

A Toolbox for Probabilistic Regression Models

Forecasts, Visualizations, Scoring Rules, and Software Infrastructure

Achim Zeileis

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

Probabilistic regression models

Classical approach: Model conditional expectation $E(y_i|\mathbf{x}_i) = \mu_i$ of a response y_i given explanatory variables \mathbf{x}_i for $i = 1, \dots, n$.

Regression model:

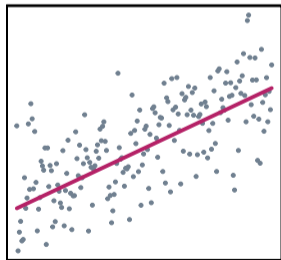
$$\mu_i = r(\mathbf{x}_i)$$

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Regression model: Linear model.

$$\mu_i = r(\mathbf{x}_i) = \beta_0 + \beta_1 \cdot x_{i,1} + \dots + \beta_k \cdot x_{i,k}$$



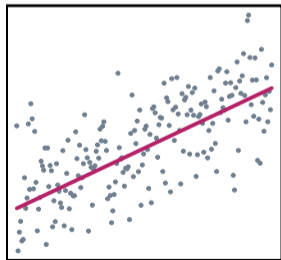
LM, GLM

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Regression model: Generalized linear model with link function $g(\cdot)$.

$$\mu_i = r(\mathbf{x}_i) = g^{-1}(\beta_0 + \beta_1 \cdot x_{i,1} + \dots + \beta_k \cdot x_{i,k})$$



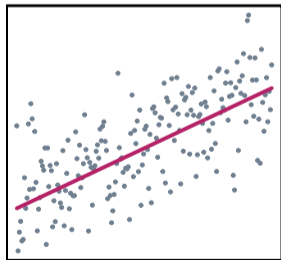
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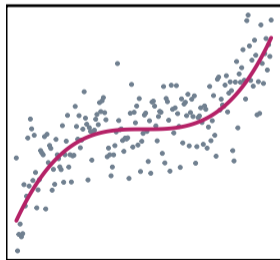
Classical approach: Model conditional expectation $E(y_i|\mathbf{x}_i) = \mu_i$ of a response y_i given explanatory variables \mathbf{x}_i for $i = 1, \dots, n$.

Regression model: Generalized additive model with link function $g(\cdot)$.

$$\mu_i = r(\mathbf{x}_i) = g^{-1}(\beta_0 + s(x_{i,1}) + \dots + s(x_{i,k}))$$



LM, GLM



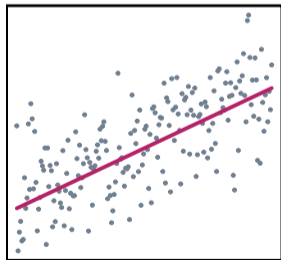
GAM

Probabilistic regression models

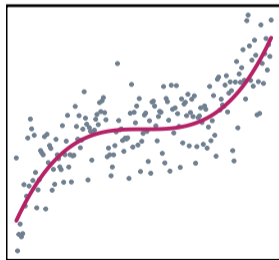
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Regression model: Algorithmic, machine learning, nonparametric, ...

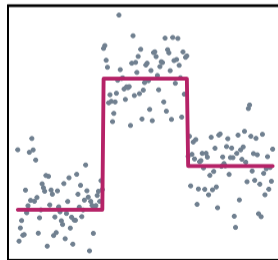
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LM, GLM



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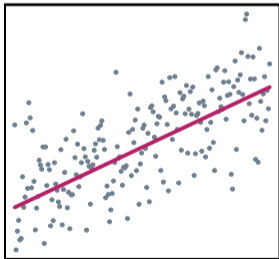
Regression tree

Probabilistic regression models

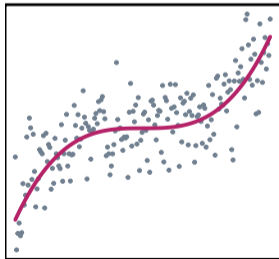
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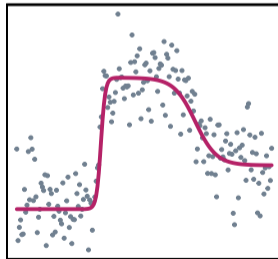
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LM, GLM



GAM



Random forest

Probabilistic regression models

Often: Further assumptions are made beyond the mean specification, especially for estimation and inference.

- Constant variance for least squares.
- Higher moments may co-vary with expectation μ_j , e.g., in exponential family (Poisson, binomial, ...)
- Full distribution for maximum likelihood or Bayesian MCMC, etc.

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But typically: Focus is on conditional means.

- *Forecasting:* $\hat{\mu}_i = \hat{r}(\mathbf{x}_i)$.
- *Scores:* $(y_i - \hat{\mu}_i)^2$ or $|y_i - \hat{\mu}_i|$.
- *Inference:* Robustness/adjustments under misspecification.

Probabilistic regression models

However: Mean forecasts are often of limited interest.

- *Football:* Average goals of team A vs. team B.
- *Precipitation:* Average amount of precipitation today.

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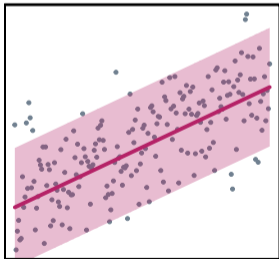
Instead: Full distribution of interest.

- *Football:* Probability for 0, 1, . . . goals, implying win/draw/lose probability.
- *Precipitation:* Probability of no/moderate/extreme precipitation.

Probabilistic regression models

Models:

- Classical models under full assumptions.

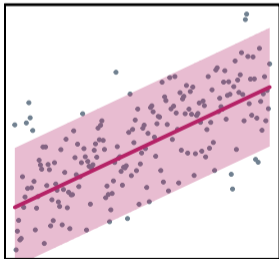


Normal (G)LM w/ constant variance

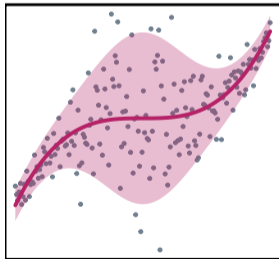
Probabilistic regression models

Models:

- Classical models under full assumptions.
- Generalized additive models for location, scale, and shape.



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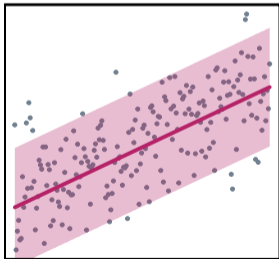


GAMLSS

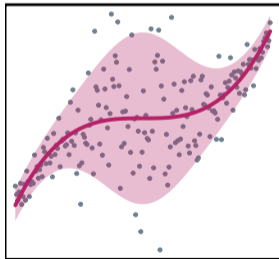
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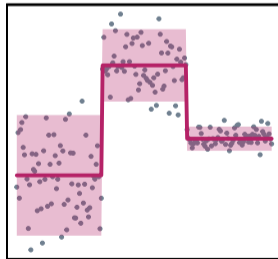
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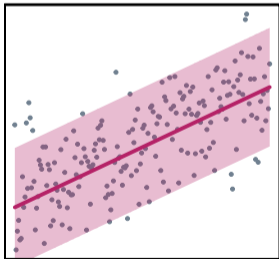


Distributional tree

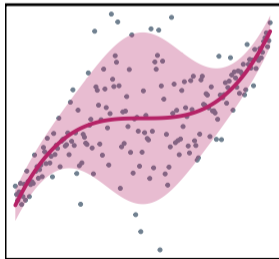
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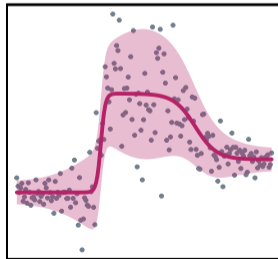
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Normal (G)LM w/ constant variance



GAMLSS



Distributional forest

Probabilistic regression models

Formally: Fit full probability distribution for each observation y_i .

Often: Assume parametric response distribution with parameter vector θ_i .

Cumulative distribution function: $F(y_i|\theta_i)$.

Probability density function: $f(y_i|\theta_i)$.

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Cumulative distribution function: $F(y_i|\theta_i)$.

Probability density function: $f(y_i|\theta_i)$.

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

- 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> head(FIFA2018)
```

	goals	team	match	type	stage	logability	difference
1	5	RUS	1	A	group	0.1531	0.8638
2	0	KSA	1	A	group	-0.7108	-0.8638
3	0	EGY	2	A	group	-0.2066	-0.4438
4	1	URU	2	A	group	0.2372	0.4438
5	3	RUS	3	A	group	0.1531	0.3597
6	1	EGY	3	A	group	-0.2066	-0.3597

Illustration: Goals in the 2018 FIFA World Cup

Model: Poisson GLM with log link.

Regression: Number of goals per team explained by ability difference.

$$\log(\hat{\lambda}_i) = \hat{\beta}_0 + \hat{\beta}_1 \cdot \text{difference}_i$$

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

```
R> lmtest::coeftest(m)
```

z test of coefficients:

	Estimate	Std. Error	z value	Pr(> z)	
(Intercept)	0.2127	0.0813	2.62	0.0088	**
difference	0.4134	0.1058	3.91	9.3e-05	***

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Illustration: Goals in the 2018 FIFA World Cup

Forecasting: In-sample for simplicity.

```
R> head(procast(m))
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                                distribution
1 Poisson distribution (lambda = 1.7680)
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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.

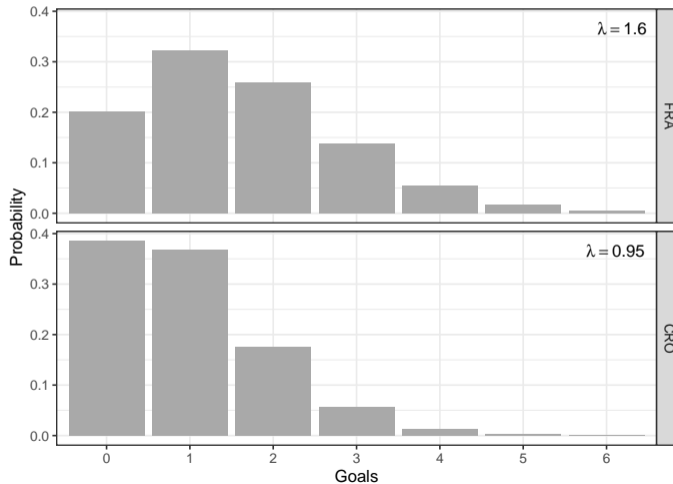


Illustration: Goals in the 2018 FIFA World Cup

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

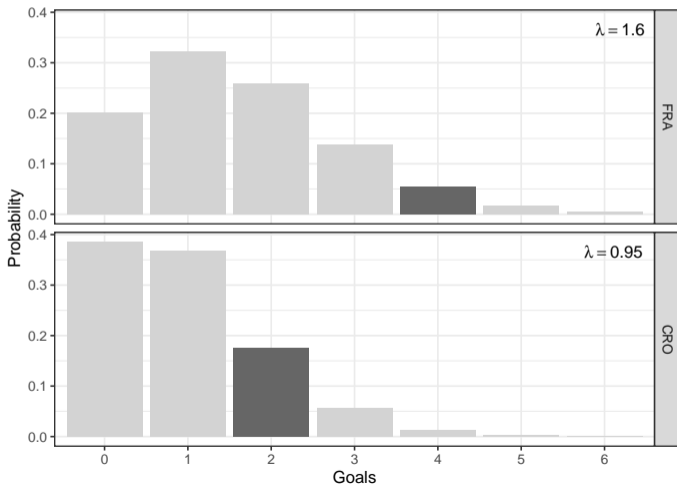


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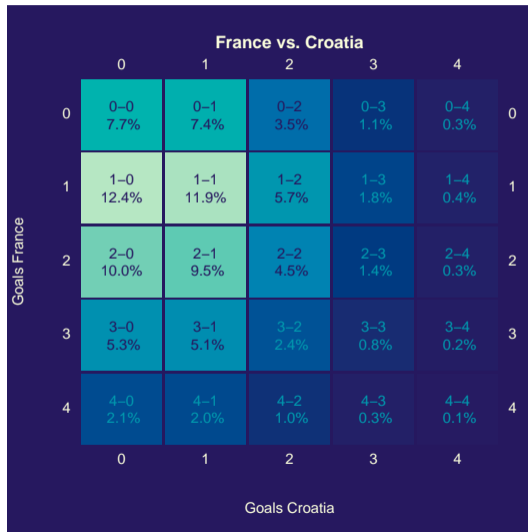


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Possible extensions:

- More observations: Fit on previous World Cups, forecast out-of-sample.
- More covariates: Previous matches, team structure, economic indicators.
- More flexible models: GAM, random forests, boosting, . . .
- More flexible distributions: Bivariate, overdispersion, zero inflation.

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Here: Focus on goodness-of-fit assessment.

In particular: Graphical assessment of model calibration.

Goodness of fit: Scoring rules

Log-score: Log-likelihood; basis for information criteria and classical inference.

$$\log f(y_i | \hat{\theta}_i)$$

Goodness of fit: Scoring rules

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(Continuous) ranked probability score: Bounded alternative to log-score.

$$\int (F(z | \hat{\theta}_i) - \mathbf{1}(z \geq y_i))^2 dz$$

Goodness of fit: Residuals

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

- Uniformly distributed if model correctly specified.
- Uniquely defined for continuous distributions.
- Otherwise consider uniform draw between $F(y_i - 1 | \hat{\theta}_i)$ and $F(y_i | \hat{\theta}_i)$.

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(Randomized) quantile residuals: $\Phi^{-1}(u_i)$.

- Map to normal scale (from uniform).
- More similar to residuals in classical linear regression.
- More emphasis on deviations in the tails of the distribution.

Goodness of fit: Graphical assessment

Ideas:

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

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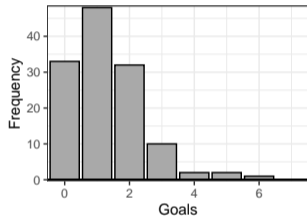
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Questions: Graphics are not new but novel unifying view.

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

Goodness of fit: Graphical assessment

Ideas: Illustrated for FIFA Poisson model.

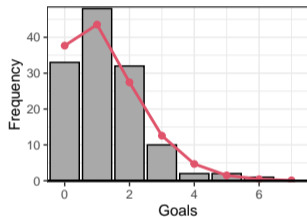


Marginal calibration:

- Observed frequencies.

Goodness of fit: Graphical assessment

Ideas: Illustrated for FIFA Poisson model.

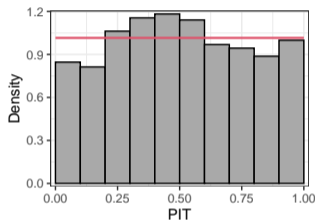
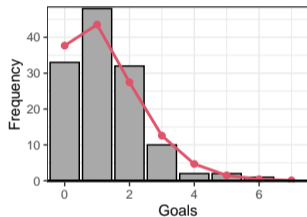


Marginal calibration:

- Observed frequencies.
- Compare: Expected.

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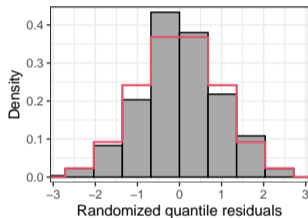
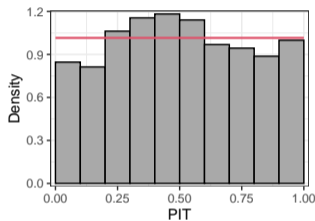
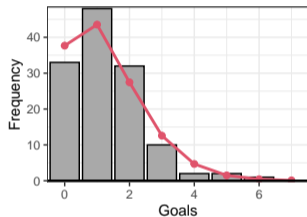
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Probabilistic calibration:

- Probability integral transform.
- Compare: Uniform.

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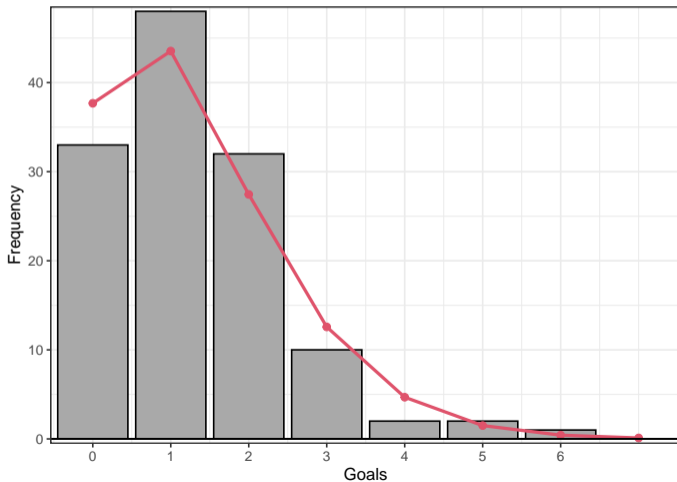
- Probability integral transform.
- Compare: Uniform.

Probabilistic calibration:

- (Randomized) quantile residuals.
- Compare: Normal

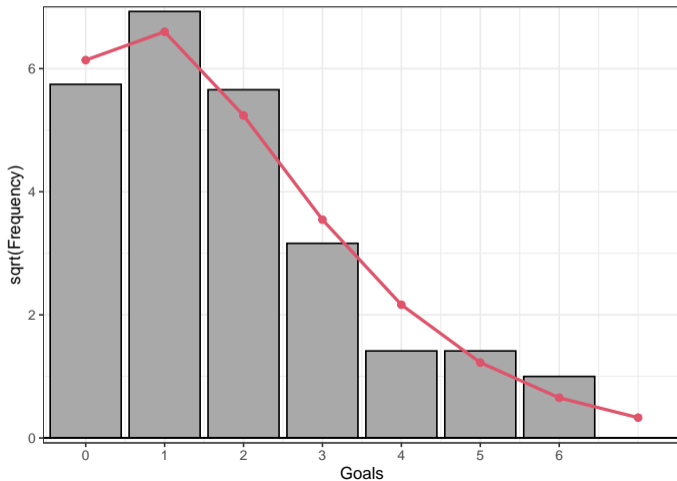
Goodness of fit: Marginal calibration

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



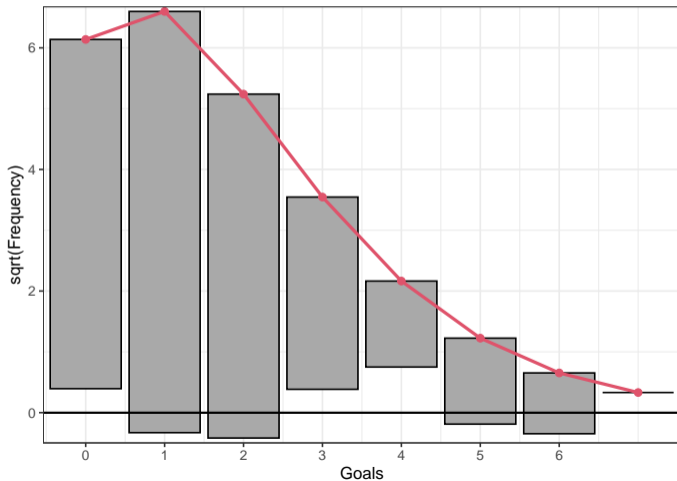
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$\sqrt{\text{Observed}}$ vs. $\sqrt{\text{expected frequencies}}$: Standing, with reference line.



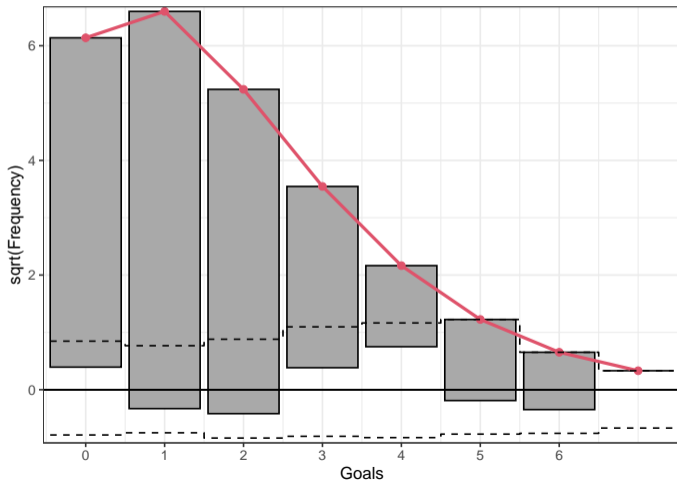
Goodness of fit: Marginal calibration

$\sqrt{\text{Observed}}$ 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.

Goodness of fit: Marginal calibration

Rootogram:

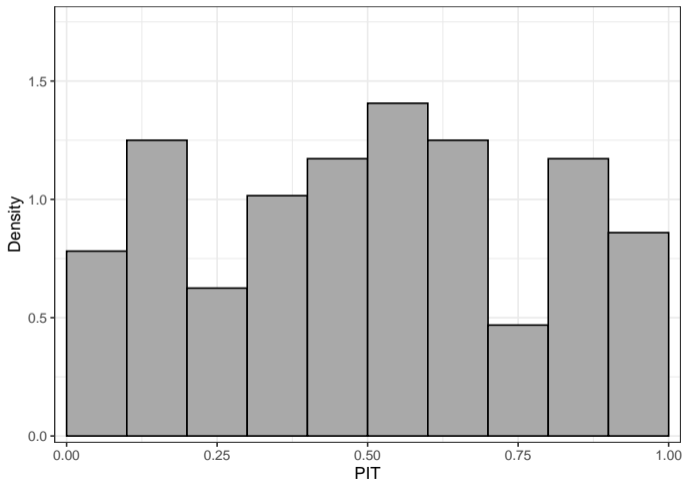
- Frequencies on raw or square-root scale.
- Hanging, standing, or suspended styled rootograms.

Overall:

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

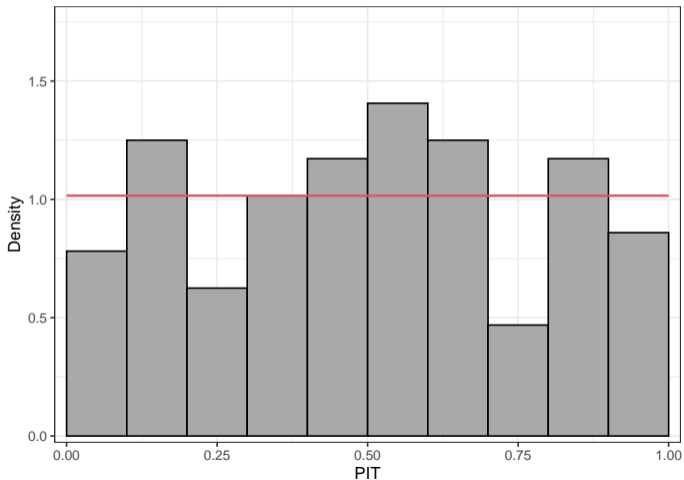
Goodness of fit: Probabilistic calibration

PIT: Randomization 1a.



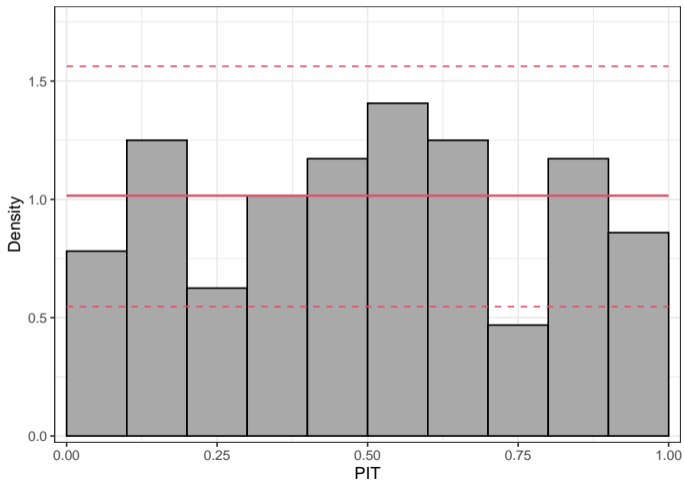
Goodness of fit: Probabilistic calibration

PIT: Randomization 1a, with reference line.



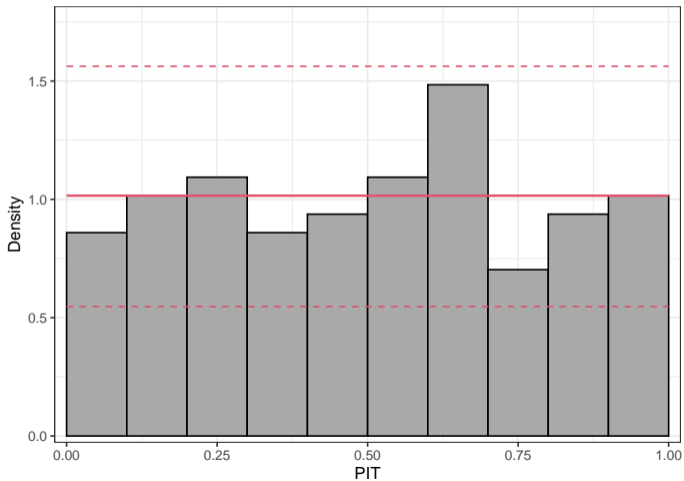
Goodness of fit: Probabilistic calibration

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



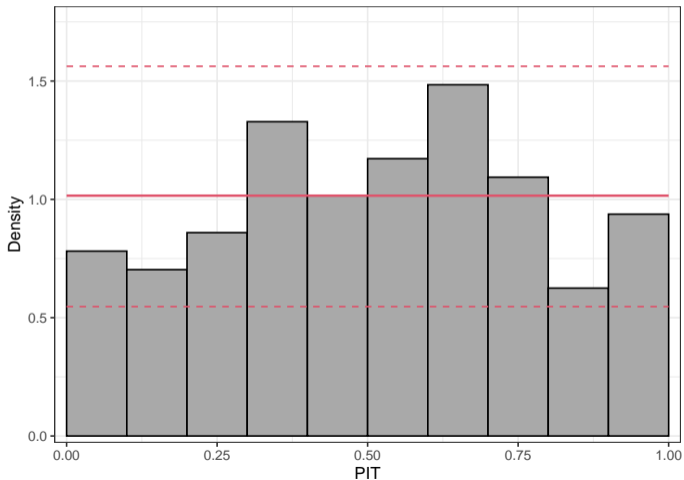
Goodness of fit: Probabilistic calibration

PIT: Randomization 1b.



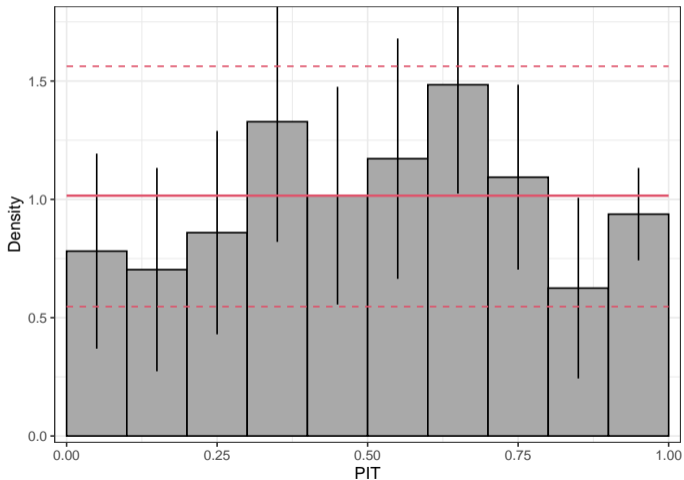
Goodness of fit: Probabilistic calibration

PIT: Randomization 1c.



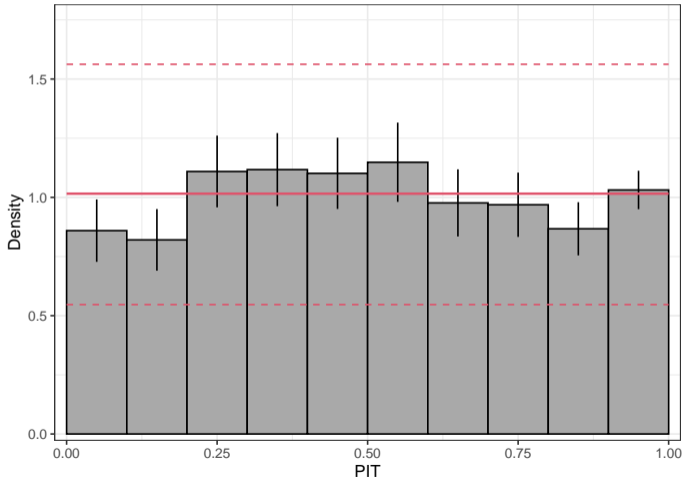
Goodness of fit: Probabilistic calibration

PIT: Randomization 1c, with simulation intervals.



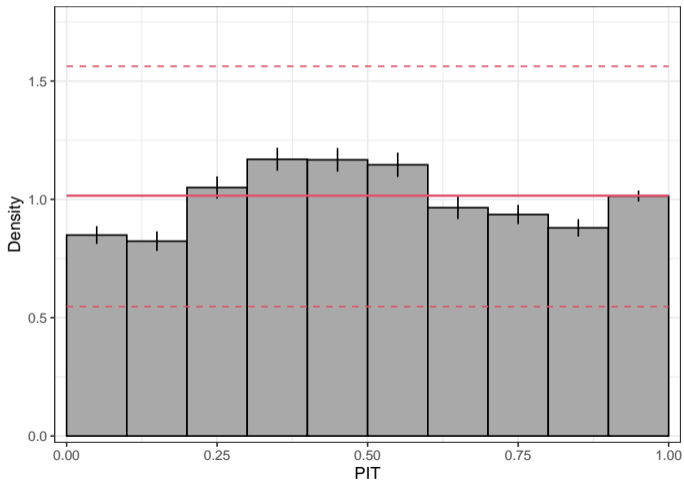
Goodness of fit: Probabilistic calibration

PIT: 10 random draws.



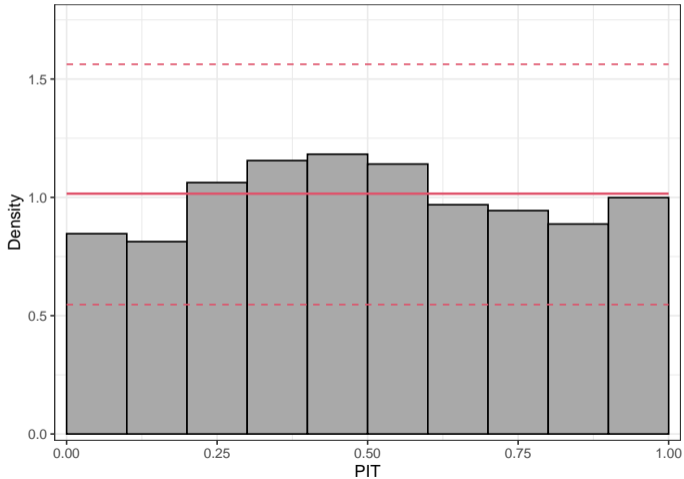
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PIT: 100 random draws.



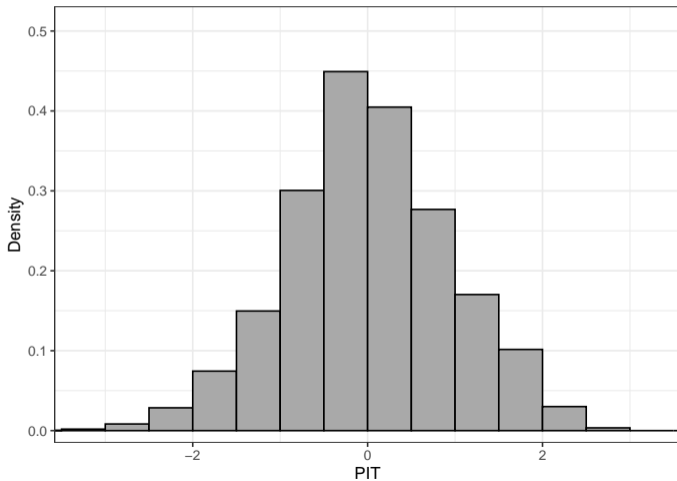
Goodness of fit: Probabilistic calibration

PIT: Expected.



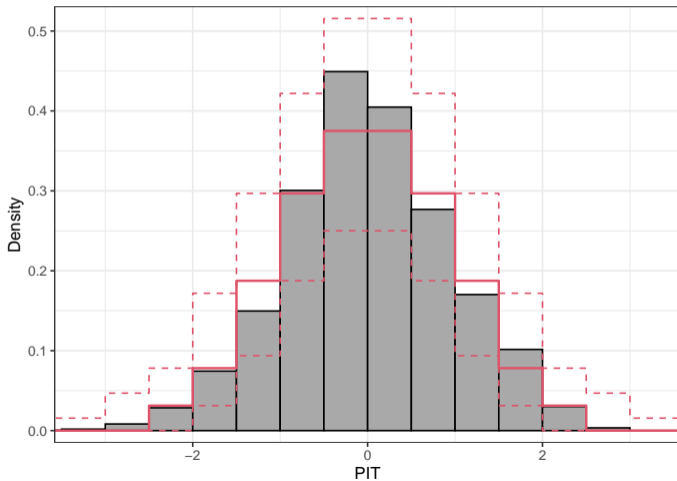
Goodness of fit: Probabilistic calibration

Randomized quantile residuals: Expected.



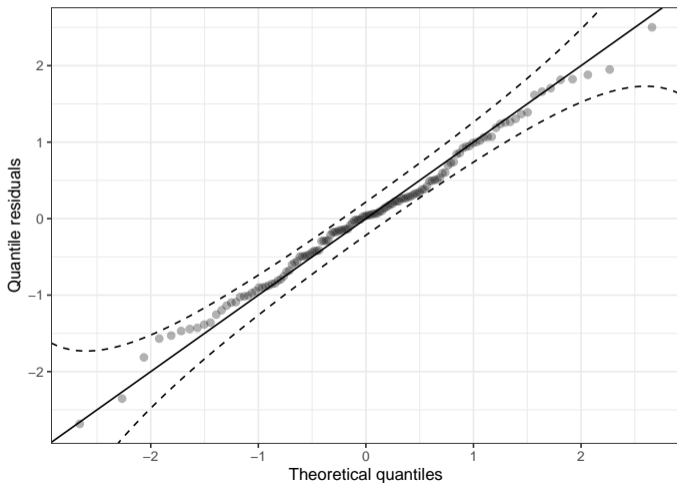
Goodness of fit: Probabilistic calibration

Randomized quantile residuals: Expected, with reference.



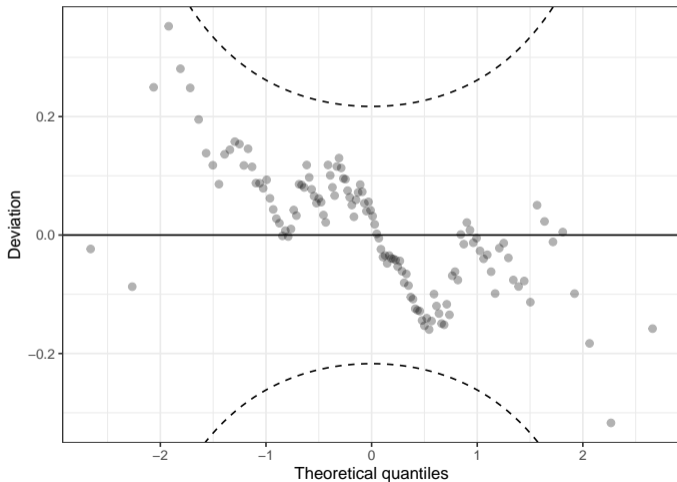
Goodness of fit: Probabilistic calibration

Observed vs. expected quantiles: Q-Q plot.



Goodness of fit: Probabilistic calibration

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



Goodness of fit: Probabilistic calibration

PIT histogram:

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

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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: Precipitation in Innsbruck

Observation data:

- 3 day-accumulated precipitation amounts over 13 years (2000–2013).
- Observation station “Innsbruck” in Austria.

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Model assumptions:

- Homoscedastic linear regression:

$$\hat{\mu}_i = \hat{\beta}_0 + \hat{\beta}_1 \cdot \text{ensmean}_i, \quad \hat{\sigma} = \text{sd}(\epsilon)$$

- Heteroscedastic censored regression with a logistic distribution assumption:

$$y_i \sim \text{Logistic}_0(\hat{\mu}_i = \hat{\beta}_0 + \hat{\beta}_1 \cdot \text{ensmean}_i, \hat{\sigma}_i = \exp(\hat{\gamma}_0 + \hat{\gamma}_1 \cdot \text{enssd}_i))$$

Illustration: Precipitation in Innsbruck

Data: Observations and numerical ensemble mean.

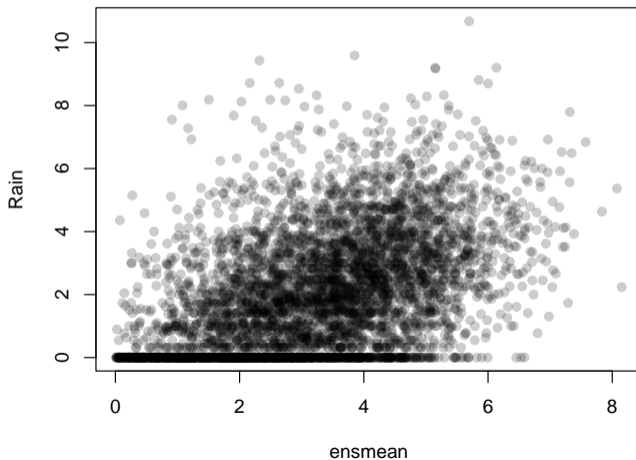


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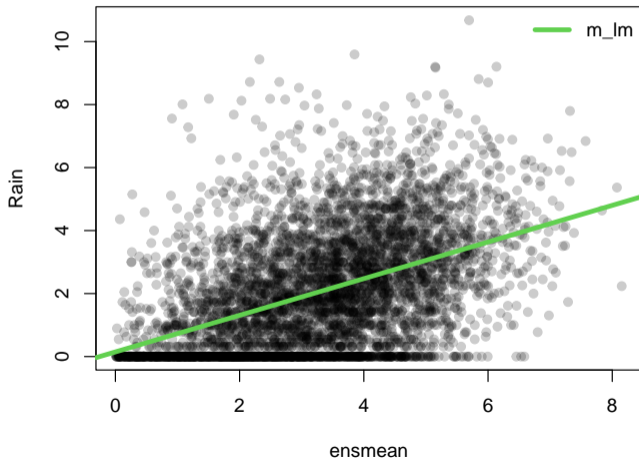


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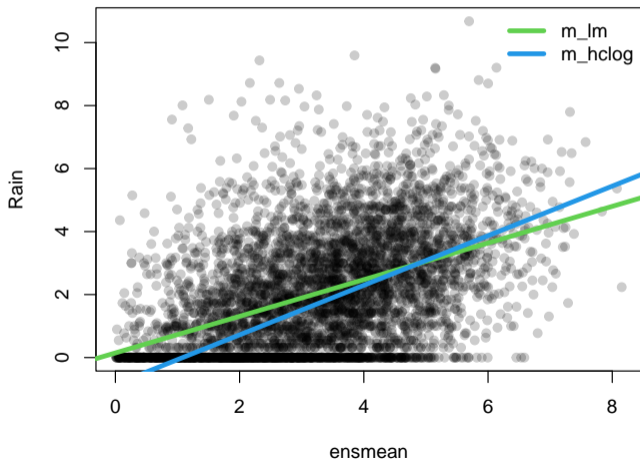


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Rootogram:

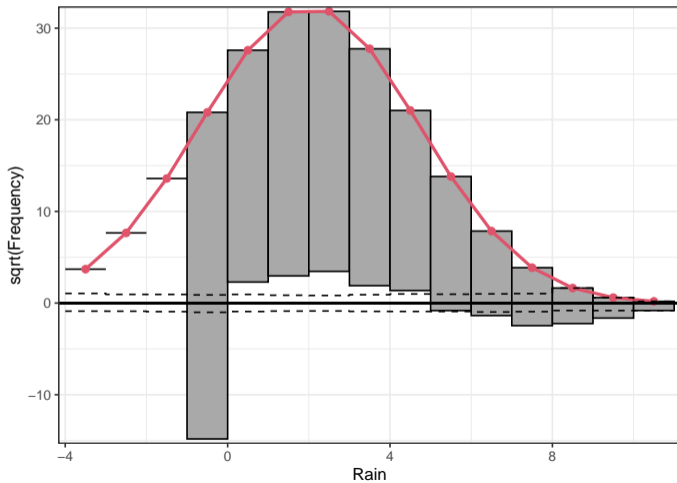


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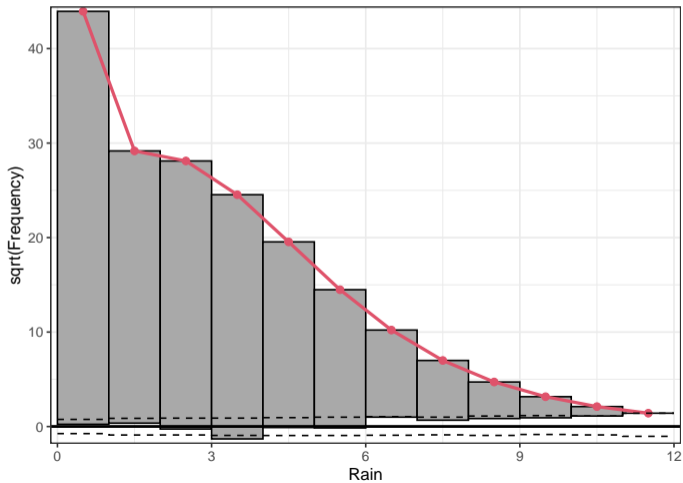


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PIT histogram:

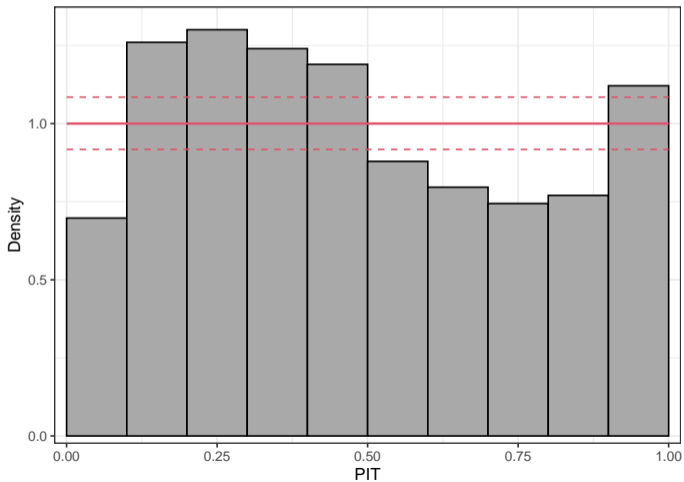


Illustration: Precipitation in Innsbruck

PIT histogram:

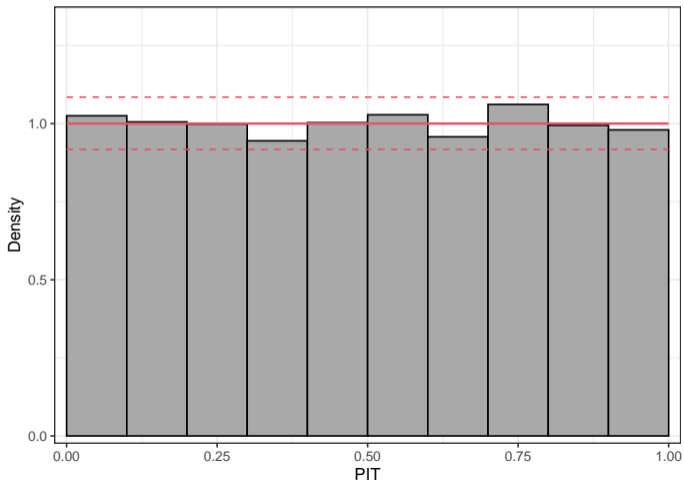


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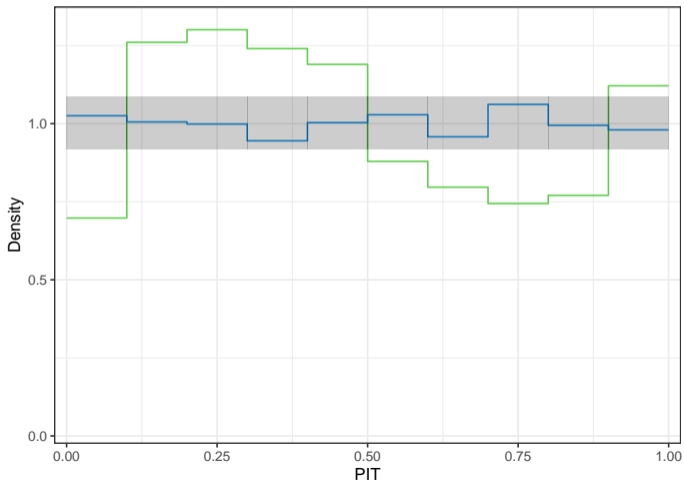


Illustration: Precipitation in Innsbruck

Q-Q residual plot:

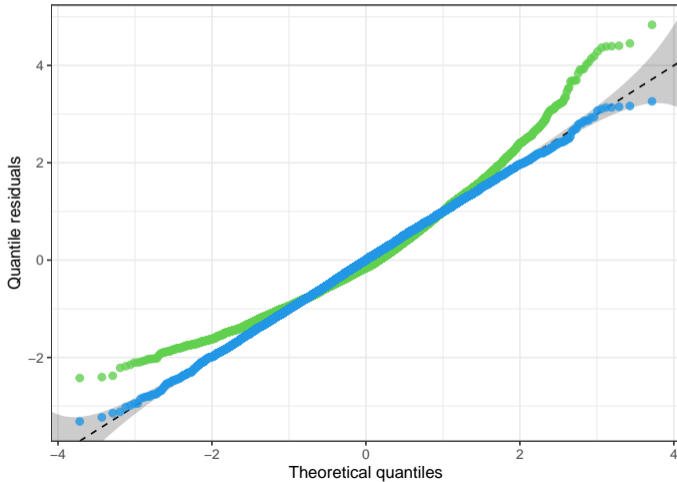
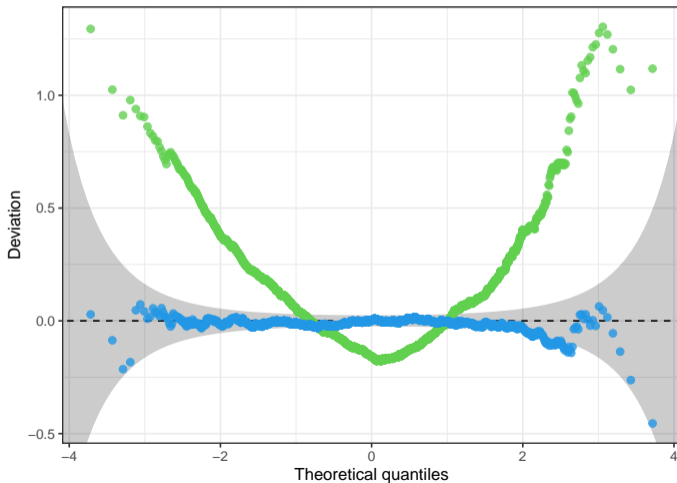


Illustration: Precipitation in Innsbruck

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.

Software: topmodels & distributions3

Probabilistic forecasts:

```
R> p <- procast(m)
```

```
R> head(p, 3)
```

```
                                distribution
1 Poisson distribution (lambda = 1.7680)
2 Poisson distribution (lambda = 0.8655)
3 Poisson distribution (lambda = 1.0297)
```

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For final:

```
R> p_final <- tail(p$distribution, 2)
```

```
R> pdf(p_final, 0:4)
```

```
      d_0    d_1    d_2    d_3    d_4
127 0.2010 0.3225 0.2587 0.13836 0.05550
128 0.3853 0.3675 0.1752 0.05572 0.01329
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Software: topmodels & distributions3

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```

Scoring rules:

```
R> proscore(m, type = c("LogS", "CRPS", "MSE"), aggregate = TRUE)
```

```
      LogS  CRPS  MSE
1 -1.388 0.562 1.162
```

References

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