

# Package ‘xvm’

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**Title** Read, Parse and Visualize 'XVG'/'XPM' Files from Molecular Dynamics

**Version** 0.0.2

## Description

Provides tools for reading, parsing and visualizing simulation data stored in 'xvg'/'xpm' file formats (commonly generated by 'GROMACS' molecular dynamics software). Streamlines post-processing and analysis of molecular dynamics ('MD') simulation outputs, enabling efficient exploration of molecular stability and conformational changes. Supports import of trajectory metrics ('RMSD', energy, temperature) and creation of publication-ready visualizations through integration with 'ggplot2'.

**URL** <https://github.com/RightSZ/xvm>, <https://rightsz.github.io/xvm/>

**BugReports** <https://github.com/RightSZ/xvm/issues>

**Encoding** UTF-8

**Roxygen** list(markdown = TRUE)

**RoxygenNote** 7.3.2

**Imports** ggplot2, tidyr, ggnewscale, plotly

**Suggests** knitr, rmarkdown, ggpubr, stringr

**VignetteBuilder** knitr

**License** GPL (>= 3)

**NeedsCompilation** no

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export_xvg	<i>export xvg data</i>
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## Description

write the data component of an xvg\_data object (or multiple objects) to a delimited text file, controlled via the sep parameter rather than file extension detection.

## Usage

```
export_xvg(xvg_data, file, sep = "\t", row.names = FALSE, merge = FALSE, ...)
```

## Arguments

xvg_data	An object of class xvg_data, or a list of xvg_data objects, as returned by read_xvg().
file	Path to the output file (any extension is acceptable).
sep	Field separator (e.g., "\t" for TSV, "," for CSV). Default is "\t".
row.names	Logical, whether to write row names. Default is FALSE.
merge	Logical, whether to merge multiple xvg_data objects before exporting. Default is FALSE.
...	Additional arguments passed to write.table().

## Value

Invisibly returns the path to the written file.

## Examples

```
## Not run:
xvg <- read_xvg(system.file("extdata/rmsd.xvg", package = "xvm"))
# Export as TSV
export_xvg(xvg, "rmsd.tsv", sep = "\t")
# Export as CSV
export_xvg(xvg, "rmsd.csv", sep = ",")

## End(Not run)
```

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format_text	<i>format special text</i>
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**Description**

processes special formatting in xvg files, converting super/subscripts to ggplot2-compatible expressions. Specifically handles:

- Superscripts: Converts \S...\N to ^ notation
- Subscripts: Converts \s...\N to [] notation

**Usage**

```
format_text(text)
```

**Arguments**

text	character string containing xvg-formatted text
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**Value**

formatted character string with ggplot2-compatible expressions

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merge_xvg_data	<i>merge multiple xvg data objects</i>
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**Description**

combines multiple xvg data objects into a single structure, preserving metadata from the first object and adding a group identifier to track the source of each data point.

**Usage**

```
merge_xvg_data(xvg_data)
```

**Arguments**

xvg_data	a list of xvg data objects, each containing 'data' and 'metadata' components
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**Value**

a merged xvg data object with:

- data - Combined data frame with an additional 'group' column identifying the source
- metadata - Metadata from the first object in the list

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parse_xpm	<i>parse xpm (X PixMap) file content into structured data</i>
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### Description

This function parses xpm file content, extracts metadata, color mappings, and matrix data, returning a structured list for further processing.

### Usage

```
parse_xpm(xpm_content)
```

### Arguments

xpm\_content      a character string containing the xpm file content.

### Value

a list with the following components:

- data - Data frame containing matrix values with coordinates
- title - Chart title extracted from xpm
- legend - Legend text extracted from xpm
- x\_label - X-axis label extracted from xpm
- y\_label - Y-axis label extracted from xpm
- color\_map - Named list mapping color codes to hex values
- color\_values - Named list mapping color codes to numeric values

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parse_xvg	<i>parse xvg File Content</i>
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### Description

parses content from a single GROMACS-generated xvg file

### Usage

```
parse_xvg(lines, skip_comments = TRUE)
```

### Arguments

lines              character vector of text lines from xvg file  
 skip\_comments    logical indicating whether to skip comment lines (default: TRUE)

### Value

list containing xvg data and metadata with following structure:

- data - Data frame containing numerical data
- metadata - List containing title, axis labels, legends and their formatted versions

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plot_xpm	<i>plot xpm data</i>
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**Description**

plot xpm data using ggplot2

**Usage**

```
plot_xpm(xpm_data, interpolate = FALSE)
```

**Arguments**

xpm_data	a xpm object returned by read_xpm
interpolate	logical indicating whether to use raster interpolation (TRUE) or discrete tiles (FALSE). Default is FALSE.

**Value**

a ggplot2 object

**Examples**

```
library(xvm)
xpm_file_path <- system.file("extdata/gibbs.xpm", package = "xvm")
xpm_data <- read_xpm(xpm_file_path)
plot_xpm(xpm_data) # plot the xpm data using plot_xpm() function
```

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plot_xpm_3d	<i>generate 3d scatter plot from xpm Data</i>
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**Description**

creates 3d visualization of xpm data with scatter plot.

**Usage**

```
plot_xpm_3d(xpm_data, reversescale = FALSE, point_size = 2)
```

**Arguments**

xpm_data	a xpm object (from <a href="#">read_xpm()</a> ) or list containing parsed objects.
reversescale	whether to reverse the color scale; default is FALSE
point_size	the size of the points in the scatter plot; default is 2

**Value**

a plotly object

## Examples

```
library(xvm)
xpm_file_path <- system.file("extdata/gibbs.xpm", package = "xvm")
xpm_data <- read_xpm(xpm_file_path)
plot_xpm_3d(xpm_data) # plot 3D scatter plot from xpm file
```

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plot_xpm_facet	<i>generate faceted plots from xpm Data</i>
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## Description

creates dual-panel visualizations of xpm data with scatter or area plots.

## Usage

```
plot_xpm_facet(xpm_data, plot_type = "scatter")
```

## Arguments

xpm_data	a xpm object (from <a href="#">read_xpm()</a> ) or list containing parsed objects.
plot_type	visualization type: "scatter" (default) or "area".

## Value

a ggplot2 object with:

- Dual facets showing x/y axis relationships
- Automatic data transformation for visualization
- NULL if invalid plot\_type specified

## Examples

```
library(xvm)
xpm_file_path <- system.file("extdata/gibbs.xpm", package = "xvm")
xpm_data <- read_xpm(xpm_file_path)
plot_xpm_facet(xpm_data) # plot pseudo-3D from xpm file
```

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`plot_xvg`*plot xvg data*

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## Description

plot xvg data using ggplot2

## Usage

```
plot_xvg(  
  xvg_data,  
  merge = FALSE,  
  title = NULL,  
  subtitle = NULL,  
  use_color_scale = NULL,  
  ...  
)
```

## Arguments

<code>xvg_data</code>	xvg data object returned by <code>read_xvg</code>
<code>merge</code>	logical; if TRUE and multiple datasets provided, merge them (default: FALSE)
<code>title</code>	chart title (default uses xvg file's title)
<code>subtitle</code>	chart subtitle (default uses xvg file's subtitle)
<code>use_color_scale</code>	custom color scale function (e.g., <code>ggsci::scale_color_bmj</code> ) to override default colors
<code>...</code>	additional parameters passed to <code>ggplot2::geom_line</code>

## Value

a ggplot2 object

## Examples

```
library(xvm)  
rmsd_file_path <- system.file("extdata/rmsd.xvg", package = "xvm")  
rmsd_data <- read_xvg(rmsd_file_path)  
plot_xvg(rmsd_data) # plot the xvg data using plot_xvg() function
```

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`read_xpm`*read xpm files*

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## Description

This function reads xpm (X PixMap) files, validates their existence, and returns parsed data structures in a list format.

## Usage

```
read_xpm(xpm_files)
```

## Arguments

`xpm_files`          a character vector containing paths to one or more xpm files.

## Details

The function performs the following operations:

1. Validates input type (must be character vector)
2. Checks for file existence and filters missing files with warnings
3. Reads valid files and parses them using [parse\\_xpm\(\)](#)
4. Returns aggregated results in a named list

## Value

list with the following components:

- `data` - Data frame containing matrix values with coordinates
- `title` - Chart title extracted from xpm
- `legend` - Legend text extracted from xpm
- `x_label` - X-axis label extracted from xpm
- `y_label` - Y-axis label extracted from xpm
- `color_map` - Named list mapping color codes to hex values
- `color_values` - Named list mapping color codes to numeric values

## Examples

```
library(xvm)
# Retrieve the path to the example file included in the package
xpm_file_path <- system.file("extdata/gibbs.xpm", package = "xvm")
xpm_data <- read_xpm(xpm_file_path) # read the xpm file using read_xpm() function
names(xpm_data)
```



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read_xvg	<i>read xvg files</i>
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**Description**

read one or more GROMACS-generated xvg files

**Usage**

```
read_xvg(xvg_files, skip_comments = TRUE)
```

**Arguments**

xvg_files	character vector of xvg file paths
skip_comments	logical indicating whether to skip comment lines (default: TRUE)

**Value**

Named list containing xvg data, using filenames (without extension) as keys

**Examples**

```
library(xvm)
# Retrieve the path to the example file included in the package:
rmsd_file_path <- system.file("extdata/rmsd.xvg", package = "xvm")
rmsd_data <- read_xvg(rmsd_file_path) # read the xvg file using read_xvg() function
names(rmsd_data)
```

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summary_xvg	<i>summarize xvg Data</i>
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**Description**

compute basic summary statistics (mean, sd, min, median, max) for each variable in one or more xvg\_data objects.

**Usage**

```
summary_xvg(xvg_data, merge_results = FALSE)
```

**Arguments**

xvg_data	a list of class 'xvg_data' or a list containing multiple 'xvg_data' objects, as returned by read_xvg().
merge_results	logical, whether to combine results from multiple objects (default: FALSE). When TRUE, results will include a 'group' column identifying the source.

**Value**

a data.frame with columns:

**group** (Optional) Source identifier when processing multiple objects with merge\_results=TRUE.

**variable** Name of the variable (column) in the xvg data.

**mean** Arithmetic mean of that variable.

**sd** Standard deviation.

**min** Minimum value.

**median** Median value.

**max** Maximum value.

**Examples**

```
path <- system.file("extdata/rmsd.xvg", package = "xvm")
xvg <- read_xvg(path)
summary_xvg(xvg)
```

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