

# Package ‘pressuRe’

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**Type** Package

**Title** Imports, Processes, and Visualizes Biomechanical Pressure Data

**Version** 0.2.5

**Description** Allows biomechanical pressure data from a range of systems to be imported and processed in a reproducible manner. Automatic and manual tools are included to let the user define regions (masks) to be analyzed. Also includes functions for visualizing and animating pressure data. Example methods are described in Shi et al., (2022) <[doi:10.1038/s41598-022-19814-0](https://doi.org/10.1038/s41598-022-19814-0)>, Lee et al., (2014) <[doi:10.1186/1757-1146-7-18](https://doi.org/10.1186/1757-1146-7-18)>, van der Zward et al., (2014) <[doi:10.1186/1757-1146-7-20](https://doi.org/10.1186/1757-1146-7-20)>, Najafi et al., (2010) <[doi:10.1016/j.gaitpost.2009.09.003](https://doi.org/10.1016/j.gaitpost.2009.09.003)>, Cavanagh and Rodgers (1987) <[doi:10.1016/0021-9290\(87\)90255-7](https://doi.org/10.1016/0021-9290(87)90255-7)>.

**License** MIT + file LICENSE

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**Depends** R (>= 2.10)

**NeedsCompilation** no

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animate_pressure	<i>Animate pressure</i>
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---

### Description

Produces animation (gif) of pressure data

### Usage

```
animate_pressure(
  pressure_data,
  plot_colors = "default",
  fps,
  dpi = 96,
  file_name
)
```

### Arguments

pressure_data	Array. A 3D array covering each timepoint of the measurement. z dimension represents time
plot_colors	String
fps	Numeric. Number of frames per second in animation
dpi	Numeric. Resolution of gif
file_name	Name (including path) of export file

**Value**

Animation in gif format

**Examples**

```
emed_data <- system.file("extdata", "emed_test.lst", package = "pressuRe")
pressure_data <- load_emed(emed_data)
animate_pressure(pressure_data, fps = 10, file_name = "pli_gif.gif")
```

---

arch_index	<i>Calculate Arch Index.</i>
------------	------------------------------

---

**Description**

Calculate Arch Index.

**Usage**

```
arch_index(pressure_data, plot = TRUE)
```

**Arguments**

pressure_data	List. Includes a 3D array covering each timepoint of the measurement. z dimension represents time
plot	Logical. Not implemented yet

**Value**

Numeric. Arch index value

**Examples**

```
emed_data <- system.file("extdata", "emed_test.lst", package = "pressuRe")
pressure_data <- load_emed(emed_data)
arch_index(pressure_data)
```

---

auto_detect_side	<i>Detect foot side</i>
------------------	-------------------------

---

**Description**

Detects which foot plantar pressure data is from (left or right), usually would only be needed for barefoot pressure plate data. Generally reliable but may be thrown off by severe deformities or abnormal walking patterns

**Usage**

```
auto_detect_side(pressure_data)
```

**Arguments**

pressure\_data List. First item should be a 3D array covering each timepoint of the measurement. z dimension represents time

**Value**

String. "LEFT" or "RIGHT"

**Examples**

```
emed_data <- system.file("extdata", "emed_test.lst", package = "pressuRe")
pressure_data <- load_emed(emed_data)
auto_detect_side(pressure_data)
```

---

cop	<i>Center of pressure</i>
-----	---------------------------

---

**Description**

Generates xy coordinates for center of pressure during each frame of measurement

**Usage**

```
cop(pressure_data)
```

**Arguments**

pressure\_data List. First item is a 3D array covering each timepoint of the measurement. z dimension represents time

**Value**

Data frame with x and y coordinates of COP throughout trial

**Examples**

```
emed_data <- system.file("extdata", "emed_test.lst", package = "pressuRe")
pressure_data <- load_emed(emed_data)
cop(pressure_data)
```

---

cpei

*CPEI*

---

**Description**

Determine Center of Pressure Excursion Index (CPEI) for footprint pressure data

**Usage**

```
cpei(pressure_data, foot_side, plot_result = TRUE)
```

**Arguments**

**pressure\_data** List. First item is a 3D array covering each timepoint of the measurement. Not currently available for pedar.

**foot\_side** String. "right" or "left". Required for automatic detection of points

**plot\_result** Logical. Plots pressure image with COP and CPEI overlaid

**Value**

Numeric. CPEI value

**Author(s)**

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**Examples**

```
emed_data <- system.file("extdata", "emed_test.lst", package = "pressuRe")
pressure_data <- load_emed(emed_data)
cpei(pressure_data, foot_side = "auto", plot_result = FALSE)
```

---

create_mask_auto	<i>Automatically mask pressure footprint</i>
------------------	--

---

## Description

Automatically creates mask for footprint data

## Usage

```
create_mask_auto(
  pressure_data,
  masking_scheme,
  foot_side = "auto",
  res_value = c(20000, 20000, 1e+05, 20000),
  plot = TRUE,
  template_mask = NULL
)
```

## Arguments

pressure_data	List. First item is a 3D array covering each timepoint of the measurement. z dimension represents time
masking_scheme	String. "automask_simple", "automask_novel", "pedar_mask1", "pedar_mask2", "pedar_mask3", "template". "simple_automask" applies a simple 3 part mask (hindfoot, midfoot, forefoot) "automask_novel" attempts to apply a 9-part mask (hindfoot, midfoot, mets, hallux, lesser toes), similar to the standard novel automask "pedar_mask1" splits the insole into 4 regions using sensel boundaries: hindfoot, midfoot, forefoot, and toes- <a href="https://www.ncbi.nlm.nih.gov/pmc/articles/PMC9470545/">https://www.ncbi.nlm.nih.gov/pmc/articles/PMC9470545/</a> "pedar_mask2" splits the insole into 4 regions using percentages: hindfoot, forefoot, hallux, and lesser toes- <a href="https://jfootankleres.biomedcentral.com/articles/10.1186/1757-1146-7-18">https://jfootankleres.biomedcentral.com/articles/10.1186/1757-1146-7-18</a> "pedar_mask3" splits the foot into 9 regions using sensel boundaries: medial hindfoot, lateral hindfoot, medial midfoot, lateral midfoot, MTPJ1, MTPJ2-3, MTPJ4-5, hallux, and lesser toes- <a href="https://jfootankleres.biomedcentral.com/articles/10.1186/1757-1146-7-20">https://jfootankleres.biomedcentral.com/articles/10.1186/1757-1146-7-20</a>
foot_side	String. "RIGHT", "LEFT", or "auto". Auto uses auto_detect_side function
res_value	Numeric vector. Adjusting these values can help if the heel, midfoot, toe, and hallux lines aren't correct. Default values are c(10000, 10000, 100000, 10000). These lines are calculated using a least cost function and the parameter essentially adjusts the resistance of the pressure value for that algorithm
plot	Logical. Whether to play the animation
template_mask	Data frame. Mask to be used if "template_mask" is selected as the masking scheme

**Value**

List. Masks are added to pressure data variable

- `pressure_array`. 3D array covering each timepoint of the measurement. z dimension represents time
- `pressure_system`. String defining pressure system
- `sens_size`. Numeric vector with the dimensions of the sensors
- `time`. Numeric value for time between measurements
- `masks`. List
- `events`. List

**Examples**

```
emed_data <- system.file("extdata", "emed_test.lst", package = "pressuRe")
pressure_data <- load_emed(emed_data)
pressure_data <- create_mask_auto(pressure_data, "automask_novel",
res_value = c(20000, 20000, 100000, 20000), foot_side = "auto", plot = FALSE)
```

---

create\_mask\_manual      *Create masking*

---

**Description**

Allows user to manually define mask regions

**Usage**

```
create_mask_manual(
  pressure_data,
  mask_definition = "by_vertices",
  n_masks = 1,
  n_verts = 4,
  n_sens = 4,
  threshold = 0.005,
  plot_existing_masks = TRUE,
  mask_names = "default",
  plot = TRUE
)
```

**Arguments**

`pressure_data` List. First item is a matrix covering each timepoint of the measurement.

`mask_definition`

String. "by\_vertices" or "by\_sensors". The first option let's you draw a shape around the area you want to select, the second allows you to define this area by clicking on specific sensors

n_masks	Numeric. Number of masks to add
n_verts	Numeric. Number of vertices in mask
n_sens	Numeric. Number of sensors mask will contain
threshold	Numeric. Distance between adjacent mask vertices before sharing vertex coordinates
plot_existing_masks	Logical. Show existing masks
mask_names	List. Mask names. Default is "custom_mask#"
plot	Logical. Show new maks on pressure image

### Value

List. Mask(s) are added to pressure data variable

- pressure\_array. 3D array covering each timepoint of the measurement. z dimension represents time
- pressure\_system. String defining pressure system
- sens\_size. Numeric vector with the dimensions of the sensors
- time. Numeric value for time between measurements
- masks. List
- events. List
- sensor\_polygons. Data frame with corners of sensors
- max\_matrix Matrix with maximum image

### Examples

```

emed_data <- system.file("extdata", "emed_test.lst", package = "pressuRe")
pressure_data <- load_emed(emed_data)
pressure_data <- create_mask_manual(pressure_data, mask_definition = "by_vertices",
n_masks = 1, n_verts = 4)
pressure_data <- create_mask_manual(pressure_data, mask_definition = "by_sensors",
n_masks = 1, n_sens = 8)

```

---

edit\_mask

*Edit mask*

---

### Description

Allows user to manually adjust mask vertices



**Usage**

```
edit_mask(
  pressure_data,
  n_edit,
  threshold = 0.002,
  edit_list = seq(1, length(pressure_data[[5]])),
  image = "max"
)
```

**Arguments**

pressure_data	List. First item is a 3D array covering each timepoint of the measurement.
n_edit	Numeric. Number of vertices to edit
threshold	Numeric. Distance between point clicked and vertex that is selected
edit_list	List. Mask numbers that want to be edited. (Default is to load all masks so that adjacent masks with shared coordinates are modified together)
image	String. "max" = footprint of maximum sensors. "mean" average value of sensors over time

**Value**

List. Edited mask is added to the pressure data variable

- pressure\_array. 3D array covering each timepoint of the measurement. z dimension represents time
- pressure\_system. String defining pressure system
- sens\_size. Numeric vector with the dimensions of the sensors
- time. Numeric value for time between measurements
- masks. List
- events. List
- sensor\_polygons. Data frame with corners of sensors
- max\_matrix Matrix with maximum image

**Examples**

```
emed_data <- system.file("extdata", "emed_test.lst", package = "pressuRe")
pressure_data <- load_emed(emed_data)
pressure_data <- create_mask_auto(pressure_data, "automask_novel",
  foot_side = "auto", plot = FALSE)
pressure_data <- edit_mask(pressure_data, n_edit = 1, threshold = 0.002,
  image = "max")
```

---

footprint	<i>Footprint</i>
-----------	------------------

---

**Description**

Determines footprint of pressure data

**Usage**

```
footprint(pressure_data, variable = "max", frame = NULL, plot = FALSE)
```

**Arguments**

pressure_data	List. Includes a 3D array covering each timepoint of the measurement. z dimension represents time
variable	String. "max" = maximum value of each sensor across full dataset. "mean" = average value of sensors over full dataset. "frame" = an individual pressure frame. "meanmax" average max values across cycles ( currently just for pedar)
frame	Integer. Only used if variable = "frame".
plot	Logical. Display pressure image

**Value**

Matrix. Maximum or mean values for all sensors

**Examples**

```
emed_data <- system.file("extdata", "emed_test.lst", package = "pressuRe")
pressure_data <- load_emed(emed_data)
footprint(pressure_data, plot = FALSE)
```

---

load_emed	<i>Load emed data</i>
-----------	-----------------------

---

**Description**

Imports and formats .lst files collected on emed system and exported from Novel software

**Usage**

```
load_emed(pressure_filepath, trim_active = FALSE)
```

**Arguments**

pressure_filepath	String. Filepath pointing to emed pressure file
trim_active	Logical. Restricts frames to only the first continuous foot contact

**Value**

A list with information about the pressure data.

- `pressure_array`. 3D array covering each timepoint of the measurement. z dimension represents time
- `pressure_system`. String defining pressure system
- `sens_size`. Numeric vector with the areas of the sensors
- `time`. Numeric value for time between measurements
- `masks`. List
- `events`. List
- `sensor_polygons`. Data frame with corners of sensors
- `max_matrix` Matrix with maximum image

**Examples**

```
emed_data <- system.file("extdata", "emed_test.lst", package = "pressuRe")
pressure_data <- load_emed(emed_data)
```

---

load_footscan	<i>Load footscan data</i>
---------------	---------------------------

---

**Description**

Imports and formats files collected on footscan systems (formerly RSScan)

**Usage**

```
load_footscan(pressure_filepath)
```

**Arguments**

`pressure_filepath`  
String. Filepath pointing to emed pressure file

**Value**

A list with information about the pressure data.

- `pressure_array`. 3D array covering each timepoint of the measurement. z dimension represents time
- `pressure_system`. String defining pressure system
- `sens_size`. Numeric vector with the dimensions of the sensors
- `time`. Numeric value for time between measurements
- `masks`. List
- `events`. List
- `sensor_polygons`. Data frame with corners of sensors
- `max_matrix`. Matrix

**Examples**

```

footscan_data <- system.file("extdata", "footscan_test.xls", package = "pressuRe")
pressure_data <- load_footscan(footscan_data)

```

---

load\_pedar

*Load pedar data*


---

**Description**

Imports and formats .asc files collected on pedar system and exported from Novel software

**Usage**

```
load_pedar(pressure_filepath)
```

**Arguments**

```

pressure_filepath
      String. Filepath pointing to pedar pressure file

```

**Value**

A list with information about the pressure data.

- `pressure_array`. 3D array covering each timepoint of the measurement. z dimension represents time
- `pressure_system`. String defining pressure system
- `sens_size`. String with sensor type
- `time`. Numeric value for time between measurements
- `masks`. List
- `events`. List
- `sensor_polygons`. Data frame with corners of sensors
- `max_matrix` Matrix with maximum image

**Examples**

```

pedar_data <- system.file("extdata", "pedar_example.asc", package = "pressuRe")
pressure_data <- load_pedar(pedar_data)

```

---

load_pliance	<i>Load pliance data</i>
--------------	--------------------------

---

## Description

Imports and formats .asc files collected on pliance system and exported from Novel software

## Usage

```
load_pliance(pressure_filepath)
```

## Arguments

pressure\_filepath  
String. Filepath pointing to pliance pressure file

## Value

A list with information about the pressure data.

- pressure\_array. 3D array covering each timepoint of the measurement. z dimension represents time
- pressure\_system. String defining pressure system
- sens\_size. String with sensor type
- time. Numeric value for time between measurements
- masks. List
- events. List
- sensor\_polygons. Data frame with corners of sensors
- max\_matrix. Matrix

## Examples

```
pliance_data <- system.file("extdata", "pliance_test.asc", package = "pressuRe")  
pressure_data <- load_pliance(pliance_data)
```

---

load_tekscan	<i>Load Tekscan data</i>
--------------	--------------------------

---

### Description

Imports and formats files collected on tekscan systems and exported from Tekscan software

### Usage

```
load_tekscan(pressure_filepath)
```

### Arguments

pressure\_filepath  
String. Filepath pointing to emed pressure file

### Value

A list with information about the pressure data.

- pressure\_array. 3D array covering each timepoint of the measurement. z dimension represents time
- pressure\_system. String defining pressure system
- sens\_size. Numeric vector with the dimensions of the sensors
- time. Numeric value for time between measurements
- masks. List
- events. List
- sensor\_polygons. Data frame with corners of sensors
- max\_matrix. Matrix

### Examples

```
tekscan_data <- system.file("extdata", "fscan_testL.asf", package = "pressuRe")  
pressure_data <- load_tekscan(tekscan_data)
```

---

load_xsensor	<i>Load xsensor data</i>
--------------	--------------------------

---

### Description

Imports and formats files collected on xsensor insole systems

### Usage

```
load_xsensor(pressure_filepath)
```

### Arguments

pressure\_filepath  
String. Filepath pointing to emed pressure file

### Value

A list with information about the pressure data.

- pressure\_array. 2D array covering each timepoint of the measurement. row dimension represents time
- pressure\_system. String defining pressure system
- sens\_size. Numeric vector with the dimensions of the sensors
- time. Numeric value for time between measurements
- masks. List
- events. List
- sensor\_polygons. Data frame with corners of sensors
- max\_matrix. Matrix

### Examples

```
xsensor_data <- system.file("extdata", "xsensor_data.csv", package = "pressuRe")  
pressure_data <- load_xsensor(xsensor_data)
```

---

mask_analysis	<i>Analyze masked regions of pressure data</i>
---------------	--

---

## Description

Analyze masked regions of pressure data

## Usage

```
mask_analysis(
  pressure_data,
  partial_sensors = FALSE,
  variable = "press_peak_sensor",
  pressure_units = "kPa",
  area_units = "cm2"
)
```

## Arguments

pressure_data	List. Includes a 3D array covering each timepoint of the measurement. z dimension represents time
partial_sensors	Logical Defines how sensors that do not lie wholly within mask are dealt with. If FALSE, they will be excluded; if TRUE, for relevant variables their contribution will be weighted by the proportion of the sensor that falls within the mask border
variable	String. Variable to be determined. "press_peak_sensor", "press_peak_mask", "contact_area_peak", "pti_1", "pti_2", "force_time_integral", "force_peak", "dpli", "press_peak_sensor_ts", "force_ts". Variables ending in "_ts" produce time series data
pressure_units	String. Default "kPa". Other options: "MPa", "Ncm2" (Newtons per square centimeter)
area_units	String. Default is "cm2" (square centimeters). Other options "m2" (square meters); "mm2" (square millimeters)

## Value

Data frame. Contains values for each mask plus additional information relevant to the data including cycle/step and foot side

## Examples

```
emed_data <- system.file("extdata", "emed_test.lst", package = "pressuRe")
pressure_data <- load_emed(emed_data)
pressure_data <- create_mask_auto(pressure_data, "automask_simple", plot = FALSE)
mask_analysis(pressure_data, FALSE, variable = "press_peak_sensor")
```



---

pedar\_insole\_areas     *pedar sensor size data*

---

**Description**

Sensor sizes for different pedar insoles

**Usage**

pedar\_insole\_areas

**Format**

## 'pedar\_insole\_areas' A data frame with 198 rows and 8 columns:

**u** areas for size u

**v** areas for size v

**w** areas for size w

**x** areas for size x

**y** areas for size y

**uw** areas for size uw

**xw** areas for size xw

**vw** areas for size vw

**Source**

Scott Telfer

---

pedar\_insole\_grid     *pedar sensor grid data*

---

**Description**

Sensor outline coordinates

**Usage**

pedar\_insole\_grid

**Format**

## 'pedar\_insole\_grid' A data frame with 199 rows and 8 columns:

**V1** x1 coordinate

**V2** y1 coordinate

**V3** x2 coordinate

**V4** y2 coordinate

**V5** x3 coordinate

**V6** y3 coordinate

**V7** x4 coordinate

**V8** y4 coordinate

**Source**

Scott Telfer

---

plot\_pressure

*Plot pressure*

---

**Description**

Produces visualization of pressure data

**Usage**

```
plot_pressure(  
  pressure_data,  
  variable = "max",  
  smooth = FALSE,  
  frame,  
  step_n = "max",  
  plot_COP = FALSE,  
  plot_outline = FALSE,  
  plot_masks = FALSE,  
  plot_colors = "default",  
  break_values,  
  break_colors,  
  sensor_outline = TRUE,  
  plot = TRUE,  
  legend = TRUE  
)
```

**Arguments**

pressure_data	List. Includes a 3D array covering each timepoint of the measurement. z dimension represents time
variable	String. "max" = footprint of maximum sensors. "mean" = average value of sensors over time (usually for static analyses). "frame" = an individual frame
smooth	Logical. Not implemented. If TRUE, plot will interpolate between sensors to increase data density
frame	Integer.
step_n	If numeric, the step number to plot (only for insole data). If "max", the max across complete trial, if "meanmax", the max on a per step basis
plot_COP	Logical. If TRUE, overlay COP data on plot. Default = FALSE
plot_outline	Logical. If TRUE, overlay convex hull outline on plot
plot_masks	Logical. If TRUE, overlay mask outline on plot
plot_colors	String. "default": novel color scheme; "custom": user supplied
break_values	Vector. If plot_colors is "custom", values to split colors at
break_colors	Vector. If plot_colors is "custom", colors to use. Should be one shorter than break_values
sensor_outline	Logical. Sensor outline to be shown
plot	Logical. If TRUE, plot will be displayed
legend	Logical. If TRUE, legend will be added to plot

**Value**

ggplot plot object

**Examples**

```

emed_data <- system.file("extdata", "emed_test.lst", package = "pressuRe")
pressure_data <- load_emed(emed_data)
plot_pressure(pressure_data, variable = "max", plot_COP = FALSE)
plot_pressure(pressure_data, variable = "frame", frame = 20,
              plot_colors = "custom", break_values = c(100, 200, 300),
              break_colors = c("light blue", "light green", "yellow", "pink"))

```

---

pressure\_interp      *Interpolate pressure data*

---

**Description**

Resamples pressure data over time. Useful for normalizing to stance phase, for example

**Usage**

```
pressure_interp(pressure_data, interp_to)
```

**Arguments**

- pressure\_data List. First item should be a 3D array covering each timepoint of the measurement. z dimension represents time.
- interp\_to Integer. Number of frames to interpolate to

**Value**

- pressure\_array. 3D array covering each timepoint of the measurement. z dimension represents time
- pressure\_system. String defining pressure system
- sens\_size. Numeric vector with the dimensions of the sensors
- time. Numeric value for time between measurements
- masks. List
- events. List
- sensor\_polygons. Data frame with corners of sensors
- max\_matrix. Matrix

**Examples**

```
emed_data <- system.file("extdata", "emed_test.lst", package = "pressuRe")
pressure_data <- load_emed(emed_data)
pressure_data <- pressure_interp(pressure_data, interp_to = 101)
```

---

select_steps	<i>Select steps</i>
--------------	---------------------

---

**Description**

Select steps, usually from insole data, and format for analysis

**Usage**

```
select_steps(  
  pressure_data,  
  threshold = "auto",  
  min_frames = 10,  
  n_steps = 5,  
  skip = 2  
)
```

**Arguments**

pressure_data	List. First item should be a 3D array covering each timepoint of the measurement. z dimension represents time.
threshold	Numeric. Threshold force to define start and end of step. If "auto", function will set threshold at minimum force in trial + 10N
min_frames	Numeric. Minimum number of frames that need to be in step
n_steps	Numeric. Target number of steps/cycles. User will be asked to keep selected steps until this target is reached or they run out of candidate steps
skip	Numeric. Usually the first few steps of a trial are accelerating and not representative of steady state walking so this removes them

**Value**

- pressure\_array. 3D array covering each timepoint of the measurement. z dimension represents time
- pressure\_system. String defining pressure system
- sens\_size. Numeric vector with the dimensions of the sensors
- time. Numeric value for time between measurements
- masks. List
- events. Data frame
- sensor\_polygons. Data frame with corners of sensors
- max\_matrix. Matrix

**Examples**

```
pedar_data <- system.file("extdata", "pedar_example.asc", package = "pressuRe")
pressure_data <- load_pedar(pedar_data)
pressure_data <- select_steps(pressure_data)
```

---

whole\_pressure\_curve    *Whole pressure curve*

---

**Description**

Generates vectors with option to plot for force, peak/mean pressure and area for complete measurement. Useful for checking data

**Usage**

```
whole_pressure_curve(
  pressure_data,
  variable,
  side,
  threshold = 10,
  plot = FALSE
)
```

**Arguments**

<code>pressure_data</code>	List. A 3D array covering each timepoint of the measurement. z dimension represents time
<code>variable</code>	String. "peak_pressure", "force", or "area"
<code>side</code>	For insole data only
<code>threshold</code>	Numeric. Threshold value for sensor to be considered active. Currently only applies to insole data
<code>plot</code>	Logical. If TRUE also plots data as line curve

**Value**

Numeric vector containing variable values

**Examples**

```
emed_data <- system.file("extdata", "emed_test.lst", package = "pressuRe")
pressure_data <- load_emed(emed_data)
whole_pressure_curve(pressure_data, variable = "peak_pressure", plot = FALSE)
whole_pressure_curve(pressure_data, variable = "area", plot = FALSE)
whole_pressure_curve(pressure_data, variable = "force", plot = FALSE)
```

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