Does age matter? The impact of rodent age on study outcomes

Samuel J Jackson1,*, Nick Andrews2, Doug Ball3, Ilaria Bellantuono4, James Gray3, Lamia Hachoumi5, Alan Holmes6, Judy Latcham7, Anja Petrie8, Paul Potter9, Andrew Rice10, Alison Ritchie11, Michelle Stewart12, Carol Strepka13, Mark Yeoman5 and Kathryn Chapman1

Abstract
Rodent models produce data which underpin biomedical research and non-clinical drug trials, but translation from rodents into successful clinical outcomes is often lacking. There is a growing body of evidence showing that improving experimental design is key to improving the predictive nature of rodent studies and reducing the number of animals used in research. Age, one important factor in experimental design, is often poorly reported and can be overlooked. The authors conducted a survey to assess the age used for a range of models, and the reasoning for age choice. From 297 respondents providing 611 responses, researchers reported using rodents most often in the 6–20 week age range regardless of the biology being studied. The age referred to as ‘adult’ by respondents varied between six and 20 weeks. Practical reasons for the choice of rodent age were frequently given, with increased cost associated with using older animals and maintenance of historical data comparability being two important limiting factors. These results highlight that choice of age is inconsistent across the research community and often not based on the development or cellular ageing of the system being studied. This could potentially result in decreased scientific validity and increased experimental variability. In some cases the use of older animals may be beneficial. Increased scientific rigour in the choice of the age of rodent may increase the translation of rodent models to humans.

Keywords
rodent, age, experimental design, development, senescence

The most frequently used mammals in scientific research are rodents, predominantly rats and mice. There are various experimental factors which are critical to the quality of data obtained from rodent models and these factors differ according to the...
requirements of the experiment. Previous studies have shown that reporting of animal experiments in peer-reviewed publications is poor due to omission of details of experimental design such as blinding, randomization, and details of strain, sex and age.

Significant differences exist in disease-relevant systems in young or aged animals compared with middle-aged counterparts, and these differences may affect the outcome of studies investigating basic disease biology, mechanisms of drug action and efficacy. Inconsistent choice of age in rodent models between experiments or laboratories has the potential to impact on data quality, potentially increasing variability and reducing relevance to the human disease being studied. In basic or mechanistic studies, biological pathways or mechanisms of action may be obscured by inter-animal variability arising from inappropriately aged animals. In drug development, the translation of compounds from non-clinical research to phase II clinical trials has shown that reporting of animal experiments in peer-reviewed publications is poor due to omission of details of experimental design such as blinding, randomization, and details of strain, sex and age.

The profound effect of age in models of stroke has been documented, and young animals have been used to model a disease generally seen in older humans. Alterations in blood flow and brain biochemistry with age have a significant impact in humans, but these pathologies are not reflected when modelling the disease with young rodents. Older rats have been shown to exhibit a difference in susceptibility to, or take longer to recover from, ischemic insults.

Age-related phenotypes are being uncovered in commonly used inbred strains. For example, the widely used C57BL/6J mouse has impaired glucose tolerance due to a missense mutation in nicotinamide nucleotide transhydrogenase (Nnt), which has implications for diabetic phenotypes, as well as causing age-related decline in mitochondrial functions. In addition C57BL/6J mice have a mutation in the cadherin 23 (Cdh23) gene, resulting in age-related hearing loss. This may impact on behavioural studies as the mice age.

The microbiome plays important roles in the maintenance of appropriate immune system function and response to disease. Ageing can affect the composition of the mouse microbiome, and this has been shown to modulate mouse models of allergic airway disease, potentially reducing the quality of data obtained.

Drug metabolism by the liver has a critical impact on systemically administered compounds, and is therefore of primary importance during the development of new pharmaceuticals. The gene expression of critical liver enzymes is dramatically different between young and older counterparts, potentially leading to discrepancies or errors affecting drug candidate selection or development.

A working group was convened by the National Centre for the Replacement, Refinement and Reduction of Animals in Research (NC3Rs) to explore the age of rodents used to model physiology, disease or drug efficacy, and to assess the impact on scientific outcomes, with the aim of improving selection of the age of rodent models. Here, we present literature examples illustrating the importance of a well-informed age choice, and data from a survey regarding the age of rodents used in biomedical research. In addition, two data-based case studies are included as supplemental data (Figures S2 and S3; all supplemental material can be found online at http://lan.sagepub.com). By improving decision-making around the age of rodent used in biomedical research, the number of animals used can be reduced while maintaining or improving data quality.
Methods

Survey regarding age of rodent used

A cross-sector working group, reflected in the authorship, was formed from a diverse range of disciplines with an interest in the impact of age on rodent models of disease. In order to explore the age at which rodents are currently being used in research, an online survey was formulated by the expert working group with reference to published guidance on survey design.38 The survey requested data on the type of rodent model used by the researcher, and the age chosen for the model (see supplemental Table S1). Researchers could enter information on up to five separate models. Researchers were also asked for additional information on why the specific age was chosen and evidence for age suitability of the model. A drop-down list was used to establish reasons for choice of age (see supplemental Table S1 Q4).

The survey was pilot-tested and refined within the working group, and distributed as a hyperlink in a standard text via 20 learned societies and research charities, as well as individual scientists and scientific networks. The disciplines covered cancer, cardiovascular disease, endocrinology, immunology, musculoskeletal disease, neuroscience, and respiratory disease, and encompassed academic and pharmaceutical researchers in the UK and EU. This enabled distribution to a representative sample of researchers, as described in the Results section.

The data analysis workflow is shown in supplemental Figure S1. Data from the survey was anonymized and initially analysed to assess information about respondent location and field of work, and which models were represented in the data-set. Models which could not be classified (see below) or without a justification for the age used were removed. Each model was individually classified by ‘System/Process’ (see supplemental Table S2) and ‘Disease/Condition’ (a list of 233 diseases and conditions) as defined by the Wellcome Trust (London, UK).39

Comparison of age used for the same model paradigm across laboratories. Where the same model paradigm was reported in multiple responses, these were grouped and the range of ages plotted.

Age considered ‘adult’. Where the choice of age was justified by describing the animal as ‘adult’, the range of ages was plotted as a frequency graph.

Results

Given the literature demonstrations of changes in disease modelling in rodents with age, a survey was designed to collect information on the age at which researchers used rodents to model human disease or physiology. A diagrammatic description of the data workflow followed in this publication can be found in supplemental Figure S1.

There were 297 respondents to the survey, predominantly from the academic sector (80%), with 18% from the pharmaceutical sector. Location information was provided by 108 respondents (35% of the total) which demonstrated that responses were received from 27 UK and 31 non-UK locations.

The survey allowed for up to four responses per respondent. The total number of responses received was 611, representing a cross section of disciplines including neuroscience, immunology, cancer, genetics, physiology and toxicology. The range of ages reported was wide (2–160 weeks) but was heavily clustered around the 8–12 week age range for both mice and rats.

When the same model paradigm was reported multiple times, comparison of the age used across different laboratories was possible. Figure 1 illustrates the range of ages used for each model. This demonstrates that researchers in different laboratories used different ages of animal in the same experimental paradigm.

Responses (11.5%) cited that the reason for the choice of age was the rodent being ‘adult’. However, the age range over which the term ‘adult’ was applied differed amongst laboratories. The age range was 6–20 weeks (Figure 2). This illustrates the potential discrepancy when using general descriptions of rodent age.

Ten percent of the total responses described the use of rodents over 16 weeks of age (24% of these in rats and 76% in mice). The age in these cases was justified by one of four reasons: onset of the disease in rodent did not occur until after 16 weeks had passed (39%); the researchers wanted to capture advanced disease time-points including comorbidities which occurred after 16 weeks (33%); the animal was of a sufficient size for surgery to be performed (10%); the model was of a disease of older age in humans (18%).

A drop-down list was used to capture the reasons behind the choice of age. Respondents supplied 704 responses, with the highest cited reasons for the choice of age being historical data comparability (24%), supply/availability (18%) and cost (15%). When these reasons were split between academic and industry respondents, a difference was evident (Figure 3). Academic respondents cited the cost of animals as the factor which impacted their choice of rodent age the most (18% academic versus 6% industry), while industry respondents reported that historical data comparability was the most influential factor on the choice of rodent age (28% industry versus 23% academic).
Figure 1. Age ranges of rodent reported for specific models. Where a model was described sufficiently, there was opportunity to compare the age of specific model systems across responses. Respondents described that a range of ages of rodent were used for a given model. Ages used for all models clustered around the 8–12 week range, regardless of the biology being studied. The use of some ages, particularly at the low or high extremes, was justified by a specific biological reasoning.

Figure 2. Age considered adult: models described as ‘adult’ were identified, and the ages plotted as frequencies. This illustrates that the description ‘adult’ can encompass a wide range of ages, between six and 20 weeks for mice and between eight and 16 weeks for rats. These ages encompass ongoing development in a range of systems which could adversely affect the outcomes of an experiment. This could be avoided by basing the choice of age on the development of the system under examination. In addition, precise ages should be reported rather than ambiguous terminology such as ‘adult’.
Discussion

Taken together, these data demonstrate that: (1) choosing an age of rodent relevant to the human disease being studied can have a critical impact on the performance of the model; (2) the age of rodent is often based on practical considerations or convention; and (3) the age of rodent reported as ‘adult’ is inconsistent between studies or laboratories.

Reporting

Given that reporting of animal age and the reasoning behind the choice of age in publications has generally been poor,1 age choices are often based on convention or anecdotal evidence. The NC3Rs’ ARRIVE guidelines2 set out the attributes of experimental design to be reported to increase the reproducibility and transparency of in vivo experiments, including the age of animal used. More widespread and stringent adoption of the ARRIVE guidelines will ensure that reporting of animal age is more consistent in the future.

Consistency in age choice

The majority of responses described using animals between eight and 12 weeks, with a range from two to 120 weeks of age. When the same model paradigm was described in multiple responses, these were compared; the age used varied over a range of up to 20 weeks. This discrepancy in the age of animal used between different laboratories using the same model could lead to data variability obtained between those laboratories. When rodent age is not accurately reported, it cannot be taken into account in subsequent experiments, and this may lead to results not being repeatable elsewhere. This illustrates the importance of using standardized or consensus protocols.
and designing experiments with reference to published work on the biology of the system being studied.

Age considered ‘adult’

By examining responses which indicated that the age used represented an ‘adult’, the disparity when using this terminology was uncovered. The definition of ‘adult’ in this context is likely to be related to the sexual maturity of the rodent. Indeed, rodents are sexually mature and able to breed from around five weeks of age.40–42 However, this is not a sufficient basis on which to consider the whole animal to be fully developed. Many systems are immature at this age and may take weeks or months to develop to maturity. Additionally, there may be differences in the maturity of different strains at a particular age.43,44 The maturity of the system can significantly impact the outcome of an experiment, as demonstrated in the literature examples cited in this manuscript. ‘Adult’ would be defined differently in many of these examples; therefore this or other terminology referring to a ‘stage of life’ should be replaced with the actual age used when designing, documenting and reporting an experiment.

Where animals over 16 weeks of age were detailed in the response, the range of reasons used to justify this were very narrow, either pertaining to the biology of the model animal or the human disease being modelled. This reflects the fact that older animals are only likely to be used when justified by a specific biological attribute related to ageing; this principle is rarely applied to younger rodents.

Reasons for age choice

By gathering qualitative data regarding reasons for age choice for a large range of rodent models, we have demonstrated that ‘cost’ and ‘supply’ are second only to ‘historical data compatibility’ as reasons for choosing the age of rodent for a study. The cost pressure is particularly evident when academic respondents are analysed separately from pharmaceutical company respondents. Several academic respondents commented that cost precluded modelling in aged animals and that including higher costs for aged animals would appear unattractive on a grant application. Comparison of the cost of animals at six and 12 weeks from one major commercial supplier to the UK and US markets illustrates this issue: purchase of animals at 12 weeks will result in a cost increase of at least 50% per animal, sometimes close to 100% (Table 1). Researchers who purchase young animals and age them in-house will also face increased costs for housing and husbandry during this period. In order to support more robust and predictive rodent studies, funding bodies and researchers will need to work together to develop a strategy which takes into account that use of older rodents may be beneficial to data quality and the appropriate use of animals, and where justified assess the increased animal costs on grant applications accordingly.

Comparisons between mice and humans

Some respondents to the survey indicated that they were using a certain age of rodent as it was an equivalent age to the human being modelled. Equivalent ages between rodent and human have been defined by comparing survival rates between mice and humans over their lifespans. For example, one study41 defined a mature adult C57BL/6J mouse as 3–6 months (equivalent in this analysis to 20–30 human years), a middle-aged mouse as 10–14 months (equivalent to 38–47 human years) and an old mouse as 18–24 months (equivalent to 56–69 human years). The mature adult mouse represents a stage where development has ceased, but senescence has not yet started, and is recommended as the comparative age for studies into the effects of ageing. Using this approach, 12 weeks should be the minimum for a model of adult disease. Similarly calculations to ascertain equivalent rat and human ages have been published which illustrate the discrepancy in the speed of development at different stages between rats and humans.40,42 A framework and online tool have also been developed to translate neurobiological development across multiple species, including rodents and humans.45,46

Databases and information sharing

The Mouse Phenome Database47,48 (Jackson Laboratory, Bar Harbor, ME, USA) was the most frequently cited resource in the questionnaire for assessing the appropriate age for mice. Other resources cited include the Age-Phenome Knowledgebase49 (Ben Gurion University of the Negev, Beer Sheva, Israel) which allows access to literature sources based on searches of age, phenotype, or both, in mice and humans. Similar resources for rats are not available, but the Rat Genome Database (Bioinformatics
Research Center, Milwaukee, WI, USA) is intended to encompass these data in the future. To facilitate studies on aged rodents and reduce the overall numbers of animals used, networks such as the Shared Ageing Research Models (ShARM) initiative aim to promote the use and knowledge of aged rodents as models of ageing. In the US, the Aging Rodent Colonies project at the National Institute on Aging (Bethesda, MD, USA) banks and breeds aged rodents and tissues for supply to National Institutes of Health (NIH)-funded researchers working on ageing.

Conclusion and recommendations for best practice

Use of an inconsistent age of rodents in many areas of scientific research may impact data quality. By increasing awareness of this basic experimental parameter, it may be possible to improve model reliability and reduce the number of animals used in some experiments. Use of an appropriate and consistent age choice plus an improvement in reporting of age will require engagement of researchers across the academic and industry communities. Importantly, given the cost implications of using older animals, engagement of funding bodies is required to ensure that this is taken into account in research funding applications.

Recommendations

- The age of rodent used in a study should be based on the development or cellular ageing of the system or disease under scrutiny.
- Funding bodies and grant reviewers may need to take into account the increased costs associated with using older animals, and application for increased funding for this should be considered where justified.
- Effective communication between researchers and animal facility staff/veterinarians will increase the likelihood that an appropriate age is chosen.
- Accurate and consistent reporting of age in peer-reviewed literature will allow comparison of experimental methodology and development of consensus-based best practice.

Acknowledgments

The authors would like to thank Scott McCleary and Ian Holyer (GlaxoSmithKline, Stevenage, UK) for their work in generating and compiling data included in supplemental case study 1 of this manuscript, and the staff at the University of Brighton Bioresource Centre for their contribution to supplemental case study 2.

Declaration of conflicting interests

The author(s) declared no potential conflicts of interest with respect to the research, authorship, and/or publication of this article.

Funding

The author(s) received no financial support for the research, authorship, and/or publication of this article.

References

1. Festing MF and Altman DG. Guidelines for the design and statistical analysis of experiments using laboratory animals. ILAR J 2002; 43: 244–258.
2. Kilkenny C, Browne WJ, Cuthill IC, Emerson M and Altman DG. Improving bioscience research reporting: the ARRIVE guidelines for reporting animal research. PLoS Biol 2010; 8: e1000412.
3. Landis SC, Amara SG, Asadullah K, et al. A call for transparent reporting to optimize the predictive value of preclinical research. Nature 2012; 490: 187–191.
4. Adams DJ. The Valley of Death in anticancer drug development: a reassessment. Trends Pharmacol Sci 2012; 33: 173–180.
5. Talevi A, Bellera CL, Di Ianni M, Gantner M, Bruno-Blanch LE and Castro EA. CNS drug development – lost in translation? Mini Rev Med Chem 2012; 12: 959–970.
6. Bandeira F, Lent R and Herculano-Houzel S. Changing numbers of neuronal and non-neuronal cells underlie postnatal brain growth in the rat. Proc Natl Acad Sci USA 2009; 106: 14108–14113.
7. Somerville JM, Aspden RM, Armour KE, Armour KJ and Reid DM. Growth of C57BL/6 mice and the material and mechanical properties of cortical bone from the tibia. Calcif Tissue Int 2004; 74: 469–475.
8. Carnieli DS, Yoshioka E, Silva LF, et al. Inflammation and remodeling in infantile, juvenile, and adult allergic sensitized mice. Pediatr Pulmonol 2011; 46: 650–665.
9. Astori M, Finke D, Karapetian O and Acha-Orbea H. Development of T–B cell collaboration in neonatal mice. Int Immunol 1999; 11: 445–451.
10. Downes N and Mullins P. The development of myelin in the brain of the juvenile rat. Toxicol Pathol 2013; 42: 913–922.
11. Fu Y, Rusznak Z, Herculano-Houzel S, Watson C and Paxinos G. Cellular composition characterizing postnatal development and maturation of the mouse brain and spinal cord. Brain Struct Funct 2013; 218: 1337–1354.
12. Brodt MD, Ellis CB and Silva MJ. Growing C57BL/6 mice increase whole bone mechanical properties by increasing geometric and material properties. J Bone Miner Res 1999; 14: 2159–2166.
13. Halloran BP, Ferguson VL, Simske SJ, Burghardt A, Venton LL and Majumdar S. Changes in bone structure and mass with advancing age in the male C57BL/6J mouse. J Bone Miner Res 2002; 17: 1044–1050.
14. Holsapple MP, West LJ and Landreth KS. Species comparison of anatomical and functional immune system
development. Birth Defects Res B Dev Reprod Toxicol 2003; 68: 321–334.

15. Kay MM, Mendoza J, Diven J, Denton T, Union N and Lajiness M. Age-related changes in the immune system of mice of eight medium and long-lived strains and hybrids. I. Organ, cellular, and activity changes. Mech Ageing Dev 1979; 11: 295–346.

16. Velardi A and Cooper MD. An immunofluorescence analysis of the ontogeny of myeloid, T, and B lineage cells in mouse hemopoietic tissues. J Immunol 1984; 133: 672–677.

17. Ghia P, ten Boekel E, Rolink AG and Melchers F. B-cell development: a comparison between mouse and man. Immunol Today 1998; 19: 480–485.

18. Holladay SD and Smialowicz RJ. Development of the murine and human immune system: differential effects of immunotoxins depend on time of exposure. Environ Health Perspect 2000; 108(Suppl 3): 463–473.

19. Kincade PW. Formation of B lymphocytes in fetal and adult life. Adv Immunol 1981; 31: 177–245.

20. Wauthier V, Verbeek RK and Calderon PB. The effect of ageing on cytochrome p450 enzymes: consequences for drug biotransformation in the elderly. Curr Med Chem 2007; 14: 745–757.

21. Palasz A, Wiaderkiewicz A, Wiaderkiewicz R, et al. Age-related changes in the mRNA levels of CYP1A1, CYP2B1/2 and CYP3A1 isoforms in rat small intestine. Genes Nutr 2012; 7: 197–207.

22. Greaves LC, Barron MJ, Campbell-Shiel G, Kirkwood TB and Turnbull DM. Differences in the accumulation of mitochondrial defects with age in mice and humans. Mech Ageing Dev 2011; 132: 588–591.

23. Nelson JF, Felicio LS, Osterburg HH and Finch CE. Differential contributions of ovarian and extravarian factors to age-related reductions in plasma estradiol and progestosterone during the estrous cycle of C57BL/6J mice. Endocrinology 1992; 130: 805–810.

24. Mobbs CV, Cheyney D, Sinha YN and Finch CE. Age-correlated and ovary-dependent changes in relationships between plasma estradiol and luteinizing hormone, prolactin, and growth hormone in female C57BL/6J mice. Endocrinology 1985; 116: 813–820.

25. Bouxsein ML, Myers KS, Shultz KL, Donahue LR, Rosen CJ and Beamer WG. Ovariectomy-induced bone loss varies among inbred strains of mice. J Bone Miner Res 2005; 20: 1085–1092.

26. Ankolekar S, Rewell S, Howells DW and Bath PM. The influence of stroke risk factors and comorbidities on assessment of stroke therapies in humans and animals. Int J Stroke 2012; 7: 386–397.

27. Shapira S, Sapir M, Wengier A, Grauer E and Kadar T. Aging has a complex effect on a rat model of ischemic stroke. Brain Res 2002; 925: 148–158.

28. Yager JY and Thornhill JA. The effect of age on susceptibility to hypoxic-ischemic brain damage. Neurosci Biobehav Rev 1997; 21: 167–174.

29. Brown AW, Marlowe KJ and Bjelke B. Age effect on motor recovery in a post-acute animal stroke model. Neurobiol Aging 2003; 24: 607–614.

30. Toye AA, Lippiat JD, Proks P, et al. A genetic and physiological study of impaired glucose homeostasis control in C57BL/6J mice. Diabetologia 2005; 48: 675–686.

31. Rydstrom J. Mitochondrial NADPH, transhydrogenase and disease. Biochim Biophys Acta 2006; 1757: 721–726.

32. Ghosh D, Levault KR and Brewer GJ. Relative importance of redox buffers GSH and NAD(P)H in age-related neurodegeneration and Alzheimer disease-like mouse neurons. Aging Cell 2014; 13: 631–640.

33. Johnson KR, Zheng QY and Noben-Trauth K. Strain background effects and genetic modifiers of hearing in mice. Brain Res 2006; 1091; 79–88.

34. Langille MG, Meehan CJ, Koenig JE, et al. Microbial shifts in the aging mouse gut. Microbiome 2014; 2: 50.

35. Vital M, Harkema JR, Rizzo M, Tiedje J and Brandenberger C. Alterations of the murine gut microbiome with age and allergic airway disease. J Immunol Res 2015; 2015: 892568.

36. Mori K, Blackshear PE, Lobenhofer EK, et al. Hepatic transcript levels for genes coding for enzymes associated with xenobiotic metabolism are altered with age. Toxicol Pathol 2007; 35: 242–251.

37. Pibiri M, Sulas P, Leoni VP, et al. Global gene expression profile of normal and regenerating liver in young and old mice. Age (Dordr) 2015; 37: 9796.

38. Bennett C, Khangura S, Brehaut JC, et al. Reporting guidelines for survey research: an analysis of published guidance and reporting practices. PLoS Med 2010; 8: e1001069.

39. Wellcome. Subject Classification of Wellcome Trust Grants. London: Wellcome, 2014.
48. Jackson. *Jackson Laboratories Mouse Phenome Database*. Bar Harbor, ME: Jackson Laboratory, 2014.
49. Geifman N and Rubin E. The mouse age phenome knowledgebase and disease-specific inter-species age mapping. *PLoS One* 2013; 8: e81114.
50. de la Cruz N, Bromberg S, Pasko D, et al. The Rat Genome Database (RGD): developments towards a phenome database. *Nucleic Acids Res* 2005; 33: D485–D491.
51. Duran AL, Potter P, Wells S, et al. Shared Ageing Research Models (ShARM): a new facility to support ageing research. *Biogerontology* 2013; 14: 789–794.
52. Nadon N. *Aged rodent colonies handbook*. Bethesda, MD: National Institute on Aging, 2014.