

Interpretable Machine Learning for Survival Analysis

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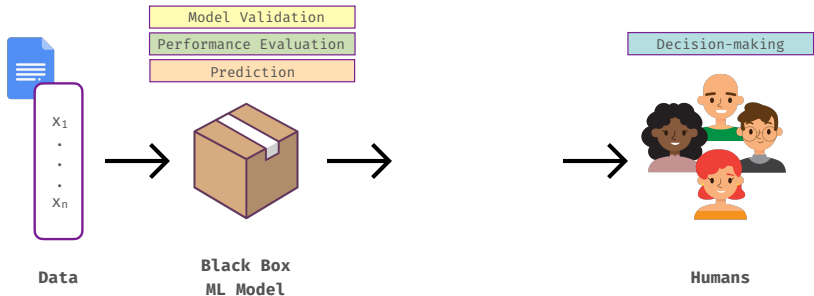
2023-09-05

Survival Analysis for Junior Researchers (SAfJR) 2023

Interpretable Machine Learning (IML)



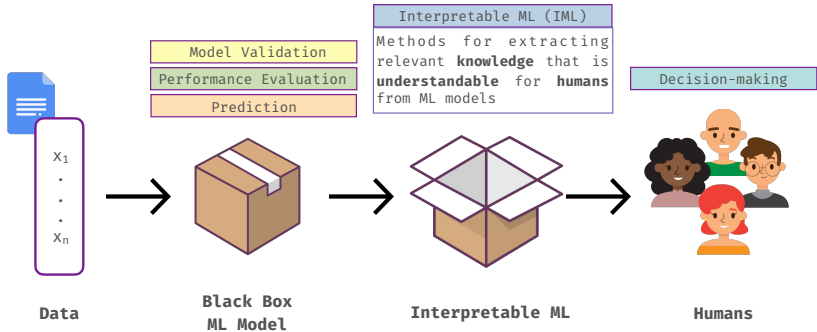
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Interpretable Machine Learning (IML)



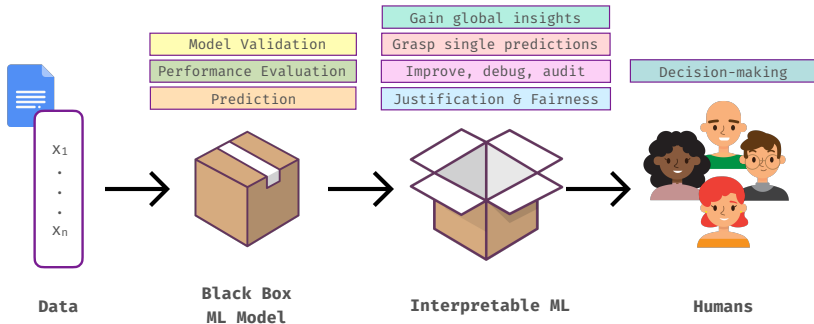
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Interpretable Machine Learning (IML)



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Interpretable Machine Learning (IML) for Survival Analysis



5

Why do we need IML specifically for Survival Analysis?

Interpretable Machine Learning (IML) for Survival Analysis



5

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1. **Lack of readily available IML methods** for functional outputs, time-to-event data, and competing risk data

Interpretable Machine Learning (IML) for Survival Analysis



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Why do we need IML specifically for Survival Analysis?

1. **Lack of readily available IML methods** for functional outputs, time-to-event data, and competing risk data
2. Applications with elevated need for **transparency, accountability, and fairness**



survex : CHEAT SHEET

Explore, Explain, and Examine Survival Models with the survex package

Survival analysis models are commonly used in medicine and other areas. Many of them are too complex to be interpreted by human. Explanation and examination is needed, but standard methods do not give a broad enough picture. **survex** provides easy-to-use methods for explaining survival models, both complex black-boxes and simpler statistical models.

Explainer

The **survex** package operates on the **explainer** objects. They can be used for calculating explanations, measuring model performance, and making predictions.

model $\xrightarrow{\text{explain()}}$ explainer

```
library(survex)
library(survival)
library(randomForestSRC)
rsf_model <- rsrc(
  Surv(time, status) ~.,
  data = veteran)
explainer <- explain(rsf_model)
```

For some models explainer are created automatically with the **explain()** function (only **model** argument is required). However, an explainer can be created for **any** survival model using the **explain_survival()** function.

Remember to:

- provide **data** as a data.frame without target column,
- provide **y** as a survival:Surv object,
- provide **predict_survival** function as a function of (model, newdata, times).

This is all you need for a fully functional explainer but you can also set:

- **predict** function,
- **predict_cumulative_hazard** function,
- **times** (for making functional predictions) on your own.

Global explanations

explain model's predictions for an entire dataset

VARIABLE IMPORTANCE

Which variables are important to the model?

```
m_parts <- model_parts(explainer)
loss_function, type, output_type
plot(m_parts)
```

Time-dependent feature importance

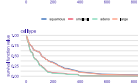
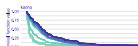


PARTIAL DEPENDENCE

How does a variable affect the average prediction?

```
m_profile <- model_profile(explainer)
variable, categorical_variables,
groups_N
plot(m_profile,
  numerical_plot_type='lines')
```

Partial dependence survival profile



Local explanations

explain model's prediction for a single observation

VARIABLE ATTRIBUTIONS

Which variables contribute to the prediction?

```
SurvSHAP()
survshap <- predict_parts(explainer,
  patient, type='survshap')
aggregation_method, B
plot(survshap)
```

SurvSHAP() model for the rsf_model



```
SurvLIME
survlime <- predict_parts(explainer,
  patient, type='survlime')
kernel_width, N
plot(survlime)
```

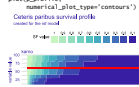


CETERIS PARIBUS

How does a variable affect the prediction?

```
p_profile <- predict_profile(explainer,
  variable, variable_type, patient)
plot(p_profile,
  numerical_plot_type='contours')
```

Ceteris paribus survival profile



Performance

examine model's quality

MODEL PERFORMANCE

How good is the model?

```
m_perf <- model_performance(explainer)
cph_perf <- model_performance(cph_exp,
  type='times')
plot(m_perf, cph_perf)
```

Model performance created for the rsf_model



```
plot(m_perf, cph_perf,
  metrics_type='scalar')
```



Prediction

explore model's predictions

MAKING PREDICTIONS

What is the model's prediction?

```
Survival function
predict(explainer, veteran,
  output_type='survival')
```

Cumulative hazard function

```
predict(explainer, veteran,
  output_type='chf')
```

Risk score/prognostic index

```
predict(explainer, veteran,
  output_type='risk')
```



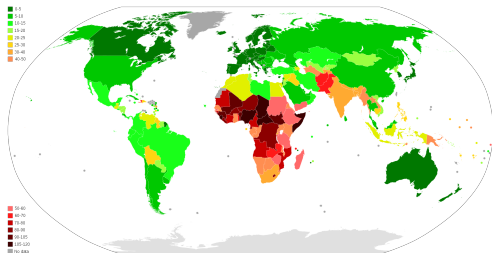
Demographic and Health Surveys (DHS) Data



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Nationally-representative household surveys to collect indicators in the areas of population, health, and nutrition from US-AID

Goal: Predict under-5 year mortality of children in Ghana in 2014 (N=7734)



Under-5 mortality rate:
World average: 36.6/1000
Ghana: 44.7/1000
Germany: 3.7/1000

Demographic and Health Surveys (DHS) Data



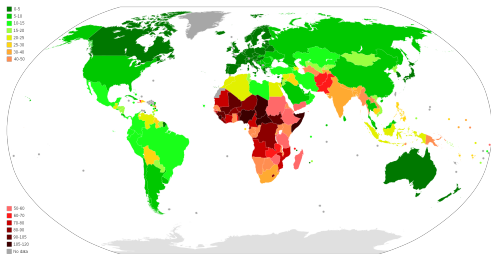
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Nationally-representative household surveys to collect indicators in the areas of population, health, and nutrition from US-AID

Goal: Predict **under-5 year mortality** of children in **Ghana** in 2014 (N=7734)

Features:

- Sex of child
- Mother place of residence
- Mother wealth index
- Mother age
- Mother education
- BMI mother



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Ghana: 44.7/1000
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Model Fitting and Model Performance



Fit ML model to make predictions:

Classic setting:

$$\hat{y} = \hat{f}(\mathbf{x}), \text{ given } (\mathbf{x}^i, y^i)_{i=1}^N$$

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Survival setting:

$$\hat{\mathbb{P}}(T > t) = \hat{S}(t|\mathbf{x}), \text{ given } (\mathbf{x}^i, \delta^i, T^i)_{i=1}^N$$

Model Fitting and Model Performance



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Fit ML model to make predictions:

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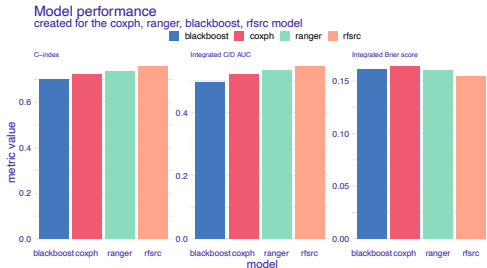
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Fitted models:

- **coxph**: CoxPH Model
- **blackboost**: Gradient boosting with regression trees survival learner
- **ranger**: Ranger random survival forest learner
- **rfsrc**: Survival random forest SRC learner



Feature Effects: Individual Conditional Expectation (ICE)



9

Local feature effects: Visualize or quantify marginal contribution of a feature w.r.t. predictions **for each observation**

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4. Use grid points to make predictions

$$\hat{f}^{(i)}(\mathbf{x}_A^{*(g)}) = \hat{f}^{(i)}(\mathbf{x}_A^{*(g)}, \mathbf{x}_{-A}^{(i)}), \quad \forall \mathbf{x}_A^{*(g)}, \forall i$$

5. Plot point pairs

$$\{(\mathbf{x}_A^{*(g)}, \hat{f}^{(i)}(\mathbf{x}_A^{*(g)}))\}_{g=1}^G$$

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$$\hat{S}^{(i)}(t|\mathbf{x}_A^{*(g)}) = \hat{S}^{(i)}(t|\mathbf{x}_A^{*(g)}, \mathbf{x}_{-A}^{(i)}), \quad \forall \mathbf{x}_A^{*(g)}, \forall i, \forall t \in \{1, \dots, T\}$$

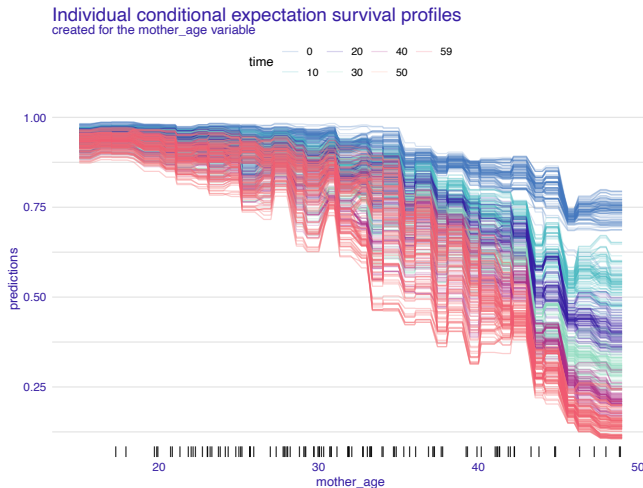
5. Plot point pairs

$$\left\{ (\mathbf{x}_A^{*(g)}, \hat{f}^{(i)}(\mathbf{x}_A^{*(g)})) \right\}_{g=1}^G \qquad \left\{ (\mathbf{x}_A^{*(g)}, \hat{S}^{(i)}(t|\mathbf{x}_A^{*(g)})) \right\}_{g=1}^G$$

ICE Curves: Age of Mother at Birth



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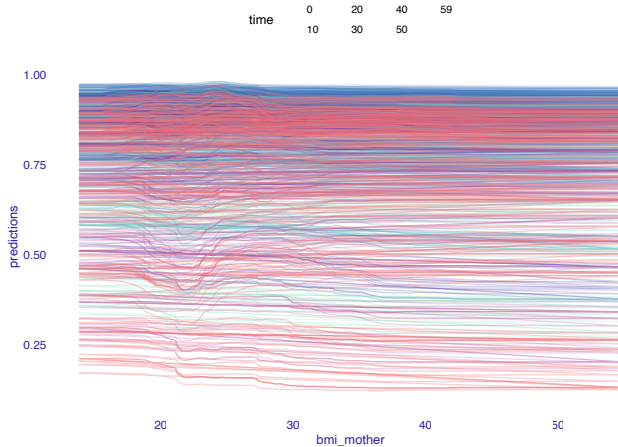
Note: These are **not** Kaplan-Meier survival curves!

ICE Curves: BMI of Mother



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Individual conditional expectation survival profiles
created for the bmi_mother variable



Feature Effects: Partial Dependence Plot (PDP)



Global feature effects: Visualize or quantify marginal contribution of a feature w.r.t. predictions **over all observations** 12

Feature Effects: Partial Dependence Plot (PDP)

Global feature effects: Visualize or quantify marginal contribution of a feature w.r.t. predictions over all observations

PD function: Expectation of prediction w.r.t. marginal distribution of \mathbf{x}_A

$$f_{PD}(\mathbf{x}_A) = \mathbb{E}_{\mathbf{x}_{-A}}(\hat{f}(\mathbf{x}_A, \mathbf{x}_{-A})) = \int_{-\infty}^{\infty} \hat{f}(\mathbf{x}_A, \mathbf{x}_{-A}) d\mathbb{P}(\mathbf{x}_{-A})$$

Estimation: For a grid value $\mathbf{x}_A^{*(g)}$, average ICE curves pointwise at $\mathbf{x}_A^{*(g)}$ over all observed $\mathbf{x}_{-A}^{*(i)}$

$$\hat{f}_{PD}(\mathbf{x}_A^{*(g)}) = \frac{1}{N} \sum_{i=1}^N \hat{f}(\mathbf{x}_A^{*(g)}, \mathbf{x}_{-A}^{*(i)})$$

Feature Effects: Partial Dependence Plot (PDP)



Global feature effects: Visualize or quantify marginal contribution of a feature w.r.t. predictions over all observations

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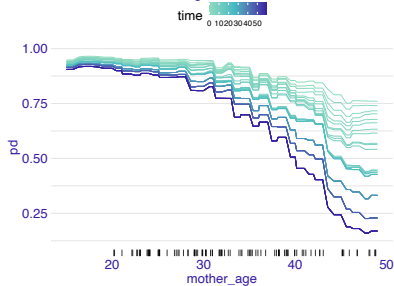
$$\hat{S}_{PD}(t|\mathbf{x}_A^{*(g)}) = \frac{1}{N} \sum_{i=1}^N \hat{S}(t|\mathbf{x}_A^{*(g)}, \mathbf{x}_{-A}^{*(i)})$$

PDP: Age of Mother at Birth

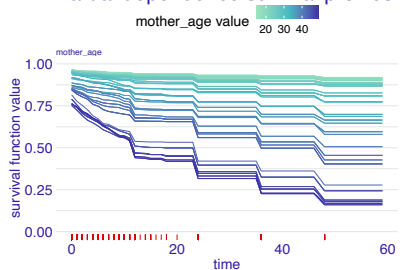


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Partial dependence survival profiles
created for the mother_age variable



Partial dependence survival profiles

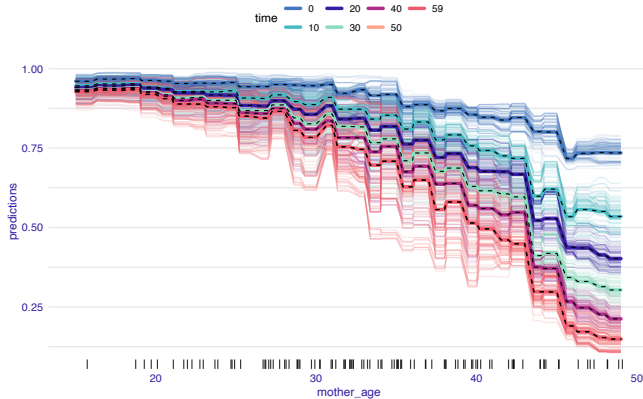


PDP+ICE: Age of Mother at Birth



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Individual conditional expectation and partial dependence survival profiles
created for the mother_age variable

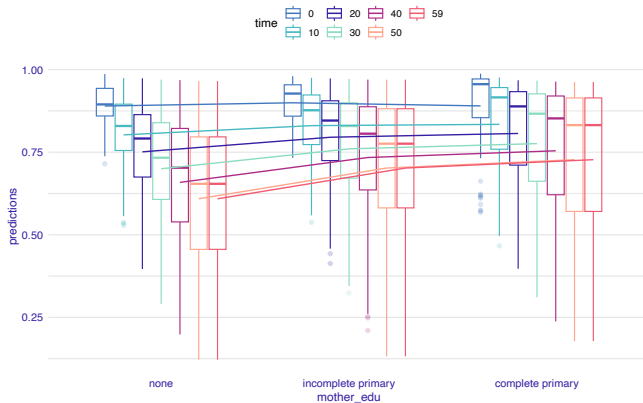


PDP+ICE: Mother's Highest Level of Education



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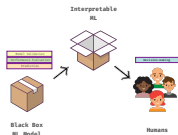
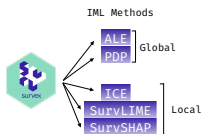
Individual conditional expectation and partial dependence survival profiles
created for the mother_edu variable



Conclusion

Why should you be interested in our research?

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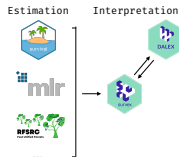


Crucial IML methods
readily available in
survex

Model
estimation and
interpretation
in **unified**
survex
interface

Pushing for
shift to
prioritize model
performance
without
sacrificing
interpretability

Promotion of
accountability,
transparency
and **fairness** in
survival model
interpretations



Thank you for your attention!

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Contact

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