

1 **Introducing *circStudio*, a Python package for preprocessing,**
2 **analyzing and modeling actigraphy data**

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14 **Abstract**

15 Actigraphy is a non-invasive and cost-effective method for monitoring behavioral rhythms under
16 real-world conditions by collecting time-resolved measurements of locomotor activity, light
17 exposure, and temperature. Although several open-source packages support specific aspects of
18 actigraphy analysis, aspects such as preprocessing, metric calculation, and mathematical modeling
19 are often distributed across separate software packages, limiting interoperability and increasing
20 programming overhead. Here we introduce *circStudio*, a Python package that unifies actigraphy
21 data processing and mathematical modeling of circadian rhythms within a single framework. Built
22 from the *pyActigraphy* codebase and integrating circadian models from the Arcascope *circadian*
23 package, *circStudio* provides flexible preprocessing tools, support for multiple actigraphy file
24 formats through adaptor classes, standalone functions for computing commonly used actigraphy
25 metrics, and implementations of several mathematical models of circadian rhythms. The package
26 enables users to move efficiently from raw wearable data to physiologically interpretable circadian
27 outputs. Ultimately, *circStudio* aims to facilitate reproducible workflows and to provide a flexible
28 foundation for research applications across circadian biology, sleep science, and digital health.

29

30 **Keywords**

31 Actigraphy, circadian rhythm, sleep, time series, wearables, computer software, signal processing,
32 chronobiology, data analysis, data visualization

33

34 **Introduction**

35 Actigraphy is a non-invasive method that uses wearable sensors to record continuous time
36 series of locomotor activity, light exposure and peripheral temperature, enabling the quantification
37 and inference of sleep-wake patterns under ambulatory conditions (Fekedulegn et al., 2020). In
38 contrast with polysomnography (PSG), the gold standard for assessing sleep but a method that is
39 expensive and often disruptive to normal sleep routines, actigraphy is well suited for large-scale
40 data collection owing to its non-invasive nature and relatively low cost (Fekedulegn et al., 2020).
41 Actigraphy time series can be used to derive commonly reported actigraphy variables or features
42 (Carpenter et al., 2025). These features include interdaily stability (IS), which measures the
43 strength and consistency of the 24-hour rest-activity pattern across days in the actigraphy
44 recording; the sleep regularity index (SRI), which quantifies the similarity of sleep-wake patterns
45 between consecutive days; and the relative amplitude (RA), which contrasts the difference between
46 the 10 most active (or illuminated) hours and the least active (or illuminated) hours of the day
47 (Carpenter et al., 2025).

48 Actigraphy time series can also be used to model the response of the circadian system to a
49 given pattern of light exposure through mathematical models of circadian rhythms (Hannay &
50 Moreno, 2020). One of the most widely used formalisms was originally developed by Kronauer
51 based on a Van der Pol (VDP) oscillator and later refined by Forger and Jewett (Forger et al., 1999;
52 Jewett et al., 1999). More recently, Hannay proposed a formalism aimed at improving the
53 physiological interpretability of model state equations (Hannay et al., 2019). In particular, the
54 Hannay framework describes how the dynamics of rhythms at the level of individual cells can give
55 rise to collective rhythm amplitude in the suprachiasmatic nucleus (SCN) of the hypothalamus
56 (Hannay et al., 2019). Other models extend the VDP formalism to incorporate non-photic input
57 (St. Hilaire et al., 2007), while others simulate the daily trajectory of melatonin across different
58 physiological compartments, such as the pineal gland, plasma and exogenous melatonin (Breslow
59 et al., 2013).

60 Over the years, several actimetry sensor brands have been developed, each producing data
61 in distinct proprietary file formats and typically accompanied by dedicated, closed-source analysis
62 software (Hammad et al., 2021). Although these tools facilitate basic processing of sensor data,
63 they often restrict the range of analytical approaches available to researchers and lead to actigraphy
64 data underutilization (Fekedulegn et al., 2020). Moreover, reliance on device-specific software and

65 closed-source processing pipelines makes it difficult to generalize results across studies (Hammad
66 et al., 2021). To address these limitations, several programming packages have been developed.
67 For instance, in the R programming language, a number of packages have been dedicated to
68 actigraphy data analysis, such as *LightLogR* (Zauner et al., 2025) for the analysis of personal light
69 exposure.

70 Critically, most analytical functionality for actigraphy data has not been integrated into a
71 single, easy-to-use package that supports preprocessing, analysis, and modeling within a unified
72 framework. In Python, one of the most notable open-source toolboxes developed for actigraphy
73 data analysis is *pyActigraphy* (Hammad et al., 2021), which also has a dedicated module for light
74 exposure analysis called *pyLight* (Hammad et al., 2024). Another Python package, *circadian*,
75 enables the calculation of such models (Tavella et al., 2023), but it is not optimized to handle noisy
76 light schedules and often fails to compute model state variables under more realistic light exposure
77 patterns. Moreover, moving between *pyActigraphy* and *circadian* typically requires additional
78 programming effort. To bridge these tools and provide an integrated workflow for actigraphy
79 analysis and circadian modeling, we introduce *circStudio*, a Python framework that combines
80 actigraphy data processing with mathematical modeling of circadian rhythms.

81

82 ***circStudio* design and architecture**

83 The *circStudio* package was developed on top of *pyActigraphy*, although it introduces
84 several architectural differences. *circStudio* is organized into two main modules: `io`, which
85 handles file input/output operations and preprocessing, and `analysis`, which provides
86 functionality for data analysis. In contrast, *pyActigraphy* distributes preprocessing (e.g., `io`, `log`,
87 `mask`) and analysis (e.g., `metrics`, `sleep`, `light`, `viz`) functionality across multiple modules. A
88 key motivation for reorganizing the codebase was to reduce code duplication and foster community
89 contributions, while providing greater flexibility for users. One example of this flexibility is the
90 segregation of data analysis functions from the `Raw` class, which is responsible for reading and
91 preprocessing actigraphy files. This architecture enables users to implement custom preprocessing
92 routines and use *circStudio* exclusively to compute actigraphy-derived metrics and/or simulate
93 mathematical models.

94 A typical workflow in *circStudio* involves preprocessing, feature extraction, mathematical
95 modeling, and visualization of actigraphy data. Raw recordings are first imported and preprocessed

96 to handle device-specific formats using adaptor classes, masking, and data cleaning. The processed
97 time series can then be used to extract commonly reported actigraphy-derived features, including
98 activity, light exposure, and sleep-related variables. The processed data may also serve as inputs
99 to mathematical models of circadian rhythms that simulate the response of the circadian system to
100 light exposure. Finally, *circStudio* provides functionality to visualize the resulting time series,
101 model outputs, and several actigraphy-derived metrics.

102

103 **Installation**

104 The latest version of *circStudio* can be retrieved directly from PyPI using:

105

106

```
107 pip install circStudio
```

108

109

110 The installation can be verified by running:

111

112

```
113 Python -c "import circstudio; print('circstudio imports')"
```

114

115

116 **Example workflows**

117 A typical use case begins by importing the package and loading an actigraphy file, followed
118 by signal preprocessing, metric computation, and optionally simulating circadian system dynamics
119 using a light intensity time series as input to mathematical models of circadian rhythms.

120

121 **Loading actigraphy data with the *Raw* class**

122 The first step in any analysis is to import *circStudio*:

123

124

```
125 import circstudio as cs
```

126

127

128 The preferred method for loading actigraphy data in *circStudio* is to create a new instance
129 of the `Raw` class. This requires the actigraphy data to be preformatted as a table containing activity
130 and light series. Unlike *pyActigraphy*, *circStudio* decouples the metric computation from the `Raw`
131 class, which is used exclusively for signal preprocessing. This design allows users to define custom
132 preprocessing pipelines and to apply *circStudio* metrics to other types of time series, such as
133 peripheral body temperature.

134 To create a `Raw` instance, users must specify the dataframe (`df`), the activity (`activity`)
135 and light (`light`) time series (set `light` to `None` if no light time series is available), as well as
136 start time (`start_time`), total duration (`period`), and sampling frequency (`frequency`):

137

138

```
139 # Create a new Raw instance  
140 raw = cs.io.Raw(  
141     df=data, # pandas.DataFrame  
142     activity=data['Activity'], # Activity time series  
143     light=data['Light'], # Light time series  
144     start_time=data.index[0], # Start time  
145     period=(data.index[-1]-data.index[0]), # Total duration  
146     frequency=data.index.freq # Sampling frequency  
147
```

148

149 **Reading actigraphy files using adaptor subclasses**

150 Although the `Raw` class is intentionally format-agnostic and flexible, *circStudio* provides
151 several adaptor subclasses for commonly used actigraphy formats. These adaptor classes, migrated
152 from *pyActigraphy*, enable the conversion of supported file types into `Raw` instances with minimal
153 effort. In the following example, a `.txt` file generated by an ActTrust (Condor Instruments)
154 device, included as a sample file, is loaded. To access the sample files distributed with *circStudio*,
155 a file path is first constructed:

156

157

```
158 fpath = os.path.join(os.path.dirname(cs.__file__))  
159
```

160

161 A `Raw` instance can then be created using the helper function `read_atr`:

162

163

```
164 raw = cs.io.read_atr(os.path.join(fpath, 'data', 'test_sample_atr.txt'))
```

165

166

167 Some ActTrust files contain an extra line above the header, (e.g., `#ActLogModel=2.0.0`)
168 that should be ignored during import. This can be handled using the `skip_rows` argument:

169

170

```
171 raw = cs.io.read_atr(os.path.join(fpath, 'data', 'test_sample_atr.txt'),  
172 skip_rows=1)
```

173

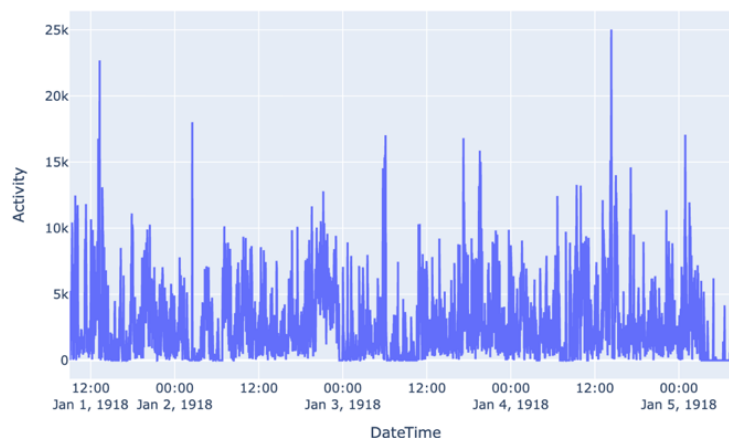
174

175 Once an original actigraphy file has been converted into a Raw object, the underlying time
176 series can be accessed directly. For example, the locomotor activity signal is stored as a
177 `pandas.Series` and can be retrieved via `raw.activity`, while the light intensity time series can
178 be accessed via `raw.light`. In both cases, the `plot()` method provides a convenient way to
179 quickly visualize activity and light recordings. For instance, the following command plots the
180 activity time series:

181

```
182 raw.plot(mode='activity', log=False)
```

183



184

Figure 1 Locomotor activity time series. Calling the `raw.plot(mode='activity')` method generates an interactive visualization of the activity signal. To get a similar visualization for the light intensity time series, use `raw.plot(mode='light')`.

185 **Preprocessing actigraphy data for subsequent analyzes**

186 **Discarding invalid sequences at the start and end of the recording**

187 To remove potentially invalid segments at the beginning and end of a recording, *circStudio*
188 follows an approach similar to that implemented in *pyActigraphy*. Briefly, users can exclude initial
189 portions of the recording by specifying `start_time` and remove terminal portions by defining the
190 duration to retain after `start_time` using the `period` argument:

191

192

```
193 raw = cs.io.read_awd(os.path.join(fpath, 'data', 'example_01.AWD'),  
194                     start_time='1918-01-24 08:00:00',  
195                     period='9 days'  
196
```

197

198 **Discarding invalid sequences during the recording**

199 *circStudio* allows users to mask spurious periods of inactivity within a recording by
200 marking them as `NaN`. Some downstream analyses, however, require continuous time series, such
201 as fitting a Cosinor model or simulating the response of the circadian system to a given light time
202 series. To address this, *circStudio* provides an imputation procedure that fills missing values using
203 the mean value from the corresponding times across other days in the recording. All masking
204 operations, including data imputation, are handled via the `apply_filters()` method, allowing
205 multiple filters to be applied with a single command.

206 A mask can be created for inactivity periods lasting at least 120 min:

207

208

```
209 raw.create_inactivity_mask(duration='120min')  
210
```

211

212 Custom masks can also be added from a CSV file containing *Mask*, *Start_time*, *Stop_time*, and
213 an optional label for each inactivity segment to be masked:

214

```
215 # Add mask periods from a CSV file  
216 raw.add_mask_periods(os.path.join(fpath, 'data', 'example_masklog.csv'))  
217  
218
```

219

220 The mask can then be applied and the resulting missing values imputed:

221

222

```
223 # Apply mask and impute NaN using the mean  
224 raw.apply_filters(apply_mask=True, impute_nan=True)
```

225

226

227 Additionally, `apply_filters()` also enables the signal to be resampled and binarized. For
228 instance, to binarize:

229

230

```
231 # Apply binarize data using a custom threshold of 4 activity counts  
232 raw.apply_filters(binarize=True, threshold=4)
```

233

234

235 To remove all filters and restore the original signal, the `reset_filters()` method can be used:

236

237

```
238 # Clear all the filters  
239 raw.reset_filters()
```

240

241

242 For some applications, such as using mathematical models of circadian rhythms, smoothing
243 may be useful to reduce the effect of extreme fluctuations in the input signal. Thus, *circStudio*
244 provides tools to smooth the activity and light series:

245

246

```
247 ts_smoothed = flm.smooth_timeseries(raw.activity, method='scott',  
248 verbose=True)
```

249

250

251 **Computing actigraphy-derived variables**

252 After completing preprocessing, a broad range of actigraphy variables can be calculated.
253 Examples include total average daily activity (ADAT), interdaily stability (IS), and the average

254 activity during the 10 most active hours of the day (M10). In contrast to *pyActigraphy*, metrics are
255 implemented as standalone functions requiring a data argument, which should be a *pandas.Series*
256 index by date and time. Of note, the time series used for these calculations should be preprocessed
257 beforehand, for example using `apply_filters()`.

258

259

```
260 adat(data=raw.activity)  
261 IS(data=raw.activity)  
262 m10(data=raw.activity)  
263
```

264

265 **Simulating mathematical models of circadian rhythms**

266 *circStudio* implements a suite of mathematical models of circadian rhythms developed to
267 predict human circadian physiology: `Forger` (Forger et al., 1999), `Jewett` (Jewett et al., 1999),
268 `HannaySP` and `HannayTP` (Hannay et al., 2019), `Hilaire07` (St. Hilaire et al., 2007), `Breslow13`
269 (Breslow et al., 2013), and `Skeldon23` (Skeldon et al., 2023). The model equations and parameters
270 were retrieved from the circadian package developed by the Arcascope team (Tavella et al., 2023).
271 The general workflow for these models parallels that used for metric computation: a light intensity
272 time series is first imported and preprocessed, after which a model instance is created. For example,
273 a new instance of the Forger model can be initialized using the following syntax:

274

275

```
276 # Create a new instance of the Forger model using light data  
277 forger = cs.Forger(data=raw.light)  
278
```

279

280 With the exception of `Hilaire07` and `Skeldon23`, model initialization syntax is consistent across
281 implementations:

282

283

```
284 # Jewett model  
285 jewett = cs.Jewett(data=raw.light)  
286  
287 # HannaySP model  
288 hannaysp = cs.HannaySP(data=raw.light)
```

```
289
290 # HannayTP model
291 hannaytp = cs.HannayTP(data=raw.light)
292
293 # Breslow13 model (melatonin model)
294 breslow13 = cs.Breslow13(data=raw.light)
295
```

296

297 The `Hilaire07` model requires a light time series and a sleep schedule as inputs. By
298 default, the sleep schedule is inferred using the Roenneberg algorithm, although a user-defined
299 schedule may also be provided via the `sleep_algo` argument. Any externally supplied sleep
300 schedule must share the same datetime index as `raw.light`. This argument can also be used to
301 select alternative sleep algorithms, such as Crespo:

302

303

```
304 # Hilaire07 model
305 hilaire07 = cs.Hilaire07(data=raw.light, sleep_algo='Roenneberg')
306
```

307

308 Before extracting phase markers, it is often useful to estimate new initial conditions from
309 the observed light time series. Because these models are formulated as systems of differential
310 equations, their numerical solution depends on the initial state. `circStudio` provides default initial
311 conditions to allow immediate use, but improved estimates can be obtained by iteratively looping
312 over the observed light time series and identifying a stable model state that yields consistent phase
313 markers, in this case, stable Dim Light Melatonin Onset (DLMO) predictions across successive
314 iterations. This can be achieved by specifying the schedule and the number of iterations:

315

316

```
317 # Loop over the provided light trajectory to determine new initial state
318 forger.get_initial_conditions(data=raw.light, loop_number=50)
319
```

320

321 Daily predictions of DLMO can then be extracted as:

322

323

324

```
325 # Obtain array of daily DLM0s
326 forger.dlmos() % 24
327
```

328

329 Model states can also be visualized using the `plot()` method:

330

331

```
332 forger.plot(states=True)
333
```

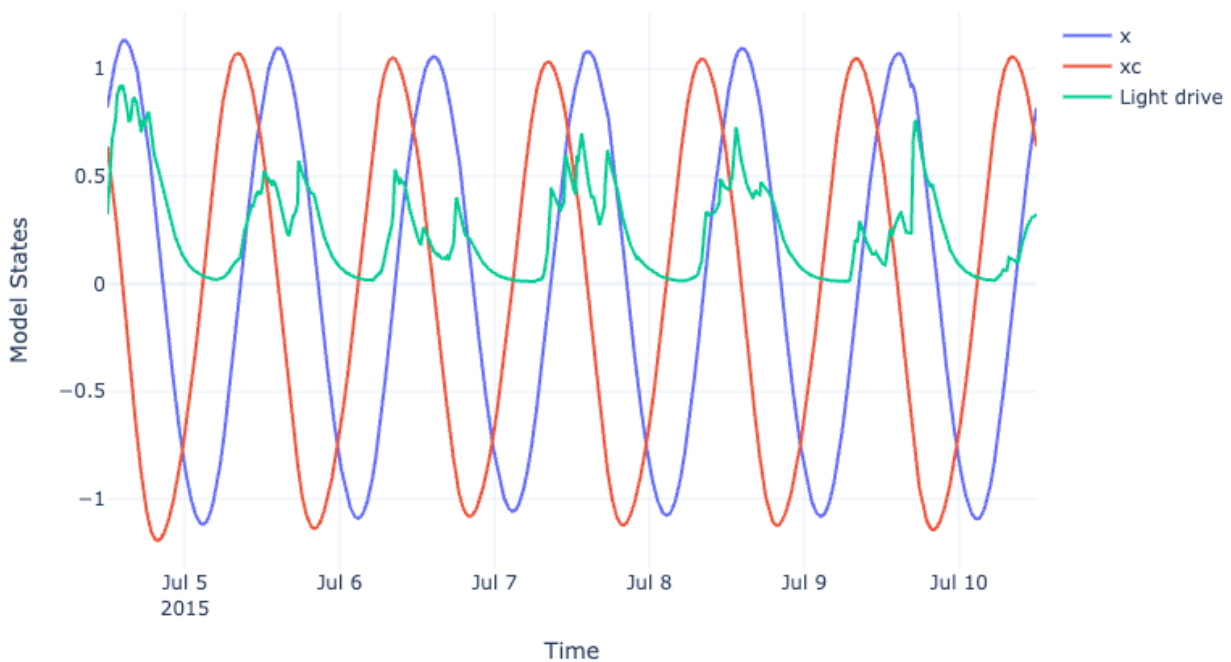


Figure 2 Interactive visualization of the three *Forger* model states (x , x_c and light drive) generated using the `plot()` method.

334

This is particularly useful for models whose state variables have physiological meaning, such as `breslow13`:

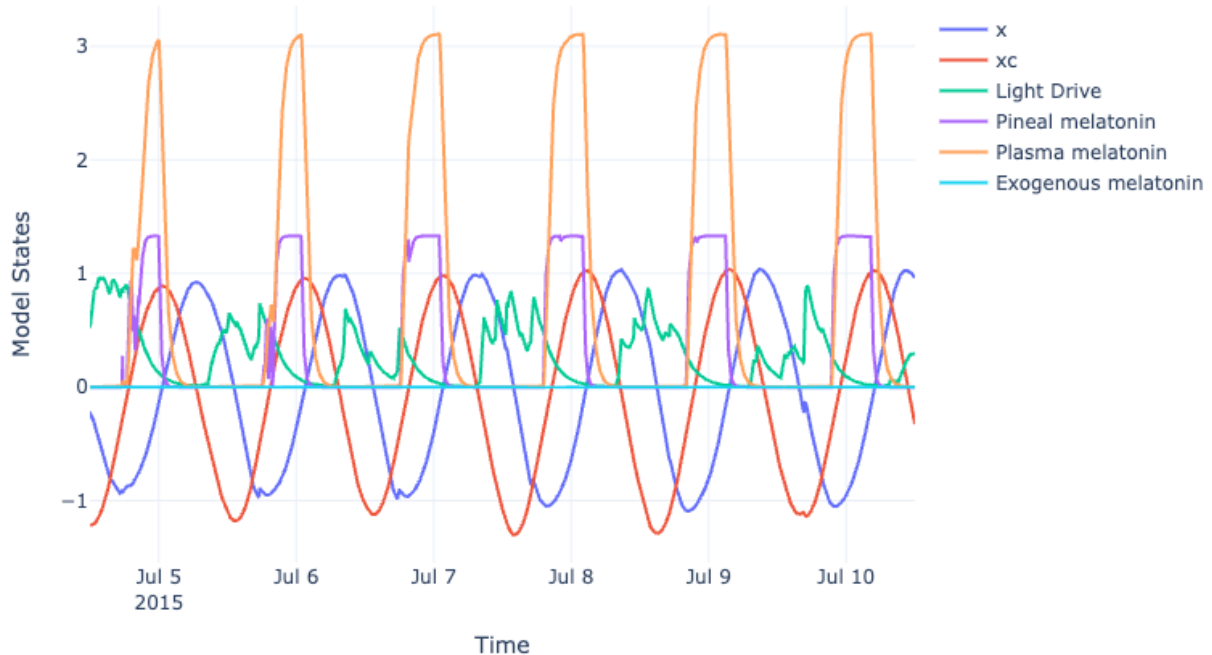
335

336

337

```
338 breslow13.plot(states=True)
339
```

340



341

342 **Figure 3 Interactive visualization of *Breslow13* model states.** This model is able to predict the melatonin trajectory in three
343 physiological compartments: pineal melatonin, plasma melatonin and exogenous melatonin.

344

345 **Predicting sleep pressure trajectory using the *Skeldon23* model**

346 The *Skeldon23* implementation follows a different workflow from the other models
347 because it includes feedback between continuous and discrete components. Briefly, *Skeldon23*
348 evolves four continuous state variables (x, x_c, n, h) through numerical integration, while the
349 sleep/wake state s is updated iteratively through a switching rule based on the current homeostatic
350 and circadian state. In practice, the sleep state is held constant within each time interval, the
351 continuous equations are integrated over that interval, and the sleep state is then updated at the
352 interval boundary. Given the nature of this implementation, *circStudio* separates model
353 instantiation from model execution in *Skeldon23*: creating the object defines parameters and
354 inputs, whereas calling `run()` performs the numerical simulation and populates the predicted
355 trajectories:

356

357

```
358 # Create new instance of Skeldon23 model  
359 skeldon23 = cs.Skeldon23(data=raw.light)
```

360

```
361 # Obtain new initial conditions based on the given light schedule
```

```
362 initial_conditions = skeldon23.get_initial_conditions(data=raw.light,  
363 loop_number=20)  
364  
365 # Run the model  
366 skeldon23.run(initial_state=initial_conditions)  
367  
368 # Plot the results  
369 skeldon23.plot(states=True)  
370
```

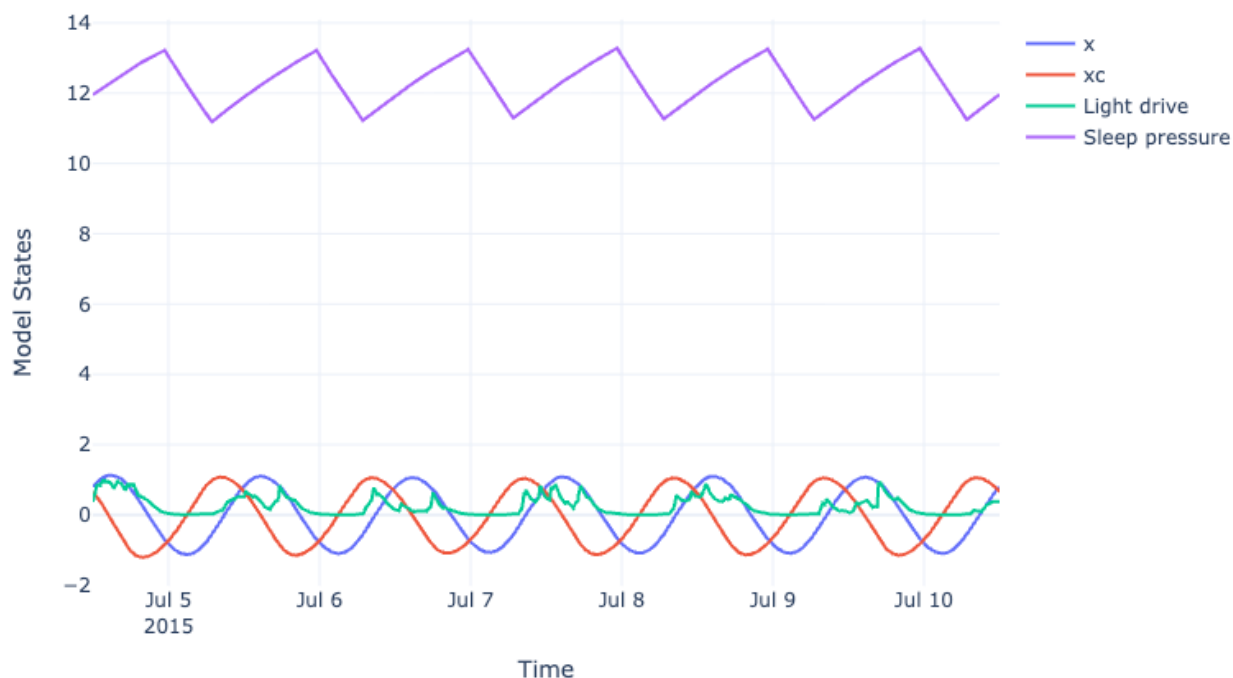


Figure 4 Visualization of model states of the *Skeldon23* model. This model is able to predict a trajectory corresponding to sleep pressure.

371
372 **Additional documentation and GitHub repository**
373 In addition to the workflows described above, circStudio provides further functionality for
374 Cosinor analysis, Singular Spectrum Analysis, (Multi-Fractal) Detrended Fluctuation Analysis and
375 Functional Linear Modeling. These methods were migrated from *pyActigraphy* and improved
376 documentation was written, including interactive examples:

377
378 **Table 1** Online resources available for *circStudio*, including tutorials, documentation, and the source code repository.

Resource	Link
Installation	https://djmarques.github.io/circStudio/tutorial_0.html

Loading actigraphy files	https://djmarques.github.io/circStudio/tutorial_1.html
Preprocessing actigraphy data	https://djmarques.github.io/circStudio/tutorial_2.html
Computing common metrics	https://djmarques.github.io/circStudio/tutorial_3.html
Mathematical models of circadian rhythms	https://djmarques.github.io/circStudio/tutorial_4.html
Cosinor analysis	https://djmarques.github.io/circStudio/tutorial_5.html
Singular spectrum analysis	https://djmarques.github.io/circStudio/tutorial_6.html
(Multi-fractal) Detrended Fluctuation Analysis	https://djmarques.github.io/circStudio/tutorial_7.html
Functional linear modelling	https://djmarques.github.io/circStudio/tutorial_8.html
API reference	https://djmarques.github.io/circStudio/api.html
Source code (GitHub repository)	https://github.com/djmarques/circStudio

379

380 **Applying *circStudio* to a practical case: submarine study**

381 To illustrate the use of *circStudio*, we considered a biological question derived from our
382 previous work on a submarine mission (Marques et al., 2025). In that study, we identified the Sleep
383 Regularity Index (SRI) and relative amplitude (RA) as candidate behavioral markers of shift-
384 specific organization in a military submarine environment. Building on this, we asked whether
385 reduced sleep regularity during time spent aboard the submarine was associated with lower relative
386 amplitude. This hypothesis was motivated by the expectation that irregular sleep-wake patterns
387 increase locomotor activity during the habitual sleep period, as measured by the average activity
388 levels during the least five hours of the day, thereby reducing the contrast between rest and activity
389 that underlies RA.

390 To answer this question, we first specify the directory containing the raw actigraphy files
391 corresponding to the submarine mission period (here denoted `source`). We then iterate over all
392 files in this directory:

393

```
394  
395 # Iterate over raw actigraphy recordings from submarine mission  
396 for filename in os.listdir(source):  
397     fpath = os.path.join(source, filename)  
398  
399
```

400 Within this loop, circadian metrics can be computed for each individual. It is possible to
401 obtain RA, L5 and SRI as follows:

402

```
403     ra = cs.ra(data=raw.activity)  
404     sri = cs.SleepRegularityIndex(data=raw.activity)  
405     l5 = cs.l5(data=raw.activity)[1]
```

407

408

409 The resulting metrics were stored and converted into a tabular structure for downstream
410 analysis. We then assessed the association between SRI and RA using Spearman's rank correlation
411 coefficient (SciPy), observing a moderate positive correlation ($\rho = 0.422$, $p = 0.025$). In addition,
412 SRI was negatively correlated with L5 ($\rho = -0.514$, $p = 0.025$), consistent with increased activity
413 during the least active period in individuals with more irregular sleep patterns. These findings are
414 consistent with the hypothesis that reduced sleep regularity may be associated with diminished
415 circadian rhythm robustness, potentially mediated by elevated nocturnal activity. However, given
416 the observational nature of the analysis, these results should be interpreted cautiously and primarily
417 as hypothesis-generating. This example illustrates how *circStudio* can be used to rapidly explore
418 biologically meaningful relationships in actigraphy data and to guide the formulation of testable
419 hypotheses in real-world settings.

420

421 **Discussion**

422 A major contribution of *circStudio* is the provision of a unified platform for actigraphy analysis
423 and data-driven simulation of circadian physiology within a single Python package. By combining
424 flexible preprocessing tools, established actigraphy metrics, and several mathematical models of
425 circadian rhythms, the package enables users to move efficiently from actimetry and wearable data
426 to physiologically interpretable circadian outputs. This unification facilitates the development of
427 health-oriented applications by providing direct access to research-grade analytical tools, while
428 also offering a flexible foundation for building custom analysis pipelines for researchers in
429 biological rhythms and neuroscience.

430 This integration of the actigraphy analysis capabilities of *pyActigraphy* with advanced
431 circadian modeling tools from the *circadian* package developed by Arcascope is consistent with
432 the original vision of *pyActigraphy*. Even though the *circStudio* codebase follows a different

433 architecture, this redesign contributes to the broader goal of establishing a comprehensive toolkit
434 for actigraphy analysis, and some functionalities developed in *circStudio* may be backported to
435 *pyActigraphy* in future releases. Future *circStudio* versions will also introduce standard algorithms
436 to automatically detect spurious periods of inactivity, thereby simplifying the identification and
437 removal of non-wear periods.

438 One potential limitation of *circStudio* is its specialization in actigraphy analysis. A possible
439 future direction would therefore be the development of a more general package for biological time
440 series analysis, providing a broader toolkit applicable to diverse time-resolved datasets in biology.
441 Additionally, even though the syntax is intentionally concise, the development of a graphical user
442 interface could further broaden the accessibility of *circStudio* across circadian and sleep research.

443

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