



Beta Regression in R

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Overview

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Motivation

Goal: Model dependent variable $y \in (0, 1)$, e.g., rates, proportions, concentrations etc.

Common approach: Model transformed variable \tilde{y} by a linear model, e.g., $\tilde{y} = \text{logit}(y)$ or $\tilde{y} = \text{probit}(y)$ etc.

Disadvantages:

- Model for mean of \tilde{y} , not mean of y (Jensen's inequality).
- Data typically heteroskedastic.

Idea: Model *y* directly using suitable parametric family of distributions plus link function.

Specifically: Maximum likelihood regression model using alternative parametrization of beta distribution (Ferrari & Cribari-Neto 2004).

Beta regression

Beta distribution: Continuous distribution for 0 < y < 1, typically specified by two shape parameters p, q > 0.

Alternatively: Use mean $\mu = p/(p+q)$ and precision $\phi = p+q$.

Probability density function:

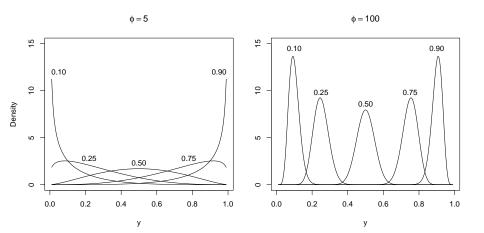
$$f(y) = \frac{\Gamma(p+q)}{\Gamma(p) \Gamma(q)} y^{p-1} (1-y)^{q-1}$$
$$= \frac{\Gamma(\phi)}{\Gamma(\mu\phi) \Gamma((1-\mu)\phi)} y^{\mu\phi-1} (1-y)^{(1-\mu)\phi-1}$$

where $\Gamma(\cdot)$ is the gamma function.

Properties: Flexible shape. Mean $E(y) = \mu$ and

$$Var(y) = \frac{\mu (1 - \mu)}{1 + \phi}.$$

Beta regression



Beta regression

Regression model:

- Observations i = 1, ..., n of dependent variable y_i .
- Link parameters μ_i and ϕ_i to sets of regressor x_i and z_i .
- Use link functions g_1 (logit, probit, ...) and g_2 (log, identity, ...).

$$g_1(\mu_i) = \mathbf{x}_i^{\top} \boldsymbol{\beta},$$

 $g_2(\phi_i) = \mathbf{z}_i^{\top} \boldsymbol{\gamma}.$

Inference:

- ullet Coefficients eta and γ are estimated by maximum likelihood.
- The usual central limit theorem holds with associated asymptotic tests (likelihood ratio, Wald, score/LM).

Implementation in R

Model fitting:

- Package betareg with main model fitting function betareg().
- Interface and fitted models are designed to be similar to glm().
- Model specification via formula plus data.
- Two part formula, e.g., y ~ x1 + x2 + x3 | z1 + z2.
- Log-likelihood is maximized numerically via optim().
- Extractors: coef(), vcov(), residuals(), logLik(),...

Inference:

- Base methods: summary(), AIC(), confint().
- Methods from Imtest and car: lrtest(), waldtest(), coeftest(), linearHypothesis().
- Moreover: Multiple testing via multcomp and structural change tests via strucchange.

Data: From Smithson & Verkuilen (2006).

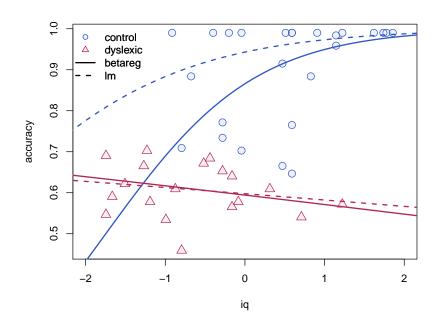
- 44 Australian primary school children.
- Dependent variable: Score of test for reading accuracy.
- Regressors: Indicator dyslexia (yes/no), nonverbal iq score.

Analysis:

- OLS for transformed data leads to non-significant effects.
- OLS residuals are heteroskedastic.
- Beta regression captures heteroskedasticity and shows significant effects.

```
R> data("ReadingSkills", package = "betareg")
R> rs_ols <- lm(qlogis(accuracy) ~ dyslexia * iq,
+ data = ReadingSkills)
R> coeftest(rs_ols)
t test of coefficients:
          Estimate Std. Error t value Pr(>|t|)
(Intercept) 1.60107 0.22586 7.0888 1.411e-08 ***
dyslexia -1.20563 0.22586 -5.3380 4.011e-06 ***
iq 0.35945 0.22548 1.5941 0.11878
Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
R> bptest(rs_ols)
studentized Breusch-Pagan test
data: rs_ols
BP = 21.692, df = 3, p-value = 7.56e-05
```

```
R> rs_beta <- betareg(accuracy ~ dyslexia * iq | dyslexia + iq,
   data = ReadingSkills)
R> coeftest(rs_beta)
z test of coefficients:
             Estimate Std. Error z value Pr(>|z|)
(Intercept)
             1.12323
                      0.14283 7.8638 3.725e-15 ***
             dyslexia
            iq
dyslexia:iq
          (phi)_(Intercept) 3.30443
                      0.22274 14.8353 < 2.2e-16 ***
(phi)_dyslexia 1.74656
                      0.26232 6.6582 2.772e-11 ***
(phi)_iq 1.22907
                      0.26720 4.5998 4.228e-06 ***
Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
```



Extensions: Partitions and mixtures

So far: Reuse standard inference methods for fitted model objects.

Now: Reuse fitting functions in more complex models.

Model-based recursive partitioning: Package party.

- Idea: Recursively split sample with respect to available variables.
- Aim: Maximize partitioned likelihood.
- Fit: One model per node of the resulting tree.

Latent class regression, mixture models: Package flexmix.

- Idea: Capture unobserved heterogeneity by finite mixtures of regressions.
- Aim: Maximize weighted likelihood with k components.
- Fit: Weighted combination of *k* models.

Beta regression trees

Partitioning variables: dyslexia and further random noise variables.

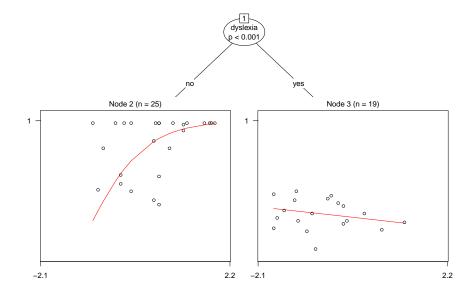
```
R> set.seed(1071)
R> ReadingSkills$x1 <- rnorm(nrow(ReadingSkills))
R> ReadingSkills$x2 <- runif(nrow(ReadingSkills))
R> ReadingSkills$x3 <- factor(rnorm(nrow(ReadingSkills)) > 0)
```

Fit beta regression tree: In each node accuracy's mean and precision depends on iq, partitioning is done by dyslexia and the noise variables x1, x2, x3.

```
R> rs_tree <- betatree(accuracy ~ iq | iq,
+ ~ dyslexia + x1 + x2 + x3,
+ data = ReadingSkills, minsplit = 10)
R> plot(rs_tree)
```

Result: Only relevant regressor dyslexia is chosen for splitting.

Beta regression trees



Setup:

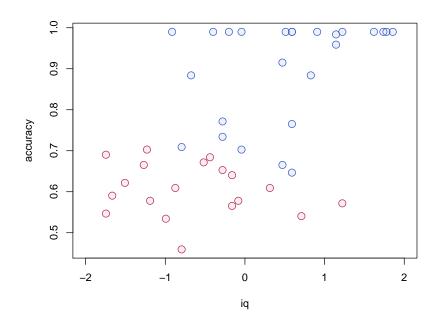
- No dyslexia information available.
- Look for k = 3 clusters: Two different relationships of type
 accuracy ~ iq, plus component for ideal score of 0.99.

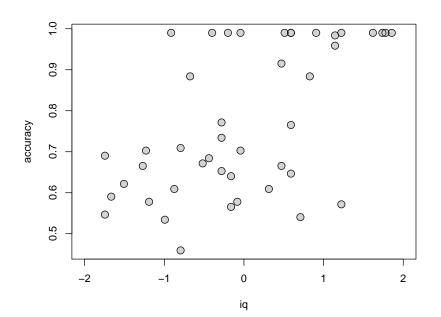
Fit beta mixture regression:

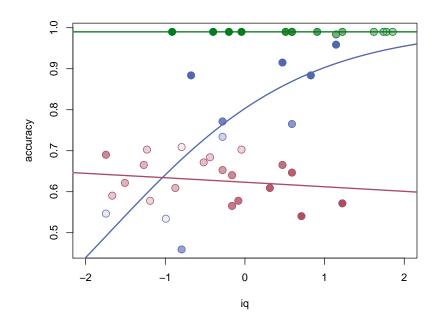
```
R> rs_mix <- betamix(accuracy ~ iq, data = ReadingSkills, k = 3,
+    nstart = 10, extra_components = extraComponent(
+    type = "uniform", coef = 0.99, delta = 0.01))</pre>
```

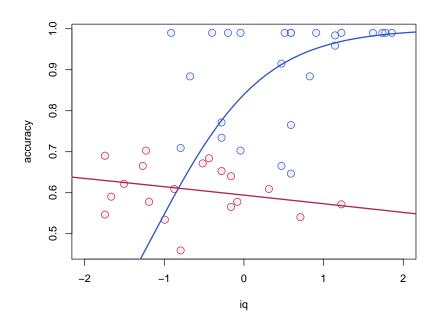
Result:

- Dyslexic children separated fairly well.
- Other children are captured by mixture of two components: ideal reading scores, and strong dependence on iq score.









Summary

Beta regression and extensions:

- Flexible regression model for proportions, rates, concentrations.
- Can capture skewness and heteroskedasticity.
- R implementation betareg, similar to glm().
- Due to design, standard inference methods can be reused easily.
- Fitting functions can be plugged into more complex fitters.
- Convenience interfaces available for: Model-based partitioning, finite mixture models.

References

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