

Visualizing Goodness of Fit of Probabilistic Regression Models

Achim Zeileis

<https://topmodels.R-Forge.R-project.org/>

Probabilistic regression models

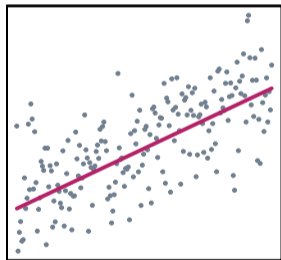
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Regression model: $\mu_i = r(\mathbf{x}_i)$

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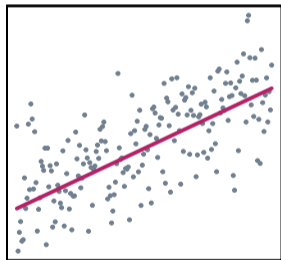


LM, GLM

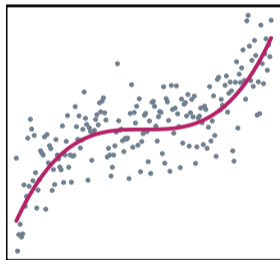
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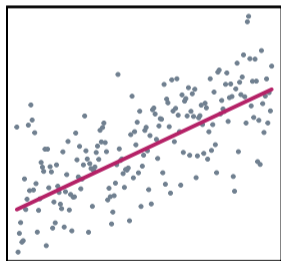


GAM

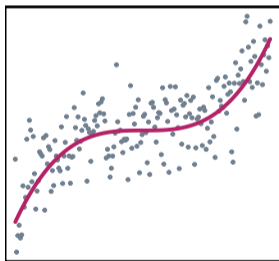
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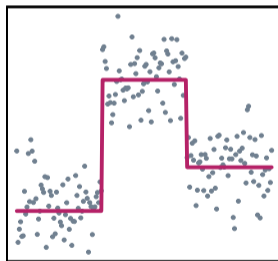
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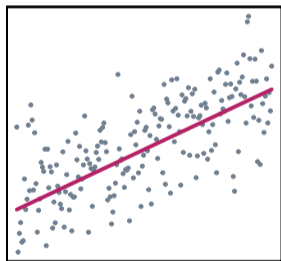


Regression tree

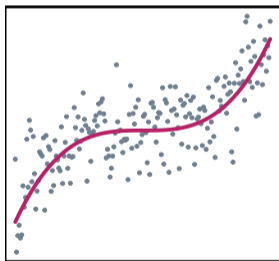
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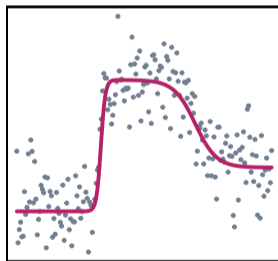
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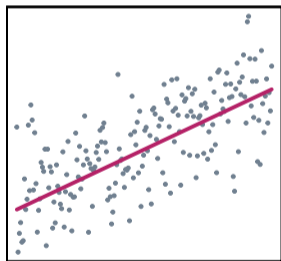
Random forest

Probabilistic regression models

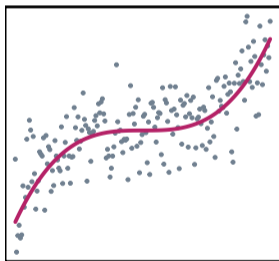
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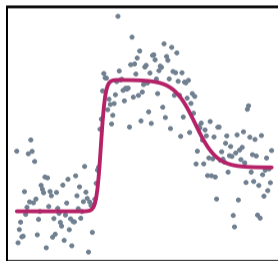
Often: Full conditional probability distribution is of interest.



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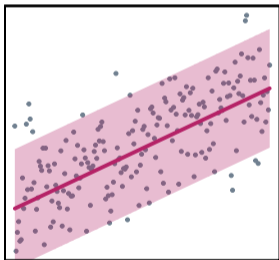
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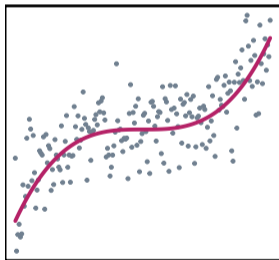
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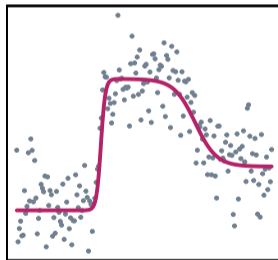
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Normal (G)LM w/ constant variance



GAM



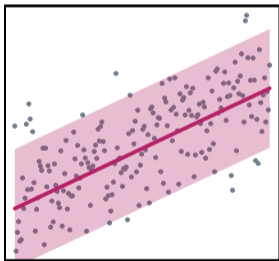
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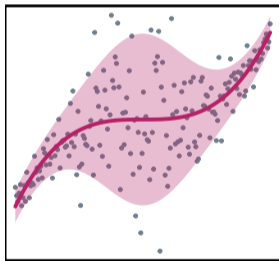
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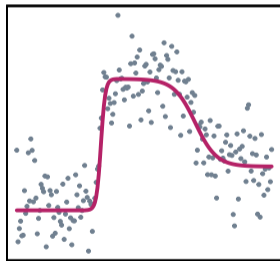
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GAMLSS



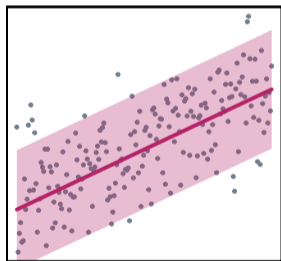
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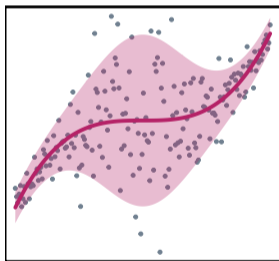
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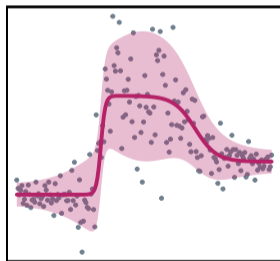
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Distributional forest

Probabilistic regression models

Formally: Fit distribution with cumulative distribution function $F(y_i|\theta_i)$ and parameter vector θ_i for each observation y_i .

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Forecasting: $\hat{\theta}_i = \hat{r}(\mathbf{x}_i)$.

- Model fit typically yields distribution parameters.
- Implies all other aspects of the distribution $F(\cdot|\theta_i)$.
- Thus: Moments, quantiles, probabilities, ...

Illustration: Goals in the 2018 FIFA World Cup

Response: Goals scored by the two teams in all 64 matches.

Covariates: Basic match information and prediction of team (log-)abilities (based on bookmakers odds).

```
R> data("FIFA2018", package = "distributions3")
```

```
R> tail(FIFA2018, 2)
```

	goals	team	match	type	stage	logability	difference
127	4	FRA	64	Final	knockout	0.8866	0.629
128	2	CRO	64	Final	knockout	0.2576	-0.629

Model: Poisson GLM with mean λ_i using log link.

Illustration: Goals in the 2018 FIFA World Cup

In R:

```
R> m <- glm(goals ~ difference, data = FIFA2018, family = poisson)
```

Forecasting: In-sample for simplicity.

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R> tail(procast(m), 2)
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Implies:

- Probabilities for match results (assuming independence of goals).
- Corresponding probabilities for win/draw/lose.

Illustration: Goals in the 2018 FIFA World Cup

Example: Probabilities for final France-Croatia.

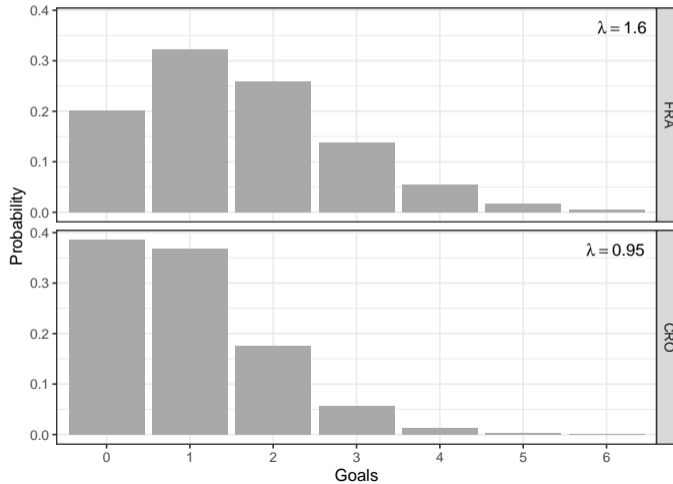
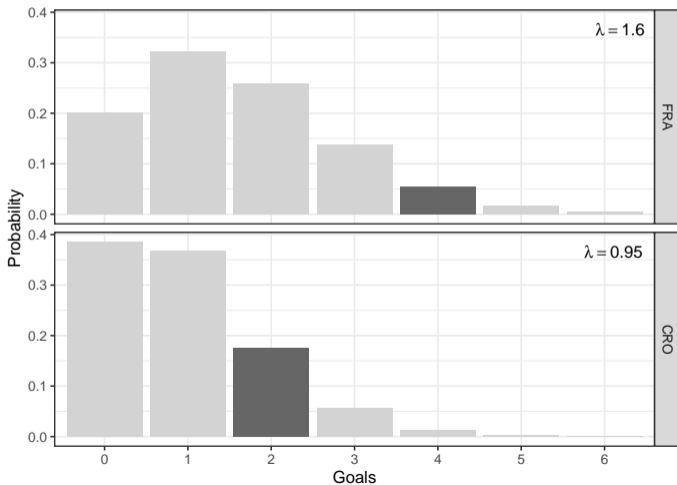


Illustration: Goals in the 2018 FIFA World Cup

Example: Probabilities for final France-Croatia. Result 4-2.



Goodness of fit

Idea:

- Use visualizations instead of just summing up scores.
- Gain more insights graphically.
- Reveal different types of model misspecification.

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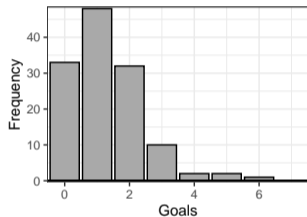
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Questions: Graphics are not new but novel unifying view.

- What are useful elements of such graphics?
- What are relative (dis)advantages?

Goodness of fit

Ideas: Illustrated for FIFA Poisson model.

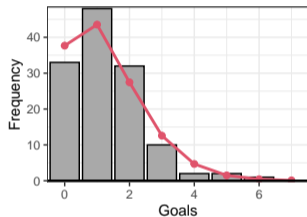


Marginal calibration:

- Observed frequencies.

Goodness of fit

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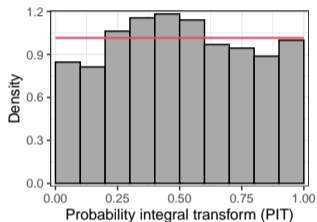
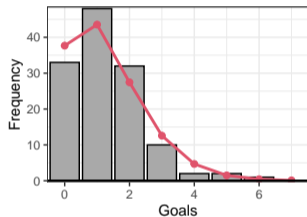


Marginal calibration:

- Observed frequencies.
- Compare: Expected.

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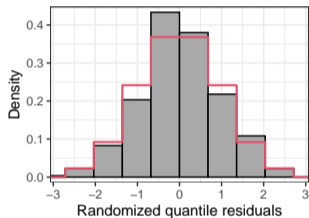
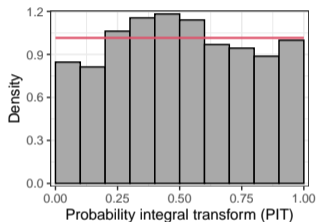
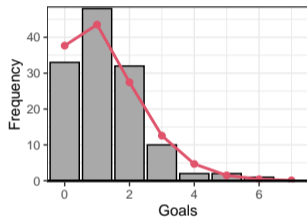
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Probabilistic calibration:

- Probability integral $u_i = F(y_i | \hat{\theta}_i)$.
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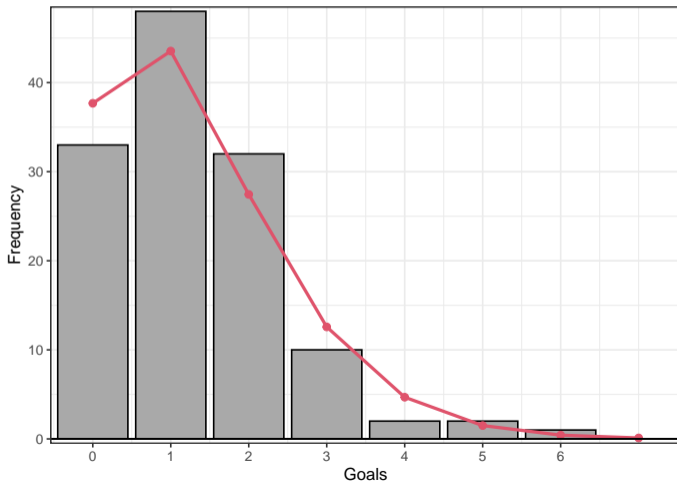
- Probability integral $u_i = F(y_i | \hat{\theta}_i)$.
- Compare: Uniform.

Probabilistic calibration:

- Quantile residuals $\Phi^{-1}(u_i)$.
- Compare: Normal

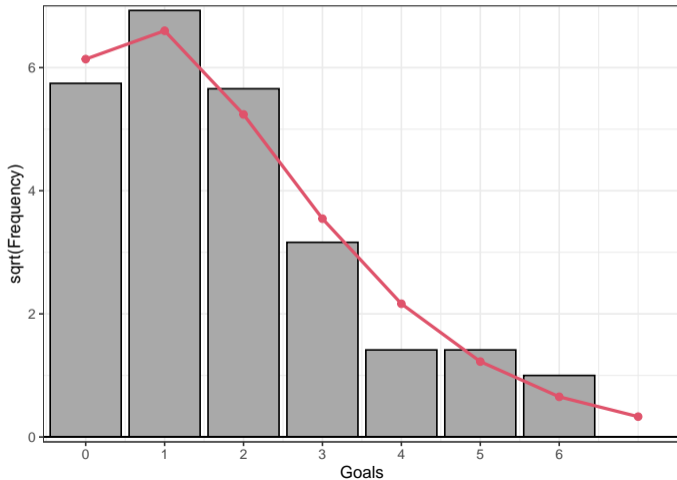
Goodness of fit: Marginal calibration

Observed vs. expected frequencies: Standing, with reference line.



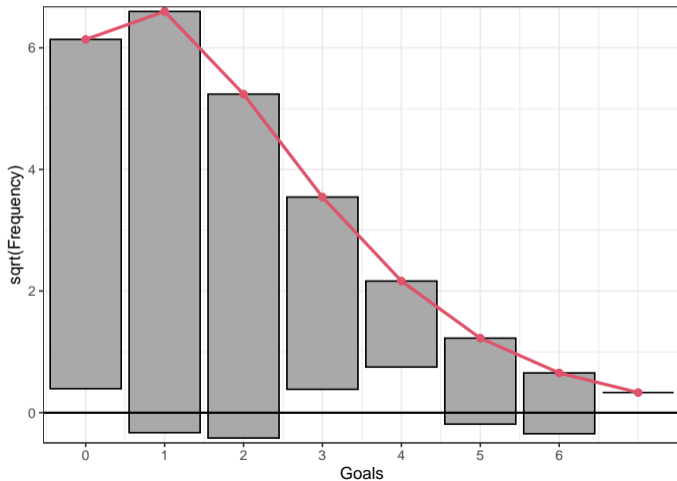
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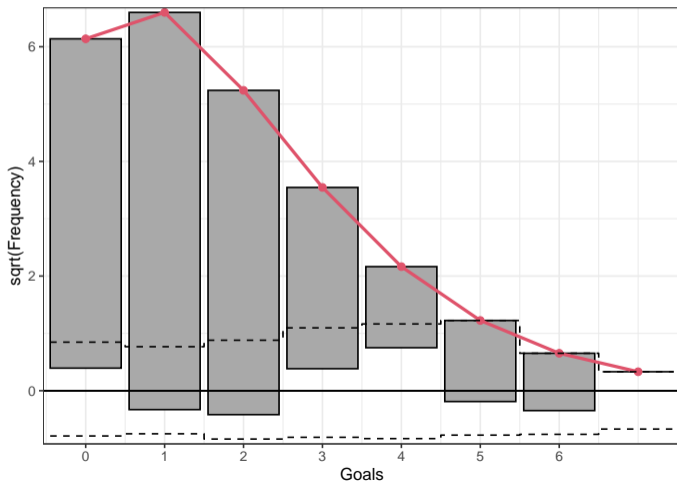
Goodness of fit: Marginal calibration

$\sqrt{\text{Observed}}$ vs. $\sqrt{\text{expected frequencies}}$: Hanging.



Goodness of fit: Marginal calibration

$\sqrt{\text{Observed}}$ vs. $\sqrt{\text{expected frequencies}}$: Hanging, with confidence interval.



Goodness of fit: Marginal calibration

Rootogram:

- Frequencies on raw or square-root scale.
- Hanging, standing, or suspended styled rootograms.

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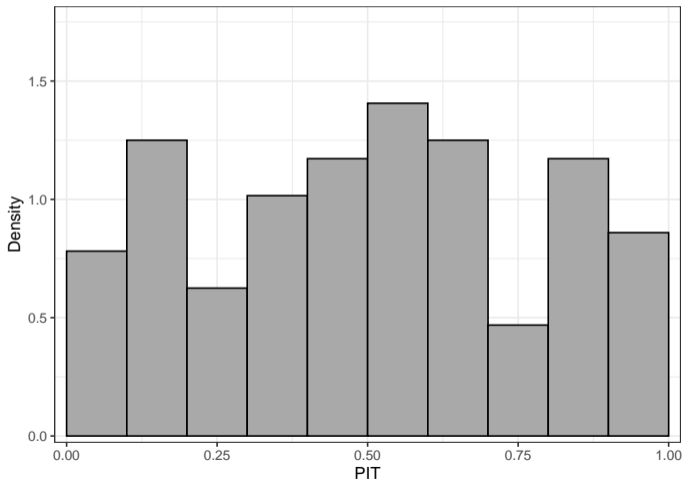
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Overall:

- *Advantage:* Scale of observations is natural, direct interpretation.
- *Disadvantage:* Needs to be compared with a combination of distributions.

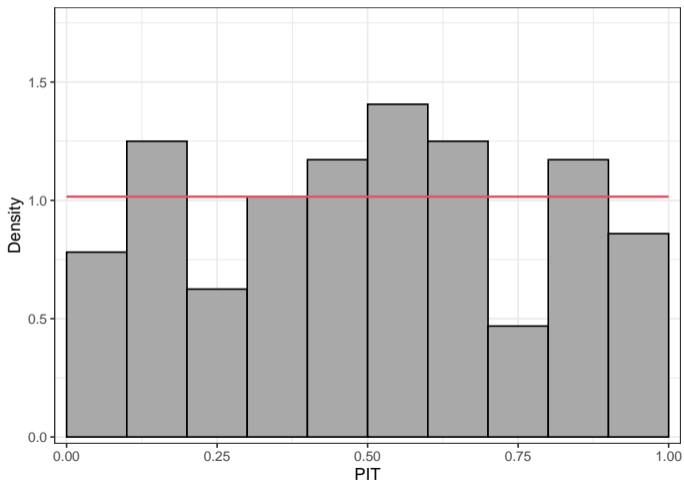
Goodness of fit: Probabilistic calibration

PIT: Randomization 1a.



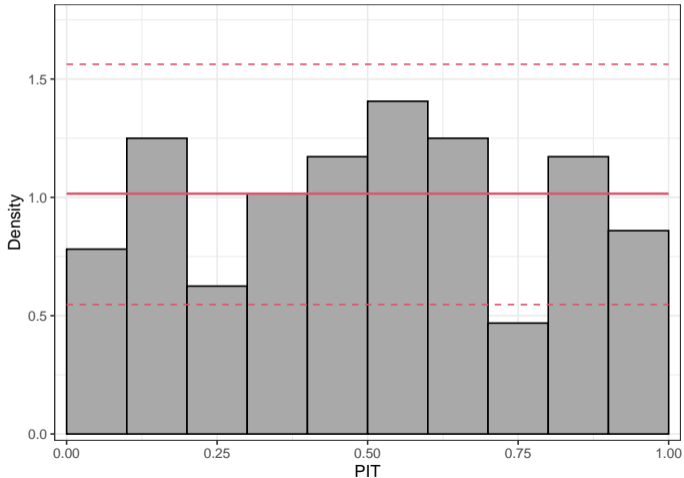
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PIT: Randomization 1a, with reference line.



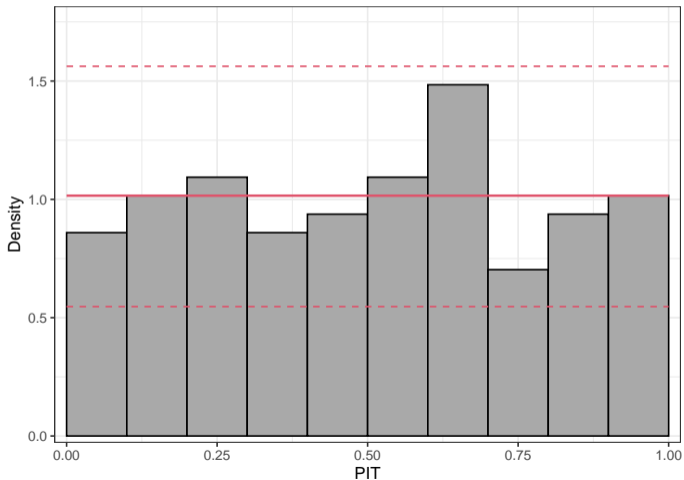
Goodness of fit: Probabilistic calibration

PIT: Randomization 1a, with reference line and confidence interval.



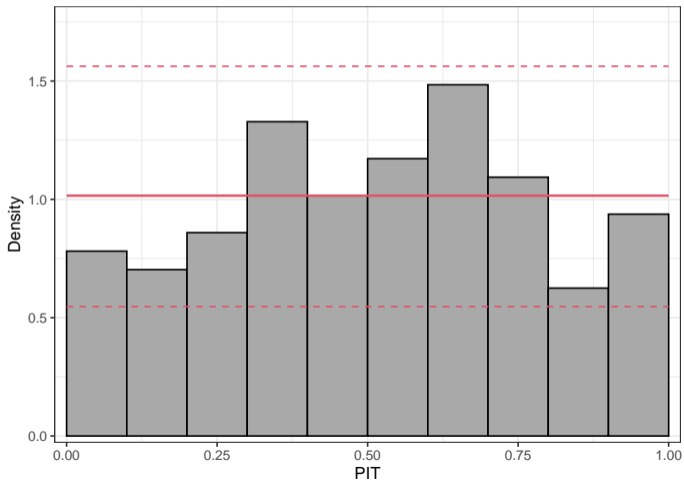
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PIT: Randomization 1b.



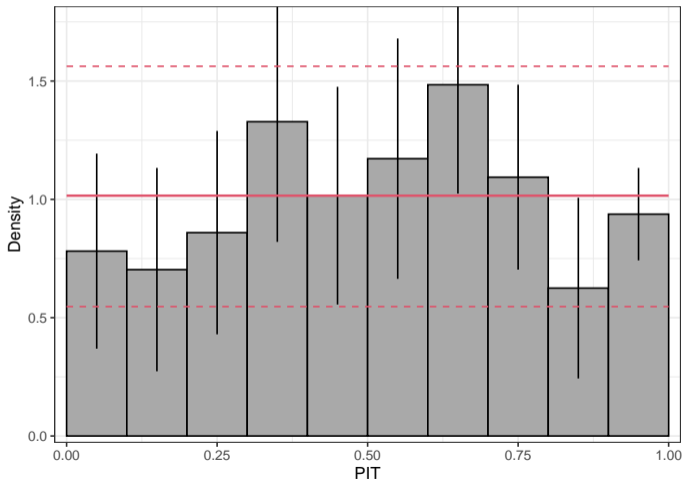
Goodness of fit: Probabilistic calibration

PIT: Randomization 1c.



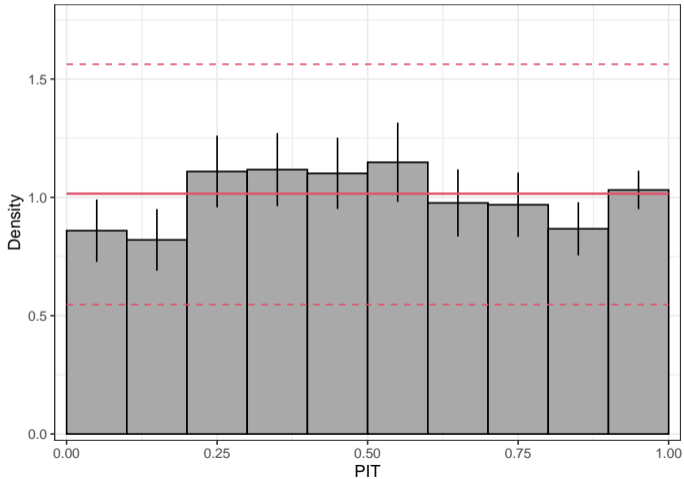
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PIT: Randomization 1c, with simulation intervals.



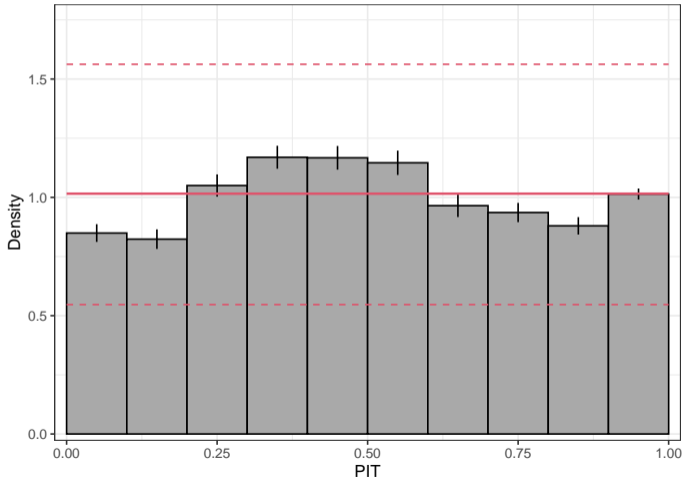
Goodness of fit: Probabilistic calibration

PIT: 10 random draws.



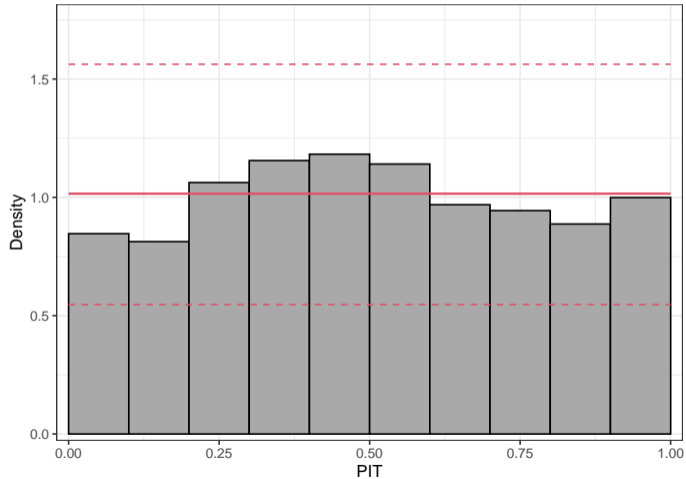
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PIT: 100 random draws.



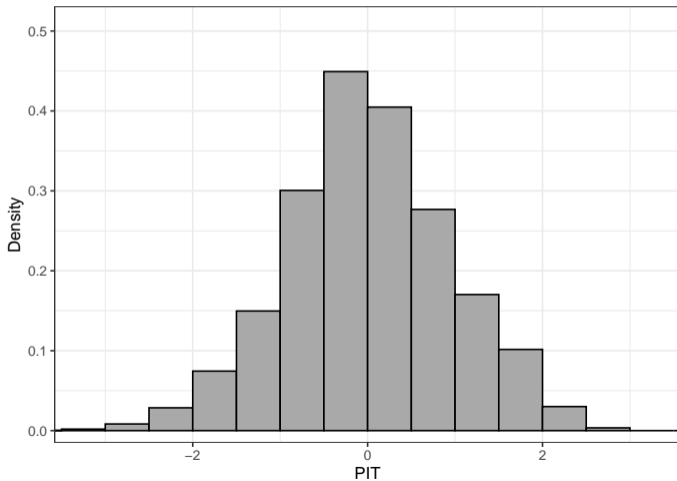
Goodness of fit: Probabilistic calibration

PIT: Expected.



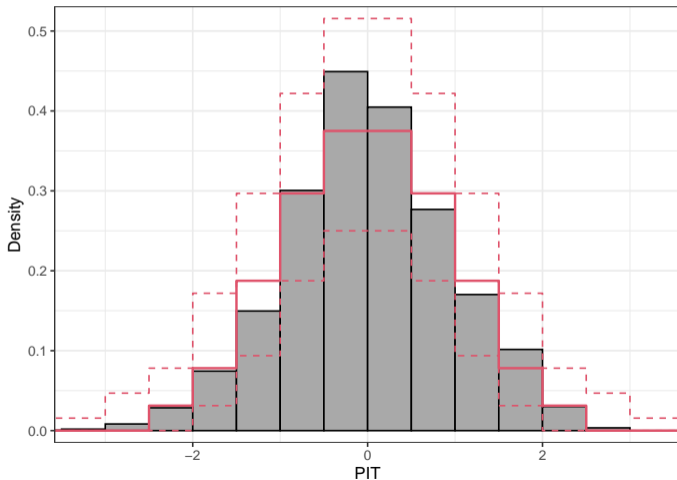
Goodness of fit: Probabilistic calibration

Randomized quantile residuals: Expected.



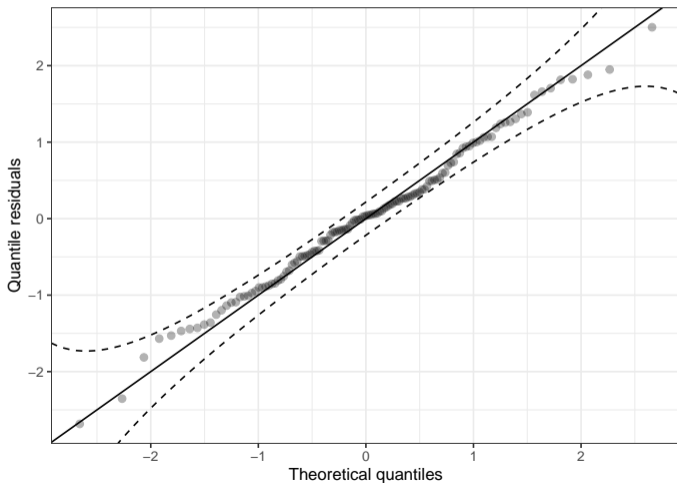
Goodness of fit: Probabilistic calibration

Randomized quantile residuals: Expected, with reference.



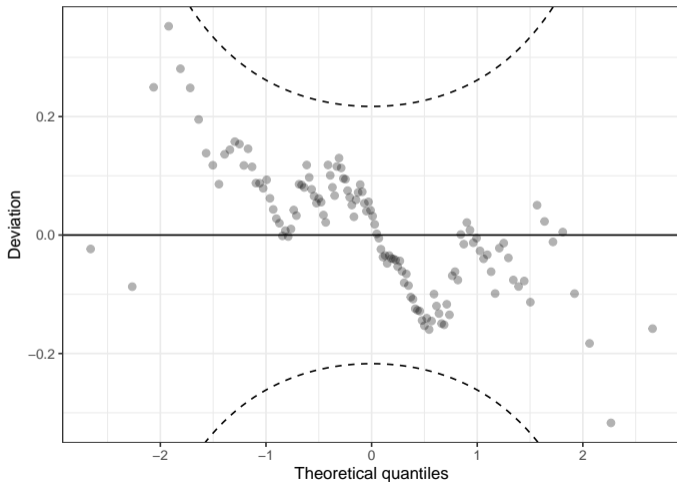
Goodness of fit: Probabilistic calibration

Observed vs. expected quantiles: Q-Q plot.



Goodness of fit: Probabilistic calibration

Observed vs. expected quantiles: Detrended Q-Q plot (worm plot).



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PIT histogram:

- Probability scale or transformed to normal scale.
- Randomized or expected for discrete distributions.

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- Normal or uniform scale.
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Q-Q residuals plot:

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Overall:

- *Advantage:* Comparison with only one distribution (uniform or normal).
- *Disadvantages:* Scale is not so natural. May require randomization.

Illustration: Loss aversion in adolescents

Experiment: Behaviour of adolescents (mostly 11–19).

- *Setup:* Nine rounds of a lottery with positive expectation.
- *Response:* Proportion of invested points across all rounds.
- *Covariates:* Arrangement (single vs. team), gender, age.

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Goodness of fit: Similar fitted means but rather different distributions.

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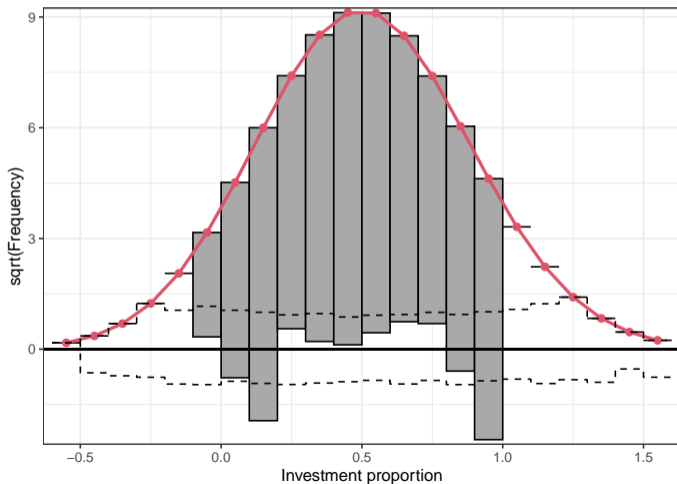


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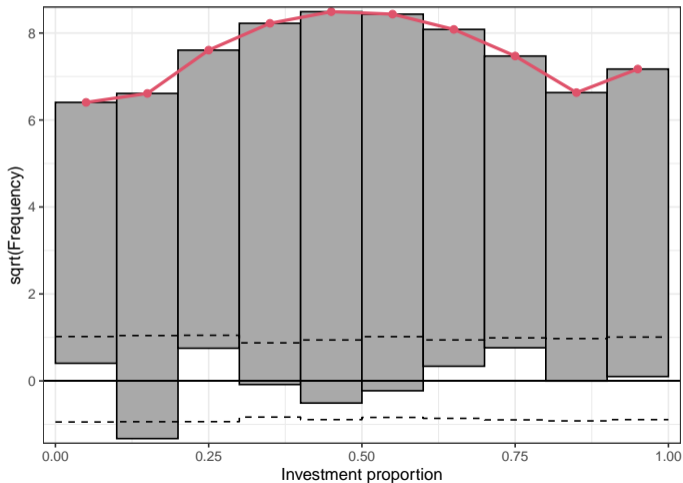


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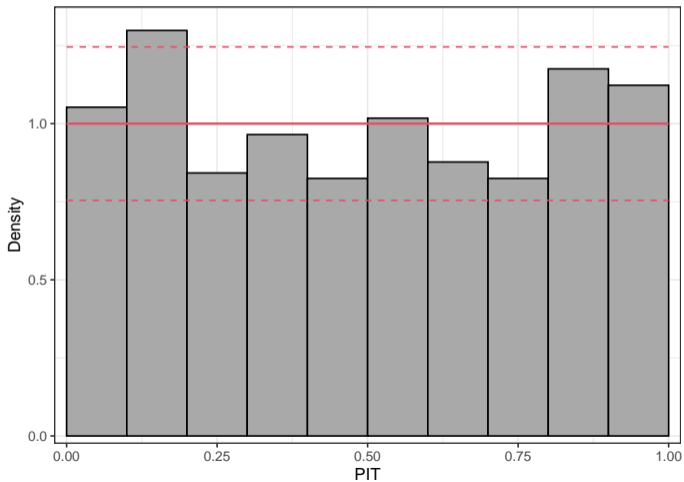


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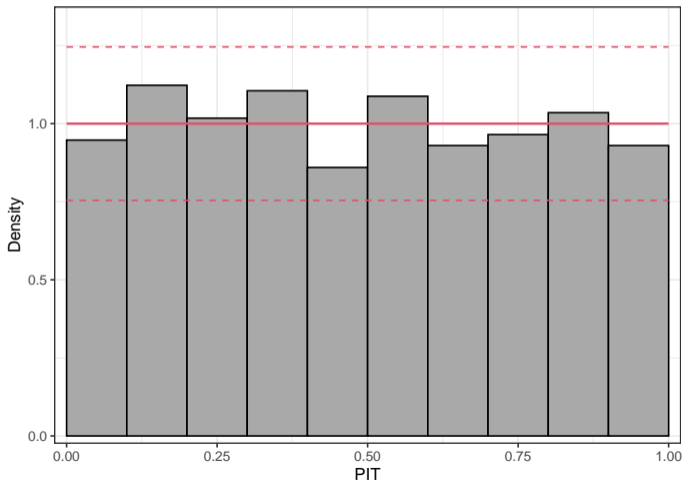


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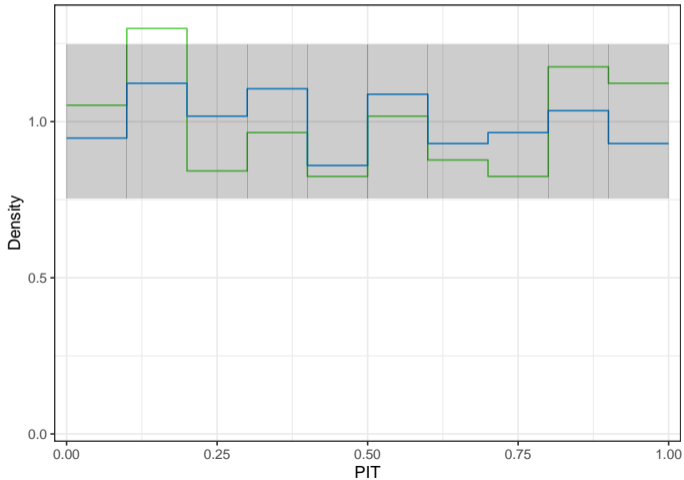


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Q-Q residual plot:

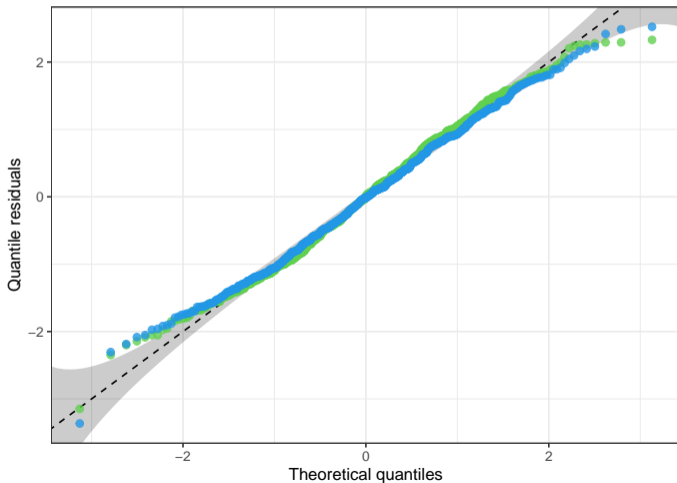
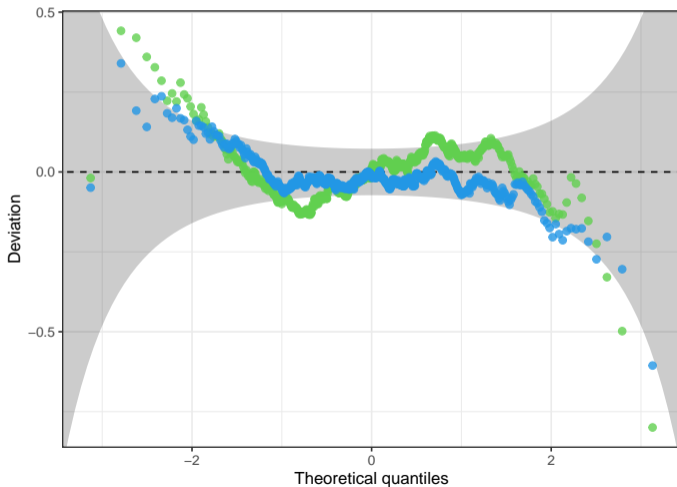


Illustration: Loss aversion in adolescents

Q-Q residual plot: Detrended.



Software: topmodels

R package: *topmodels*. Forecasting and assessment of probabilistic models.

Not yet on CRAN: <https://topmodels.R-Forge.R-project.org/>

Visualizations:

<code>rootogram()</code>	Rootograms of observed and fitted frequencies
<code>pithist()</code>	PIT histograms
<code>qqrplot()</code>	Q-Q plots for quantile residuals
<code>wormplot()</code>	Worm plots for quantile residuals
<code>reliagram()</code>	(Extended) reliability diagrams

Software: topmodels

Numeric quantities:

<code>procast()</code>	Probabilistic forecasts (probabilities, quantiles, etc.)
<code>proscore()</code>	Evaluate scoring rules for procasts
<code>pitresiduals()</code>	Probability integral transform (PIT) residuals
<code>qresiduals()</code>	(Randomized) quantile residuals

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Object orientation:

- Work with distribution objects (vectorized) from *distributions3*.
- Model classes like `lm`, `glm`, `gamlss`, `bamlss`, `hurdle`, `zeroinfl`, ...
- New model classes can be easily added if distribution can be extracted.

References

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