

# Package: tmle3 (via r-universe)

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**Title** The Extensible TMLE Framework

**Version** 0.2.1

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**Description** A general framework supporting the implementation of targeted maximum likelihood estimators (TMLEs) of a diverse range of statistical target parameters through a unified interface. The goal is that the exposed framework be as general as the mathematical framework upon which it draws.

**Depends** R (>= 3.6.0), graphics

**Imports** sl3 (>= 1.4.5), delayed, data.table, assertthat, R6, uuid, methods, ggplot2, stats, foreach, mvtnorm, magrittr, stringr, digest

**Suggests** testthat, knitr, origami (>= 1.0.3), tableone, speedglm, rmarkdown, Rsolnp, npls, tmle, tmle3shift, future, future.apply, xgboost

**Remotes** github::tlverse/sl3@fix-tests

**Additional\_repositories** <https://tlverse.r-universe.dev>

**License** GPL-3

**URL** <https://tlverse.org/tmle3>

**BugReports** <https://github.com/tlverse/tmle3/issues>

**Encoding** UTF-8

**LazyData** yes

**LazyLoad** yes

**VignetteBuilder** knitr

**RoxygenNote** 7.3.2

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**Repository** <https://tlverse.r-universe.dev>

**RemoteUrl** <https://github.com/tlverse/tmle3>

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all_ancestors	<i>Helper functions for the NPSEM</i>
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## Description

all\_ancestors returns a list of all\_ancestors of the specified node. time\_ordering attempts to find a time\_ordering for the variables.

## Usage

```
all_ancestors(node_name, npsem)
```

```
time_ordering(npsem)
```

## Arguments

node_name	the node to search for ancestors of
npsem	the NPSEM, defined by a list of <a href="#">tmle3_Node</a> objects.

---

bound	<i>Bound (Truncate) Likelihoods</i>
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**Description**

Bound (Truncate) Likelihoods

**Usage**

```
bound(x, bounds)
```

**Arguments**

x	the likelihood values to bound
bounds	Either a length two vector of c(lower,upper) or a lower bound, where the upper is then 1 - lower

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CF_Likelihood	<i>Counterfactual Likelihood</i>
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**Description**

Represents a counterfactual likelihood where one or more likelihood factors has been replaced with an intervention as specified by `intervention_list`. Inherits from [Likelihood](#). Other factors (including their updates) are taken from an underlying `observed_likelihoood` estimated from observed data.

**Usage**

```
make_CF_Likelihood(...)
```

**Arguments**

...	Passes all arguments to the constructor. See documentation for the Constructor below.
-----	---

**Format**

[R6Class](#) object.

**Value**

Likelihood object

**Constructor**

make\_CF\_Likelihood(observed\_likelihood, intervention\_list, ...)

observed\_likelihood [Likelihood](#) object specifying the relevant factors of the observed likelihood

intervention\_list A list of objects inheriting from [LF\\_base](#), representing the intervention.

... Not currently used.

**Fields**

observed\_likelihood [Likelihood](#) object specifying the relevant factors of the observed likelihood

intervention\_list A list of objects inheriting from [LF\\_base](#), representing the intervention.

**See Also**

Other Likelihood objects: [LF\\_base](#), [LF\\_derived](#), [LF\\_emp](#), [LF\\_fit](#), [LF\\_known](#), [LF\\_static](#), [LF\\_targeted](#), [Likelihood](#), [Targeted\\_Likelihood](#), [define\\_lf\(\)](#)

---

define\_lf

*Define a Likelihood Factor*

---

**Description**

Define a Likelihood Factor

**Usage**

define\_lf(LF\_class, ...)

**Arguments**

LF\_class the class of likelihood factor. Should inherit from [LF\\_base](#)

... arguments that define the likelihood factor. See the constructor for the specified LF\_class.

**See Also**

Other Likelihood objects: [CF\\_Likelihood](#), [LF\\_base](#), [LF\\_derived](#), [LF\\_emp](#), [LF\\_fit](#), [LF\\_known](#), [LF\\_static](#), [LF\\_targeted](#), [Likelihood](#), [Targeted\\_Likelihood](#)

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define_param	<i>Define a Parameter</i>
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**Description**

Define a Parameter

**Usage**

```
define_param(Param_class, ...)
```

**Arguments**

Param_class	the class of the Parameter. Should inherit from <a href="#">Param_base</a>
...	arguments that define the parameter See the constructor for the specified Parameter.

**See Also**

Other Parameters: [Param\\_ATC](#), [Param\\_ATE](#), [Param\\_ATT](#), [Param\\_MSM](#), [Param\\_TSM](#), [Param\\_base](#), [Param\\_delta](#), [Param\\_mean](#), [Param\\_stratified](#), [Param\\_survival](#), [tmle3\\_Fit](#)

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delta_param_ATE	<i>PAR = Linear Contrast EY1-EY0</i>
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**Description**

PAR = Linear Contrast EY1-EY0

**Usage**

```
delta_param_ATE
```

**Format**

An object of class list of length 4.

---

delta_param_OR	<i>Odds Ratio odds(Y1)/odds(Y0)</i>
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---

**Description**

Odds Ratio odds(Y1)/odds(Y0)

**Usage**

delta\_param\_OR

**Format**

An object of class list of length 5.

---

delta_param_PAF	<i>PAF = 1 - (1/RR(EY/E0))</i>
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---

**Description**

PAF = 1 - (1/RR(EY/E0))

**Usage**

delta\_param\_PAF

**Format**

An object of class list of length 5.

---

delta_param_PAR	<i>PAR = Linear Contrast EY-EY0</i>
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---

**Description**

PAR = Linear Contrast EY-EY0

**Usage**

delta\_param\_PAR

**Format**

An object of class list of length 4.

---

delta_param_RR	<i>Risk Ratio EY1/EY0</i>
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---

**Description**

Risk Ratio EY1/EY0

**Usage**

delta\_param\_RR

**Format**

An object of class list of length 5.

---

density_formula	<i>Get and Plot Propensity Scores</i>
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**Description**

Get and Plot Propensity Scores

**Usage**

```
density_formula(tmle_task, node = "A")
get_propensity_scores(likelihood, tmle_task, node = "A")
propensity_score_plot(likelihood, tmle_task, node = "A")
propensity_score_table(likelihood, tmle_task, node = "A")
```

**Arguments**

tmle_task	a tmle_task data object
node	a character specifying which node to use
likelihood	a fitted likelihood object



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discretize\_variable     *Discretize Continuous Variable*

---

**Description**

Converts a data.table column from continuous to a discrete factor

**Usage**

```
discretize_variable(data, variable, num_cats, breakpoints = NULL)
```

**Arguments**

data	data.table, containing the column to change
variable	character, the name of the column to change
num_cats	integer, the number of bins to generate
breakpoints	numeric vector, the breakpoints to use. If NULL, these will be quantiles.

**Value**

the updated data.table, modified in place

---

ED\_from\_estimates     *Get Empirical Mean of EIFs from Estimates*

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**Description**

Get Empirical Mean of EIFs from Estimates

**Usage**

```
ED_from_estimates(estimates)
```

**Arguments**

estimates	a list of estimates objects
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LF\_base

*Base Class for Defining Likelihood Factors***Description**

A Likelihood factor models a conditional density function. The conditioning set is defined as all parent nodes (defined in [tmle3\\_Task](#)). In the case of a continuous outcome variable, where a full density isn't needed, this can also model a conditional mean. This is the base class, which is intended to be abstract. See below for a list of possible likelihood factor classes.

**Format**

[R6Class](#) object.

**Value**

LF\_base object

**Constructor**

```
define_lf(LF_base, name, ..., type = "density")
```

name character, the name of the factor. Should match a node name in the nodes specified by [tmle3\\_Task\\$npsem](#)

... Not currently used.

type character, either "density", for conditional density or, "mean" for conditional mean

**Methods**

get\_density(tmle\_task) Get conditional density values for for the observations in tmle\_task.

- tmle\_task: [tmle3\\_Task](#) to get likelihood values for

get\_mean(tmle\_task) Get conditional mean values for for the observations in tmle\_task.

- tmle\_task: [tmle3\\_Task](#) to get likelihood values for

**Fields**

name character, the name of the factor. Should match a node name in the nodes specified by [tmle3\\_Task\\$npsem](#)

type character, either "density", for conditional density or, "mean" for conditional mean

variable\_type [variable\\_type](#) object, specifying the data type of the outcome variable. Only available after Likelihood training.

values Possible values of the outcome variable, retrieved from the variable\_type object. Only available after Likelihood training.

**See Also**

Other Likelihood objects: [CF\\_Likelihood](#), [LF\\_derived](#), [LF\\_emp](#), [LF\\_fit](#), [LF\\_known](#), [LF\\_static](#), [LF\\_targeted](#), [Likelihood](#), [Targeted\\_Likelihood](#), [define\\_lf\(\)](#)

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LF_derived	<i>Derived Likelihood Factor Estimated from Data + Other Likelihood values, using sl3.</i>
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**Description**

Uses an sl3 learner to estimate a likelihood factor from data. Inherits from [LF\\_base](#); see that page for documentation on likelihood factors in general.

**Format**

[R6Class](#) object.

**Value**

LF\_base object

**Constructor**

```
define_lf(LF_fit, name, learner, ..., type = "density")
```

name character, the name of the factor. Should match a node name in the nodes specified by [tmle3\\_Task\\$npsem](#)

learner An sl3 learner to be used to estimate the factor

... Not currently used.

type character, either "density", for conditional density or, "mean" for conditional mean

**Fields**

learner The learner or learner fit object

**See Also**

Other Likelihood objects: [CF\\_Likelihood](#), [LF\\_base](#), [LF\\_emp](#), [LF\\_fit](#), [LF\\_known](#), [LF\\_static](#), [LF\\_targeted](#), [Likelihood](#), [Targeted\\_Likelihood](#), [define\\_lf\(\)](#)

---

 LF\_emp

*Likelihood Factor Estimated using Empirical Distribution*


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### Description

Uses the empirical probability distribution (puts mass  $1/n$  on each of the observations, or uses weights if specified) to estimate a marginal density. Inherits from [LF\\_base](#); see that page for documentation on likelihood factors in general. Only compatible with marginal likelihoods (no parent nodes). Only compatible with densities (no conditional means). The type argument will be ignored if specified.

### Format

[R6Class](#) object.

### Value

LF\_base object

### Constructor

```
define_lf(LF_emp, name, ...)
```

name character, the name of the factor. Should match a node name in the nodes specified by [tmle3\\_Task\\$npsem](#)

... Not currently used.

### See Also

Other Likelihood objects: [CF\\_Likelihood](#), [LF\\_base](#), [LF\\_derived](#), [LF\\_fit](#), [LF\\_known](#), [LF\\_static](#), [LF\\_targeted](#), [Likelihood](#), [Targeted\\_Likelihood](#), [define\\_lf\(\)](#)

---

 LF\_fit

*Likelihood Factor Estimated from Data using sl3.*


---

### Description

Uses an `sl3` learner to estimate a likelihood factor from data. Inherits from [LF\\_base](#); see that page for documentation on likelihood factors in general.

### Format

[R6Class](#) object.

### Value

LF\_base object

**Constructor**

```
define_lf(LF_fit, name, learner, ..., type = "density")
```

name character, the name of the factor. Should match a node name in the nodes specified by [tmle3\\_Task\\$npsem](#)

learner An sl3 learner to be used to estimate the factor

... Not currently used.

type character, either "density", for conditional density or, "mean" for conditional mean

**Fields**

learner The learner or learner fit object

**See Also**

Other Likelihood objects: [CF\\_Likelihood](#), [LF\\_base](#), [LF\\_derived](#), [LF\\_emp](#), [LF\\_known](#), [LF\\_static](#), [LF\\_targeted](#), [Likelihood](#), [Targeted\\_Likelihood](#), [define\\_lf\(\)](#)

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LF\_known

*Known True Likelihood Factor*

---

**Description**

Incorporate existing knowledge about the likelihood Inherits from [LF\\_base](#); see that page for documentation on likelihood factors in general.

**Format**

[R6Class](#) object.

**Value**

LF\_base object

**Constructor**

```
define_lf(LF_fit, name, mean_fun, density_fun, ..., type = "density")
```

name character, the name of the factor. Should match a node name in the nodes specified by [tmle3\\_Task\\$npsem](#)

mean\_fun A function that takes a sl3 regression task and returns true conditional means

density\_fun A function that takes a sl3 regression task and returns true conditional densities

... Not currently used.

type character, either "density", for conditional density or, "mean" for conditional mean

**See Also**

Other Likelihood objects: [CF\\_Likelihood](#), [LF\\_base](#), [LF\\_derived](#), [LF\\_emp](#), [LF\\_fit](#), [LF\\_static](#), [LF\\_targeted](#), [Likelihood](#), [Targeted\\_Likelihood](#), [define\\_lf\(\)](#)

---

LF\_static

*Static Likelihood Factor*

---

**Description**

Likelihood factor for a variable that only has one value with probability 1. This is used for static interventions. Inherits from [LF\\_base](#); see that page for documentation on likelihood factors in general.

**Format**

[R6Class](#) object.

**Value**

LF\_base object

**Constructor**

```
define_lf(LF_static, name, type, value, ...)
```

name character, the name of the factor. Should match a node name in the nodes specified by [tmle3\\_Task\\$npsem](#)

type character, either "density", for conditional density or, "mean" for conditional mean

value the static value

... Not currently used.

**Fields**

value the static value.

**See Also**

Other Likelihood objects: [CF\\_Likelihood](#), [LF\\_base](#), [LF\\_derived](#), [LF\\_emp](#), [LF\\_fit](#), [LF\\_known](#), [LF\\_targeted](#), [Likelihood](#), [Targeted\\_Likelihood](#), [define\\_lf\(\)](#)

---

LF_targeted	<i>Use a likelihood factor from an existing targeted likelihood</i>
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**Description**

Uses an `s13` learner to estimate a likelihood factor from data. Inherits from [LF\\_base](#); see that page for documentation on likelihood factors in general.

**Format**

[R6Class](#) object.

**Value**

LF\_base object

**Constructor**

```
define_lf(LF_fit, name, learner, ..., type = "density")
```

`name` character, the name of the factor. Should match a node name in the nodes specified by `tmle3_Task$npsem`

`learner` An `s13` learner to be used to estimate the factor

`...` Not currently used.

`type` character, either "density", for conditional density or, "mean" for conditional mean

**Fields**

`learner` The learner or learner fit object

**See Also**

Other Likelihood objects: [CF\\_Likelihood](#), [LF\\_base](#), [LF\\_derived](#), [LF\\_emp](#), [LF\\_fit](#), [LF\\_known](#), [LF\\_static](#), [Likelihood](#), [Targeted\\_Likelihood](#), [define\\_lf\(\)](#)

Likelihood

*Class for Likelihood***Description**

This object represents an estimate of the relevant factors of the likelihood estimated from data, or based on *a priori* knowledge where appropriate. That is, it represents some subset of  $P_n$ . This object inherits from `Lrrnr_base`, and so shares some properties with `s13` learners. Specifically, to fit a likelihood object to data, one calls `likelihood$train(tmle3_task)`. Each likelihood factor is represented by an object inheriting from `LF_base`.

**Usage**

```
make_Likelihood(...)
```

**Arguments**

... Passes all arguments to the constructor. See documentation for the Constructor below.

**Format**

`R6Class` object.

**Value**

Likelihood object

**Constructor**

```
make_Likelihood(factor_list, ...)
```

`factor_list` A list of objects inheriting from `LF_base`, representing the individual relevant factors.

... Not currently used.

**Methods**

`validate_task(tmle_task)` Ensure that this likelihood is compatible with a particular `tmle3_Task`, in that the factor names must match the `tmle_task$npsem` names.

- `tmle_task`: the `tmle3_Task` to validate.

`get_initial_likelihoods(tmle_task, nodes=NULL)` Gets initial (i.e. before any TMLE updates) likelihood values for the specified nodes (or all nodes if none are specified) for the observations in `tmle_task`.

- `tmle_task`: `tmle3_Task` to get likelihood values for
- `nodes`: character vectors, the list of nodes to get likelihood values for. If missing, values will be provided for all nodes.



`get_likelihoods(tmle_task, nodes=NULL)` Gets updated (i.e. after all TMLE updates) likelihood values for the specified nodes (or all nodes if none are specified) for the observations in `tmle_task`.

- `tmle_task`: [tmle3\\_Task](#) to get likelihood values for
- `nodes`: character vectors, the list of nodes to get likelihood values for. If missing, values will be provided for all nodes.

`get_possible_counterfactuals(nodes)` Gets all possible combination of counterfactual values for a set of nodes. This is useful for marginalizing over a node. Returns a `data.frame` with one row per possibility.

- `nodes`: character vectors, the list of nodes to get counterfactual values for. If missing, values will be provided for all nodes.

### Fields

`factor_list` The list of [LF\\_base](#) objects specifying the relevant likelihood factors

`observed_values` The likelihood values for the observed data. These are cached, as they are used in many places in TMLE

`update_list` A list of `tmle_updates` that have been calculated for this likelihood

### See Also

Other Likelihood objects: [CF\\_Likelihood](#), [LF\\_base](#), [LF\\_derived](#), [LF\\_emp](#), [LF\\_fit](#), [LF\\_known](#), [LF\\_static](#), [LF\\_targeted](#), [Targeted\\_Likelihood](#), [define\\_lf\(\)](#)

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Likelihood_cache	<i>Cache Likelihood values, update those values</i>
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---

### Description

Cache Likelihood values, update those values

---

Param_ATT	<i>Additive Effect of Treatment Among the Treated</i>
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---

### Description

Parameter definition for the Additive Effect of Treatment Among the Treated (ATT). Currently supports multiple static intervention nodes. Does yet not support dynamic rule or stochastic interventions.

### Format

[R6Class](#) object.

**Value**

Param\_base object

**Current Issues**

- clever covariates doesn't support updates; always uses initial (necessary for iterative TMLE, e.g. stochastic intervention)
- doesn't integrate over possible counterfactuals (necessary for stochastic intervention)
- clever covariate gets recalculated all the time (inefficient)

**Constructor**

`define_param(Param_ATT, observed_likelihood, intervention_list, ..., outcome_node)`

`observed_likelihood` A [Likelihood](#) corresponding to the observed likelihood

`intervention_list_treatment` A list of objects inheriting from [LF\\_base](#), representing the treatment intervention.

`intervention_list_control` A list of objects inheriting from [LF\\_base](#), representing the control intervention.

`...` Not currently used.

`outcome_node` character, the name of the node that should be treated as the outcome

**Fields**

`cf_likelihood_treatment` the counterfactual likelihood for the treatment

`cf_likelihood_control` the counterfactual likelihood for the control

`intervention_list_treatment` A list of objects inheriting from [LF\\_base](#), representing the treatment intervention

`intervention_list_control` A list of objects inheriting from [LF\\_base](#), representing the control intervention

**See Also**

Other Parameters: [Param\\_ATE](#), [Param\\_ATT](#), [Param\\_MSM](#), [Param\\_TSM](#), [Param\\_base](#), [Param\\_delta](#), [Param\\_mean](#), [Param\\_stratified](#), [Param\\_survival](#), [define\\_param\(\)](#), [tmle3\\_Fit](#)

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Param_ATE	<i>Average Treatment Effect</i>
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---

**Description**

Parameter definition for the Average Treatment Effect (ATE).

**Format**

[R6Class](#) object.

**Value**

Param\_base object

**Constructor**

`define_param(Param_ATT, observed_likelihood, intervention_list, ..., outcome_node)`

`observed_likelihood` A [Likelihood](#) corresponding to the observed likelihood

`intervention_list_treatment` A list of objects inheriting from [LF\\_base](#), representing the treatment intervention.

`intervention_list_control` A list of objects inheriting from [LF\\_base](#), representing the control intervention.

`...` Not currently used.

`outcome_node` character, the name of the node that should be treated as the outcome

**Fields**

`cf_likelihood_treatment` the counterfactual likelihood for the treatment

`cf_likelihood_control` the counterfactual likelihood for the control

`intervention_list_treatment` A list of objects inheriting from [LF\\_base](#), representing the treatment intervention

`intervention_list_control` A list of objects inheriting from [LF\\_base](#), representing the control intervention

**See Also**

Other Parameters: [Param\\_ATC](#), [Param\\_ATT](#), [Param\\_MSM](#), [Param\\_TSM](#), [Param\\_base](#), [Param\\_delta](#), [Param\\_mean](#), [Param\\_stratified](#), [Param\\_survival](#), [define\\_param\(\)](#), [tmle3\\_Fit](#)

Param\_ATT

*Additive Effect of Treatment Among the Treated***Description**

Parameter definition for the Additive Effect of Treatment Among the Treated (ATT). Currently supports multiple static intervention nodes. Does yet not support dynamic rule or stochastic interventions.

**Format**

[R6Class](#) object.

**Value**

Param\_base object

**Current Issues**

- clever covariates doesn't support updates; always uses initial (necessary for iterative TMLE, e.g. stochastic intervention)
- doesn't integrate over possible counterfactuals (necessary for stochastic intervention)
- clever covariate gets recalculated all the time (inefficient)

**Constructor**

```
define_param(Param_ATT, observed_likelihood, intervention_list, ..., outcome_node)
```

`observed_likelihood` A [Likelihood](#) corresponding to the observed likelihood

`intervention_list_treatment` A list of objects inheriting from [LF\\_base](#), representing the treatment intervention.

`intervention_list_control` A list of objects inheriting from [LF\\_base](#), representing the control intervention.

`...` Not currently used.

`outcome_node` character, the name of the node that should be treated as the outcome

**Fields**

`cf_likelihood_treatment` the counterfactual likelihood for the treatment

`cf_likelihood_control` the counterfactual likelihood for the control

`intervention_list_treatment` A list of objects inheriting from [LF\\_base](#), representing the treatment intervention

`intervention_list_control` A list of objects inheriting from [LF\\_base](#), representing the control intervention

**See Also**

Other Parameters: [Param\\_ATC](#), [Param\\_ATE](#), [Param\\_MSM](#), [Param\\_TSM](#), [Param\\_base](#), [Param\\_delta](#), [Param\\_mean](#), [Param\\_stratified](#), [Param\\_survival](#), [define\\_param\(\)](#), [tmle3\\_Fit](#)

---

 Param\_base

*Base Class for Defining Parameters*


---

**Description**

A parameter is a function of the likelihood. Once given a [Likelihood](#) object, a parameter will a value. These objects also contain information about the efficient influence function (EIF) of a parameter, as well as its clever covariate(s).

**Format**

[R6Class](#) object.

**Value**

Param\_base object

**Constructor**

`define_param(Param_base, observed_likelihoood, ..., outcome_node)`

`observed_likelihoood` A [Likelihood](#) corresponding to the observed likelihood

`...` Not currently used.

`outcome_node` character, the name of the node that should be treated as the outcome

**Methods**

`clever_covariates(tmle_task = NULL)` Get the clever covariates for an TMLE update step.

- `tmle_task`: [tmle3\\_Task](#) to get clever covariate values for. If NULL, the `tmle_task` used to train the observed likelihood will be used

`estimates(tmle_task = NULL)` Get the parameter estimates and influence curve values.

- `tmle_task`: [tmle3\\_Task](#) to get clever covariate values for. If NULL, the `tmle_task` used to train the observed likelihood will be used

**Fields**

`observed_likelihoood` the observed likelihood

`outcome_node` character, the name of the outcome node

**See Also**

Other Parameters: [Param\\_ATC](#), [Param\\_ATE](#), [Param\\_ATT](#), [Param\\_MSM](#), [Param\\_TSM](#), [Param\\_delta](#), [Param\\_mean](#), [Param\\_stratified](#), [Param\\_survival](#), [define\\_param\(\)](#), [tmle3\\_Fit](#)

---

Param_delta	<i>Delta Method Parameters</i>
-------------	--------------------------------

---

**Description**

These parameters are smooth functionals of one or more other params They are not fit directly with tmlle, but are estimated using the delta method todo: better docs They do not return have clever covariates

**See Also**

Other Parameters: [Param\\_ATC](#), [Param\\_ATE](#), [Param\\_ATT](#), [Param\\_MSM](#), [Param\\_TSM](#), [Param\\_base](#), [Param\\_mean](#), [Param\\_stratified](#), [Param\\_survival](#), [define\\_param\(\)](#), [tmlle3\\_Fit](#)

---

Param_mean	<i>Mean of Outcome Node</i>
------------	-----------------------------

---

**Description**

Parameter for marginal mean of Y:  $\Psi = E[Y]$ . No TMLE update needed, but can be used in delta method calculations. Useful for example, in calculating attributable risks.

**Format**

[R6Class](#) object.

**Value**

Param\_base object

**Constructor**

`define_param(Param_TSM, observed_likelihood, intervention_list, ..., outcome_node)`

`observed_likelihood` A [Likelihood](#) corresponding to the observed likelihood

`...` Not currently used.

`outcome_node` character, the name of the node that should be treated as the outcome

**Fields**

`cf_likelihood` the counterfactual likelihood for this treatment

`intervention_list` A list of objects inheriting from [LF\\_base](#), representing the intervention

**See Also**

Other Parameters: [Param\\_ATC](#), [Param\\_ATE](#), [Param\\_ATT](#), [Param\\_MSM](#), [Param\\_TSM](#), [Param\\_base](#), [Param\\_delta](#), [Param\\_stratified](#), [Param\\_survival](#), [define\\_param\(\)](#), [tmlle3\\_Fit](#)

Param\_MSM

*Stratified Parameter Estimates via MSM***Description**

Stratified Parameter Estimates via MSM

**Format**

R6Class object.

**Value**

Param\_base object

**Current Issues**

- clever covariates doesn't support updates; always uses initial (necessary for iterative TMLE, e.g. stochastic intervention)
- clever covariate gets recalculated all the time (inefficient)

**Constructor**

```
define_param(Param_MSM, observed_likelihoood, strata_variable, ...)
```

observed\_likelihoood A [Likelihood](#) corresponding to the observed likelihood  
msm form of the MSM. Default is "A + V", consistent with the default of treatment\_node and strata\_name.

weight "Cond.Prob.", "Unif." or custom input function. Note that custom function should support vector input. Default is "Cond.Prob."

... Not currently used.

covariate\_node character, the name of the node that should be treated as the covariate

treatment\_node character, the name of the node that should be treated as the treatment

outcome\_node character, the name of the node that should be treated as the outcome

**Fields**

cf\_likelihoood the counterfactual likelihood for this treatment

**See Also**

Other Parameters: [Param\\_ATC](#), [Param\\_ATE](#), [Param\\_ATT](#), [Param\\_TSM](#), [Param\\_base](#), [Param\\_delta](#), [Param\\_mean](#), [Param\\_stratified](#), [Param\\_survival](#), [define\\_param\(\)](#), [tmle3\\_Fit](#)

---

Param_stratified	<i>Stratified Parameter Estimates</i>
------------------	---------------------------------------

---

**Description**

Stratified Parameter Estimates

**Format**

[R6Class](#) object.

**Value**

Param\_base object

**Current Issues**

- clever covariates doesn't support updates; always uses initial (necessary for iterative TMLE, e.g. stochastic intervention)
- doesn't integrate over possible counterfactuals (necessary for stochastic intervention)
- clever covariate gets recalculated all the time (inefficient)

**Constructor**

```
define_param(Param_TSM, observed_likelihood, intervention_list, ..., outcome_node)
```

`observed_likelihood` A [Likelihood](#) corresponding to the observed likelihood

`intervention_list` A list of objects inheriting from [LF\\_base](#), representing the intervention.

`...` Not currently used.

`outcome_node` character, the name of the node that should be treated as the outcome

**Fields**

`cf_likelihood` the counterfactual likelihood for this treatment

`intervention_list` A list of objects inheriting from [LF\\_base](#), representing the intervention

**See Also**

Other Parameters: [Param\\_ATC](#), [Param\\_ATE](#), [Param\\_ATT](#), [Param\\_MSM](#), [Param\\_TSM](#), [Param\\_base](#), [Param\\_delta](#), [Param\\_mean](#), [Param\\_survival](#), [define\\_param\(\)](#), [tmle3\\_Fit](#)



---

Param_survival	<i>Survival Curve</i>
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---

**Description**

Survival Curve

**Format**

[R6Class](#) object.

**Value**

Param\_base object

**Constructor**

`define_param(Param_survival, observed_likelihood, intervention_list, ..., outcome_node)`

`observed_likelihood` A [Likelihood](#) corresponding to the observed likelihood

`intervention_list` A list of objects inheriting from [LF\\_base](#), representing the intervention.

... Not currently used.

`outcome_node` character, the name of the node that should be treated as the outcome

**Fields**

`cf_likelihood` the counterfactual likelihood for this treatment

`intervention_list` A list of objects inheriting from [LF\\_base](#), representing the intervention

**See Also**

Other Parameters: [Param\\_ATC](#), [Param\\_ATE](#), [Param\\_ATT](#), [Param\\_MSM](#), [Param\\_TSM](#), [Param\\_base](#), [Param\\_delta](#), [Param\\_mean](#), [Param\\_stratified](#), [define\\_param\(\)](#), [tmle3\\_Fit](#)

---

Param_TSM	<i>Treatment Specific Mean</i>
-----------	--------------------------------

---

**Description**

Parameter definition for the Treatment Specific Mean (TSM):  $E_W[E_Y|A(Y|A=a|W)]$ . Currently supports multiple static intervention nodes. Does yet not support dynamic rule or stochastic interventions.

**Format**

[R6Class](#) object.

**Value**

Param\_base object

**Current Issues**

- clever covariates doesn't support updates; always uses initial (necessary for iterative TMLE, e.g. stochastic intervention)
- doesn't integrate over possible counterfactuals (necessary for stochastic intervention)
- clever covariate gets recalculated all the time (inefficient)

**Constructor**

define\_param(Param\_TSM, observed\_likelihood, intervention\_list, ..., outcome\_node)

observed\_likelihood A [Likelihood](#) corresponding to the observed likelihood

intervention\_list A list of objects inheriting from [LF\\_base](#), representing the intervention.

... Not currently used.

outcome\_node character, the name of the node that should be treated as the outcome

**Fields**

cf\_likelihood the counterfactual likelihood for this treatment

intervention\_list A list of objects inheriting from [LF\\_base](#), representing the intervention

**See Also**

Other Parameters: [Param\\_ATC](#), [Param\\_ATE](#), [Param\\_ATT](#), [Param\\_MSM](#), [Param\\_base](#), [Param\\_delta](#), [Param\\_mean](#), [Param\\_stratified](#), [Param\\_survival](#), [define\\_param\(\)](#), [tmle3\\_Fit](#)

---

plot\_vim

*Plot results of variable importance analysis*

---

**Description**

Plot results of variable importance analysis

**Usage**

plot\_vim(vim\_results)

**Arguments**

vim\_results Object produced by invoking `tmle3_vim`.

---

point\_tx\_npsem      *Helper Functions for Point Treatment*

---

**Description**

Handles the common W (covariates), A (treatment/intervention), Y (outcome) data structure

**Usage**

```
point_tx_npsem(node_list, variable_types = NULL)

point_tx_task(data, node_list, variable_types = NULL, ...)

point_tx_likelihood(tmle_task, learner_list)
```

**Arguments**

node_list	a list of character vectors, listing the variables that comprise each node
variable_types	a list of variable types, one for each node. If missing, variable types will be guessed
data	a data.frame, or data.table containing data for use in estimation
...	extra arguments.
tmle_task	a <a href="#">tmle3_Task</a> as constructed via point_tx_task
learner_list	a list of sl3 learners, one for A and one for Y to be used for likelihood estimation

---

process\_missing      *Preprocess Data to Handle Missing Variables*

---

**Description**

Process data to account for missingness in preparation for TMLE

**Usage**

```
process_missing(
  data,
  node_list,
  complete_nodes = c("A", "Y"),
  impute_nodes = NULL,
  max_p_missing = 0.5
)
```

**Arguments**

data	data.table, containing the missing variables
node_list	list, what variables comprise each node
complete_nodes	character vector, nodes we must observe
impute_nodes	character vector, nodes we will impute
max_p_missing	numeric, what proportion of missing is tolerable? Beyond that, the variable will be dropped from the analysis

**Details**

Rows where there is missingness in any of the complete\_nodes will be dropped. Then, missingness will be median-imputed for the variables in the impute\_nodes. Indicator variables of missingness will be generated for these nodes.

Then covariates will be processed as follows:

1. any covariate with more than max\_p\_missing missingness will be dropped
2. indicators of missingness will be generated
3. missing values will be median-imputed

**Value**

list containing the following elements:

- data, the updated dataset
- node\_list, the updated list of nodes
- n\_dropped, the number of observations dropped
- dropped\_cols, the variables dropped due to excessive missingness

---

submodel_logit	<i>Logistic Submodel Fluctuation</i>
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---

**Description**

Logistic Submodel Fluctuation

**Usage**

```
submodel_logit(eps, X, offset)
```

**Arguments**

eps	...
X	...
offset	...

---

```
summary_from_estimates
```

*Summarize Estimates*

---

**Description**

Generates a `data.table` summarizing results with inference

**Usage**

```
summary_from_estimates(
  task,
  estimates,
  param_types = NULL,
  param_names = NULL,
  init_psi = NULL,
  simultaneous_ci = FALSE
)
```

**Arguments**

<code>task</code>	<code>tmle3_Task</code> containing the observed data of interest; the same as that passed to <code>tmle3</code>
<code>estimates</code>	<code>list</code> , TMLE estimates of parameter and ICs from <code>tmle3_Fit\$estimates</code>
<code>param_types</code>	the types of the parameters being estimated
<code>param_names</code>	the names of the parameters being estimated
<code>init_psi</code>	the names of the parameters being estimated
<code>simultaneous_ci</code>	if <code>TRUE</code> , calculate simultaneous confidence intervals

**Value**

`data.table` summarizing results

---

```
survival_tx_npsem
```

*Helper Functions for Survival Analysis*

---

**Description**

Handles the `W` (covariates), `A` (treatment/intervention), `T_tilde` (time-to-event), `Delta` (censoring indicator), `t_max` (the maximum time to estimate) survival data structure

**Usage**

```
survival_tx_npsem(node_list, variable_types = NULL)

survival_tx_task(data, node_list, variable_types = NULL, ...)

survival_tx_likelihood(tmle_task, learner_list)
```

**Arguments**

<code>node_list</code>	a list of character vectors, listing the variables that comprise each node
<code>variable_types</code>	a list of variable types, one for each node. If missing, variable types will be guessed
<code>data</code>	a <code>data.frame</code> , or <code>data.table</code> containing data for use in estimation
<code>...</code>	extra arguments.
<code>tmle_task</code>	a <code>tmle3_Task</code> as constructed via <code>survival_tx_task</code>
<code>learner_list</code>	a list of <code>sl3</code> learners, one for A and one for Y to be used for likelihood estimation

---

Targeted\_Likelihood    *Targeted Likelihood*

---

**Description**

Represents a likelihood where one or more likelihood factors has been updated to target a set of parameter(s)

**Format**

`R6Class` object.

**Value**

Likelihood object

**Constructor**

```
make_Likelihood(factor_list, ...)
```

`factor_list` A list of objects inheriting from `LF_base`, representing the individual relevant factors.

`...` Not currently used.

## Methods

`validate_task(tmle_task)` Ensure that this likelihood is compatible with a particular `tmle3_Task`, in that the factor names must match the `tmle_task$npsem` names.

- `tmle_task`: the `tmle3_Task` to validate.

`get_initial_likelihoods(tmle_task, nodes=NULL)` Gets initial (i.e. before any TMLE updates) likelihood values for the specified nodes (or all nodes if none are specified) for the observations in `tmle_task`.

- `tmle_task`: `tmle3_Task` to get likelihood values for
- `nodes`: character vectors, the list of nodes to get likelihood values for. If missing, values will be provided for all nodes.

`get_likelihoods(tmle_task, nodes=NULL)` Gets updated (i.e. after all TMLE updates) likelihood values for the specified nodes (or all nodes if none are specified) for the observations in `tmle_task`.

- `tmle_task`: `tmle3_Task` to get likelihood values for
- `nodes`: character vectors, the list of nodes to get likelihood values for. If missing, values will be provided for all nodes.

`get_possible_counterfactuals(nodes)` Gets all possible combination of counterfactual values for a set of nodes. This is useful for marginalizing over a node. Returns a `data.frame` with one row per possibility.

- `nodes`: character vectors, the list of nodes to get counterfactual values for. If missing, values will be provided for all nodes.

## Fields

`factor_list` The list of `LF_base` objects specifying the relevant likelihood factors

`observed_values` The likelihood values for the observed data. These are cached, as they are used in many places in TMLE

`update_list` A list of `tmle_updates` that have been calculated for this likelihood

## See Also

Other Likelihood objects: `CF_Likelihood`, `LF_base`, `LF_derived`, `LF_emp`, `LF_fit`, `LF_known`, `LF_static`, `LF_targeted`, `Likelihood`, `define_lf()`

---

tmle3

*TMLE from a tmle3\_Spec object*

---

## Description

Using a `tmle3_Spec` object, fit a TMLE

## Usage

```
tmle3(tmle_spec, data, node_list, learner_list = NULL)
```

**Arguments**

tmle_spec	<a href="#">tmle3_Spec</a> , defines the TMLE
data	<code>data.frame</code> , the raw data
node_list	list, defines which variables are which nodes
learner_list	list, defines which learners are used to fit which likelihood factors

**Value**

A [tmle3\\_Fit](#) object

---

tmle3_Fit	<i>TMLE fit object</i>
-----------	------------------------

---

**Description**

A `tmle_fit` object, containing initial and updated estimates, as well as data about the fitting procedure. TMLE updates are calculated when the object is constructed.

**Usage**

```
fit_tmle3(...)
```

**Arguments**

... Passes all arguments to the constructor. See documentation for the Constructor.

**Format**

[R6Class](#) object.

**Value**

`Param_base` object

**Constructor**

```
fit_tmle3(tmle_task, likelihood, tmle_params, updater, max_it=100, ...)
```

`tmle_task` A [tmle3\\_Task](#) object defining the data and NP-SEM

`likelihood` A [Likelihood](#) object defining the factorized likelihood

`tmle_params` A list of parameters inheriting from [Param\\_base](#) defining the parameter(s) of interest

`updater` A [tmle3\\_Update](#) object defining the update procedure, including submodel and loss function

`maxit` integer, maximum number of TMLE iterations

... Not currently used.



**Methods**

`set_timings(start_time, task_time, likelihood_time, params_time, fit_time)` Provide the timings for the different steps of the TMLE procedure, for later reporting to the user

- `tmle_task`: [tmle3\\_Task](#) to get clever covariate values for. If NULL, the `tmle_task` used to train the observed likelihood will be used

`estimates(tmle_task = NULL)` Get the parameter estimates and influence curve values.

- `tmle_task`: [tmle3\\_Task](#) to get clever covariate values for. If NULL, the `tmle_task` used to train the observed likelihood will be used

**Fields**

`tmle_task` A [tmle3\\_Task](#) object defining the data and NP-SEM

`likelihood` A [Likelihood](#) object defining the factorized likelihood

`tmle_params` A list of parameters inheriting from [Param\\_base](#) defining the parameter(s) of interest

`tmle_names` A list of parameter names, obtained by calling `param$name` on each parameter

`updater` A [tmle3\\_Update](#) object defining the update procedure, including submodel and loss function

`steps` integer, the number of steps until TMLE converged

`ED` vector, the mean of the EIF for all the parameters

`initial_psi` vector, the initial parameter estimates

`estimates` list, final parameter estimates and ICs

`summary` `data.table`, summary of results

`timings` `data.frame`, timings for each step (provided by `tmle3_Fit$set_timings`)

**See Also**

Other Parameters: [Param\\_ATC](#), [Param\\_ATE](#), [Param\\_ATT](#), [Param\\_MSM](#), [Param\\_TSM](#), [Param\\_base](#), [Param\\_delta](#), [Param\\_mean](#), [Param\\_stratified](#), [Param\\_survival](#), [define\\_param\(\)](#)

---

tmle3\_Node

*A Node (set of variables) in an NPSEM*

---

**Description**

This class defines a node in an NPSEM

**Usage**

`define_node(...)`

**Arguments**

... Passes all arguments to the constructor. See documentation for the Constructor below.

**Format**

R6Class object.

**Value**

tmle3\_Node object

**Constructor**

make\_tmle3\_task(name, variables, parents = c(), variable\_type = NULL)

name character, the name of node

variables character vector, the names of the variables that comprise the node

parents character vector, the names of the parent nodes. If censoring, node is assumed to have no parents.

variable\_type [variable\\_type](#) object, specifying the data type of this variable. If censoring, variable\_type will be guessed later from the data.

**Methods**

guess\_variable\_type(variable\_data) Guesses the [variable\\_type](#) from the provided data. This will be called by the [tmle3\\_Task](#) constructor if no variable\_type was provided.

- variable\_data: the observed variable data.

**Fields**

name character, the name of node

variables character vector, the names of the variables that comprise the node

parents character vector, the names of the parent nodes. If censoring, node is assumed to have no parents.

variable\_type [variable\\_type](#) object, specifying the data type of this variable.

---

tmle3_Spec	<i>Defines a TML Estimator (except for the data)</i>
------------	--

---

**Description**

Current limitations: pretty much tailored to Param\_TSM

---

tmle3_Spec_ATC	<i>Defines a TML Estimator (except for the data)</i>
----------------	--

---

**Description**

Defines a TML Estimator (except for the data)

---

tmle3_Spec_ATE	<i>Defines a TML Estimator (except for the data)</i>
----------------	--

---

**Description**

Defines a TML Estimator (except for the data)

---

tmle3_Spec_ATT	<i>Defines a TML Estimator (except for the data)</i>
----------------	--

---

**Description**

Defines a TML Estimator (except for the data)

---

tmle3_Spec_MSM	<i>Defines a Stratified TML Estimator with MSM (except for the data)</i>
----------------	--

---

**Description**

Defines a Stratified TML Estimator with MSM (except for the data)

---

tmle3_Spec_OR	<i>Defines a TML Estimator for the Odds Ratio</i>
---------------	---

---

**Description**

Current limitations: pretty much tailored to Param\_TSM see TODOs for places generalization can be added

---

tmle3_Spec_PAR	<i>Defines a tmle (minus the data)</i>
----------------	--

---

**Description**

Current limitations: pretty much tailored to Param\_TSM see TODOs for places generalization can be added

---

tmle3\_Spec\_RR      *Defines a TML Estimator for the Risk Ratio*

---

**Description**

Current limitations: pretty much tailored to Param\_TSM see TODOs for places generalization can be added

---

tmle3\_Spec\_stratified      *Defines a Stratified TML Estimator (except for the data)*

---

**Description**

Defines a Stratified TML Estimator (except for the data)

---

tmle3\_Spec\_survival      *Defines a TML Estimator (except for the data)*

---

**Description**

Defines a TML Estimator (except for the data)

---

tmle3\_Spec\_TSM\_all      *Defines a TML Estimator (except for the data)*

---

**Description**

Current limitations: pretty much tailored to Param\_TSM See TODOs for places generalization can be added

tmle3\_Task

*Class for Storing Data and NPSEM for TMLE***Description**

This class inherits from [sl3\\_Task](#). In addition to all the methods supported by [sl3\\_Task](#), it supports the following.

**Usage**

```
make_tmle3_Task(...)
```

**Arguments**

... Passes all arguments to the constructor. See documentation for the Constructor below.

**Format**

[R6Class](#) object.

**Value**

tmle3\_Task object

**Constructor**

```
make_tmle3_task(data, npsem, ...)
```

`data` A `data.frame` or `data.table` containing the underlying data

`npsem` A list of [tmle3\\_Node](#) objects, where each is created using [define\\_node](#). These specify the NPSEM. See examples.

... Other arguments passed to the constructor of [sl3\\_Task](#). **NB:** Support for these is currently limited.

**Methods**

`get_tmle_node(node_name, bound = FALSE)` Gets the data associated with a `tmle_node`. Bounds the data if requested.

- `node_name`: character, the name of the node to get.
- `bound`: logical, if true the data is transformed to be in (0,1) based on pre-specified bounds.

`get_regression_task(target_node, bound = FALSE)` Gets a [sl3\\_Task](#) suitable for fitting the conditional likelihood factor with the `target_node` as the outcome.

- `target_node`: character, the name of the node to get.

`generate_counterfactual_task(uuid, new_data)` Generates a new `tmle_Task` where some nodes are overridden to have counterfactual values.

- `uuid`: A unique identifier for the counterfactual task, as generated by `UUIDgenerate`
- `new_data`: A `data.frame` or `data.table` with the counterfactual values. Column names must refer to node names in the `npsem` for this task.

## Fields

`npsem` The list of `tmle3_Node` objects specifying the NPSEM

---

<code>tmle3_Update</code>	<i>Defines an update procedure (submodel+loss function)</i>
---------------------------	---

---

## Description

Current Limitations: loss function and submodel are hard-coded (need to accept arguments for these)

## Constructor

`define_param(maxit, cvtmle, one_dimensional, constrain_step, delta_epsilon, verbose)`

`maxit` The maximum number of update iterations

`cvtmle` If TRUE, use CV-likelihood values when calculating updates.

`one_dimensional` If TRUE, collapse clever covariates into a one-dimensional clever covariate scaled by the mean of their EIFs.

`constrain_step` If TRUE, step size is at most `delta_epsilon` (it can be smaller if a smaller step decreases the loss more).

`delta_epsilon` The maximum step size allowed if `constrain_step` is TRUE.

`convergence_type` The convergence criterion to use: (1) "scaled\_var" corresponds to  $\sqrt{\text{Var}(D)/n}/\log n$  (the default) while (2) "sample\_size" corresponds to  $1/n$ .

`fluctuation_type` Whether to include the auxiliary covariate for the fluctuation model as a covariate or to treat it as a weight. Note that the option "weighted" is incompatible with a multi-epsilon submodel (`one_dimensional = FALSE`).

`use_best` If TRUE, the final updated likelihood is set to the likelihood that minimizes the ED instead of the likelihood at the last update step.

`verbose` If TRUE, diagnostic output is generated about the updating procedure.

---

tmle3\_Update\_survival *Defines an update procedure (submodel+loss function) for survival data*

---

### Description

Current Limitations: loss function and submodel are hard-coded (need to accept arguments for these)

### Constructor

```
define_param(maxit, cvtmle, one_dimensional, constrain_step, delta_epsilon, verbose)
```

maxit The maximum number of update iterations

cvtmle If TRUE, use CV-likelihood values when calculating updates.

one\_dimensional If TRUE, collapse clever covariates into a one-dimensional clever covariate scaled by the mean of their EIFs.

constrain\_step If TRUE, step size is at most delta\_epsilon (it can be smaller if a smaller step decreases the loss more).

delta\_epsilon The maximum step size allowed if constrain\_step is TRUE.

convergence\_type The convergence criterion to use: (1) "scaled\_var" corresponds to  $\sqrt{\text{Var}(D)/n}/\log n$  (the default) while (2) "sample\_size" corresponds to  $1/n$ .

fluctuation\_type Whether to include the auxiliary covariate for the fluctuation model as a covariate or to treat it as a weight. Note that the option "weighted" is incompatible with a multi-epsilon submodel (one\_dimensional = FALSE).

verbose If TRUE, diagnostic output is generated about the updating procedure.

---

tmle3\_vim *Compute Variable Importance Measures (VIM) with any given parameter*

---

### Description

Compute Variable Importance Measures (VIM) with any given parameter

### Usage

```
tmle3_vim(
  tmle_spec,
  data,
  node_list,
  learner_list = NULL,
  adjust_for_other_A = TRUE
)
```

**Arguments**

tmle\_spec      [tmle3\\_Spec](#), defines the TMLE  
 data            data.frame, the raw data  
 node\_list      list, defines which variables are which nodes  
 learner\_list   list, defines which learners are used to fit which likelihood factors  
 adjust\_for\_other\_A      Whether or not to adjust for other specified intervention nodes.

---

tmle_ATC	<i>All Treatment Specific Means</i>
----------	-------------------------------------

---

**Description**

O=(W,A,Y) W=Covariates A=Treatment (binary or categorical) Y=Outcome (binary or bounded continuous)

**Usage**

```
tmle_ATC(treatment_level, control_level)
```

**Arguments**

treatment\_level      the level of A that corresponds to treatment  
 control\_level      the level of A that corresponds to a control or reference level

---

tmle_ATE	<i>All Treatment Specific Means</i>
----------	-------------------------------------

---

**Description**

O=(W,A,Y) W=Covariates A=Treatment (binary or categorical) Y=Outcome (binary or bounded continuous)

**Usage**

```
tmle_ATE(treatment_level, control_level)
```

**Arguments**

treatment\_level      the level of A that corresponds to treatment  
 control\_level      the level of A that corresponds to a control or reference level



---

tmle_ATT	<i>All Treatment Specific Means</i>
----------	-------------------------------------

---

**Description**

O=(W,A,Y) W=Covariates A=Treatment (binary or categorical) Y=Outcome (binary or bounded continuous)

**Usage**

```
tmle_ATT(treatment_level, control_level)
```

**Arguments**

treatment\_level            the level of A that corresponds to treatment  
control\_level            the level of A that corresponds to a control or reference level

---

tmle_MSM	<i>Make MSM version of Stratified TML estimator class</i>
----------	---

---

**Description**

O=(W,A,Y) W=Covariates A=Treatment (binary or categorical) Y=Outcome (binary or bounded continuous)

**Usage**

```
tmle_MSM(weight = "Cond.Prob.", n_samples = 30)
```

**Arguments**

weight            h(A, V)  
n\_samples            number of samples to draw for each observation if A is continuous

---

tmle_OR	<i>Odds Ratio</i>
---------	-------------------

---

### Description

O = (W, A, Y) W = Covariates A = Treatment (binary or categorical) Y = Outcome (binary or bounded continuous)

### Usage

```
tmle_OR(baseline_level, contrast_level)
```

### Arguments

baseline\_level The baseline risk group.

contrast\_level The contrast risk group.

---

tmle_PAR	<i>PAR and PAF</i>
----------	--------------------

---

### Description

O=(W,A,Y) W=Covariates A=Treatment (binary or categorical) Y=Outcome (binary or bounded continuous)

### Usage

```
tmle_PAR(baseline_level)
```

### Arguments

baseline\_level the baseline risk group

---

tmle_RR	<i>Risk Ratio</i>
---------	-------------------

---

**Description**

O = (W, A, Y) W = Covariates A = Treatment (binary or categorical) Y = Outcome (binary or bounded continuous)

**Usage**

```
tmle_RR(baseline_level, contrast_level)
```

**Arguments**

baseline\_level The baseline risk group.

contrast\_level The contrast risk group.

---

tmle_stratified	<i>Stratified version of TML estimator from other Spec classes</i>
-----------------	--

---

**Description**

O=(W,A,Y) W=Covariates A=Treatment (binary or categorical) Y=Outcome (binary or bounded continuous)

**Usage**

```
tmle_stratified(base_spec, base_estimate = TRUE)
```

**Arguments**

base\_spec An underlying spec to stratify.

base\_estimate Indicate whether to report base parameter.

---

tmle_survival	<i>Treatment Specific Survival</i>
---------------	------------------------------------

---

**Description**

See the associated handbook chapter

**Usage**

```
tmle_survival(treatment_level, control_level, target_times = NULL, ...)
```

**Arguments**

treatment_level	the level of A that corresponds to treatment
control_level	the level of A that corresponds to a control or reference level
target_times	the time points to be targeted at during the TMLE adjustment
...	others args passed to spec

---

tmle_TSM_all	<i>All Treatment Specific Means</i>
--------------	-------------------------------------

---

**Description**

O=(W,A,Y) W=Covariates A=Treatment (binary or categorical) Y=Outcome (binary or bounded continuous)

**Usage**

```
tmle_TSM_all()
```

---

train_lf	<i>Manually Train Likelihood Factor The internal training process for likelihood factors is somewhat obtuse, so this function does the steps to manually train one, which is helpful if you want to use a likelihood factor independently of a likelihood object</i>
----------	--

---

**Description**

Manually Train Likelihood Factor The internal training process for likelihood factors is somewhat obtuse, so this function does the steps to manually train one, which is helpful if you want to use a likelihood factor independently of a likelihood object

**Usage**

```
train_lf(lf, tmle_task)
```

**Arguments**

lf	the likelihood factor to train
tmle_task	the task to use for training

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