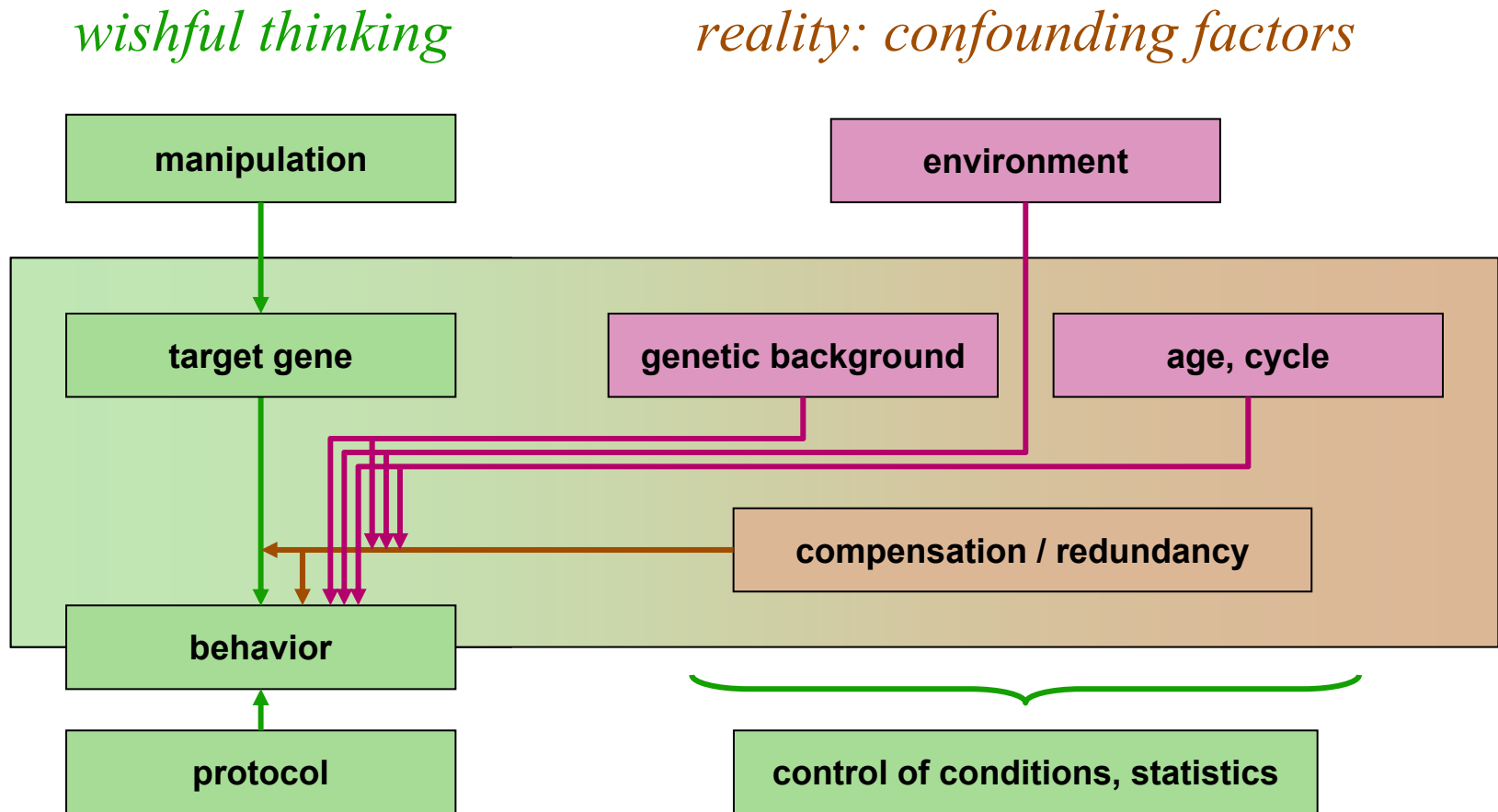

6th TT Meeting 2005, Barcelona, Spain, 11-13 Sept 2005

Choice of Strains, Strain Genetic-Differences, Modifiers



David P. Wolfer M.D.
Institute of Anatomy, University of Zurich
dpwolfer@anatom.unizh.ch

Gene targeting and behavior

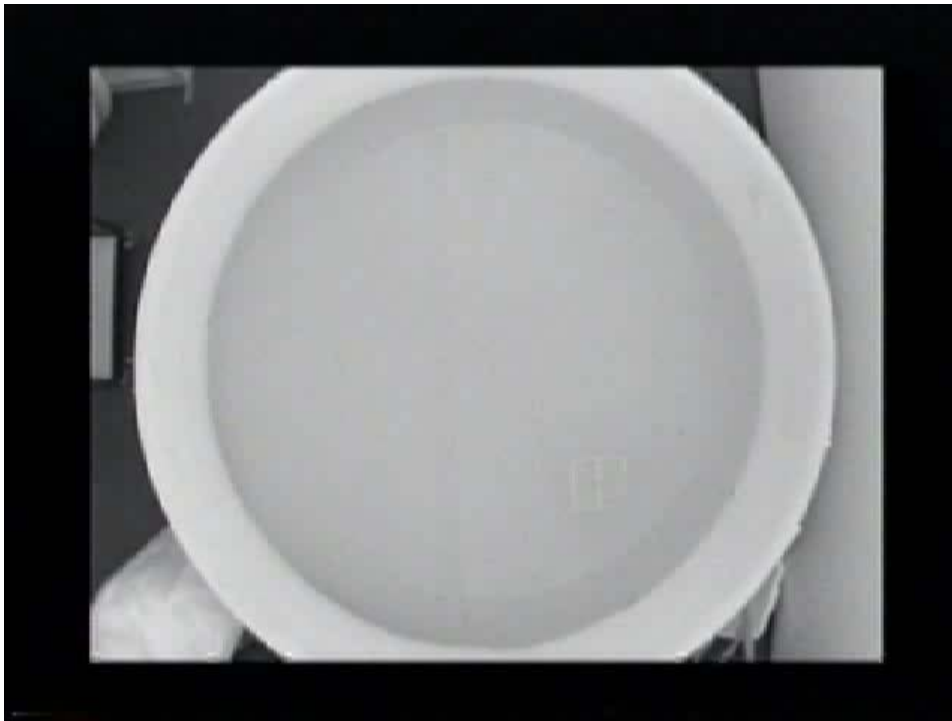


Genetic background:

Does it need consideration? YES!

Is it a hopeless issue? NO!

The place navigation task: testing spatial memory in a swimming pool



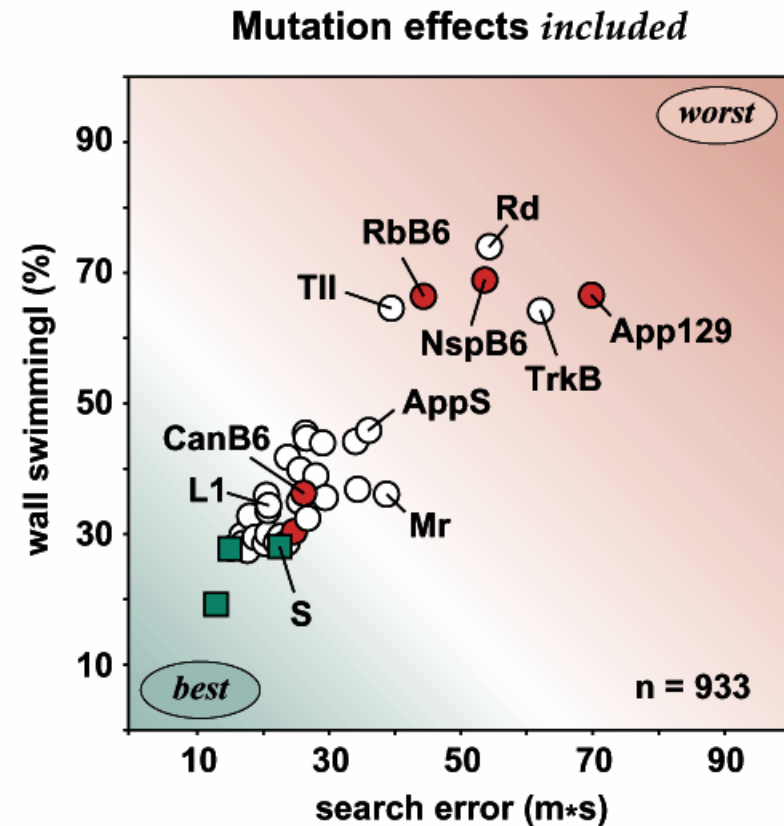
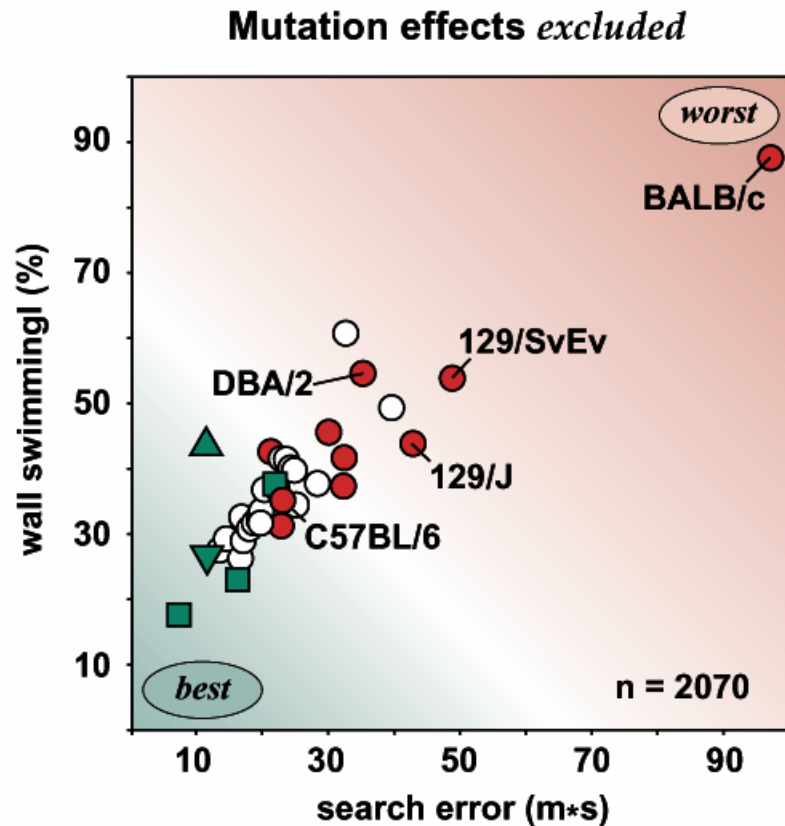
Morris RGM,
Learn Motiv, 12:239-260, 1981

Rats learn to swim to a hidden platform using cues located outside the pool.

Morris RGM et al.,
Nature, 297:681-683, 1982

Lesions of the hippocampus disrupt spatial navigation but not the ability to swim.

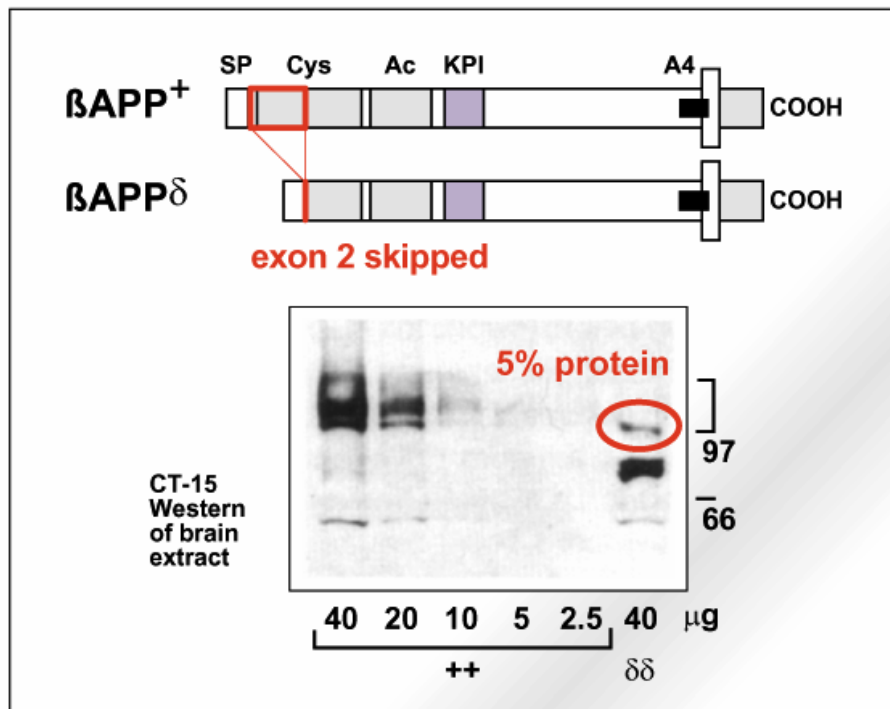
Place navigation in the water-maze: training performance of non/mutant mice



Genetic background effects

- Inconsistent or false negative results due to mutation-independent effects of genetic background:
 - baseline shift
 - ceiling/floor effects, non-performance of control animals
- Inconsistent results due to background x mutation interactions:
 - strain differences in ability to compensate
- False negative results due to large variability:
 - genetic noise in genetically inhomogeneous samples
- False positive results due to systematic differences in genetic background of mutant and control groups:
 - breeding of separate mutant and control lines
 - insertion effects
 - genetic linkage (“flanking gene problem”)

β APP-deficient mice



Phenotype of β APP $\delta\delta$ mice has been characterized in different genetic backgrounds:

Reduced body weight, impaired place navigation, abnormal openfield exploration, frequent agenesis of corpus callosum.

Müller et al. Cell 79:755-65, 1994

Retardation of neurological, behavioral, and somatic postnatal development. Permanent reduction of grip strength.

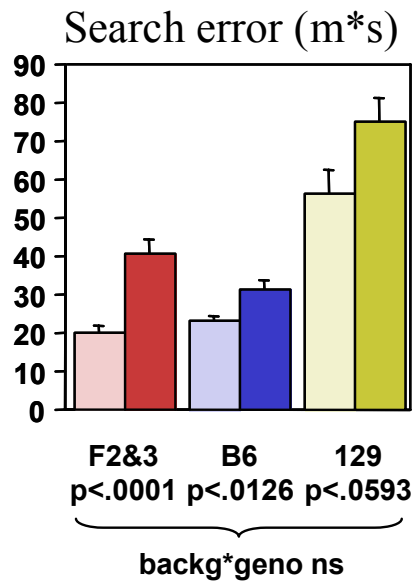
Tremml et al. BBR 95:65-76, 1998

Reduction of brain weight and forebrain commissure size. 100% corpus callosum agenesis in 129 background.

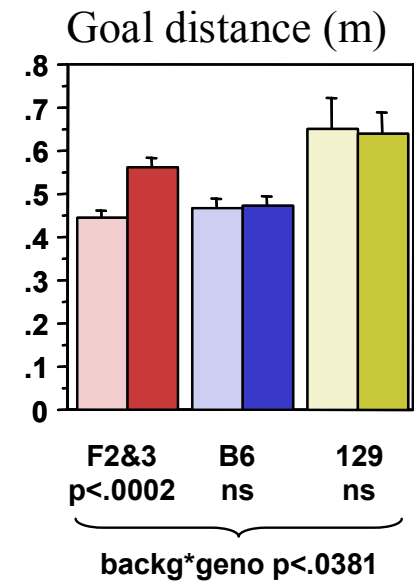
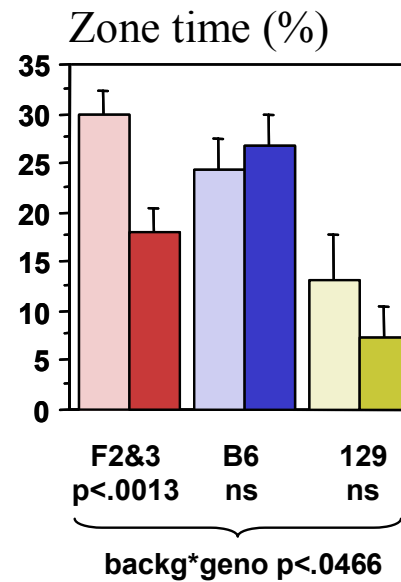
Magara et al. PNAS 96:4656-61, 1999


Genetic background modifies the impairment of β APP-deficient mice in the water-maze task


Acquisition



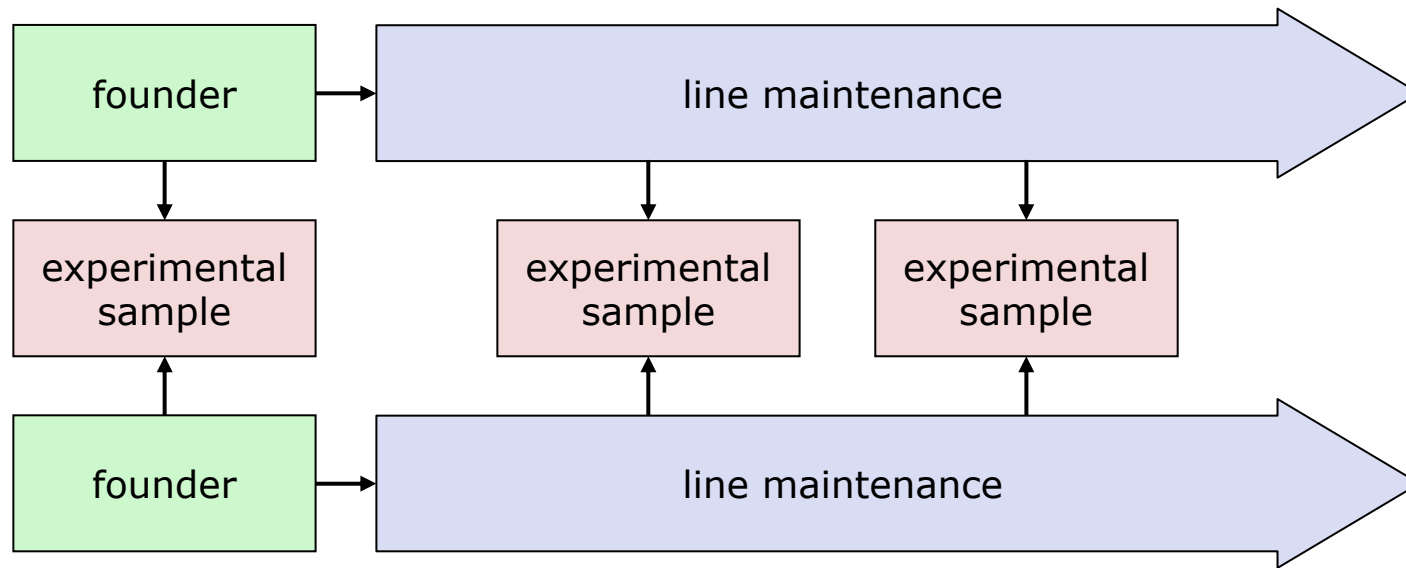
Transfer test







 62 wild-type
 65 APP δ/δ


 70 F2&3 C57BL/6 x 129Sv/Ev
 40 C57BL/6
 17 129Sv/Ev

Line generation, maintenance and analysis



Different genetic backgrounds may be appropriate for

-  generation of founder animals
-  long-term maintenance of mutant lines
-  animals used for experimental investigation

Common choices of genetic background

Founder

Strain determined by requirements of transgenic technique:

- homologous recombination: 129- or C57BL/6- derived ES-cells
- random insertion transgenics: F1-hybrid oocytes

Line maintenance

Goal: efficient breeding, easy transfer between labs, repeated generation of experimental samples with same genetic background:

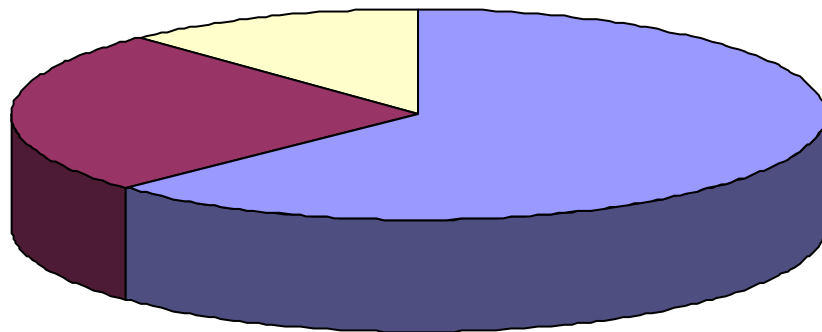
- backcross to ES-cell donor strain: *co-isogenic* line
- backcross to other commonly available strain: *congenic* line
- propagation of first test sample without control of background

Experimental samples

Goals: rapid and reproducible results, low variability, no ceiling or floor effects, suitability for wide range of investigations:

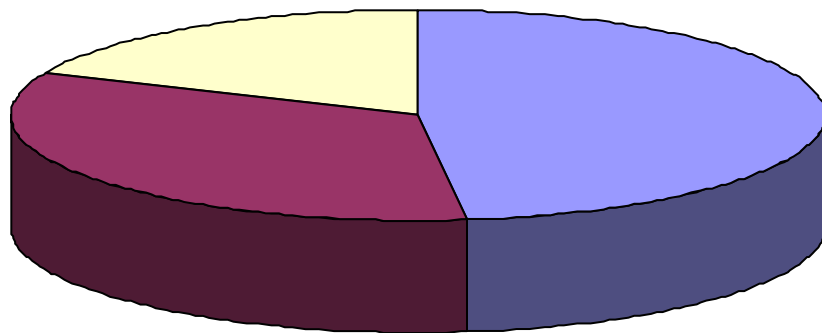
- F2/F3 crosses, inbred strains, F1 crosses

Genetic background of mouse lines tested in Zurich



1987-2004, 7167 mice

63% ■ F2 / mixed
25% ■ inbred
12% ■ F1 hybrid

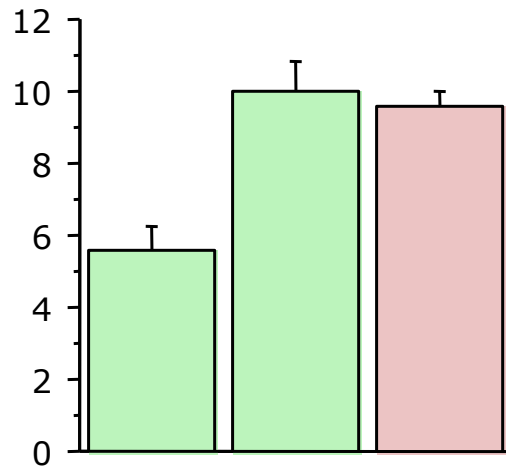


2002-2004, 2194 mice

48% ■ F2 / mixed
34% ■ inbred
18% ■ F1 hybrid

Inbreeding and variability of water-maze learning

Water-maze, training
SD search error (m,SE)

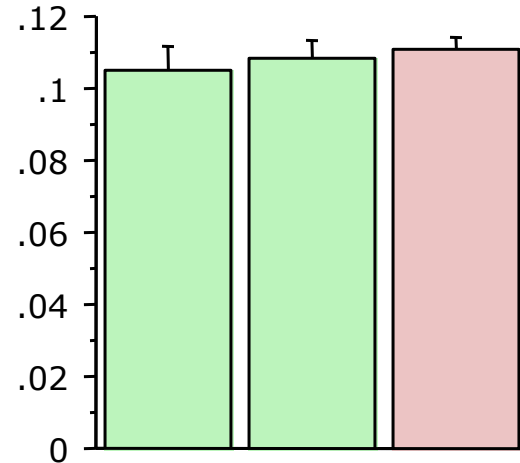


hybrid n=23 inbred n=52 mixed n=147

p<.0005 ns
p<.0004

2989 mice,
average group
size 13.5
(non-mutants)

Water-maze, probe test
SD goal distance (m,SE)







hybrid n=23 inbred n=52 mixed n=147


ns ns
ns

Minimal requirements for genetic background

Check that genetic background in experiment is...

-  ① ...free of bias between control and mutant groups
-  ② ...well documented and easy to reproduce
-  ③ ...compatible with all experimental procedures, with controls producing a baseline that prevents ceiling or floor effects

 *Maintain* mutation as congenic or co-isogenic line, e.g. by backcrossing to C57BL/6

 *Analyze* mutation in inbred, F1 hybrid or F2 samples using littermate controls (at least for behavioral experiments)