Electrocardiography Abnormalities in Macaques after Infection with Encephalitic Alphaviruses

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Abstract: Eastern (EEEV) and Venezuelan (VEEV) equine encephalitis viruses (EEVs) are related, (+)ssRNA arboviruses that can cause severe, sometimes fatal, encephalitis in humans. EEVs are highly infectious when aerosolized, raising concerns for potential use as biological weapons. No licensed medical countermeasures exist; given the severity/rarity of natural EEV infections, efficacy studies require animal models. Cynomolgus macaques exposed to EEV aerosols develop fever, encephalitis, and other clinical signs similar to humans. Fever is nonspecific for encephalitis in macaques. Electrocardiography (ECG) metrics may predict onset, severity, or outcome of EEV-attributable disease. Macaques were implanted with thermometry/ECG radiotransmitters and exposed to aerosolized EEV. Data was collected continuously, and repeated-measures ANOVA and frequency-spectrum analyses identified differences between courses of illness and between pre-exposure and post-exposure states. EEEV-infected macaques manifested widened QRS-intervals in severely ill subjects post-exposure. Moreover, QT-intervals and RR-intervals decreased during the febrile period. VEEV-infected macaques suffered decreased QT-intervals and RR-intervals with fever onset. Frequency-spectrum analyses revealed differences in the fundamental frequencies of multiple metrics in the post-exposure and febrile periods compared to baseline and confirmed circadian dysfunction. Heart rate variability (HRV) analyses revealed diminished variability post-exposure. These analyses support using ECG data alongside fever and clinical laboratory findings for evaluating medical countermeasure efficacy.

Keywords: alphavirus; vaccine; arbovirus; animal models; nonhuman primates; electrocardiography; ECG; aerosol; encephalitis; equine

1. Introduction
Equine encephalitis viruses (EEVs) constitute a distinct portion of the alphavirus genus of Family Togaviridae. Alphaviruses are grouped into New World and Old World viruses, which generally correspond with signs of clinical disease; New World alphaviruses are more likely to cause encephalitis while Old World viruses tend to cause arthralgia [1-4]. These mosquito-borne viruses were the first alphaviruses to be isolated through the 1930s, and VEEV underwent research in later decades by the United States and the Union of Soviet Socialist Republics as a potential biological warfare agent after it was realized that VEEV was highly infectious when aerosolized [5-10]. These pathogens remain important medical and veterinary pathogens to this day [11].

In the clinical context, encephalitis comprises infection or inflammation of brain parenchyma that may produce signs of diagnosable disease [12]. EEVs can produce a febrile course, the eponymous encephalitis, and a variety of neurological presentations including but not limited to seizures, tremor, and photophobia [1,12]. Other symptoms seen in EEV-induced disease includes nausea, vomiting, diarrhea, focal neurological deficits, or altered mental status [13-18]. VEEV and western equine encephalitis virus are only rarely fatal in naturally-transmitted human cases, and
then typically only in very young or elderly patients [15,19-24]. North American isolates of EEEV
have a high case fatality rate, estimated to range from 30-70%, although sub-clinical disease may be
underreported [13,25]. Survivors of virulent infections can develop persistent neurologic sequelae
that affect motor function. Pathological evaluation can confirm or rule out encephalitis by
examination of cellular inclusion bodies, lymphocytic infiltrates, and immunohistochemical
staining; however, the post-mortem nature of these procedures limits their clinical applicability [26].
Apart from a thorough patient history, confirmation of EEVs as the cause of disease requires
detection of EEV-specific IgM or IgG in plasma or cerebrospinal fluid (CSF) [13,18,27].

No consensus governs the case management of EEEV infections and no licensed treatments
or vaccines are available to mitigate an outbreak or intentional release [1,28]. Because aerosol
dissemination is not a natural route of transmission, determining efficacy of countermeasures
against aerosolized EEVs in a clinical trial is neither logistically nor ethically possible. Licensure of
such countermeasures can proceed only through the FDA’s Animal Rule, which allows for pivotal
efficacy studies to be conducted in well-characterized animal models that meet appropriate criteria
[29]. These criteria require that the disease and pathophysiology in the animal model must resemble
what is known to occur in humans. To evaluate the efficacy of a therapeutic treatment against viral
encephalitis, biomarkers that correspond with viral penetration into the CNS and/or neurological
disease severity are needed. For EEVs, infection of the cynomolgus macaque closely resembles the
disease seen in humans [30,31]. Fever was the predominant physiological sign observed in past EEEV
studies, but it is not clear if this corresponds with a response to peripheral viral replication or viral
penetration into the CNS [18,32-34]. In VEEV infection of humans and macaques, the febrile
response is biphasic; the consensus in the field has been that the first fever likely represents
peripheral viral replication while the second fever coincides with viral penetration into the CNS. In
macaques, the second febrile period is associated with neurological signs indicative of encephalitis
[35]. In contrast, EEEV infections do not have a fever in the initial period seen with VEEV but fever
onset roughly corresponds with the second fever peak for VEEV and onset of neurological signs in
macaques [36]. Previous work with macaques found that heart rate increased in association with
fever onset after infection with EEEV [35,36]. Exploration of a wider range of electrocardiographic
metrics leverages the advantages inherent in continuous radiofrequency telemetry monitoring
equipment. In this study, we hypothesized that physiological changes encountered in macaques
following infection with EEEV or VEEV would produce measurable changes in electrocardiographic
metrics that would precede or accompany the onset of febrile or encephalitic disease. We performed
repeated measures ANOVA and multiple comparisons of electrocardiographic metrics between
pre-exposure, post-exposure, febrile periods for macaques that developed fever, and recovery
periods for macaques that survived febrile courses. Heart rate variability (HRV) was analyzed
between days from the baseline period through the post-infection period. Finally, frequency-spectrum analysis was performed on the electrocardiography metrics. Findings indicated
that changes in QRS complexes, QT-Interval, and RR-Interval, among other metrics, could serve as
sentinels of EEEV-induced disease.

2. Results

2.1. ECG Changes in EEEV-infection of Macaques

To explore the utility of monitoring ECG changes associated with EEEV-induced viral
encephalitis caused by New World-alphaviruses, four macaques were exposed to small-particle
(≤5μm mass mean aerodynamic diameter) aerosols containing EEEV. The outcome and the febrile
response after infection of the macaques with EEEV are shown in Table 1. Macaques exposed to
EEEV manifested either severe or non-severe courses of disease (Figure 1a). Severe disease was
defined as a course that precipitated moribund status requiring euthanasia. Aerosol exposure to
EEEV produced severe disease courses in the two macaques (M161-16, M163-16) that received more
than 1.1*10^8 PFU. In contrast, the two macaques (M160-16, M162-16) that received doses below this
amount survived without fever.
Table 1. EEV Courses of Infection in Macaque Cohort. Macaques received aerosols of either EEV or VEEV. *Doses in log_{10} pfu. Neur = neurological signs (tremors, seizures, photophobia, head pressing), \( \delta \)TTD = Time to death, S = Survived to 28 days post infection. \( \gamma \)Fever data. \( \gamma \)Max =
maximum difference from predicted; Onset = onset of fever duration > 8 hrs; Duration = fever duration, in hours; Fever-hours = summation of all significant differences in temperature from predicted.

<table>
<thead>
<tr>
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<th>Macaque</th>
<th>Sex</th>
<th>Dose*</th>
<th>TTD( ^{\delta} )</th>
<th>Neuro( ^{\gamma} )</th>
<th>( \Delta )Max( ^{\gamma} )</th>
<th>Onset( ^{\gamma} )</th>
<th>Duration( ^{\gamma} )</th>
<th>Fever-hours( ^{\gamma} )</th>
</tr>
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<td>6</td>
<td>Y</td>
<td>4.2</td>
<td>3</td>
<td>70.8</td>
<td>144.2</td>
</tr>
<tr>
<td></td>
<td>163-16</td>
<td>M</td>
<td>7.5</td>
<td>6</td>
<td>Y</td>
<td>4.4</td>
<td>3</td>
<td>65.0</td>
<td>154.9</td>
</tr>
<tr>
<td></td>
<td>Mean</td>
<td></td>
<td></td>
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<td>3</td>
<td>67.9</td>
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</tr>
<tr>
<td></td>
<td>160-16</td>
<td>M</td>
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<td>S</td>
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<td>14.3</td>
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</tr>
<tr>
<td></td>
<td>162-16</td>
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<td>N</td>
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<td>41.5</td>
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<tr>
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<td>18.5</td>
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<td>S</td>
<td>Y</td>
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<td>1</td>
<td>187.8</td>
<td>171.6</td>
</tr>
<tr>
<td></td>
<td>165-16</td>
<td>F</td>
<td>6.9</td>
<td>S</td>
<td>Y</td>
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<td>1</td>
<td>260.5</td>
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<td></td>
<td>170-16</td>
<td>M</td>
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<td>S</td>
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<td>180.5</td>
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<td></td>
<td>171-16</td>
<td>M</td>
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<td>S</td>
<td>Y</td>
<td>3.3</td>
<td>1</td>
<td>238.0</td>
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<td>Mean</td>
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<td></td>
<td>3.0</td>
<td>1</td>
<td>216.7</td>
<td>228.4</td>
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</tbody>
</table>

In the macaques with severe EEV disease (M161-16, M163-16), the disease course replicated with high fidelity what was reported previously with a different EEV strain [36]. Post-challenge, the macaques became febrile shortly after the 3-day mark, culminating in a febrile plateau of 40-41°C around 4.0 days post-infection (dpi) (Figure 1b,c). This plateau was followed by a terminal decline in temperature beginning approximately 5 dpi, heralding a moribund state occurring at approximately 6 dpi. For macaques with severe disease, 0-3 dpi defined the post-infection period and the time following 3 dpi denoted the febrile period. None of the EEV-infected macaques with severe disease survived, so a recovery period was not defined. For the two macaques that survived lower doses of EEV there was no febrile response post-infection (Figure 1d,e) nor any changes in behavior or activity or any other clinically apparent sequelae.

In the immediate post-infection period, the profile of QRS Complex data show no readily apparent changes between 0 dpi to 3 dpi, in either the directly observable trace (Figure 2a) or in daily aggregates of the QRS complex (Figure 2b). Beginning with fever onset on 3 dpi, the QRS complex exhibited progressive increases in severely diseased EEV-infected macaques which continued until euthanasia criteria were met (Table 2, p<0.05). Macaques exposed to the lower doses of EEV displayed no significant deviations from their baseline QRS complexes (Figure 2b).

Similar to the QRS-Interval, there are no visually apparent changes for any of the EEV-infected macaques in the QT-interval trace (Figure 3a) or the daily QT-interval aggregate (Figure 3b) in the post-infection period until 3 dpi. With the onset of fever on 3 dpi, the two macaques with severe EEV disease exhibited shortened QT-intervals and a loss of diurnal variation. This continued until the macaques were moribund (Table 3, p<0.05). A false positive result surfaced from the repeated measures ANOVA for M160-16, one of the macaques without severe disease, but this result is attributable to data loss during baseline collection; (Figure 3a) this effect disappears once the days containing missing baseline data are censored. The missing diurnal baseline data for 160-16 rendered an apparent increase in QT-interval on the initial analysis. In the macaques exposed to
Figure 1. EEEV disease in macaques. Shown are models of disease for (a) EEEV-exposed macaques with non-severe and severe courses of alphavirus disease over days post-infection (x-axis). Deep blue denotes pre-infection baseline period, vertical magenta line denotes time of infection. Green bar denotes post-infection period in animals without severe/fatal disease. For animals with severe/fatal disease, powder blue defines post-infection period before the onset of fever, and red signifies the febrile post-infection period. Skull and crossbones mark the mean time to death. Actual and forecasted temperature profiles for two macaques with severe disease requiring euthanasia: (b) M161-16, and (c) M163-16, compared to macaques without severe disease: (d) M160-16 and (e) M162-16. Red: Actual Temperature (°C), Blue: Forecasted Temperature (°C) from ARIMA modeling, Black: Residual Values (°C) (deviations from forecasted baseline). Gray Lines: upper/lower bounds for residuals Magenta: aerosol challenge.
lower doses of EEEV that did not develop clinical disease, no reductions or loss of diurnal variation in QT-Interval were seen in the study period (Figure 3b).

Figure 2. Increase in QRS Complex during the febrile period in macaques with severe EEEV. Shown are the profiles (a) of QRS complex derived from 1-minute average measurements for four individual macaques infected with EEEV. Black: QRS Complex (ms), Gray: Temperature (°C), and Magenta: aerosol challenge. Blue Bars: Neurological score on corresponding day. (b) Average daily QRS Complex measurements, from repeated measures ANOVA, depicted in with boxes delimited by interquartile range and center line signifying median value. For (a) and (b), Top Left: M160-16, Bottom Left: M161-16, Top Right: M162-16, Bottom Right: M163-16.

Table 2. Repeated Measures ANOVA Statistics for QRS Complexes. Statistics and p-values for QRS complex for EEEV and VEEV-infected macaque cohorts. *Doses in log_{10} pfu. **TTD = Time to death. *P-Value lower than significance level α=0.05.

<table>
<thead>
<tr>
<th>Virus</th>
<th>Macaque</th>
<th>Sex</th>
<th>Dose</th>
<th>TTD</th>
<th>F-Statistic</th>
<th>P-Value</th>
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<td>M161-16</td>
<td>M</td>
<td>8.2</td>
<td>6</td>
<td>4.780</td>
<td>0.0346*</td>
</tr>
<tr>
<td></td>
<td>M163-16</td>
<td>M</td>
<td>7.5</td>
<td>6</td>
<td>4.034</td>
<td>0.0490*</td>
</tr>
<tr>
<td></td>
<td>M160-16</td>
<td>M</td>
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<td>S</td>
<td>3.833</td>
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</tr>
<tr>
<td></td>
<td>M162-16</td>
<td>M</td>
<td>5.7</td>
<td>S</td>
<td>3.046</td>
<td>0.331</td>
</tr>
<tr>
<td>VEEV</td>
<td>M164-16</td>
<td>F</td>
<td>7.1</td>
<td>S</td>
<td>29.55</td>
<td>&lt;0.00001*</td>
</tr>
<tr>
<td></td>
<td>M165-16</td>
<td>F</td>
<td>6.9</td>
<td>S</td>
<td>164.7</td>
<td>&lt;0.00001*</td>
</tr>
<tr>
<td></td>
<td>M170-16</td>
<td>M</td>
<td>6.3</td>
<td>S</td>
<td>48.40</td>
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</tr>
<tr>
<td></td>
<td>M171-16</td>
<td>M</td>
<td>6.0</td>
<td>S</td>
<td>84.17</td>
<td>&lt;0.00001*</td>
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RR-intervals in macaques infected with EEEV follow a similar pattern to that seen with the QT-Interval (Figure 4a). From challenge to 3 dpi, there was no visual change in RR-Interval in any of the macaque. On 3 dpi, the severely-infected macaques had decreases in both the trace and the median RR-interval in aggregate data (Figure 4b). No changes in RR-Interval were observed for macaques with nonsevere courses of disease (Table 4, p<0.05). Heart rate variability was plotted as ordered pairs of the RR-interval data collected at time n (RRL_n) and RR-interval data collected at time n+1 (RRL_{n+1}), wherein n represents an integer value in corresponding to a minute on the macaque’s time scale. In aggregate, when these scatter plots are superimposed and color-coded by pre-infection, post-infection, and febrile disease periods (Figure 5a), there is a visually notable contraction in the distribution of ordered pairs during the febrile period in the severely diseased macaques. When decomposed into Poincaré plots to showcase the day to day distribution of heart rate variability, with the central tendency of the plots represented by the geometric mean (Supplemental Figures A1 and A2), the disease courses of the severely diseased macaques demonstrate (Figure 5b) a statistically significant decrease in the geometric mean of the
Figure 3. Reduction in QT-Interval during the febrile period in macaques infected with severe EEEV. Shown are the profiles (a) of QT-Interval derived from 1-minute average measurements for four individual macaques infected with EEEV. Black: QT-Interval (ms), Gray: Temperature (°C), and Magenta: aerosol challenge. Blue Bars: Neurological score on corresponding day. (b) Average daily QT-Interval measurements, from repeated measures ANOVA, depicted in with boxes delimited by interquartile range and center line signifying median value. For (a) and (b), Top Left: M160-16, Bottom Left: M161-16, Top Right: M162-16, Bottom Right: M163-16.

Table 3. Repeated Measures ANOVA Statistics for QT-Interval. Statistics and p-values for QT-interval for EEEV and VEEV-infected macaque cohorts. *Doses in log_{10} pfu. TTD = Time to death. *P-Value lower than significance level α=0.05.

<table>
<thead>
<tr>
<th>Virus</th>
<th>Macaque</th>
<th>Sex</th>
<th>Dose*</th>
<th>TTD*</th>
<th>F-Statistic</th>
<th>P-Value</th>
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<td>6</td>
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<tr>
<td></td>
<td>M163-16</td>
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<td>6</td>
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<td>82.35</td>
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Figure 4. Reduction in RR-Interval during the febrile period in macaques infected with EEEV. Shown are the profiles (a) of RR-Interval derived from 1-minute average measurements for four individual macaques infected with EEEV. Black: RR-Interval (ms), Gray: Temperature (°C), and Magenta: aerosol challenge. Blue Bars: Neurological score on corresponding day. (b) Average daily RR-Interval measurements, from repeated measures ANOVA, depicted in with boxes delimited by interquartile range and center line signifying median value. For (a) and (b), Top Left: M160-16, Bottom Left: M161-16, Top Right: M162-16, Bottom Right: M163-16.
distribution (p<0.05) with an accompanying diminishment of variability, which recapitulates the markedly reduced heart rate variability roughly corresponding to the results seen in Figure 4.

Table 4. Repeated Measures ANOVA Statistics for RR-Interval. Statistics and p-values for RR-Interval for EEEV and VEEV-infected macaque cohorts. \(^d\)Doses in log_{10} pfu. \(^\delta\)TTD = Time to death.

<table>
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<th>Virus</th>
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Figure 5. Heart Rate Variability decreases during the febrile period in macaques infected with EEEV. (a) Distributions of heart rate variability of each EEEV-infected macaque are plotted in aggregate EEEV and color-coded by disease period (Cyan: Pre-Infection, Black: Post-Infection, Red: Febrile if applicable). (b) Repeated measures ANOVA of HRV data for macaques with nonsevere and severe EEEV infection (p<0.05). Central tendency of macaques with severe disease (Red) significantly diminishes after 3 dpi, compared to macaques with nonsevere disease (Blue).

2.2. ECG Changes in VEEV-infection of Macaques

Aerosol parameters of VEEV-infected macaques are summarized in Table 1. Exposure to VEEV produced a characteristic biphasic febrile illness in all exposed macaques with the first phase beginning within 18-24 hours post-infection and the second phase within 1-3 days after the first phase had resolved (Figure 6a). None of the macaques challenged with VEEV died from the disease, and all recovered from their fevers. Post-exposure, macaques became febrile after a latency period of a day, with a febrile peak of 40-41°C at approximately 1.3 dpi (Figure 6b-e). The temperature returned to normal by 1.5 dpi. A second fever occurred at 2.5 dpi and lasted approximately four days until 6.5-7.0 dpi. Subsequently, the macaques’ temperatures returned to normal, with the exception of hypothermia that occurred in M165-16. For analysis of VEEV-infected macaques, the post-infection period was defined as 0-1 dpi (inclusive), the febrile period as 2-7 dpi (inclusive), and the recovery period as 8+ dpi.
Figure 6. VEEV disease in macaques. Shown is the model of disease for (a) VEEV-exposed macaques over days post-infection (x-axis); all macaques developed febrile encephalitic disease. Deep blue denotes pre-infection baseline period, vertical magenta line denotes time of infection, powder blue defines post-infection period before the onset of fever, red signifies the febrile post-infection period, and yellow signifies recovery period after febrile illness subsides. Actual and forecasted temperature profiles for female macaques: (b) M164-16, (c) M165-16, and male macaques: (d) M170-16 and (e) M171-16. Red: Actual Temperature (°C), Blue Forecasted Temperature (°C) from ARIMA modeling, Black: Residual Values (°C) (deviations from forecasted baseline). Gray Lines: upper/lower bounds for residuals Magenta: aerosol challenge.
QT-intervals of the VEEV-infected macaques decreased during the post-infection period, and remained decreased during the febrile period; examination of QT-interval traces (Figure 7a) reveals that in two macaques (M165-16, M171-16) the QT-interval remained decreased during the recovery period, but for the other two in the cohort (M164-16, M170-16), the QT-interval returned to baseline during recovery. The daily QT-interval aggregate (Figure 7b) demonstrates that the QT-interval exhibited a sustained decrease beginning from the immediate post-infection period through the febrile period, until the beginning of the designated recovery period at 8 dpi; repeated measures ANOVA indicated statistically significant departures from baseline QT-interval in all four macaques (Table 3, p<0.05).

Figure 7. Reduction in QT-Interval during the febrile period in macaques infected with VEEV. Shown are the profiles (a) of QT-Interval derived from 1-minute average measurements for four individual macaques infected with VEEV. Black: QT-Interval (ms), Gray: Temperature (°C), and Magenta: aerosol challenge. Blue Bars: Neurological score on corresponding day. (b) Average daily QT-Interval measurements, from repeated measures ANOVA, depicted in with boxes delimited by interquartile range and center line signifying median value. For (a) and (b), Top Left: M164-16, Bottom Left: M165-16, Top Right: M170-16, Bottom Right: M171-16.

The RR-intervals of macaques infected with VEEV (Figure 8a) demonstrated decreases in all macaques beginning in the immediate post-infection, and remained persistently low throughout the recovery period. This effect is visible in the median RR-interval in aggregate data (Figure 8b), in

Figure 8. RR-Interval decreases and remains decreased in macaques with VEEV encephalitis. Shown are the profiles (a) of RR-Interval derived from 1-minute average measurements for four individual macaques infected with VEEV. Black: RR-Interval (ms), Gray: Temperature (°C), and Magenta: aerosol challenge. Blue Bars: Neurological score on corresponding day. (b) Average daily RR-Interval measurements, from repeated measures ANOVA, depicted in with boxes delimited by interquartile range and center line signifying median value. For (a) and (b), Top Left: M164-16, Bottom Left: M165-16, Top Right: M170-16, Bottom Right: M171-16.
which the RR-interval can be seen to trend downward beginning in the immediate post-infection period at
0 or 1 dpi (Table 4, p<0.05).

The HRV for the VEEV-infected macaques show in aggregate that when scatter plots of daily HRV
distributions are superimposed and color-coded by pre-infection, post-infection, febrile, and recovery
periods (Figure 9), the distribution of ordered pairs contracts, and remains contracted well after the febrile
period. When decomposed into Poincaré plots by day post infection, representative of each disease period
(Supplemental Figure A3), the persistence of the decrease in heart rate variability is even more apparent.
As with the related RR-interval, all macaques experienced this effect.

Figure 9. Heart Rate Variability Decreases during the febrile period in macaques infected with
EEEV. Distributions of heart rate variability of each EEEV-infected macaque are plotted in aggregate
EEEV and color-coded by disease period (Cyan: Pre-Infection, Black: Post-Infection, Red: Febrile,
Yellow: Recovery).

2.3. Frequency Spectrum Analysis

Using Fast Fourier transformation (FFT), pre-/post-infection and febrile periods were
distinguishable from one another in the macaque subjects with severe courses of disease. The
fundamental frequency, defined as the first local maximum in the frequency spectrum, of each
metric was computed in cycles per day (cpd). Patterns of change in the frequency spectra varied
little between ECG metrics, especially between those that co-varied; for instance, metrics
associated with heart rate, such as RR-interval, exhibited similar trends with respect to how the
fundamental frequencies changed between pre-, post-, and febrile periods. The amplitude of the
frequency spectrum plots was not significant, as magnitude squared (ms² in this case)
determined by the number of points fed into the FFT and the amplitude of the original signal in
the time domain. Fundamental frequencies for the RR-interval, as a representation of the overall
trend of frequency changes, are tabulated in Table 5.

Prior to infection, the fundamental frequency for RR-interval was different between
macaques; however, most fell into a fairly narrow range. Macaques infected with EEEV that did
not develop severe disease saw no significant changes in RR-I fundamental frequencies
post-infection (Table 5). Changes in RR-I fundamental frequencies were much larger in severely
diseased macaques. In the EEEV-infected macaques with severe disease, the post-infection
frequency increased to 0.879 and 0.769 cpd, then decreased to 0.571 and 0.549 cpd in the febrile
period although this was still elevated compared to baseline. Values for VEEV were larger due
to the extended time periods used for analysis. Examination of fundamental frequencies
identified relatively high cyclic rates in the VEEV cohort with differences based on sex: 5.97 cpd
and 11.25 cpd for female and male monkeys, respectively, at pre-infection. For both females and
males, the fundamental frequencies increased after VEEV infection and then dropped below
baseline levels in the febrile period with the sharpest decline in the male macaques. With the
onset of the designated recovery period in VEEV, the fundamental frequency of the latter
period, when body temperatures largely returned to normal, the fundamental frequencies in
The VEEV-infected macaques remained lower than pre-infection although the greatest reduction was seen in the males.

Table 5. Fundamental Frequencies of EEEV and VEEV Courses of Infection. This table documents fundamental frequencies observed in the ECG metric of RR-Interval. Disease periods are separated into pre-infection, post-infection (outside of any febrile period), and febrile periods for EEEV VEEV. Recovery periods were also included when applicable, specifically in the case of the VEEV cohort. Fundamental frequencies are displayed in cycles per day (cpd). Values for VEEV are larger due to the extended time periods used for analysis; the values in parentheses are normalized to the first harmonic value for the purposes of comparison to EEEV frequency values. Italicized values represent animals with fatal courses of disease.

<table>
<thead>
<tr>
<th>Virus</th>
<th>Macaque</th>
<th>Sex</th>
<th>Pre-Infection Frequency</th>
<th>Post-Infection Frequency</th>
<th>Febrile Period Frequency</th>
<th>Recovery Period Frequency</th>
</tr>
</thead>
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<tr>
<td>EEEV</td>
<td>161-16</td>
<td>M</td>
<td>0.395</td>
<td>0.879</td>
<td>0.571</td>
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<tr>
<td></td>
<td>163-16</td>
<td>M</td>
<td>0.242</td>
<td>0.769</td>
<td>0.549</td>
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<tr>
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<td>160-16</td>
<td>M</td>
<td>0.351</td>
<td>0.242</td>
<td></td>
<td></td>
</tr>
<tr>
<td></td>
<td>162-16</td>
<td>M</td>
<td>0.307</td>
<td>0.351</td>
<td></td>
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</tr>
<tr>
<td>VEEV</td>
<td>164-16</td>
<td>F</td>
<td>5.974 (0.373)</td>
<td>11.070 (0.275)</td>
<td>4.393 (0.275)</td>
<td>3.514 (0.220)</td>
</tr>
<tr>
<td></td>
<td>165-16</td>
<td>F</td>
<td>5.974 (0.373)</td>
<td>11.070 (0.286)</td>
<td>4.568 (0.286)</td>
<td>3.514 (0.220)</td>
</tr>
<tr>
<td></td>
<td>170-16</td>
<td>M</td>
<td>11.245 (0.351)</td>
<td>13.881 (0.434)</td>
<td>4.217 (0.132)</td>
<td>3.163 (0.099)</td>
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<tr>
<td></td>
<td>171-16</td>
<td>M</td>
<td>11.245 (0.351)</td>
<td>13.705 (0.428)</td>
<td>5.271 (0.165)</td>
<td>3.163 (0.099)</td>
</tr>
</tbody>
</table>

3. Discussion

3.1. QRS Complexes

As the QRS interval corresponds to ventricular depolarization in the heart and therefore the speed with which ventricular contraction occurs, a widened QRS complex could point to electrolyte imbalances such as hypokalemia. Clinical observations noted that both severely diseased EEEV-infected macaques had decreased water intake during the febrile period and signs of dehydration. However, blood chemistry analysis obtained at baseline, during aerosol, and at necropsy did not bear out this hypothesis, as potassium levels were within normal limits (3.5-5 mEq), leaving the question open as to whether autonomic dysregulation is responsible for the pathological phenomenon observed. The lack of blood chemistry between infection and necropsy means that the possibility of electrolyte balance dysregulation cannot be completely ruled out. Recent investigation has suggested that autonomic modulation of heart rate regulation can lead to morphological changes in the QRS complex, with shorter QRS complexes representing a shift towards sympathetic predominance [37]. However, in the macaques in which this occurred, the finding was more likely a covariate of an increased heart rate due to febrile illness.

3.2. QT-Intervals

The QT-interval represents the interval of time between the Q-Wave (beginning of ventricular depolarization) and the T-Wave (ventricular repolarization) of the ECG. The finding of decreased QT-intervals in the EEEV-exposed macaques with severe disease, again, represents an anomaly in that though the QT-interval is decreased, the QRS complex, which comprises a segment of the QT-interval, is increased. The QT-interval is of particular interest for the testing of therapeutic compounds, due to the extent to which certain small molecule drug candidates can affect the QT-interval. Ion channels in the myocardium can be affected by off-target effects, and this represents concern for potential pharmacological intervention [38]. However, pre-clinical studies in uninfected
animal models should mitigate the potential for such an eventuality. Perhaps the most
tought-provoking aspect of the QT-interval effects resides in the alterations to diurnal variation
during and sometimes after the period of febrile illness, a finding mirrored in the upcoming
discussion of the RR-interval and heart rate variability.

3.3. RR-Interval, Heart Rate Variability, and Frequency Spectrum Analysis

The RR-interval denotes the interval of time between two heartbeats, while heart rate variability
(HRV) represents the variation in the time interval between heartbeats, expressed in milliseconds
[39]. The notion that RR-interval and HRV can illustrate the circadian rhythm of autonomic
regulation has been well characterized [40-42]. Changes in HRV have comprised the subject of study
by multiple authors who have associated alterations in the RR-interval and HRV with pathology
such as diabetes mellitus and its associated diabetic neuropathy, heart failure, myocardial infarction,
and other chronic conditions [40,43,44]. Additionally, more recent studies have suggested a link not
only between inflammatory markers and HRV [45], but also between changes in the RR-interval and
heart rate variability in a number of diseases of infectious origin [46-48].

For both VEEV-infected and severely EEEV-infected macaques, there was a decrease in the
RR-Interval in the febrile period. For severe EEEV-infection, the decrease was greater, with a more
complete loss in diurnal variation. Subsequent work with macaques challenged with EEEV (not
published) demonstrates that the EEEV macaques with no overt clinical signs of disease are indeed
infected, as determined by plaque assay and antibody neutralization titers from plasma obtained
from serial blood draws following the aerosol challenge. Therefore, as the EEEV-challenged
macaques with nonsevere disease also have limited to no changes in RR-interval as depicted in daily
Poincaré plots, this effect can therefore distinguish severe from nonsevere courses of disease.
RR-interval remained persistently low in all four VEEV-infected macaques even through the
recovery period. Despite recovery from encephalitic disease, the results suggest that infection with
VEEV may have altered the homeostatic set point of autonomic regulation, as previously reported in
the literature [12,18,49].

The fundamental frequency represents the frequency with which the original time domain
signal (the ECG metric) repeats, typically multiple times per day due to idiosyncrasies in the
autonomic regulation of cardiac function. Although the fundamental frequency is what was used as
the benchmark, subsequent local maxima are registered as nth harmonics, or subsequent echoes, of
the fundamental frequency. Frequency spectrum analysis reveals similar trends over different
periods of EEV disease courses; post-infection, the fundamental frequencies of the ECG metrics
typically increase, then decrease during the febrile period in individuals with severe disease.

In the context of frequency spectrum analysis, for macaques with severe courses of EEEV, the
fundamental frequencies of the RR-interval increased between pre-infection, the latency period, and
febrile disease, whereas the fundamental frequencies of macaques with nonsevere courses of EEEV
remain relatively unchanged. This result corroborates the value of RR-interval in distinguishing
between severe and nonsevere cases of EEEV and highlights the capacity of frequency spectrum
analysis to perform this distinguishing function. For VEEV, the fundamental frequencies of the
RR-interval increased post-infection during the short latency period, then decreased during the
onset of febrile illness. During the recovery period, the fundamental frequency proceeded to
decrease again as the permanence of the decreased RR-interval, plotted in Poincaré plots in
Supplemental Figure A4, and diminished heart rate variability for the VEEV-exposed cohort is
recapitulated in Table 5 by the frequency spectrum analysis, suggesting a real physiological rather
than artifactual basis for these results. The existence of post-infection changes in the fundamental
frequencies in subjects who survived EEEV challenge suggests that neurological sequelae could
occur even without a strong febrile response or obvious clinical signs, sequelae that could have long
term health consequences.
3.4. Limitations and Future Work

We do acknowledge that there is a shortfall in investigating the electrocardiographic implications of the dose-response relationship due to the small sample sizes, a limitation partially offset by the sheer volume of data obtained from each subject. The potential confounding effect of subject sex likewise presents a limitation; the EEEV cohort was entirely male, while the VEEV cohort was split between the sexes. Baseline metrics varied significantly between individual macaques, although all baseline measures fell within normal limits. At this juncture, it is not clear whether the results reported here are applicable to other viral encephalitides. Direct application of these analyses in a human clinical context may suffer from limited generalizability, based on the variability in baseline ECG metrics between individual macaques and baseline ECG data would not be available for most human patients.

There are a number of remaining questions, including the following: do sex-based differences in electrocardiographic signatures precipitate effects on the response to infection by aerosol route, and if so, how does the periodicity of hormonal changes, for example, modify the outcome, especially with respect to the loss of circadian variation seen in all signals? Perhaps body weight or animal size may exert an effect on the electrocardiographic metrics not only in baseline data, but also in the post-infection period response, through the febrile period. A decrease in RR-Interval during the febrile period appears to be a generalizable phenomenon to other severe, acute infectious diseases (D.S. Reed, manuscript in preparation) although it is not clear if these other severe infections penetrate the CNS particularly when the route of exposure is also aerosol. The other parameters noted here (QT-Interval, QRS) may be specific to viral encephalitis. Finally, it may be helpful to determine whether a dose-dependent relationship exists between the manifestation of electrocardiographic changes and the dose of virus received, or between the immune response observed, or whether the ECG effects observed are binary once exposure exceeds a threshold dose. The results described appear to support the latter theory; while these results may not be specific to viral encephalitides and may not supplant fever as a biomarker for the initiation of treatment, they may serve as useful predictors of outcome. Further investigation with electroencephalography can better determine whether the differences observed between sexes are responsible for the dichotomies observed in the results, and can help to establish whether the observed ECG effects stem from a basis in autonomic dysregulation, from direct viral infection, the immune response to infection, or a combination of these factors.

Future work may improve the utility of ECG data to provide specific profiles and prognosticators of disease. Dimension reduction can be carried out among covarying metrics for the purposes of principal component analysis to provide a more robust form of prognostication. In conjunction with analysis of electroencephalography and other telemetric modalities, such methods could prove a powerful tool, followed in real time or quasi-real-time by lab staff, for investigating both viral pathogenesis and developing therapeutics and countermeasures against EEV aerosol infection. Whether EEVs directly affect cardiac responses as captured by electrocardiography metrics with respect to damage to the myocardial syncytium versus the disruption of autonomic outflow requires further study and remains an area of continuing investigation.

4. Materials and Methods

4.1 Statement on Rationale, Use, and Care of Animals

This study was approved by the University of Pittsburgh IACUC (Protocol #16026773) and was fully compliant with rules, regulations, and recommendations stipulated by the Animal Welfare Act Regulations and the Guide for the Care and Use of Laboratory Animals, [50]. The University of Pittsburgh, which includes a regional biocontainment laboratory (RBL), is accredited by the Association for Assessment and Accreditation of Laboratory Animal Care (AAALAC). The rationale for the development of a nonhuman primate (NHP) model of inhalational EEV disease fulfills the necessity criteria outlined by the FDA Animal Rule [29,51,52]. The cynomolgus macaque (Macaca fascicularis) has served as a model organism for EEVs since the discovery of these viruses in the 1930s.
Infection in the context of mg/kg cefazolin (1st-generation cephalosporin) for infection control and 5%v/v isopropyl alcohol was then sterilized with chlorhexidine antiseptic and administered for continuous anesthesia with 4.0 mg/kg ketoprofen for analgesia, and the incision site, 5 cm superior to the anterior superior iliac spine, was shaved. Macaques were anesthetized with 10mg/kg ketamine with continuous IV administration of 3% normal saline via the greater saphenous vein. Isoflurane (2%v/v) was then administered for continuous anesthesia with 4.0-4.5L/min of O2. The incision site was draped and sterilized with chlorhexidine antiseptic and 70%v/v isopropyl alcohol. A 5 cm incision was made parallel to the midline, into which the implant was inserted. ECG leads were sutured in place, one under the right pectoralis muscle and one parallel to the left inguinal area, approximating Lead II placement in a 12-lead ECG [59]. The incision was closed with a simple interrupted stitch. Post-operatively, each macaque was eligible for 3 days’ administration (PRN) of buprenorphine for analgesia and received 5 days’ administration (BID) of cefazolin for infection control.

4.2 Animal Model Telemetry Implantation

The macaques used in this study were implanted intra-abdominally with Data Sciences International (DSI, St. Paul, MN) radiofrequency transmitters (DSI Model No. M01) by a DSI veterinarian at Covance Laboratories (Princeton, NJ). In preparation for surgery, the macaques were each administered 25mg/kg cefazolin (1st-generation cephalosporin) for infection control and 1mg/kg ketoprofen for analgesia, and the incision site, 5 cm superior to the anterior superior iliac spine, was shaved. Macaques were anesthetized with 10mg/kg ketamine with continuous IV administration of 3% normal saline via the great saphenous vein. Isoflurane (2%v/v) was then administered for continuous anesthesia with 4.0-4.5L/min of O2. The incision site was draped and sterilized with chlorhexidine antiseptic and 70%v/v isopropyl alcohol. A 5 cm incision was made parallel to the midline, into which the implant was inserted. ECG leads were sutured in place, one under the right pectoralis muscle and one parallel to the left inguinal area, approximating Lead II placement in a 12-lead ECG [59]. The incision was closed with a simple interrupted stitch. Post-operatively, each macaque was eligible for 3 days’ administration (PRN) of buprenorphine for analgesia and received 5 days’ administration (BID) of cefazolin for infection control.

4.3 Virus Culture and Dose Determination

Virus cultures were conducted as described in the literature [60]. The EEEV and VEEV isolates used in this study were single passages of infectious clones generated from human isolates: V105 (EEEV) and INH-9813 (VEEV-IC). For both alphaviruses, stocks were generated through construction of cDNA libraries; capped, infectious viral RNAs were generated by in vitro RNA synthesis from linearized cDNA plasmid template genomes that were electroporated into baby hamster kidney (BHK-21, ATCC) cells (Klimstra, manuscript in preparation). The supernatant was clarified by centrifugation between 18-24 hours post-electroporation, and single-use aliquots were stored at -80°C as zero-passage (p0) stocks. The p0 stocks were titered by standard plaque assays on BHK-21 cells and used to infect Vero cells (ATCC CCL-81) at a multiplicity of infection of 10 in roller bottles. At 18-24 hours post-infection, the supernatant was clarified by centrifugation and subjected to sucrose purification. The supernatant was layered over a 20/60% sucrose cushion and ultracentrifuged. The interface between the 20/60% sucrose cushion was collected and diluted in 10mM Tris, 1mM EDTA, 100mM NaCl–STE 10X NaCl (TNE) buffer. The interface was then layered over 20% sucrose and again ultracentrifuged to pellet the virus. The pelleted virus was resuspended in Opti-MEM® Reduced Serum Medium (ThermoFisher, Catalog No. 31985-070) and single-use aliquots were stored at -80°C. The virus stock (passage 1), nebulizer samples, and aerosol samples...
were titered using standard plaque assays on BHK-21 cells and the aerosol LD₅₀ confirmed in mice before use in macaque studies.

### 4.4 Aerosol Exposure of Macaques to EEVs

For each exposure, aerosol generation was performed in a modified Class III biosafety cabinet [61]. Aerosol exposures were performed under the control of the Aero3G aerosol management platform (Biaera Technologies, Hagerstown, MD) as previously described [62]. Macaques were anesthetized with 6 mg/kg Telazol® (Tiletamine HCl / Zolazepam HCl). Each macaque was then weighed, bled, and transported to the modified Class III biosafety cabinet using a mobile transport cart. The macaque’s head was placed inside an acrylic head-only exposure chamber. Jacketed External Telemetry Respiratory Inductive Plethysmography (JET-RIP; DSI) belts were placed around the abdomen and chest of the macaque and calibrated to a pneumotach. This allowed monitoring and recording of respiratory function via the Ponemah software platform (DSI) during the aerosol [63]. EEV aerosols were generated using an Aeroneb vibrating mesh nebulizer (Aerogen, Chicago, IL) as previously described [64]. Exposures were 10 minutes in duration. To determine inhaled dose, aerosol sampling was performed during each exposure with an all-glass impinger (AGI; Ace Glass, Vineland, NJ). Particle size was measured once during each exposure at 5 minutes using an Aerodynamic Particle Sizer (TSI, Shoreview, MN). A 5-minute air wash followed each aerosol before the macaque was removed from the cabinet, and transported back to its cage and observed until fully recovered from anesthesia.

### 4.5 Electrocardiography Data Collection

Electrocardiography, activity, and temperature data were collected continuously from implanted macaques for at least two days before aerosol exposure and for the duration of the disease course post-exposure. An example of a raw ECG trace is shown in Supplemental Figure A4. A computer connected to a communication link controller (CLC; DSI) collected signals from the implanted macaques by way of transceivers mounted in the room. Video cameras were positioned to record macaque behavior and were programmed to record continuously, synchronously, alongside the telemetry data. Because of the amount of telemetry and video data generated daily, the system had to be manually stopped and restarted every morning. The Ponemah software package (v.5.2; DSI) provides a graphic user interface that presents a unified, multichannel display of the data collected from the M01 implants’ sensors. For analysis, data were read either in raw form (.RAW file extension) or reviewed in common spreadsheet, statistical analysis, and plotting programs. Body temperature and electrocardiographic measures including heart rate were sampled at 10 Hz and recorded every 5 seconds [65]. For fever, a Box-Jenkins ARIMA model was used to predict 15-minute average body temperature from baseline data; significant deviations (3 times the square root of the residual sum of squares) of actual from predicted temperatures were scored as fever. Fever duration was calculated as the number of fever points divided by 4 (the number of points per hour); fever severity was measured as ‘fever-hours’, the summation of residual elevated temperatures divided by 4. ARIMA analysis was done using Number Cruncher Statistical Systems (NCSS) 2007 software. Electrocardiography metrics studied are displayed in Supplemental Table A1. Standard deviations of these metrics were measured and recorded at the same sampling frequencies as well.

### 4.6 Electrocardiography Analysis Methods

Disease courses were defined by the appearance of fever during the clinical observation period; for each cohort of EEV-infected macaques, periods were established based on the timing of infection and fever: pre-infection, post-infection (prior to fever onset), febrile period, and recovery, when applicable. These periods were constructed according to the markers for fever explained above. Analyses of electrocardiography metrics were carried out by considering the median daily electrocardiographic metric of interest or by considering the aggregate median electrocardiographic metric of interest over the periods previously defined.

Electrocardiographic metrics were sampled from the raw data at a rate of one sample per minute for 1440 data points per day to reduce the computational burden on the analytical software
and to reduce the impact of noise on producing artificially significant results. Statistical analyses were performed in two stages: preliminary analysis of temperature and activity were performed alongside the collection of data, with limited analysis of the raw ECG traces. A more comprehensive statistical analysis was performed with a within-subjects repeated measures ANOVA approach. Due to significant variability in baseline metrics, the baseline period for each macaque served as its own control for this study. All significance levels utilized were α<0.05. Heart rate variability was mapped as described by Golinska, and Poincaré plots were constructed according to guidelines described by Henriques, et al [66,67].

To examine circadian changes in the electrocardiographic data, frequency spectrum analysis was performed by use of the fast Fourier transform (FFT). Time-series of the electrocardiographic metrics were decomposed into linear combinations of basic trigonometric functions. The frequencies of these trigonometric functions produces a discrete spectrum, with each frequency point representing a component frequency of the original time-series. The fundamental frequency, or the first defining frequency peak (local maximum) in the frequency spectrum window, in units of cycles per day, explored the differences in pre-/post-infection, febrile, and recovery periods. Subtle changes in the number of cycles per day, or the subcomponents of the normal cyclic rates of electrocardiographic metrics, were quantified and visualized. These procedures were carried out in MATLAB 2018b (Mathworks, Natick, MA) with the use of native frequency analysis algorithms.

5. Conclusions

This work was performed to further develop and characterize the macaque model for aerosol exposure to EEVs. In agreement with what we saw previously, fever was a prominent indicator of EEV disease. Across both EEEV and VEEV, it is notable that the macaques who succumbed to severe disease had a maximum temperature difference from predicted of 4.0°C or higher while all of those that survived infection had maximum temperature differences less than 4.0°C from predicted. The goal of these studies was to determine whether other physiological parameters might be suitable as biomarkers or indicators for future efficacy studies. The augmented animal model might suffice to evaluate therapeutic compounds under the FDA Animal Rule. Although these are encephalitic viruses, heart function is controlled by the nervous system so we surmised that ECG metrics would be useful for this objective. Indeed, the data suggests that ECG metrics are useful for this purpose, but the ECG metric chosen will be dependent on the particular alphavirus used and will likely need to be paired with body temperature to paint a complete picture of the disease and outcome in macaques. The most prominent set of significant electrocardiographic metrics that distinguish and characterize both EEEV and VEEV disease courses consist of QRS complexes, QT-interval, and RR-interval.
Supplementary Materials: The following are available online, Figure S1: Representative Poincaré Plots for Nonsevere EEEV Infection Cohort, Figure S2: Representative Poincaré Plots for Severe EEEV Infection Cohort, Figure S3: Representative Poincaré Plots for VEEV Infection Cohort, Figure S4: Example ECG Trace, Table S1: Electrocardiographic Metrics.


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Conflicts of Interest: The authors declare no conflict of interest. The funders had no role in the design of the study; in the collection, analyses, or interpretation of data; in the writing of the manuscript, or in the decision to publish the results. None of the submitted work was carried out in the presence of any personal, professional, or financial relationships that could be potentially construed as a conflict of interest; none of the authors listed owns controlling interest in any of the vendors or suppliers listed in this manuscript.
Appendix A – Supplementary Figures and Tables

Supplemental Figure A1. Representative Poincaré Plots for Nonsevere EEEV Infection Cohort.
Representative daily Poincaré plots for a macaque with nonsevere EEEV infection (M160-16).

Supplemental Figure A2. Representative Poincaré Plots for Severe EEEV Infection Cohort.
Distributions of heart rate variability of each EEEV-infected macaque are plotted in aggregate EEEV and color-coded by disease period (Cyan: Pre-Infection, Black: Post-Infection, Red: Febrile, Yellow: Recovery).
Supplemental Figure A3. Representative Poincaré Plots for VEEV Infection Cohort.

Representative daily (from -1, +1, +3, and +14 dpi) Poincaré plots for each macaque in the VEEV infection cohort (top to bottom: M164-16, M165-16, M170-16, and M171-16) show diminished range of HRV following infection that persists into the recovery period.

Supplemental Figure A4. Example ECG Trace. Shown is a sample ECG trace from a macaque in this study. The electrocardiogram generated by a single heartbeat provides the PQRST profile (marked) of an atrial and ventricular contraction and repolarization. Different segments of the waveform serve as key metrics that may inform upon underlying cardiac anatomical and physiological status. The P wave corresponds to the depolarization of the atria, the QRS complex corresponds to both ventricular depolarization and atrial repolarization, and the T wave represents ventricular repolarization. QRS complex is defined by the time period transited by the peak between points Q and S.
### Table A1. Electrocardiographic Metrics. Abridged list of metrics under study.

<table>
<thead>
<tr>
<th>Electrocardiography Metric</th>
<th>Abbreviation</th>
<th>Significance</th>
<th>Units</th>
</tr>
</thead>
<tbody>
<tr>
<td>Arrhythmic Beat Detection</td>
<td>BAD</td>
<td>The number of arrhythmic beats detected during a logging period.</td>
<td>Count</td>
</tr>
<tr>
<td>Heart Rate</td>
<td>HR</td>
<td>Number of heartbeats per minute.</td>
<td>bpm</td>
</tr>
<tr>
<td>Heart Rate Variability</td>
<td>HRV</td>
<td>RR-Interval matched with RR-Interval, plotted as an ordered pair.</td>
<td>ms, Unitless</td>
</tr>
<tr>
<td>Maximum Voltage Derivative of R-Wave</td>
<td>MxdV</td>
<td>Maximum rate of change of ECG trace at R-Wave.</td>
<td>mv/ms</td>
</tr>
<tr>
<td>Noise in ECG Trace</td>
<td>Noise</td>
<td>Approximation of noise level in ECG cycle; root-mean-square value of derivative in a single ECG cycle.</td>
<td>Unitless</td>
</tr>
<tr>
<td>P-Wave Count</td>
<td>PCt</td>
<td>Number of P-waves counted in one sample.</td>
<td>Count</td>
</tr>
<tr>
<td>P-Wave Height</td>
<td>P-H</td>
<td>Height of P-Wave from isoelectric level.</td>
<td>mV</td>
</tr>
<tr>
<td>PR-Interval</td>
<td>PR-I</td>
<td>Interval of time between beginning of P-Wave to the beginning of the R-Wave. Can detect atrial conduction block.</td>
<td>ms</td>
</tr>
<tr>
<td>P-Wave Width</td>
<td>P-Width</td>
<td>Interval of time from the P-Wave to the end of the P-Wave.</td>
<td>ms</td>
</tr>
<tr>
<td>QaT</td>
<td>QATN</td>
<td>Interval of time between the Q-Wave to the peak of the T-Wave.</td>
<td>ms</td>
</tr>
<tr>
<td>QR-Interval</td>
<td>QR-I</td>
<td>Interval of time between the Q-Wave to the peak of the R-Wave. Indicator of ventricular depolarization and/or contraction.</td>
<td>ms</td>
</tr>
<tr>
<td>QRS complex</td>
<td>QRS</td>
<td>Interval of time between the Q-Wave to the beginning of the S-Wave. Indicator of ventricular depolarization and/or contraction.</td>
<td>ms</td>
</tr>
<tr>
<td>QRS Amplitude</td>
<td>QRSA</td>
<td>Amplitude of QRS complex from the isoelectric level. Indicator of ventricular contraction.</td>
<td>mV</td>
</tr>
<tr>
<td>QT-Interval</td>
<td>QT-I</td>
<td>Interval of time between the Q-Wave to the end of the T-Wave. Indicator of electrolyte status as well as speed of total cardiac repolarization.</td>
<td>ms</td>
</tr>
<tr>
<td>R-Wave Height</td>
<td>R-H</td>
<td>Height of the R-Wave from the isoelectric level.</td>
<td>mV</td>
</tr>
<tr>
<td>RR-Interval</td>
<td>RR-I</td>
<td>Interval from one R-Wave peak to the next R-Wave peak.</td>
<td>ms</td>
</tr>
<tr>
<td>ST-Elevation</td>
<td>ST-E</td>
<td>Height of the T-Wave at algorithmically-determined point between S-Wave and T-Wave, to detect myocardial infarction.</td>
<td>mV</td>
</tr>
<tr>
<td>ST-Interval</td>
<td>ST-I</td>
<td>Interval of time between the S-Wave and the end of the T-Wave.</td>
<td>ms</td>
</tr>
<tr>
<td>T-Wave Height</td>
<td>T-H</td>
<td>Highest point between the end of the S-Wave and end of the T-Wave.</td>
<td>mV</td>
</tr>
<tr>
<td>Peak of T-Wave</td>
<td>T-P</td>
<td>Peak of T-Wave relative to isoelectric level, between end of S-Wave and end of the T-Wave.</td>
<td>mV</td>
</tr>
<tr>
<td>T-Wave Peak-to-End</td>
<td>T-Pe</td>
<td>Time between the peak of the T-Wave to the end of the T-Wave; indicator of ventricular reperfusion.</td>
<td>ms</td>
</tr>
</tbody>
</table>
References


59. Einthoven, W.; Fahr, G.; De Waart, A. On the direction and manifest size of the variations of potential in the human heart and on the influence of the position of the heart on the form of the electrocardiogram. *Am Heart J* 1950, 40, 163-211.


