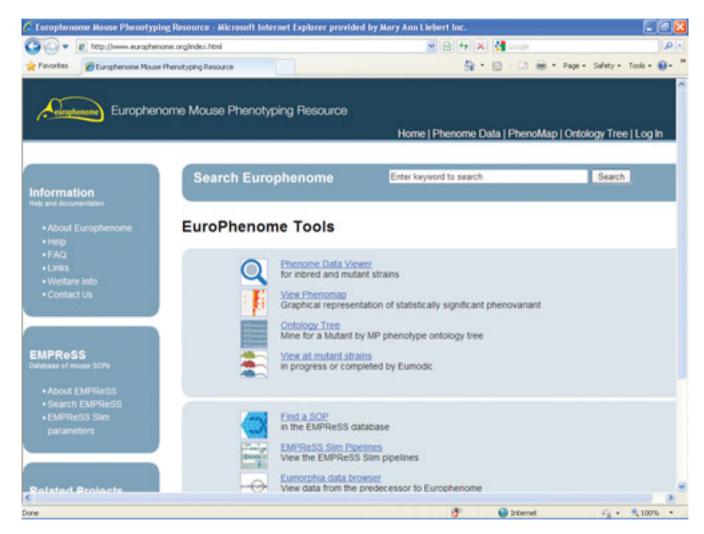




### **Data presented in EuroPhenome**



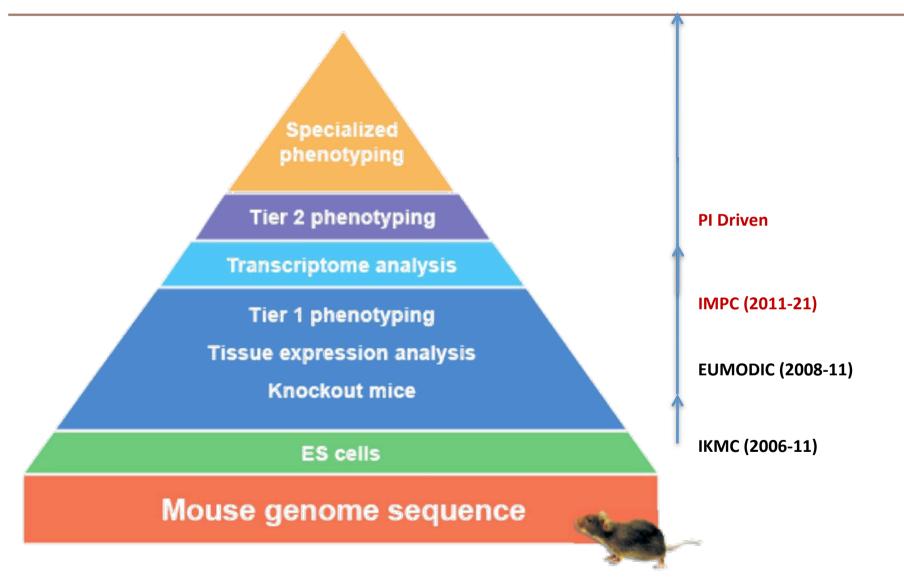






# IMPC: International Mouse Phenotyping Consortium







# Vision For Next 10 Years... MRC Harwell

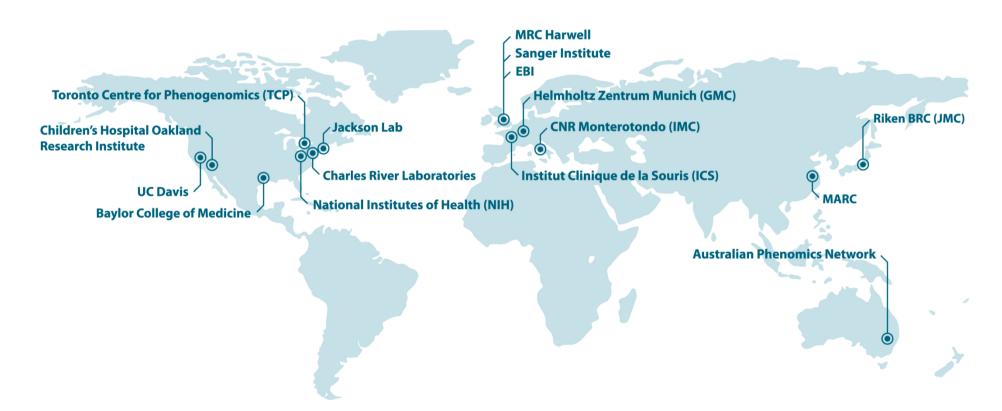


- An Encyclopaedia of Mammalian Gene Function
  - Create KO mice for ALL GENES
  - Database with associated primary phenotype info
  - Discover unforeseen gene function
  - Free access to MICE
  - Free thousands of researchers from tool generation;
  - A rich seam for future hypothesis driven research, with the potential for breakthrough discoveries
  - A transformative project that will underpin the future of biomedical science and the biology of disease systems.









22 Partners, 13 Production Centres, 9 Countries

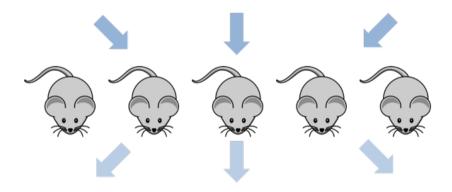
# www.mousephenotype.org



## **Harwell part in IMPC**



ES cells (EUCOMM, KOMP) => Mice, Archive Embryos, Breed Cohorts



Lethality & Fecundity Test, LacZ Expression Profile (embryo/adult), Extended EUMODIC

- 100 Lines per year (funding secured)
- Node for mouse distribution



# Genetically engineered animals and their use in understanding disease



#### Targeting the genome

- Gene targeting technology + Genome sequence
- Publicly available allele collections
- Quality control of the EUCOMM library

## Large scale phenotyping

- EUMODIC: Large scale phenotyping proof of principle
- IMPC

#### Aging screen

- ENU mutagenesis
- Screens





## **Chemical mutagenesis**



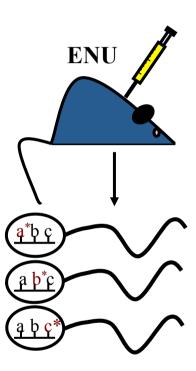
## Large scale ENU mutagenesis programmes:

## Gene driven screens: ENU DNA/Sperm archive

>10000 DNA and sperm samples, allowing the identification of point mutations for all coding and non-coding sequences in mouse genome

#### Phenotype driven screens

generating mouse models screening for a phenotype of interest





## **Building on Success**



Recessive Screens	Pedigrees screened	Mice screened	Pedigrees containing Phenodeviants	Confirmed phenotypes
ADULT				
Dysmorphology	223	7520	39	25
Imprinting/Fertility	26	648	7	7
Circadian Rhythm	67	1296	24	12
Memory/Learning	45	967	13	3
Deafness	154	4485	26	8
Bone and Mineral Disorders	35	1163	4	4
Innate Immunity	146	954	23	4
Vision	137	3965	12	7
Brain Histology	19	331	1	1
Totals	395	9122	150 (39% of pedigrees)	72
DEVELOPMENT				
Development	187	1287	79	8
L-R development	135	615	39	13
Cardiac Development	76	2626	44	22
Totals	398	2626	162 (41% of pedigrees)	43



## **Phenotype Driven Screens**



## Makes no assumptions about genetic basis of disease

- Potential to identify novel genes or pathways involved in disease
- Bias screens to search models of human disease
- Employ challenges or genetic modifications for a sensitised screen
- Screens need to be large (1000s of mice)

#### **Point mutations**

- Easier to map than QTLs
- Hyper-, hypo- and neomorphic mutations
- Generate novel models of human disease



## **Late Onset/Age-Related Disease**



## Concept

- Let the mice grow old: look at apparition of phenotype
- Would need to be a consortium of researchers
- Makes use of existing expertise and technologies
- Harwell well positioned to carry out such a screen

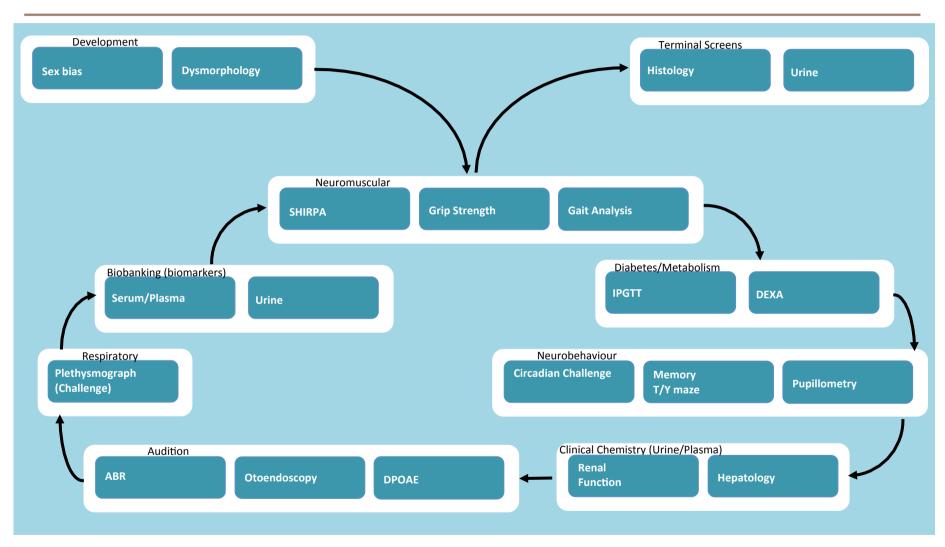
## Importance

- Ageing society Increasing pressure on society from age related diseases
- Some models are acute/early onset which does not truly reflect human disease
- Focus of funding bodies



## **Screening Pipeline**

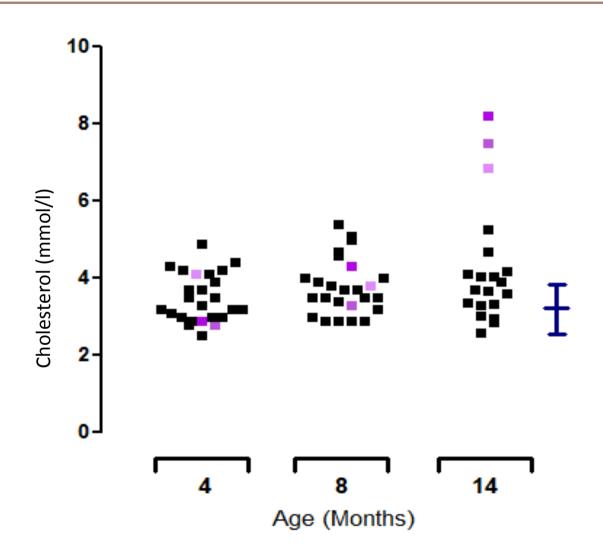






## Ageing Mutants MP-86 Hypercholesterolaemia

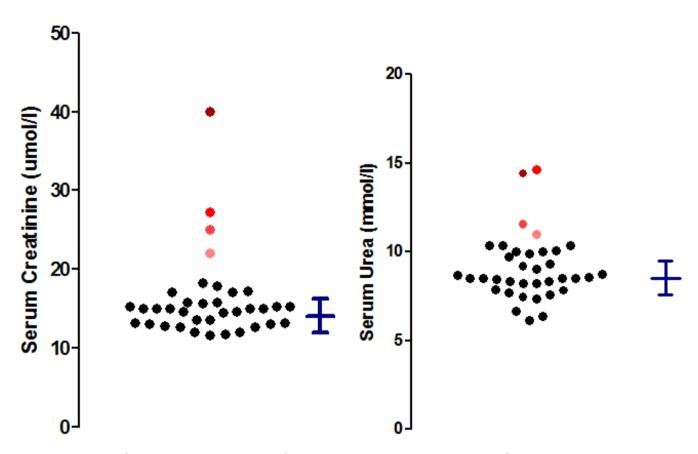






# **Ageing Mutants MPC-46 Reduced Renal Function?**





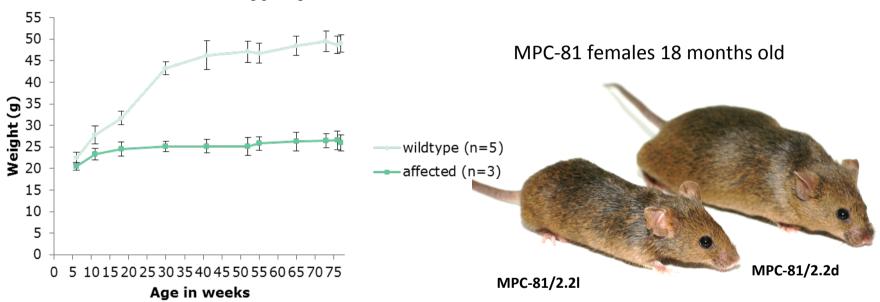
Increased serum urea and creatinine at 12 months



## MPC-81 Lean mice



## MPC-81A weights (affected and wildtypes)



Currently undergoing detailed metabolic phenotyping – activity, metabolism, food intake...



## **Mapping**



## **Conventional Mapping**

- Large pedigrees/multiple rounds of screening
  - Confidence in phenotype
  - Enough affected individuals for mapping
  - NGS to identify associated mutation



## **Acknowledgements**



#### Mary Lyon Centre, MRC Harwell

• Director: T Weaver

• FESA: M Fray

• Phenotyping: S Wells

## Mammalian Genetics Unit, MRC Harwell

• Director: S Brown

Aging programme: P Potter







## consortiaCNR (Monterotondo)

Helmhotz Zentrum (Munich)

**EUCOMM (WP7) & EUMODIC** 

- ICS (Strasbourg)
- WTSI (Hinxton)



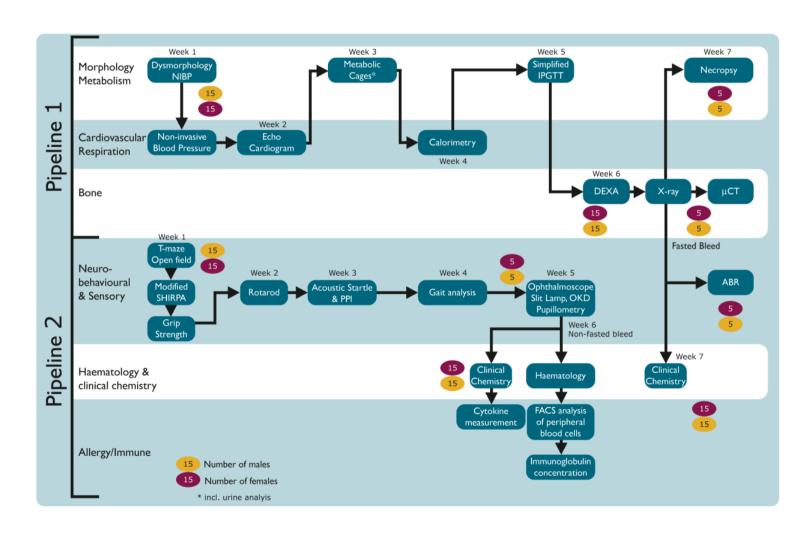






## **EUMODIC Phenotyping**

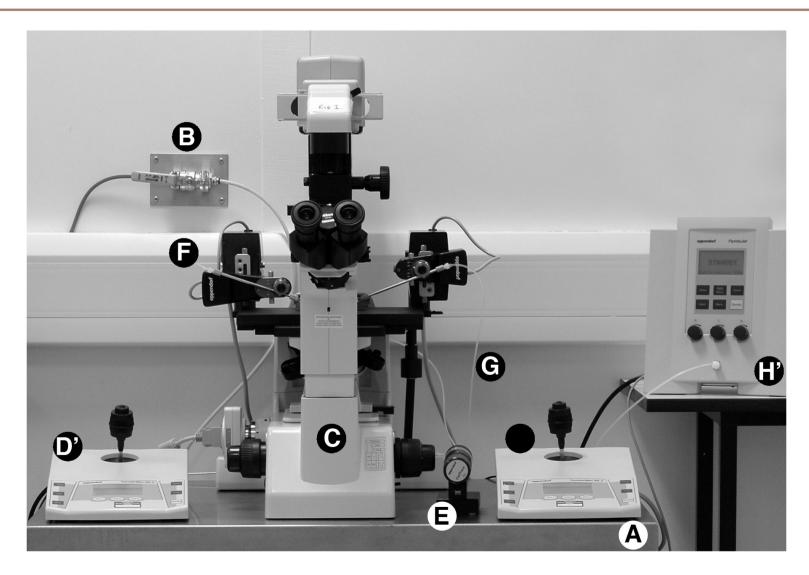






## **Principles of gene targeting**

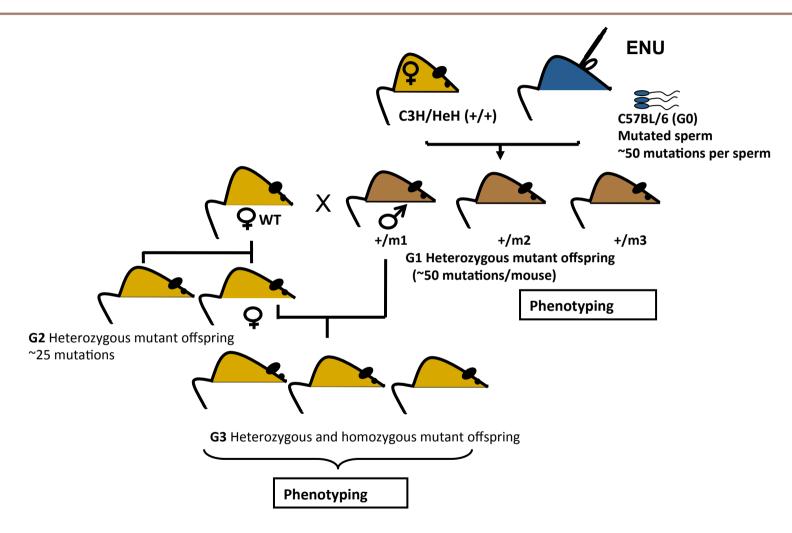






## Dominant and Recessive phenotype driven screens







## Mapping



- Conventional Mapping
  - Large pedigrees/multiple rounds of screening
    - Confidence in phenotype
    - Enough affected individuals for mapping
    - NGS to identify associated mutation
- Early (mild) disease/biomarkers
  - Early indicators of late onset disease
  - Map on mild disease/biomarker