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1 **Comprehensive ECG reference intervals in C57BL/6N substrains provide a**  
2 **generalizable guide for cardiac electrophysiology studies in mice**

3

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## 1 **Introduction**

2 Reference ranges are a powerful tool for diagnostic decision-making in clinical medicine  
3 and their use has become increasingly common<sup>1</sup>. Reference ranges are derived intervals  
4 containing a defined subset of values from a large and comparable population dataset.  
5 These values, designed to delineate the expected range of a given parameter, are used  
6 clinically to identify outlier values. Individuals presenting with values outside of a clinically  
7 defined reference range are considered abnormal and flagged for follow up clinical  
8 investigation.

9 Looking beyond clinical applications, reference ranges are of enormous value in pre-  
10 clinical, basic scientific research using *in vivo* modelling<sup>1</sup>. They are used to define  
11 “normality” for a given genetic background, sex, and age of animals, such as inbred  
12 mouse strains. To our knowledge, there are no published reference ranges for  
13 electrocardiography (ECG) in the laboratory mouse. Such reference ranges would  
14 provide the research community with the information necessary to evaluate the  
15 consequences of pharmacological, environmental, or genetic perturbations, the latter  
16 opening up the opportunity to uncover genotype\*phenotype associations.

17 We used ECG data collected under the auspices of the International Mouse Phenotyping  
18 Consortium (IMPC<sup>2</sup>, <https://www.mousephenotype.org>) to generate the first mouse-  
19 specific cardiac physiology reference ranges. Here, data were collected from over 26,000  
20 conscious or anesthetized C57BL/6N wildtype control mice stratified by sex and age. The  
21 unprecedented scale of this data resource yields a robust reference range for a broad  
22 and commonly studied set of ECG parameters that are clinically important to assess  
23 myocardial electrical processes and cardiac function.



## 1 **Results**

2 ECG data collected by IMPC contributing centers (data release, DR, 15.0) were available  
3 from 26,706 wildtype control mice, stratified as presented in **Table 1** and summarized  
4 below. All the mice were from a C57BL/6N inbred substrain. ECG was performed on  
5 conscious mice, or mice anesthetized with either isoflurane or tribromoethanol. The  
6 majority of mice (90.6% or 24,194) were tested at a mean age of 12 weeks (designated  
7 as “early adult” or EA), while the remaining 9.4% (2,512) of mice were tested at a mean  
8 age of 62 weeks (designated as “late adult” or LA). Sex was evenly distributed at both EA  
9 and LA time points. Raw data can be downloaded using the following link:  
10 <https://www.mousephenotype.org/data/previous-releases/15.0>. The total number of  
11 reported parameters varied slightly between mice and can be accessed in **Supplemental**  
12 **Table 1**.

## 13 **Variability Assessment**

14 A panel of 15 output parameters were collected from ECG, namely heart rate (HR), RR-,  
15 PR-, PQ-, ST-, and QT-interval, and QT corrected (QTc) using the Mitchell formula<sup>4</sup>, QRS  
16 complex, coefficient of variation of R-R intervals (CV), heart rate variability (HRV), pNN50,  
17 rMSSD (Root Mean Sum of Squared Distance), mean R-amplitude, mean SR-amplitude  
18 and QT corrected (QTc) dispersion (parameter definition in **Supplemental Table 2**).

19 In multi-center, large-scale, high-throughput programs such as the IMPC, variability in the  
20 measured values was to be expected. However, the extent of this variability dictates the  
21 sensitivity and robustness of each parameter.

1 Variability testing was performed on all DR 15.0 ECG data from the IMPC, independently  
2 of anesthetic agent in this analysis. For each sex, individual ECG parameters were tested  
3 for variability in EA and LA populations. The following standard metrics for assessing  
4 distribution variability were calculated:

5 1) Coefficient of variation (COV) ( $100 \times \text{standard deviation} / \text{mean}$ ) assumes a parametric  
6 distribution and normalizes the variability to the most typical score (mean) but is sensitive  
7 to outliers. To support the parametric COV test, we applied a 2) “Quartile-based CV”  
8 (QCV), defined as interquartile range (IQR) (75-25%) relative to the median  
9 ( $100 \times \text{IQR} / \text{median}$ ). QCV is a similar metric to COV but uses non-parametric measures of  
10 variability, therefore makes no assumptions of normality but is still readily influenced by  
11 outliers<sup>3,4</sup>.

12 Based on this analysis, exclusion criteria were defined as any parameter with acceptable  
13 variability based on Eurachem guidelines  
14 (<https://www.eurachem.org/index.php/publications/guides>) of  $\geq 30$  for COV (**Figure 1**) and  
15 a QCV  $\geq 30$  for EA and LA mice (**Supplemental Figure 1**). Figure 1 shows that the  
16 retained parameters are all clustered closely together, however the excluded parameters  
17 show a wide range of variability. Specifically, seven ECG-parameters (CV, HRV, pNN50,  
18 rMSSD, mean R-amplitude, mean SR-amplitude and QTc dispersion) exceeded the  
19 variability criteria in both sexes (male and female) and ages (EA and LA) and were  
20 excluded from further analysis (**Figure 1**). The variability threshold was exceeded least  
21 for QTc dispersion in EA and mean R- and SR-amplitude for LA, however, for the  
22 remaining parameters that were excluded, variability was in excess of 2-7 times the  
23 threshold.

1 A PQ-interval reference range is provided for conscious EA and LA mice (**Supplemental**  
2 **Figure 2**) however, PQ-interval was excluded from further analysis in this study because  
3 data points were only captured in EA and LA mice from one of the ten data contributing  
4 centers. The remaining seven ECG-parameters [heart rate (HR), RR-, PR-, ST- and QT-  
5 interval, QRS complex, and QT corrected (QTc) using the Mitchell formula<sup>5</sup>] consistently  
6 presented with low variability across the whole IMPC dataset thereby giving high  
7 confidence to establish robust, generalizable reference ranges for EA and LA populations  
8 on the C57BL/6N inbred genetic background.

9 Despite the exclusion of several parameters, the electrical conduction phases of a cardiac  
10 cycle were entirely captured by the robust parameters included herein (**Figure 2**). The  
11 lengths of PR-interval and QRS complex covered the atrial and ventricular depolarization  
12 phases (e.g. contraction), whereas lengths of QT- and ST-intervals implied the ventricular  
13 repolarization (e.g. relaxation) in voltage over time.

#### 14 **Assessment of Data Distribution**

15 The distribution of data was assessed via histograms for the seven selected ECG  
16 parameters stratified by sex, age, and anesthetic regime (**Figure 3**). This visual  
17 representation of the frequency of occurrence per value in the data was useful for  
18 revealing conformity to- and deviations from- a normal distribution, for each parameter.  
19 Visual inspection of the histograms showed that the data appeared practically normal for  
20 parameters PR, QT and QTc Mitchell, and modestly skewed for HR, QRS, ST and RR.  
21 To assess normality mathematically, we applied the Shapiro-Wilk test which revealed  
22 statistically significant deviation from a normal distribution for some, but not all, ECG  
23 parameters. **Table 2** presents data as median and 95% reference range (2.5th and 97.5th

1 percentile) to account for the lack of normal distribution of some parameters and to  
2 provide a consistent data presentation<sup>4</sup>. For the sake of completeness, mean, standard  
3 deviation and sample size are provided for the seven selected ECG parameters stratified  
4 by sex, age, and anesthetic regime in **Supplemental Table 1**.

5 Interestingly, male, and female data showed similar distributions by visual inspection  
6 (**Figure 3**). To test the hypothesis that there is no difference between each sex, a simple  
7 two-tailed t-test was performed independently for each anesthetic regime and age group,  
8 and Cohen's *d* was calculated as an effect size measure (**Supplemental Figure 3, 4 and**  
9 **5 - Panels a and b, stratified by age**).

10 For some parameters, p-values reached significance  $<.001$ , for others we found no  
11 evidence of a difference. However, for all parameters the corresponding Cohen's *d* value  
12 revealed small to negligible effect sizes. We therefore considered the possibility that the  
13 large group sizes could be overstating the biological differences between the sexes for  
14 some parameters.

15 To address this, we applied a bootstrap analysis stratified by age group. In brief, random  
16 sampling (1000x randomized) of different sub-sample sizes, ranging from 5 to 100 mice,  
17 were applied to test the robustness of the effect for each parameter comparing females  
18 and males. The sub-sample group sizes were chosen to more closely approximate  
19 standard experimental groups. The proportion of significant t-tests ( $p<.05$ ), from the 1000  
20 comparisons, indicates the power to find the sex difference, for that sub-sample size. If  
21 the proportion of significant tests remains near 5% regardless of sub-sample size, then  
22 this indicates the influence of the Type 1, i.e. false positive, error.

1 Recordings from both conscious (**Supplemental Figure 3**) and isoflurane anesthetized  
2 mice (**Supplemental Figure 4**) show that the ECG parameters consistently have very  
3 low proportions of significant tests for sexual dimorphism, with most parameters  
4 fluctuating around 5% of tests. Tribromoethanol anesthesia (**Supplemental Figure 5**)  
5 however, reveals weak sexual dimorphism for a subset of parameters. This may be due  
6 to a bias from drawing bootstrap samples from a much smaller “population” than the other  
7 conditions, but we cannot exclude the possibility that this anesthetic has a small but  
8 significant impact on the sexes.

### 9 **Effect of Anesthetic Agent**

10 To investigate the effect of different anesthetic agents on cardiac conduction function and  
11 ECG profiles, conscious data stratified by sex and age are displayed for comparison with  
12 those of isoflurane or tribromoethanol data (**Figure 4**). Female data are placed directly  
13 above male for ease of visualization. **Figure 4** shows distinct distribution clusters for  
14 conscious, isoflurane and tribromoethanol groups split by EA (**Figure 4 – Panels A to G**)  
15 and LA (**Figure 4 – Panels H to N**). As before, no data were available for tribromoethanol  
16 anesthesia in LA mice.

17 As expected, the physiological benchmark of highest heart rate in conscious mice  
18 compared to anesthetized animals was observed (**Figure 4 – Panels A and H**). To  
19 assess the differences between EA anesthetic states, we tested conscious versus  
20 isoflurane and conscious versus tribromoethanol groups, by a one-way ANOVA with  
21 planned comparisons, and observed highly significant differences between those groups  
22 (**Table 3**). These data clearly show differences in ECG parameters that can be attributed  
23 to the anesthetic regime; therefore, it is essential to establish reference ranges separately

1 by condition (conscious or anaesthetized) and by anesthetic (isoflurane or  
2 tribromoethanol).

### 3 **Effect of Age on ECG Parameters**

4 Two different age groups, i.e. mean of 12-weeks (minimum 8 and maximum 16 weeks)  
5 old EA and mean of 62 weeks (minimum 52 and maximum 78 weeks) old LA, have made  
6 it possible to explore the effect of age on ECG parameters in conscious and isoflurane  
7 anaesthetized mice. A two-tailed t-test was applied to test the difference between the  
8 means of EA and LA results in conscious mice (**Figure 5 - Panels a and b**). P-values  
9  $<.001$  were reached for all parameters, indicating high statistical significance and the  
10 corresponding Cohen's *d* effect size revealed negligible to medium standardized effect  
11 sizes (**Figure 5 – Panels a and b**).

12 To test the influence of unbalanced group sizes (i.e. large number of EA and smaller  
13 number of LA datasets), we applied a bootstrap analysis, this time stratified by sex  
14 (**Figure 5 – Panel c**). This bootstrap analysis demonstrated that parameters with even  
15 small to medium effect sizes required relatively large experimental group sizes to attain  
16 a conventional  $>80\%$  value for power estimates<sup>6,7</sup>, e.g. QRS and ST in conscious  
17 conditions required a group size of 50 mice to achieve  $>80\%$  power with a  $p<.05$  (**Figure**  
18 **5 – Panel c, QRS (subpanel C) and ST (subpanel F)**). As expected, for parameters with  
19 negligible Cohen's *d* effect sizes, such as HR and RR, increases in sample size do not  
20 appreciably increase power (**Figure 5 – Panel c, HR (subpanel A) and RR (subpanel**  
21 **E)**). Parameters with less than 80% power even with up to  $n=100$  animals, can be  
22 considered likely to be “similar in EA and LA” with no aging effect for most experimental

1 purposes. **Supplemental Figure 6** presents the equivalent t-test, Cohen's *d* and  
2 bootstrap analysis in EA and LA mice anesthetized with isoflurane.

3 In summary, **Figure 6** is a graphical representation of the median and 95% reference  
4 ranges (2.5th and 97.5th percentile) broken down by anesthetic regimen with the female  
5 data placed directly above equivalent male data for easy visual interpretation  
6 corresponding numeric values are presented in **Table 2**.

### 7 **Validation of Reference Ranges Using Non-IMPC Data**

8 Mice characterized by the IMPC are all substrains of one commonly used inbred genetic  
9 background, C57BL/6N. To test the validity of the reference ranges reported herein  
10 beyond C57BL/6N inbred mice, we used representative control animals from publicly  
11 available ECG data including: six founder strains from a collaborative cross study<sup>8</sup>; the  
12 Jaxwest1 project (<https://phenome.jax.org/projects/Jaxwest1>) with seven inbred strains  
13 of mice; and the Xing1: Aging study (<https://phenome.jax.org/projects/Xing1>)<sup>9</sup> with 29  
14 inbred strains of which we have included herein the 26 strains with complete ECG data.  
15 An additional dataset was included using inbred, wildtype control animals from non-IMPC  
16 studies conducted at the German Mouse Clinic where data is available upon request.  
17 Validation was also carried out for LA population by using 12- and 20-month age groups  
18 of the Xing1 study. In each non-IMPC study, where suitable we presented the data split  
19 by sex and overlaid with the sex-specific 95% reference range calculated herein for  
20 conscious mice. Due to the small sample sizes in a subset of these comparator studies,  
21 however, the combined reference ranges for females and males are summarized in  
22 **Supplementary Table 3** for further comparison. **Figure 7** shows the founder strain data  
23 from the collaborative cross study overlaid with the reference ranges split by sex whereas

1 **Supplemental Figure 7** illustrates data from the German Mouse Clinic, **Supplemental**  
2 **Figure 8** from the Jaxwest1 and **Supplemental Figures 9-13** depict LA data from the  
3 Xing1 study. Of note, HR is not presented throughout as it was not accessible for those  
4 studies yet it is indirectly visualized in the RR-interval plot due to the inverse correlation  
5 ship of HR and RR<sup>10</sup>.

6 Remarkably, and true for all ECG parameters, most non-C57BL/6N values lay within our  
7 reference values. There is a subset of outliers that fall outside of the reference ranges  
8 which is to be expected with heterogeneity of small size and phenotypic differences seen  
9 between inbred mouse strains.

10



## 1 **Discussion**

2 Reference ranges for the assessment of abnormal electrocardiograms and cardiac  
3 conduction disorders in patients have long been established and are regularly adopted  
4 by expert bodies such as the North American Society of Pacing and Electrophysiology<sup>11</sup>  
5 and the European Society of Cardiology<sup>12,13</sup>. For mouse models, however, there are no  
6 such reference ranges.

7 In this multicenter study, we have established reference ranges using an exceptionally  
8 large ECG dataset comprising more than 26,000 wildtype control mice from the  
9 International Mouse Phenotyping Consortium (IMPC). The goal of the IMPC is to extend  
10 the functional annotation of the mammalian genome via the large-scale production and  
11 phenotypic characterization of single gene knockout mouse strains for all protein coding  
12 genes. The phenotypic pipeline used to characterize these knockout strains included  
13 cardiac electrophysiology assessment using ECG. For each knockout strain  
14 characterized, age, sex and genetic background matched wildtype control animals were  
15 also assessed. The ECG data from these C57BL/6N wildtype control mice hold  
16 extraordinary value and represent the focus of the current study.

17 Thus, this study represents a large mouse data set and allows the crucial understanding  
18 of the effects of sex, age, and anesthesia on electrocardiograms in mice. To this end, we  
19 introduced a stepwise refinement of the data analysis and started with an in-depth  
20 assessment of the variability of 15 ECG parameters gathered in the IMPC. We identified  
21 seven clinically relevant ECG parameters that were highly robust and had low variability.  
22 We excluded the remaining eight ECG parameters because of the excessive level of inter-  
23 mouse variability they displayed. Five of the eight excluded parameters were direct

1 measures of heart rate variability (HRV), or represented parameters derived from HRV  
2 (HRV, pNN50, rMSSD, mean R-amplitude and mean SR-amplitude). HRV depicts the  
3 change in the time interval between successive heartbeats and is an index of the  
4 parasympathetic nervous system<sup>14,15</sup>. HRV measurement is very sensitive to  
5 experimental methods (e.g. acclimation time, ECG sampling rate, and duration of  
6 recording), and has been shown to be incompatible with a high-throughput data collection  
7 set up such as that used by the IMPC<sup>16,17</sup>. Next, CV provides an indication of the function  
8 of the parasympathetic nerve and the autonomic nervous system through the  
9 physiological phenomenon of RR variation<sup>18</sup>. Such measurements, however, require  
10 stable and prolonged measurement times to be meaningful, which, as stated above for  
11 HRV, we do not have in the context of the high throughput testing paradigm used herein.  
12 Similarly, this susceptibility to broad variability in short duration measurements also  
13 applies to the parameter QT dispersion, which is defined as the difference between the  
14 longest and shortest QT interval in one of the surface ECG leads and quantifies the spatial  
15 inhomogeneity of ventricular repolarization. Mainly for methodological reasons,  
16 parameters with high variability were excluded here, but PR interval is the exception. This  
17 parameter was only collected by one center and therefore not included in the overall  
18 evaluation, but the values were made available in full in the supplemental materials.  
19 Despite the exclusion of those parameters, the robust ECG parameters that were  
20 included entirely captured the electrical conduction phases of a cardiac cycle and  
21 provided a comprehensive ECG evaluation.

22 Understanding the sex-related impact on ECG is crucial for ensuring robust reference  
23 values. In this study, we were able to show that the values for HR, RR-, PR-, ST- and QT-

1 interval, QRS complex, and QT corrected (QTc) using the Mitchell formula<sup>5</sup> are  
2 comparable in female and male mice with negligible sexual dimorphism. There may,  
3 however, be small sex differences for some parameters depending on the anesthetic  
4 agent. This observation is of key importance, and in part consistent with previous mouse  
5 data<sup>19</sup>. Whilst sexual dimorphism was not overtly apparent in inbred mice in the absence  
6 of any environmental, pharmacological or genetic perturbations, the literature clearly  
7 supports sex differences in heart health<sup>20</sup> and therefore our recommendation is that both  
8 sexes are included in any experimental design assuming that post-treatment we may  
9 detect sex differences.

10 Anesthetics cause a dose-dependent decrease in myocardial contractile force and  
11 associated ECG alterations with the most familiar landmark of decreased HR<sup>21</sup>. Our  
12 observations are that presence of anesthesia matters, we confirm a decreased heart rate  
13 in anesthetized mice and go on to reveal distinctions in isoflurane inhalation anesthesia  
14 and intraperitoneal injected tribromoethanol induced anesthesia<sup>22,23</sup>. These distinctions  
15 are pivotal and to emphasize them we mapped the effects of three different states  
16 (conscious, isoflurane and tribromoethanol anesthesia) on seven ECG parameters in  
17 detail and present anesthesia-specific reference values.

18 HR is an important determinant of cardiovascular performance defined by the activity of  
19 the sinoatrial node, the so-called pacemaker of the heart. The dysfunction of the sinoatrial  
20 node increases with age, and HR decreases due to tissue, cellular, and molecular  
21 mechanisms that underlie the reduction in pacemaker activity with age<sup>24-26</sup>. Interestingly,  
22 we did not observe any strong age-related ECG changes in the absence of any  
23 pharmacological, environmental, or genetic challenges in inbred C57BL/6N mice. The

1 differences in the reference ranges of 12-week-old young adult mice compared to 62-  
2 week-old adult mice were negligible. Our step-by-step analysis of these data in  
3 bootstrapping showed that age-related ECG effects are more likely, if at all, to be detected  
4 in large group numbers (>50). This dependency on the group size can be used as a guide  
5 for experimental design when considering aging.

6 In the IMPC, we control for genetic diversity by using C57BL/6N inbred background  
7 substrains thereby focusing our comparison on the genetic perturbation of interest i.e. the  
8 single gene that is knocked out on this common genetic background. The transferability  
9 from the C57BL/6N background used here, however, was demonstrated by independently  
10 validating the ranges using data from a broad spectrum of non-IMPC C57BL/6N and  
11 C57BL/6J strains, and other inbred strains, including wild derived inbred strains. This  
12 validation indicates that C57BL/6N-based reference values represent a robust and  
13 comprehensive indicator of normality and can be used as a starting point for many  
14 experimental investigations of cardiac function in the mouse.

15 In summary, we have created a unique and comprehensive map of ECG reference ranges  
16 that will be foundational for future mouse studies. While based on inbred mouse  
17 substrains that are C57BL/6N in origin, these reference ranges have utility across  
18 different mouse strains and are important guides in studies of electrical conductivity  
19 disorders.

1 **Methods**

2 ***The International Mouse Phenotyping Consortium***

3 The International Mouse Phenotyping Consortium (IMPC) represents a multi-institutional  
4 and collaborative research initiative encompassing twenty-four major research  
5 organizations and funding agencies, distributed globally. The IMPC seeks to generate  
6 and phenotype a knockout mouse line for every protein-coding gene in the mouse  
7 genome ([www.mousephenotype.org](http://www.mousephenotype.org))<sup>27,2</sup>. Phenotyping is carried out under the uniform  
8 operating procedures detailed in IMPReSS (International Mouse Phenotyping Resource  
9 of Standardized Screens; [www.mousephenotype.org/impress/index](http://www.mousephenotype.org/impress/index)), which were  
10 developed and validated during the pilot programs EUMORPHIA and EUMODIC<sup>28</sup>.

11 ***IMPC Centers Contributing Electrocardiography Data***

12 IMPC data release (DR) 15.0 was used herein  
13 (<https://www.mousephenotype.org/data/previous-releases/15.0>). The following subset of  
14 ten IMPC data-contributing centers provided electrocardiography (ECG) data in DR 15.0  
15 (ethical approval details are included in parenthesis after each contributing center):

- 16 1. Baylor College of Medicine (BCM) (Institutional Animal Care and Use Committee  
17 approved license AN-5896).
- 18 2. German Mouse Clinic Helmholtz Zentrum München (GMC) (#144-10, 15-168)
- 19 3. Medical Research Council (MRC) – Harwell (HAR) (Animal Welfare and Ethical Review  
20 Body approved licenses 70/8015 and 30/3384).

1 4. Institute Clinique de la Souris, Mouse Clinical Institute (ICS) (#4789-  
2 2016040511578546v2).

3 5. The Jackson Laboratory (JAX) (Institutional Animal Care and Use Committee approved  
4 licenses 14004, 11005, and 99066. JAX AAALAC accreditation number was 000096, NIH  
5 Office of Laboratory Animal Welfare assurance number was D16-00170).

6 6. RIKEN BioResource Research Center (RBRC) (Animal Care Committee approved  
7 animal use protocols 0153, 0275, 0277, and 0279).

8 7. University of California – Davis (UCD) (Institutional Animal Care and Use Committee  
9 approved animal care and use protocol number was 19075. UCD AAALAC accreditation  
10 number is 000029, and the NIH Office of Laboratory Animal Welfare assurance number  
11 is D16-00272 # (A3433-01).

12 8. Seoul National University, Korea Mouse Phenotyping Center (KMPC) (KRIBB-AEC-  
13 19189).

14 9. Czech Centre for Phenogenomics (CCP) (AV CR 62/2016, Academy of Sci., Czech  
15 Rep.).

16 10. The Centre for Phenogenomics, Toronto (TCP) (22-0275 and 22-0279).

17 ECG data was collected from mice at one of two possible time points. For the Early Adult  
18 (EA) Pipeline data were collected at a mean of 12 weeks with the minimum of 8 and  
19 maximum of 16 weeks of age. For the Late Adult (LA) Pipeline data were collected at a  
20 mean of 62 weeks with the minimum of 52 and maximum of 78 weeks of age. Animal  
21 welfare was assessed routinely for all mice involved.

1 **Animals**

2 This study includes data collected from inbred wildtype control animals tested as part of  
3 the IMPC goals. These mice, both males and females, were on a C57BL/6N genetic  
4 background of substrains: C57BL/6NCrl (CCP, HMGU, ICS, TCP and UCD); C57BL/6NJ  
5 (JAX and BCM); C57BL/6NJcl (RBRC) and C57BL/6NTac (KMPC, HMGU, ICS and  
6 HAR). Non-IMPC mice were from four different studies: (1) The founder strains animals  
7 from a study titled “The Collaborative Cross: A Recombinant Inbred Mouse Population for  
8 the Systems Genetic Era”<sup>8</sup> with A/J, C57BL/6J, 129S1/SvImJ, NOD/ShiLtJ, NZO/HILtJ,  
9 CAST/EiJ, PWK/Ph, and WSB/EiJ inbred strains  
10 (<https://phenome.jax.org/projects/GMC13>); (2) The Jaxwest1 project, a multi-system  
11 analysis of physiology on seven inbred strains of mice: 129S1/SvImJ, A/J, BALB/cJ,  
12 C57BL/6J, DBA/2J, NOD/ShiLtJ and SJL/J (<https://phenome.jax.org/projects/Jaxwest1>);  
13 (3) Wildtype control animals from three non-IMPC studies performed at the German  
14 Mouse Clinic (<https://www.mouseclinic.de/>) with a standard sample size (20-30 control  
15 animals per study). The mouse backgrounds were: (i) An independent repeat of strain  
16 C57BL/6NJ (Jackson Laboratory strain #:005304) that is used by some of the IMPC  
17 contributing centers; (ii) C57BL/6J (JAX strain #:000664), the most commonly used inbred  
18 mouse strain and the first to have its genome sequenced; and (iii) FVB (JAX strain  
19 #:001800), a widely used multipurpose inbred line. For more information on these inbred  
20 strains, visit: <https://www.jax.org/strain>; and (4) The Xing1, Aging study:  
21 Electrocardiogram for 29 inbred strains of mice (<https://phenome.jax.org/projects/Xing1>)<sup>9</sup>.  
22 Xing1 recorded ECG characteristics in the following 26 inbred mouse strains:  
23 129S1/SvImJ, A/J, BALB/cByJ, BTBR-T+tf/J, BUB/BnJ, C3H/HeJ, C57BL/10J,

1 C57BL/6J, C57BLKS/J, C57BR/cdJ, C57L/J, CBA/J, DBA/2J, FVB/NJ, KK/HIJ, LP/J,  
2 MRL/MpJ, NOD.B10Sn-H2/J, NON/ShiLtJ, NZO/H1LtJ, NZW/LacJ, P/J, PL/J, RIIS/J,  
3 SM/J, and SWR/J. AKR/J, PWD/PhJ and SJL/J were excluded herein due to incomplete  
4 ECG data.

### 5 ***Data Collection***

6 The IMPC standard operating procedure provides an overview of the conscious and  
7 anesthetized ECG procedures used by contributing centers  
8 (<https://www.mousephenotype.org/impress/ProcedureInfo?action=list&proclD=1415&pipeID=7>). In brief, conscious ECG was collected using ECGenie equipment (Mouse  
9 Specifics, Inc.) as detailed previously by Spielmann et al.<sup>29</sup>. Based on availability of  
10 equipment and local expertise, some contributing centers opted to perform anesthetized  
11 ECG using Power Lab recording equipment and LabChart software (ADInstruments).  
12 Mice were anesthetized either with inhaled isoflurane (anesthesia was induced using 2.5-  
13 4% isoflurane in oxygen then maintained using 2-2.5% isoflurane in oxygen) or injected  
14 tribromoethanol (Sigma, stock concentration 20 mg/ml, dose calculated as 0.5 g/kg body  
15 weight). Anesthetized mice were positioned supine on a warming pad apparatus that  
16 maintained the animal's core temperature at 37°C. Needle electrodes were placed  
17 subcutaneously as follows: the negative electrode in the right forelimb; the ground  
18 electrode in the right hindlimb; and the positive electrode in the left hindlimb. ECG data  
19 was collected for up to 120 seconds and the resulting data analyzed using LabChart8  
20 software (ADInstruments). Regardless of the methodology, ECG was recorded in a dimly  
21 lit, quiet procedure room. In order to eliminate circadian influences ECG was recorded  
22 during the morning when the resting phase of a mouse begins.  
23



## 1 **Data Quality**

2 Standard protocols for ECG signal analysis were used to analyze the data. For this the  
3 software uses a so-called peak detection algorithm, which finds the peak of the R waves  
4 and calculates the heart rate (HR).

5 In anesthetized mice and for each cardiac cycle, the P, Q, R, S and T peaks were defined  
6 and from these the HR as well as the duration of the QRS complex, PQ interval, PR  
7 interval, QT interval and ST interval were automatically measured whereas in conscious  
8 ECG the peaks were automatically determined and averaged over multiple cardiac  
9 cycles. Only unique P, Q, R, S and T peaks have been included in the automatic  
10 calculations. Furthermore, the software automatically defined the end of the T-wave of  
11 each signal as the point where the signal intersects the isoelectric line. If any peaks were  
12 not selected correctly by the software, the position of this marker was corrected manually.  
13 Some subtle differences in annotation placement were observed across IMPC  
14 contributing centers. Heart rate variability (HRV) was calculated as the mean of the  
15 differences between successive heart rates for the entire set of ECG signals. The QT  
16 intervals were frequency corrected by applying the following equation (QTc) Mitchell et  
17 al.<sup>14</sup>. Noise and movement artefacts were automatically eliminated by the software.

18 Data were curated and subject to quality control at the IMPC prior to Data Release 15  
19 (August 11<sup>th</sup>, 2021) and we excluded one additional mouse from the analysis due to a  
20 biologically implausible QRS value.

## 21 **Statistical Methods**

1 Bespoke methods were developed to assess ECG reference ranges and are independent  
2 of the methodologies implemented on the IMPC portal.

3 Data analysis was conducted using R (version 4.0.4, R Core Team 2022<sup>30</sup>) with figures  
4 and tables produced in ggplot2, embedded in RMarkdown HTML files. Variability of all  
5 the data was assessed with two metrics a) coefficient of variation (COV) and b) “Quartile-  
6 based CV” (QCV), defined as interquartile range (IQR) (75-25%) relative to the median  
7 ( $100 \cdot \text{IQR} / \text{median}$ ).

8 Visual methods, as well as formal statistical tests were applied to test whether the scores  
9 of the individual parameters were normally distributed. Data were separated by age, sex  
10 and anesthesia regime and histograms for each parameter were plotted. Shapiro-Wilks  
11 tests were conducted to assess normality. Reference ranges were calculated based on  
12 median, 25th percentile and 75th percentile. In addition, the mean, standard deviation,  
13 and parameter sample size were provided to reflect the distribution of data. To reflect the  
14 distribution of each parameter, the 95% confidence intervals can be calculated by  
15  $\text{mean} \pm 1.96 \cdot \text{standard deviation}$  for each parameter.

### 16 ***Investigation of Anesthesia, Sex and Age Effects***

17 To investigate the effect of anesthesia on the different parameters, we calculated a one-  
18 way Analysis of Variance (ANOVA) with planned comparisons of “Conscious versus  
19 Isoflurane” and “Conscious versus Tribromoethanol”, separated by sex whereas  
20 “Isoflurane versus Tribromoethanol” was not tested. These planned comparisons were  
21 used to compare conscious vs unconscious. When looking for differences between

1 groups we tested the null hypothesis. P-values and F-values with degrees of freedom  
2 were calculated.

3 The effects of sex (female vs male) and age (EA vs LA) were compared using the same  
4 statistical analyses. In each case a simple two-tailed t-test was performed and the  
5 Cohen's *d* effect size calculated from the "effsize package" (R library). Due to the central  
6 limit theorem (CLT)<sup>31</sup>, the large sample sizes allowed parametric statistical testing of  
7 these effects.

8 These large group sizes provide overwhelming statistical power and may overestimate  
9 the importance of the effects. Bootstrapping tests were done to verify the biological  
10 significance of any differences in a range of more realistic experimental group sizes.

#### 11 **Data availability**

12 All data used are available to the public for download at the  
13 IMPC(<https://www.mousephenotype.org/data/previous-releases/15.0>).

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### 17 **Conflict of interest**

18 There is no conflict of interest for any of the authors listed above.

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2

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15

16



1 Table and Figure captions

2 **Table 1:** ECG data were available from a total of 26,706 mice, stratified by sex, age at  
3 testing (EA = 12 weeks of age; LA = mean of 62 weeks of age), and conscious state  
4 (conscious, anesthetized using isoflurane, or anesthetized using tribromoethanol).

5

6 **Table 2:** Median and 95% reference ranges of HR, PR-, QRS complex, QT-, RR- and  
7 ST-intervals, and QT corrected (QTc) using the Mitchell formula. Data are stratified by  
8 sex, age (EA and LA) and conscious state. **Panel a.** Conscious, isoflurane and  
9 tribromoethyl anesthetized female mice. **Panel b.** Conscious, isoflurane and  
10 tribromoethyl anesthetized male mice. Note, there were no data for LA mice anesthetized  
11 using tribromoethanol.

12

13 **Table 3:** Significant differences between the statistical comparison of conscious versus  
14 isoflurane ( $p < .001$ ) and conscious versus tribromoethanol ( $p < .001$  to  $p = .004$ ) in female  
15 (Panel a) and male (Panel b) mice for HR, PR-, QRS complex, QT-, RR-, ST-interval and  
16 QTc Mitchell. Test: p-value and F-value of one-way ANOVA with planned comparison.

17




18 **Figure 1:** Coefficient of variation (COV) analysis of data split by sex (female and male)  
19 and age (EA and LA) identified parameters with excess variability (COV >30%) that were  
20 excluded from further analysis (white bars). Parameters in blue were below the COV  
21 threshold of 30% and were retained for further analysis. These were in ascending COV  
22 percentage QTc Mitchell, PQ-, QT-, QRS-complex, ST-, RR- and PR-interval.

23

24 **Figure 2:** Representative averaged waveform of a C57BL/6N mouse in voltage over time  
25 reflecting the most commonly applied ECG annotations. Some subtle differences in  
26 annotation placement were observed across IMPC contributing centers. Figure 2 shows  
27 the depolarization and repolarization phases with prominent ECG parameters such as P,  
28 Q, R, S and T and their respective interval lengths as a function of voltage over time.

1 **Figure 3:** Histograms presenting the distribution of each selected ECG parameter for  
2 male and female mice separately. By visual inspection, no sexual dimorphism is apparent.  
3 **Panel a:** Recorded in the conscious state in early adult, EA (Sub-panels A-G) and late  
4 adult (LA) mice (Sub-panels H-N). **Panel b:** Recorded under isoflurane anesthesia in  
5 early adult, EA (Sub-panels A-G) and late adult (LA) mice (Sub-panels H-N). **Panel c:**  
6 Recorded under tribromoethanol anesthesia in early adult, EA (Sub-panels A-G). No late  
7 adult data are available for tribromoethanol anesthesia.

8

9 **Figure 4:** Comparison of the anesthetic regimes with the conscious state recordings.  
10 Distribution of the seven selected ECG parameters presented by histograms, stratified  
11 for female and male mice in early adult, EA (Sub-panels A-G) and late adult (LA)  
12 populations (Sub-panels H and N). Color code:  Conscious,  isoflurane and  tribromoethanol anesthesia.




14

15 **Figure 5:** Testing age-differences in conscious mice.

16 T-test results when comparing conscious EA versus LA data show high significance for  
17 all parameters ( $p < .001$ ) and negligible to medium Cohen's  $d$  standardized effect sizes.

18 **Panel a:** females, **Panel b:** males. **Panel c.** Bootstrap analysis of power estimates for  
19 sample sizes ranging from 5-100 mice, presented for each of the seven selected ECG  
20 parameters.

21

22 **Figure 6:** Reference ranges split by anesthetic regimen showing median, and 95%  
23 reference ranges (2.5th and 97.5th percentile). Female data are directly above the male  
24 data for EA (Sub-panels A-G) and late adult (LA) populations (Sub-panels H and N). No  
25 late adult data are available for tribromoethanol anesthesia. Color code:  Conscious,  
26  isoflurane and  tribromoethanol anesthesia.

27

1 **Figure 7:** Independent, non-IMPC study on six of the founder strain mice reported in “The  
2 Collaborative Cross: A Recombinant Inbred Mouse Population for the Systems Genetic  
3 Era”<sup>8</sup> study, including 129S1/SvImJ, A/J, C57BL/6J, NOD/ShiLtJ, NZO/HILtJ, and  
4 PWK/PhJ inbred strains show a close alignment to the reference ranges reported herein  
5 for PR-, QRS complex, QT-, RR-, and ST-interval based on multiple C57BL/6N substrains  
6 indicating good utility for those reference ranges. Data for HR was not available in this  
7 study. Mice were conscious, split by sex and ~12 weeks of age, equivalent to the IMPC  
8 EA time point. Red dotted lines depict the boundaries of the sex- specific reference ranges  
9 calculated herein, for each parameter.

10  
11 **Supplemental Table 1:** Comprehensive overview of mean, standard deviation, and  
12 sample number for each of the seven selected ECG parameters stratified by conscious  
13 state (conscious, anesthetized with isoflurane, or anesthetized with tribromoethanol), age  
14 [early (EA) and late (LA) adult time point] and sex (**Panel a.** Females; **Panel b.** Males).

15  
16 **Supplemental Table 2:** Definition and unit of measure for each ECG parameter reported.

17  
18 **Supplemental Table 3:** Comprehensive overview of median and 95% reference ranges  
19 (2.5th and 97.5th percentile) for each of the seven selected ECG parameters, stratified  
20 by conscious state (conscious, anesthetized with isoflurane, or anesthetized with  
21 tribromoethanol) and age [early (EA) and late (LA) adult time-point]. Sex is combined  
22 (females plus males) for this analysis to generate a both-sex-combined reference range.

23  
24 **Supplemental Figure 1:** “Quartile-based CV” (QCV), defined as interquartile range (IQR)  
25 (75-25%) relative to the median ( $100 \times \text{IQR} / \text{median}$ ), analysis of data split by sex (female  
26 and male) and age (EA and LA) identified parameters with excess variability (QCV  $\geq 30\%$   
27 for EA and LA time point) that were excluded from further analysis (white bars).  
28 Parameters in blue were below the QCV threshold and were retained for further analysis.

1 In the QCV analysis, but not in the COV, there were three parameters (light blue bars) in  
2 the LA population (HR, RR and PR) that marginally exceeded the limit of QCV  $\geq 30$  but  
3 were retained.

4

5 **Supplemental Figure 2:** Histograms presenting the distribution of PQ-interval data along  
6 with calculated ranges (mean  $\pm$  SD and median and 95% reference range) for conscious  
7 EA and LA mice stratified by sex. These calculations are based on data from one  
8 contributing center (German Mouse Clinic).

9

10 **Supplemental Figure 3:** Testing sex-differences in conscious mice.

11 T-test results when comparing data from conscious male and female animals for each of  
12 the seven selected ECG parameters, stratified by age [early (EA) and late (LA) adult time  
13 point]) demonstrated that some parameters show high significance ( $p < .001$ ), while for  
14 others there was no evidence of sexual dimorphism. Associated Cohen's  $d$  standardized  
15 effect sizes were negligible. **Panel a** shows EA time point, **Panel b** shows LA time point,  
16 and **Panel c** shows bootstrap analysis of power estimates for female and male samples  
17 ranging from 5-100 mice, presented for each of the seven selected ECG parameters.

18

19 **Supplemental Figure 4:** Testing sex-differences in isoflurane anesthetized mice.

20 T-test results when comparing data from isoflurane anesthetized male and female  
21 animals for each of the seven selected ECG parameters, stratified by age [early (EA) and  
22 late (LA) adult time point]) demonstrated that some parameters show significance ( $p < .05$ )  
23 and negligible to small Cohen's  $d$  standardized effect sizes. **Panel a.** shows EA time  
24 point. **Panel b.** shows LA time point, and **Panel c.** shows bootstrap analysis of power  
25 estimates for female and male samples ranging from 5-100 mice, presented for each of  
26 the seven selected ECG parameters.

27

28 **Supplemental Figure 5:** Testing sex-differences in tribromoethanol anesthetized mice.

1 T-test results when comparing data from tribromoethanol anesthetized male and female  
2 animals for each of the seven selected ECG parameters, stratified by age [early (EA) but  
3 **no** late (LA) adult time point] demonstrated that three parameters show high statistical  
4 significance ( $p < .001$ ) namely HR, PR- and RR-interval whereas medium significance for  
5 ST- ( $p = .003$ ) and QT-interval ( $p = .018$ ) and no significance for QRS complex ( $p = .522$ ) and  
6 QTc Mitchell ( $p = .283$ ) were detected with negligible to medium Cohen's *d* standardized  
7 effect sizes. **Panel a.** shows EA time point, and **Panel b.** shows bootstrap analysis of  
8 power estimates for female and male samples ranging from 5-100 mice, presented for  
9 each of the seven selected ECG parameters.

10

11 **Supplemental Figure 6:** Testing age-differences in conscious mice.

12 There is little to no aging effect by equivalent t-test, Cohen's *d* and bootstrap analysis in  
13 EA and LA mice anesthetized with isoflurane. In females, high statistical significance is  
14 detected for three parameters ( $p < .001$ ) namely HR, QRS complex and RR-interval and  
15 no significance for PR-, QT- and ST-interval and QTc Mitchell with negligible to small  
16 Cohen's *d* standardized effect sizes. In males, high statistical significance is detected for  
17 five parameters ( $p < .001$ ) namely HR, QRS complex, QT-, RR- and ST-interval and no  
18 significance for PR-interval and QTc Mitchell with negligible to small Cohen's *d*  
19 standardized effect sizes. **Panel a.** shows females. **Panel b.** shows males and **Panel c.**  
20 shows bootstrap analysis of power estimates for EA and LA sample size ranging from 5-  
21 100 mice, presented for each of the seven selected ECG parameters.

22

23 **Supplemental Figure 7:** Wildtype control animals from three non-IMPC studies tested at  
24 the German Mouse Clinic (<https://www.mouseclinic.de/>) with a standard sample size (20-  
25 30 control animals per study) on the background strains C57BL/6NJ, C57BL/6J and FVB  
26 show a close alignment to the reference ranges reported herein for HR, PR-, QRS  
27 complex, QT-, RR-, and ST-interval, and QTc Mitchell values based on multiple  
28 C57BL/6N substrains indicating good utility for those reference ranges. Mice were  
29 conscious, split by sex and ~12 weeks of age, equivalent to the IMPC EA time point. Red

1 dotted lines depict the boundaries of the sex-specific reference range calculated herein,  
2 for each parameter.

3

4 **Supplemental Figure 8:** Seven inbred strains of The Jaxwest1 project 129S1/SvImJ,  
5 A/J, BALB/cJ, C57BL/6J, DBA/2J, NOD/ShiLtJ and SJL/J show a close alignment to the  
6 reference ranges reported herein for PR-, QRS complex, QT-, RR-, and ST-interval based  
7 on multiple C57BL/6N substrains indicating good utility for those reference ranges. Mice  
8 were conscious, split by sex and ~12 weeks of age, equivalent to the IMPC EA time point.  
9 Red dotted lines depict the boundaries of the sex-specific reference range calculated  
10 herein, for each parameter.

11

12 **Supplemental Figure 9-13:** The Xing1 Aging study includes 29 inbred strains of which  
13 26 have been included here: 129S1/SvImJ, A/J, BALB/cByJ, BTBR-T+tf/J, BUB/BnJ,  
14 C3H/HeJ, C57BL/10J, C57BL/6J, C57BLKS/J, C57BR/cdJ, C57L/J, CBA/J, DBA/2J,  
15 FVB/NJ, KK/HIJ, LP/J, MRL/MpJ, NOD.B10Sn-H2/J, NON/ShiLtJ, NZO/H1LtJ,  
16 NZW/LacJ, P/J, PL/J, RIIS/J, SM/J, and SWR/J. The data show a close alignment to the  
17 reference ranges reported herein for PR-, QRS complex, QT-, RR-, and ST-interval based  
18 on multiple C57BL/6N substrains indicating good utility for those reference ranges. Mice  
19 were conscious, split by sex and ~12 months or ~20 months of age, equivalent to the  
20 IMPC LA time ranges with a minimum of 52 and a maximum of 78 weeks of age. Red  
21 dotted lines depict the sex-specific boundaries of the reference range calculated herein,  
22 for each parameter.

Table 1:

	Conscious	Isoflurane	Tribromoethanol	Sum
EA female	9,240	2,672	226	12,138
EA male	9,238	2,598	220	12,056
LA female	620	693	0	1,313
LA male	589	610	0	1,199
Sum	19,687	6,573	446	26,706

Table 2:

a FEMALE						
Parameter	Conscious		Isoflurane		Tribromoethanol	No LA data available
	EA	LA	EA	LA	EA	
Heart Rate [bpm]	760 [650;815]	748 [700;790]	415.8 [292.6;511.7]	440.5 [339.6;530.3]	440 [355.8;550.5]	
PR [ms]	28.7 [20.5;35.8]	26.3 [18.2;35]	44.5 [29.6;72.7]	46.7 [30.5;58.9]	49.1 [41.4;57.7]	
QRS [ms]	10.7 [8.4;14.8]	11.8 [8.3;15.3]	10.1 [7;17.9]	10.1 [7.8;16.6]	13.2 [10.6;17]	
QT [ms]	42 [36.4;45.8]	43.3 [37.9;46.2]	52.1 [35.4;86.9]	52 [29.5;74.2]	53.5 [40.7;75.7]	
RR [ms]	79 [73.7;92.9]	80.2 [76;86.1]	144.4 [117.3;205]	136.2 [113.2;176.7]	136.2 [108.9;166.1]	
ST [ms]	30.8 [13.1;35.8]	32.3 [27;35.3]	40 [24.3;78]	38.5 [31.9;58.3]	42.8 [28.3;63.1]	
QTc Mitchell [ms]	47.4 [41.7;51.1]	48.5 [42.5;51.5]	42.4 [29.9;65.5]	45 [25;62.8]	45.2 [36.4;60]	

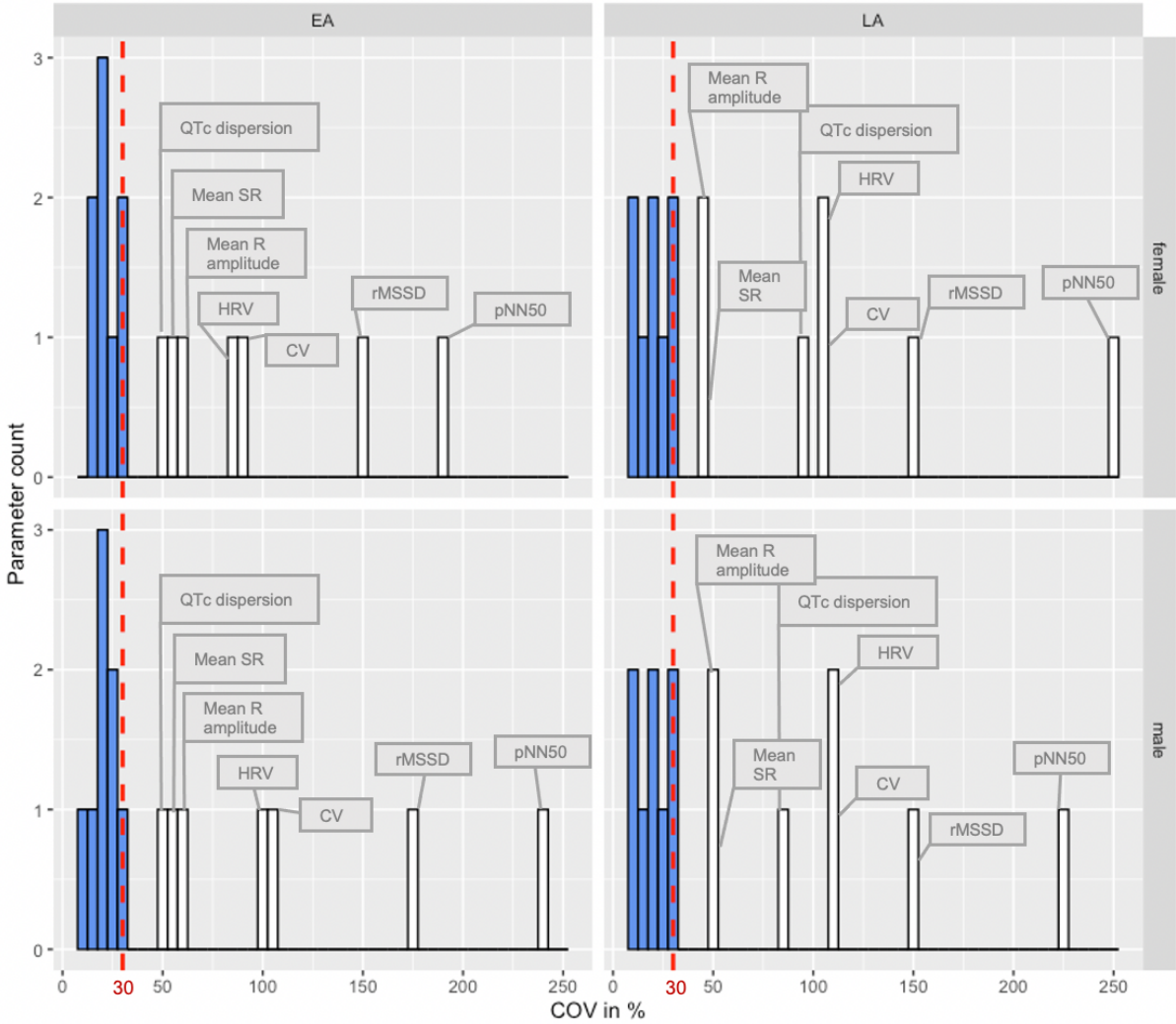
  

b MALE						
Parameter	Conscious		Isoflurane		Tribromoethanol	No LA data available
	EA	LA	EA	LA	EA	
Heart Rate [bpm]	764 [661;819.7]	751 [702.3;796.3]	422.6 [298;534.9]	443.1 [325.1;553.7]	414.1 [329.9;573.1]	
PR [ms]	28.6 [20.4;35]	25.8 [18.2;34.8]	43.2 [32.5;63.6]	44.3 [34;54.9]	47.7 [41.9;55.7]	
QRS [ms]	10.6 [8.3;14.6]	11.8 [8.6;16.1]	10 [7;19.7]	10 [7;17.3]	13.4 [10.8;16.8]	
QT [ms]	41.8 [36.6;45.5]	43.3 [38;46.6]	52.8 [37.9;89.2]	51.2 [34.6;64.8]	55.2 [42.6;76.6]	
RR [ms]	78.6 [73.2;91.4]	79.9 [75.4;86.1]	142 [112.2;201.6]	135.4 [108.4;184.6]	145 [104.7;181.7]	
ST [ms]	30.6 [12.6;35.6]	32.1 [27.6;35.3]	40.7 [26.9;79.3]	38.7 [35.8;41.9]	45.2 [32.4;65.9]	
QTc Mitchell [ms]	47.5 [41.9;51]	48.7 [42.8;51.8]	43.4 [32.6;64.9]	45.4 [32.7;58]	46 [39.3;62.8]	

Table 3:

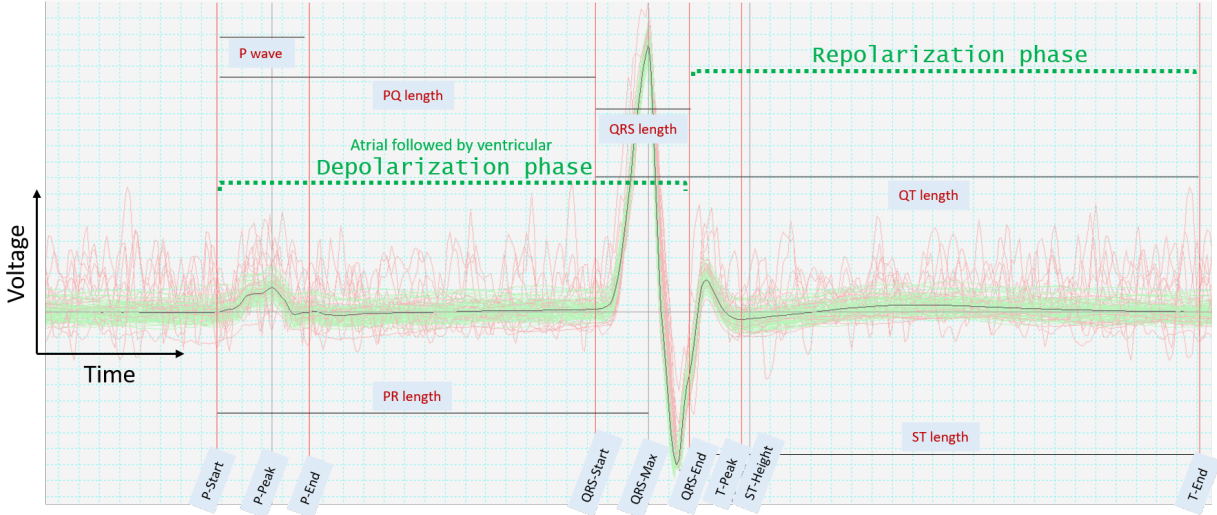
	a FEMALE		b MALE	
	Conscious vs Isoflurane	Conscious vs Tribromoethanol	Conscious vs Isoflurane	Conscious vs Tribromoethanol
Heart Rate [bpm]	F(1)=120674.4, p<.001	F(1)=2122.7, p<.001	F(1)=116569.4, p<.001	F(1)=2865.6, p<.001
PR [ms]	F(1)=21279, p<.001	F(1)=872.8, p<.001	F(1)=23952.9, p<.001	F(1)=1142.3, p<.001
QRS [ms]	F(1)=78.1, p<.001	F(1)=273.2, p<.001	F(1)=71.6, p<.001	F(1)=294.8, p<.001
QT [ms]	F(1)=1754.3, p<.001	F(1)=135, p<.001	F(1)=2067.7, p<.001	F(1)=217.7, p<.001
RR [ms]	F(1)=72292.8, p<.001	F(1)=839, p<.001	F(1)=64552.1, p<.001	F(1)=1468.3, p<.001
ST [ms]	F(1)=3278.4, p<.001	F(1)=285.3, p<.001	F(1)=4149.9, p<.001	F(1)=424.1, p<.001
QTc Mitchell [ms]	F(1)=221, p<.001	F(1)=8.3, p=.004	F(1)=88.9, p<.001	F(1)=8.3, p=.004

**Figure 1:**



**Figure 2:**

Representative averaged waveform of a C57BL/6N mouse



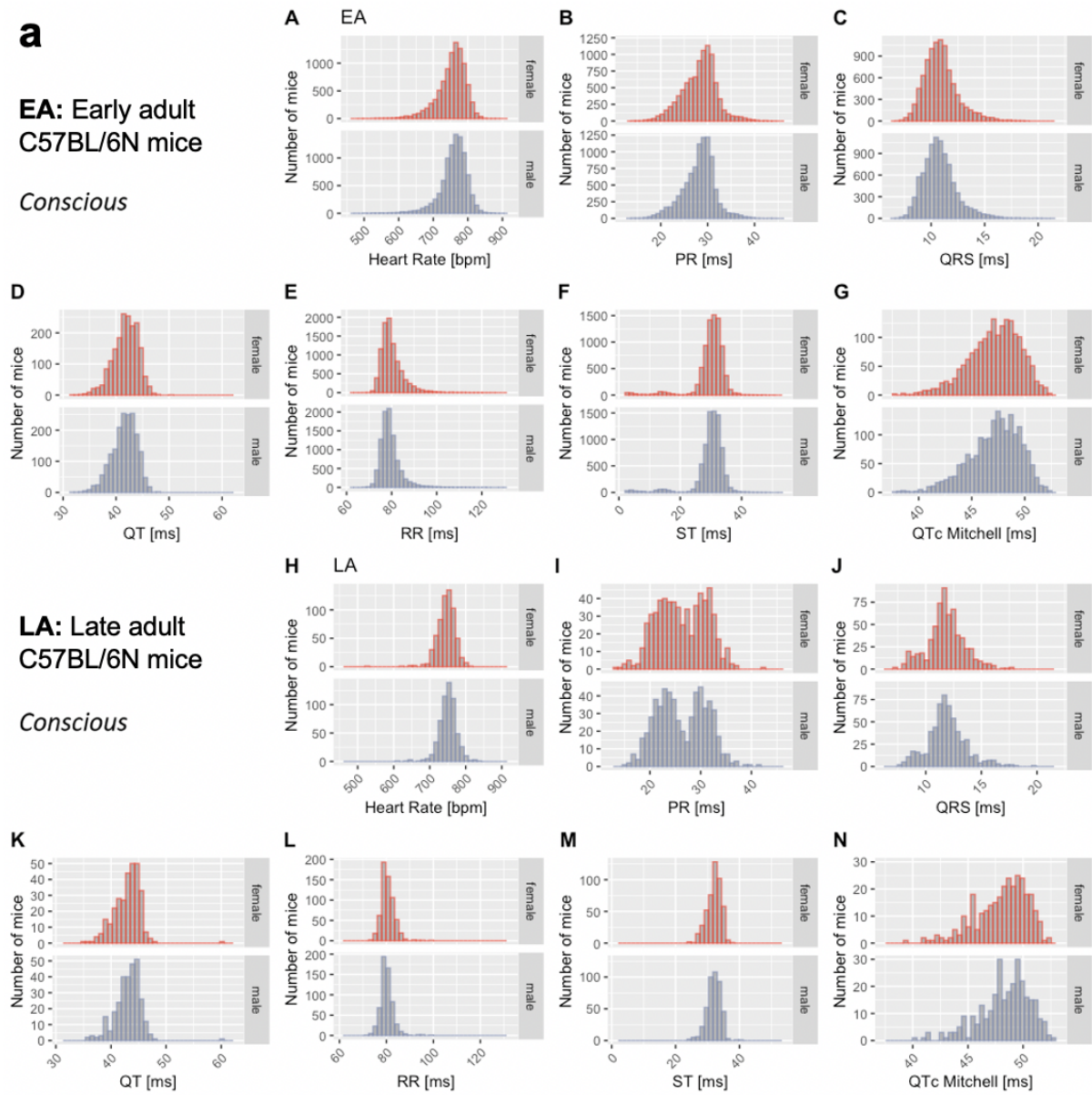


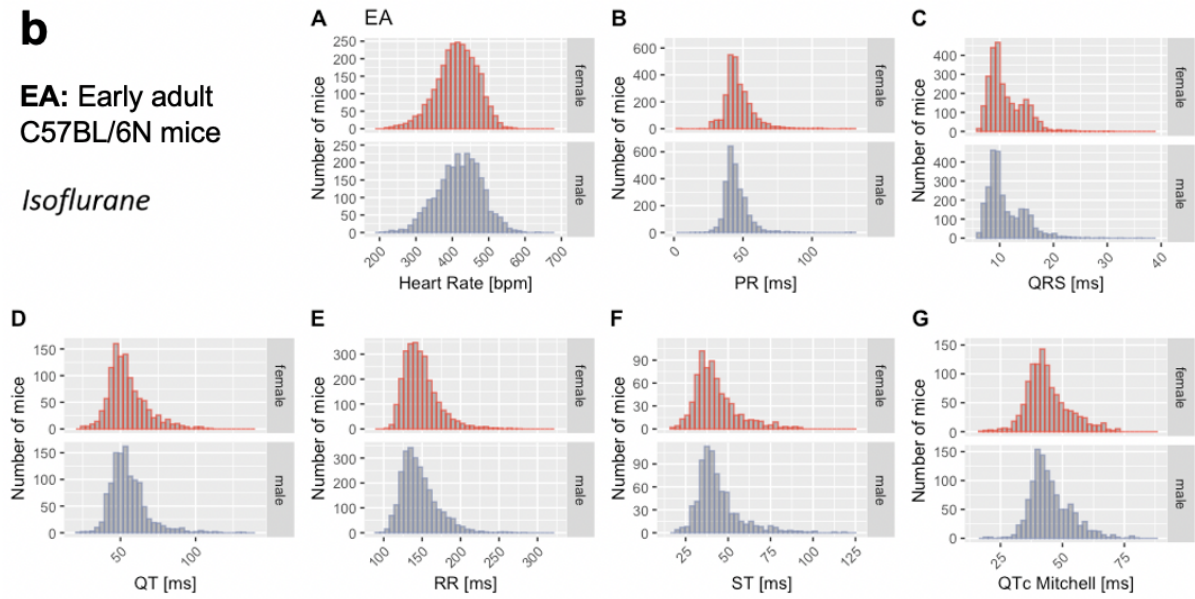
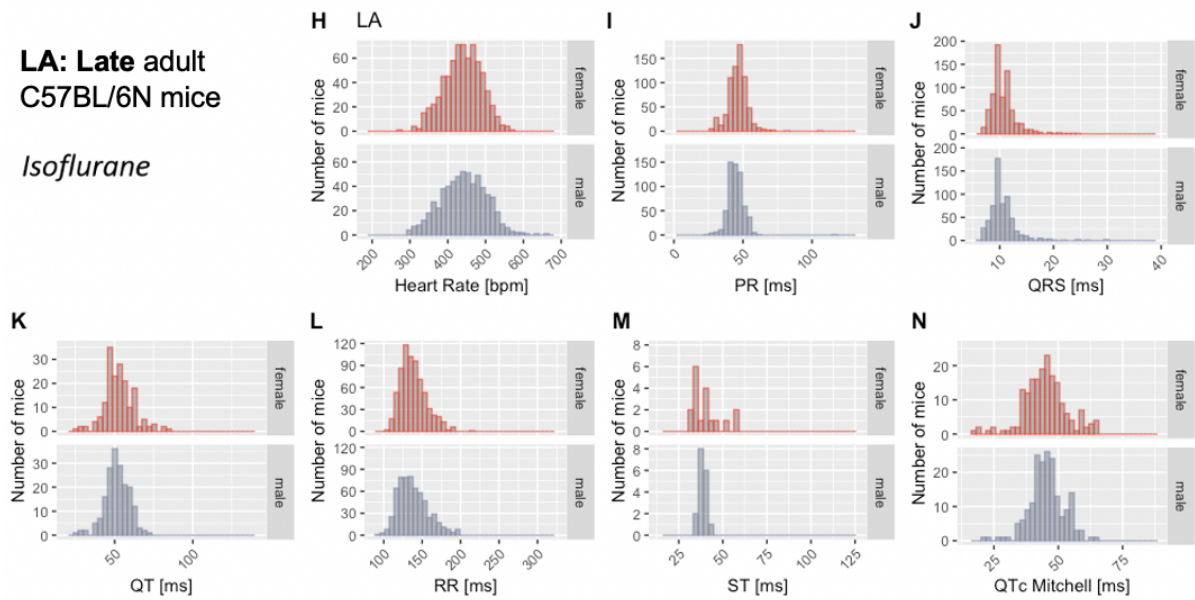
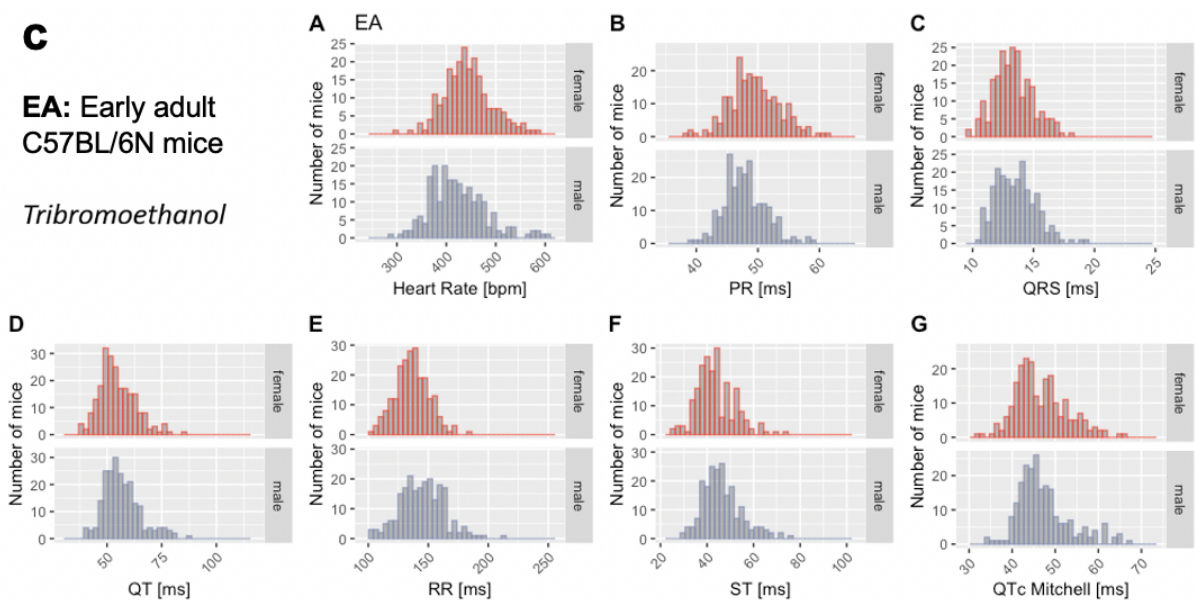
**Figure 3:**

**a**

**EA: Early adult  
C57BL/6N mice**

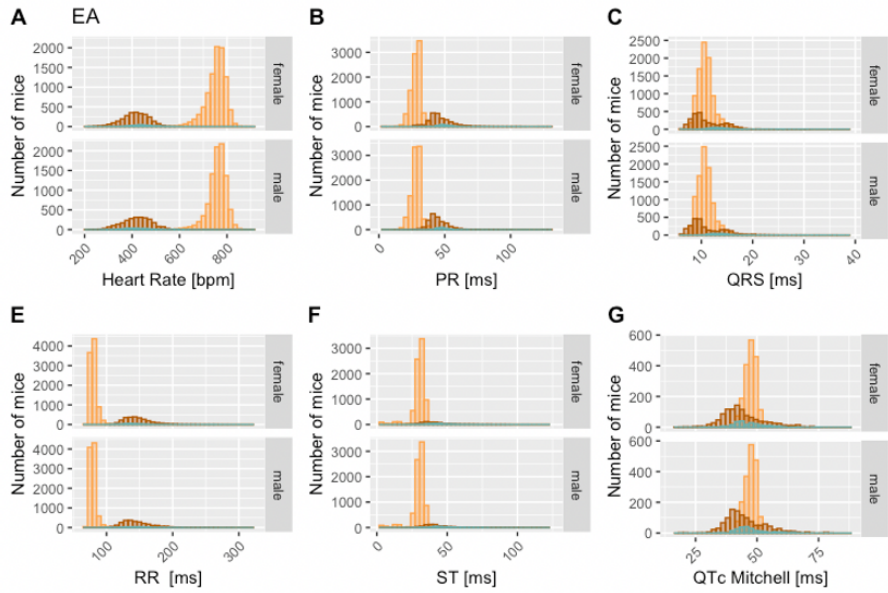
*Conscious*



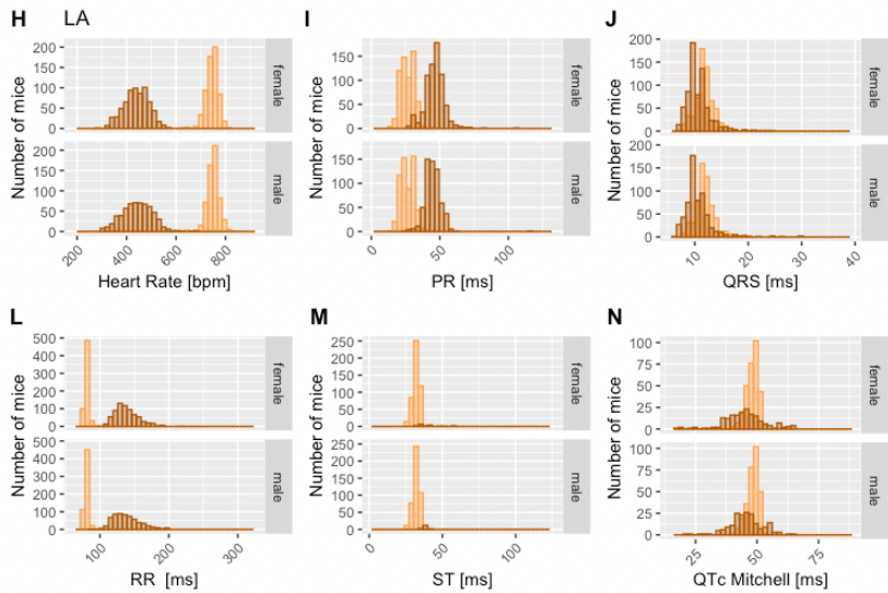
**b****EA: Early adult  
C57BL/6N mice***Isoflurane***LA: Late adult  
C57BL/6N mice***Isoflurane***C****EA: Early adult  
C57BL/6N mice***Tribromoethanol*

**Figure 4:**

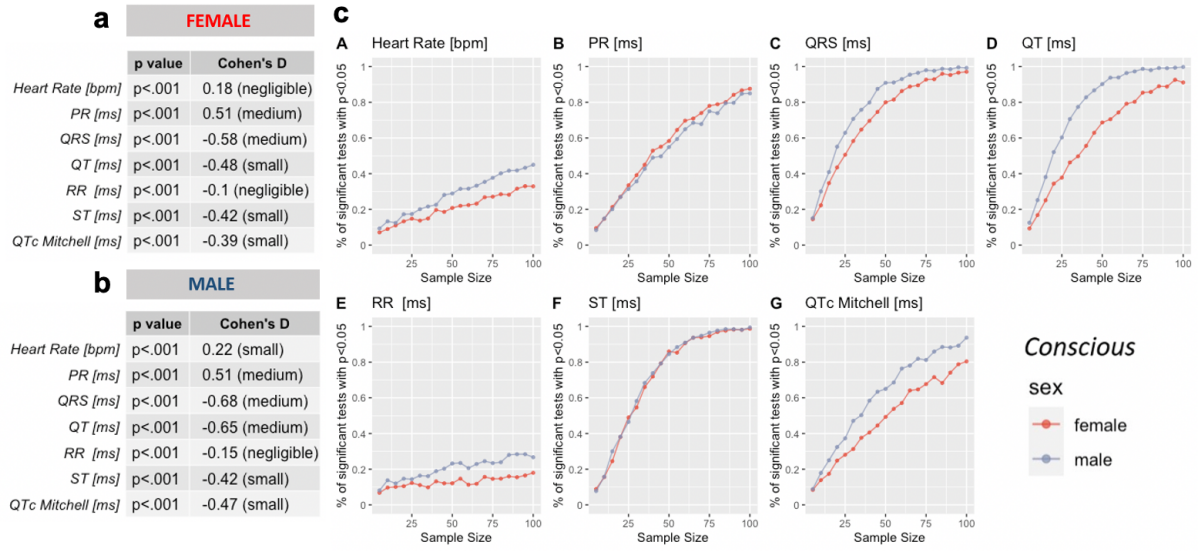
**EA: Early adult  
C57BL/6N mice**



**LA: Late adult  
C57BL/6N mice**

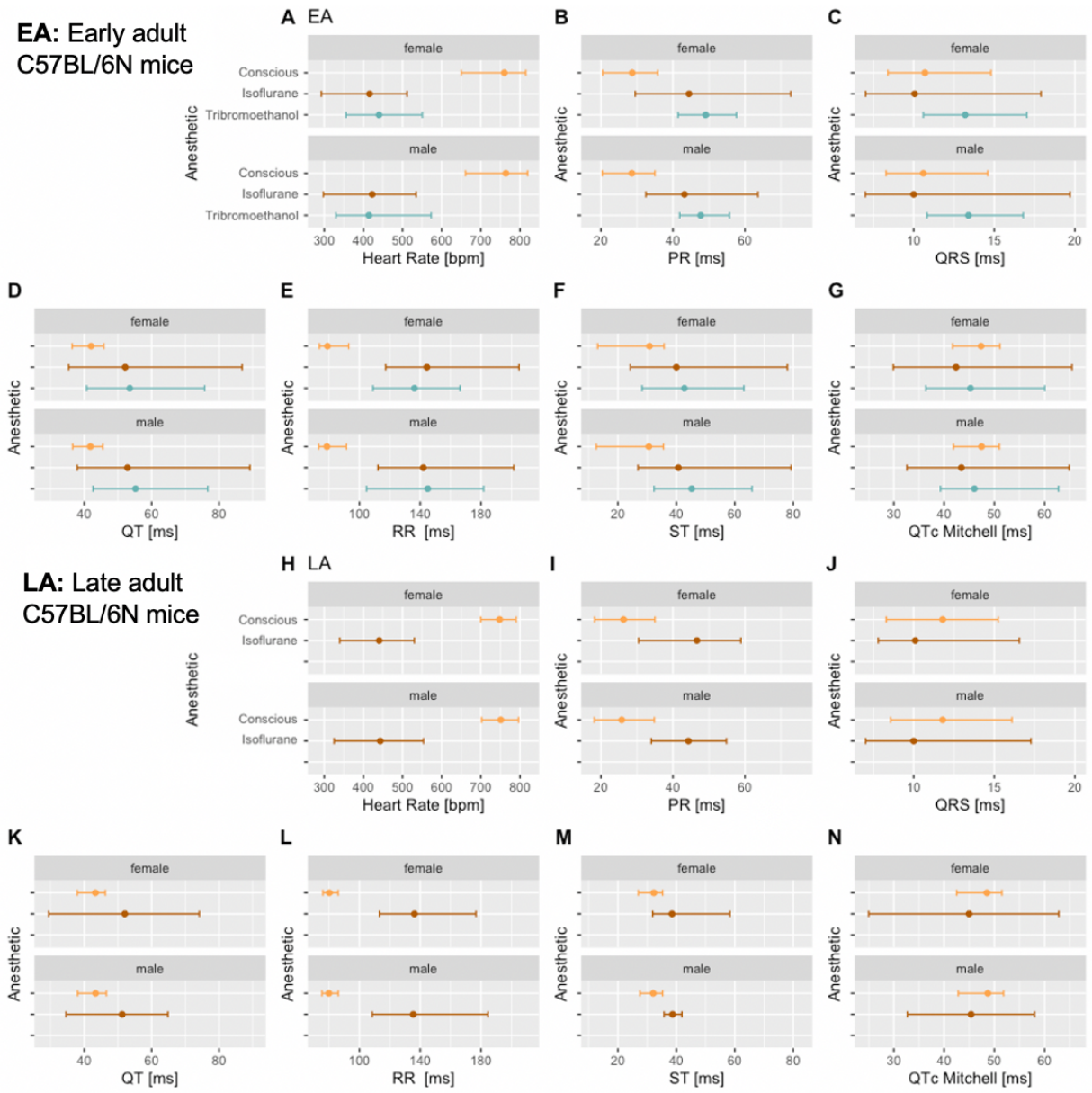


**Figure 5:**

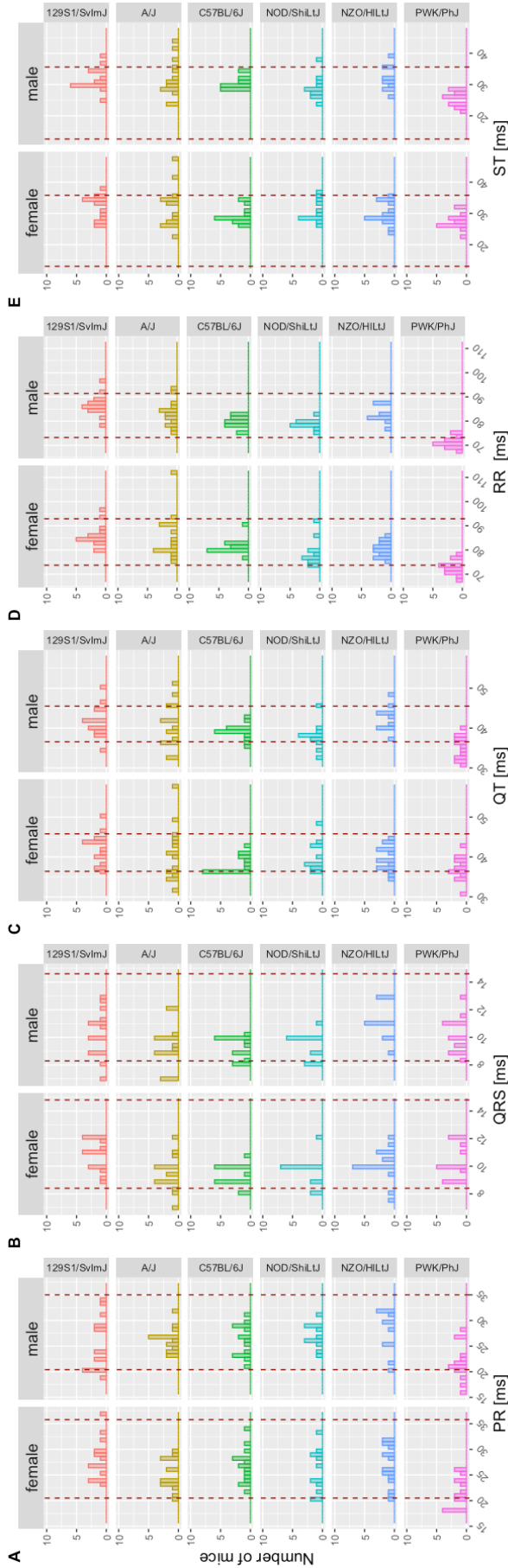




**Figure 6:**



**Figure 7:**





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**Electronic Supplementary Material**  
**Supplemental\_Table\_and\_Figures.pdf**

