

# Package ‘savvyr’

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**Type** Package

**Title** Survival Analysis for AdVerse Events with VarYing Follow-Up Times

**Version** 0.1.2

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**Description** The SAVVY (Survival Analysis for AdVerse Events with VarYing Follow-Up Times) project is a consortium of academic and pharmaceutical industry partners that aims to improve the analyses of adverse event (AE) data in clinical trials through the use of survival techniques appropriately dealing with varying follow-up times and competing events, see Stegherr, Schmoor, Beyersmann, et al. (2021) <[doi:10.1186/s13063-021-05354-x](https://doi.org/10.1186/s13063-021-05354-x)>. Although statistical methodologies have advanced, in AE analyses often the incidence proportion, the incidence density or a non-parametric Kaplan-Meier estimator are used, which either ignore censoring or competing events. This package contains functions to easily conduct the proposed improved AE analyses.

**License** Apache License 2.0

**URL** <https://openpharma.github.io/savvyr/>

**BugReports** <https://github.com/openpharma/savvyr/issues>

**Depends** R (>= 4.0)

**Imports** checkmate, etm, Rdpack

**Suggests** kableExtra, knitr, rmarkdown, testthat (>= 3.0)

**Config/Needs/documentation** roxytypes, roxylint

**RdMacros** Rdpack

**VignetteBuilder** knitr

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**NeedsCompilation** no

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aalen_johansen	<i>Aalen Johansen Estimator</i>
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Description

This function calculates the Aalen Johansen estimator of adverse events observed in  $[0, \tau]$ . Please also refer to Stegherr et al. (2021).

Usage

```
aalen_johansen(data, ce, tau)
```

Arguments

- |      |  |
|------|--|
| data | (data.frame)<br>with columns including <ul style="list-style-type: none"><li>time_to_event: Time to the first AE, death or soft competing event.</li><li>type_of_event: 0 for censored, 1 for AE, 2 for death, 3 for soft competing event.</li></ul> |
| ce   | (number)<br>code for competing event.  |
| tau  | (number)<br>milestone at which Aalen-Johansen is computed.   |

**Value**

(vector)

with the following entries:

- ae\_prob: Estimated probability of AE.
- ae\_prob\_var: Variance of that estimate.
- ce\_prob: Estimated probability of competing events.
- ce\_prob\_var: Variance of competing events.

**References**

Stegherr R, Schmoor C, Lübbert M, Friede T, Beyersmann J (2021). “Estimating and comparing adverse event probabilities in the presence of varying follow-up times and competing events.” *Pharmaceutical Statistics*, **20**(6), 1125–1146. doi:10.1002/pst.2130, <https://onlinelibrary.wiley.com/doi/abs/10.1002/pst.2130>.

**Examples**

```
set.seed(123)
dat <- generate_data(n = 5, cens = c(2, 5), haz_ae = 2, haz_death = 3, haz_soft = 5)
aalen_johansen(dat, ce = 2, tau = 4)
```

generate\_data

*Generate Example Data***Description**

This generates the dataset denoted by S1 in Table 4 of Stegherr et al. (2021), i.e. we assume constant hazards for the adverse event (AE) hazard, the hazard for the competing event of death, and the hazard for the "soft" competing events. Censoring is uniform in the given range.

**Usage**

```
generate_data(n, cens, haz_ae, haz_death, haz_soft)
```

**Arguments**

n	(count) number of patients.
cens	(numeric) minimum and maximum censoring time.
haz_ae	(number) constant hazard for AE.
haz_death	(number) constant hazard for death.
haz_soft	(number) constant hazard for soft competing event.

**Value**

(data.frame)

a data.frame with the following columns:

- id: Patient ID.
- time\_to\_event: Time to the first AE, death or soft competing event.
- type\_of\_event: 0 for censored, 1 for AE, 2 for death, 3 for soft competing event.
- cens: Censoring time.

**References**

Stegherr R, Schmoor C, Lübbert M, Friede T, Beyersmann J (2021). “Estimating and comparing adverse event probabilities in the presence of varying follow-up times and competing events.” *Pharmaceutical Statistics*, **20**(6), 1125–1146. doi:10.1002/pst.2130, <https://onlinelibrary.wiley.com/doi/abs/10.1002/pst.2130>.

**Examples**

```
set.seed(123)
generate_data(n = 5, cens = c(2, 5), haz_ae = 2, haz_death = 3, haz_soft = 5)
```

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inc\_prop

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*Computing the Incidence Proportion*


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**Description**

Computing the Incidence Proportion

**Usage**

```
inc_prop(data, tau)
```

**Arguments**

data	(data.frame) with columns including: <ul style="list-style-type: none"> <li>• time_to_event: Time to the first AE, death or soft competing event.</li> <li>• type_of_event: 0 for censored, 1 for AE, 2 for death, 3 for soft competing event.</li> </ul>
tau	(numeric) milestone at which incidence proportion is computed.

**Value**

(vector)  
with the following entries:

- `ae_prob`: Estimated probability of AE.
- `ae_prob_var`: Variance of that estimate.

**Examples**

```
set.seed(123)
dat <- generate_data(
  n = 5,
  cens = c(2, 5),
  haz_ae = 2,
  haz_death = 3,
  haz_soft = 5
)

inc_prop(dat, tau = 4)
```

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one\_minus\_kaplan\_meier

*One Minus Kaplan-Meier*

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**Description**

This function calculates the one minus Kaplan-Meier estimator of adverse events (while censoring all competing events) observed in  $[\emptyset, \tau]$ . Please also refer to formula (4) in Stegherr et al. (2021).

**Usage**

```
one_minus_kaplan_meier(data, tau)
```

**Arguments**

<code>data</code>	(data.frame) with columns including <ul style="list-style-type: none"> <li>• <code>time_to_event</code>: Time to the first AE, death or soft competing event.</li> <li>• <code>type_of_event</code>: 0 for censored, 1 for AE, 2 for death, 3 for soft competing event.</li> </ul>
<code>tau</code>	(number) milestone at which One Minus Kaplan-Meier is computed.

**Value**

(vector)

with the following entries:

- ae\_prob: Estimated probability of AE.
- ae\_prob\_var: Variance of that estimate.

**References**

Stegherr R, Beyersmann J, Jehl V, Rufibach K, Leverkus F, Schmoor C, Friede T (2021). “Survival analysis for Adverse events with VarYing follow-up times (SAVVY): Rationale and statistical concept of a meta-analytic study.” *Biometrical Journal*, **63**(3), 650-670. doi:10.1002/bimj.201900347, <https://onlinelibrary.wiley.com/doi/pdf/10.1002/bimj.201900347>, <https://onlinelibrary.wiley.com/doi/abs/10.1002/bimj.201900347>.

**Examples**

```
set.seed(123)
dat <- generate_data(n = 5, cens = c(2, 5), haz_ae = 2, haz_death = 3, haz_soft = 5)
one_minus_kaplan_meier(dat, tau = 4)
```

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prop_trans_inc_dens	<i>Computing the Probability Transform Incidence Density</i>
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**Description**

Computing the Probability Transform Incidence Density

**Usage**

```
prop_trans_inc_dens(data, tau)
```

**Arguments**

data	(data.frame) with columns including <ul style="list-style-type: none"> <li>• time_to_event: Time to the first AE, death or soft competing event.</li> <li>• type_of_event: 0 for censored, 1 for AE, 2 for death, 3 for soft competing event.</li> </ul>
tau	(number) milestone at which Probability Transform Incidence Density is computed.

**Value**

(vector)

with the following entries:

- ae\_prob: Estimated probability of AE.
- ae\_prob\_var: Variance of that estimate.

**Examples**

```
set.seed(123)
dat <- generate_data(n = 5, cens = c(2, 5), haz_ae = 2, haz_death = 3, haz_soft = 5)
prop_trans_inc_dens(dat, tau = 4)
```

---

prop\_trans\_inc\_dens\_ce

*Probability Transform Incidence Density Accounting for Competing Events*

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**Description**

This function calculates the incidence density of both adverse events and specified competing events observed in  $[0, \tau]$  and then combines and transforms the incidence densities on a probability scale. Please also refer to formulas (4) and (5) in Stegherr et al. (2021).

**Usage**

```
prop_trans_inc_dens_ce(data, ce, tau)
```

**Arguments**

data	(data.frame) with columns including <ul style="list-style-type: none"> <li>time_to_event: Time to the first AE, death or soft competing event.</li> <li>type_of_event: 0 for censored, 1 for AE, 2 for death, 3 for soft competing event.</li> </ul>
ce	(number) code for competing event.
tau	(number) milestone at which Probability Transform Incidence Density is computed.

**Value**

(vector)  
with the following entries:

- ae\_prob: Estimated probability of AE.
- ae\_prob\_var: Variance of that estimate.

**References**

Stegherr R, Schmoor C, Beyersmann J, Rufibach K, Jehl V, Brückner A, Eisele L, Künzel T, Kupas K, Langer F, Leverkus F, Loos A, Norenberg C, Voss F, Friede T (2021). “Survival analysis for Adverse events with VarYing follow-up times (SAVVY)—estimation of adverse event risks.” *Trials*, 22(1), 420. ISSN 1745-6215, doi:[10.1186/s1306302105354x](https://doi.org/10.1186/s1306302105354x).

**Examples**

```
set.seed(123)
dat <- generate_data(n = 5, cens = c(2, 5), haz_ae = 2, haz_death = 3, haz_soft = 5)
prop_trans_inc_dens_ce(dat, ce = 2, tau = 4)
```



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