

Paired Comparison Preference Models

The prefmod Package: Part I Some Examples

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Paired Comparison Models

General Considerations:

the paired comparison models we consider in this talk are based on the Bradley-Terry Model (Biometrika, 1952)

- and will be used as statistical models describing and explaining data
- the models are similar to regression models (ANOVA) other types of dependent variables (multivariate, nominal) [Generalised Linear Models](#)
- but will **not** be considered as measurement models in this case one should use another approach e.g. taken by Winkelmaier in the **R** package `eba`

Paired Comparisons

- method of data collection
- given a set of J items



- individuals are asked to judge pairs of objects

j preferred to k   k preferred to j

- aim is to rank objects into a preference order
- obtain an overall ranking of the objects

Overview

- [LLBT models](#): loglinear Bradley-Terry models
- Basic LLBT
- [Extended LLBT](#)
undecided response
subject covariates
object specific covariates
- [Pattern Models](#)
Paired comparison \rightarrow pattern models
Ranking \rightarrow pattern models
Rating \rightarrow pattern models



The Basic Bradley-Terry Model (BT)

for the each comparison (jk) of object j to object k we observe:

- $n_{(j>k)}$... the number of times j is preferred to k
- $n_{(k>j)}$... the number of times k is preferred to j

$$N_{(jk)} = n_{(j>k)} + n_{(k>j)} \quad \text{total number of responses to comparison } (jk)$$

the probability that j is preferred to k in comparison (jk)

$$P(j \succ k) = \frac{\pi_j}{\pi_j + \pi_k} \quad \begin{array}{l} \pi\text{'s are called } \textit{worth parameters} \\ \text{and are non-negative numbers} \\ \text{describing the location of the objects} \end{array}$$



The Basic Loglinear BT Model (LLBT)

the model can be formulated as a log-linear model following the usual Multinomial / Poisson - equivalence.

the expected value $m_{(j>k)}$ of $n_{(j>k)}$ is $m_{(j>k)} = N_{(jk)}p_{(j>k)}$

$$P(j \succ k) = \frac{\pi_j}{\pi_j + \pi_k} = c_{(jk)} \frac{\sqrt{\pi_j}}{\sqrt{\pi_k}} \quad \text{where } c_{(jk)} \text{ is constant for a given comparison}$$

then our basic paired comparison model for one comparison is

$$\ln m_{(j>k)} = \mu_{(jk)} + \lambda_j - \lambda_k \quad \begin{array}{l} \lambda\text{'s are the object parameters} \\ \mu\text{'s are nuisance parameters} \end{array}$$

this model formulation is feasible for further extensions



terms and relations

- relation between π and λ :

$$\lambda_j = \ln \sqrt{\pi_j}$$

$$\pi_j = \exp 2\lambda_j$$

- identifiability of π s is obtained by the restriction $\pi_J = 1$ via $\lambda_J = 0$
- the worth parameters are calculated by

$$\pi_j = \frac{\exp(2\lambda_j)}{\sum_j \exp(2\lambda_j)}, \quad j = 1, 2, \dots, J$$

where $\sum_j \pi_j = 1$



LLBT Design Structure

- for 3 objects we have 3 comparisons

PC	decision	counts	μ	λ_1	λ_2	λ_3
(12)	O_1	$n_{(1>2)}$	1	1	-1	0
(12)	O_2	$n_{(2>1)}$	1	-1	1	0
(13)	O_1	$n_{(1>3)}$	2	1	0	-1
(13)	O_3	$n_{(3>1)}$	2	-1	0	1
(23)	O_2	$n_{(2>3)}$	3	0	1	-1
(23)	O_3	$n_{(3>2)}$	3	0	-1	1

the **design structure** consists of counts (dependent variable) and the **design matrix X** with:

μ which is a factor (dummies for μ_1, μ_2, μ_3) and variates for the objects O_1, O_2, O_3

LLBT Model Structure

model formula for the first line is:

$$\ln m_{(1>2)} = \mu_1 + \lambda_1 - \lambda_2$$

general matrix notation:

$$\begin{pmatrix} \ln m_{(1>2)} \\ \ln m_{(2>1)} \\ \ln m_{(1>3)} \\ \ln m_{(3>1)} \\ \ln m_{(2>3)} \\ \ln m_{(3>2)} \end{pmatrix} = \begin{pmatrix} 1 & 0 & 0 & 1 & -1 & 0 \\ 1 & 0 & 0 & -1 & 1 & 0 \\ 0 & 1 & 0 & 1 & 0 & -1 \\ 0 & 1 & 0 & -1 & 0 & 1 \\ 0 & 0 & 1 & 0 & 1 & -1 \\ 0 & 0 & 1 & 0 & -1 & 1 \end{pmatrix} \begin{pmatrix} \mu_1 \\ \mu_2 \\ \mu_3 \\ \lambda_1 \\ \lambda_2 \\ \lambda_3 \end{pmatrix}$$

WINE example (fictitious)

4 sicilian red wines have been tasted pairwise by 1000 people (all of them prefer red to white wines)

aim of the study:

- preference order for different wines

data:

- PC-responses about the choices of 4 selected wines
- the wines are:
 - O1 = Nero D'**A**vola (also called 'Calabrese')
 - O2 = **G**aglioppo (a red of Calabrian origin frequently grown in Sicily)
 - O3 = **P**erricone (Pignatello - esoteric, robust red)
 - O4 = **N**erello (Mascalese - strong red)

Data preparation

- 4 objects (items) A,G,P,N
- number of comparisons is $4 \cdot 3/2 = 6$
- for data input the ordering of comparisons is important

wines	wine first in comparison			
	A ₁	G ₂	P ₃	N ₄
A ₁	-			
G ₂	V1	-		
P ₃	V2	V3	-	
N ₄	V4	V5	V6	-

V1	V2	V3	V4	V5	V6
(12)	(13)	(23)	(14)	(24)	(34)
(A,G)	(A,P)	(G,P)	(A,N)	(G,N)	(P,N)

Coding

One possible coding for each comparison (V1, ... V6) is

$$(jk) = \begin{cases} 1 & \text{if first object is preferred to second object } (j > k) \\ -1 & \text{if second object is preferred to first object } (k > j) \end{cases}$$

First respondent gave the following responses:

V1	V2	V3	V4	V5	V6
(12)	(13)	(23)	(14)	(24)	(34)
(A,G)	(A,P)	(G,P)	(A,N)	(G,N)	(P,N)
1	1	1	1	-1	-1

NOTE:

- missing responses should be coded with NA
- if the coding is **not** (1, -1) but any other two numbers, the smaller number means the first object is preferred e.g. a coding with (0, 1) ⇒ 0 means first object preferred



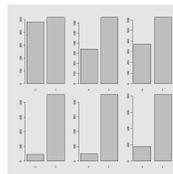
Wine data

- preparations

```
> library(prefmod)
> load("wine.Rdata")
> wnames <- c("A", "G", "P", "N")
```

produce barplots for all comparisons

```
> par(mfrow = c(2, 3))
> for (i in 1:6) barplot(table(wine[, i]))
> par(mfrow = c(1, 1))
```

Function: `llbtPC.fit()`

- preparations

```
> library(prefmod)
> load("wine.Rdata")
> wnames <- c("A", "G", "P", "N")
```

- fit simple model

```
> m1 <- llbtPC.fit(wine, nitems = 4, obj.names = wnames)
> m1
```



```
> m1
Call:
```

```
gnm(formula = formula, eliminate = elim, family = poisson, data = dfr)
```

Coefficients of interest:

	A	G	P	N
	1.118	1.073	0.794	NA

```
Deviance:          2.1059
Pearson chi-squared: 2.1101
Residual df:       3
```



- model fit was done by the package `gnm()`
- We see (parameter) estimates for $\lambda_1, \lambda_2, \lambda_3$ and $\lambda_4 = NA$ (is set to zero because of restriction for estimation)
- We do not see parameter estimates for $\mu_1, \mu_2, \mu_3, \mu_4$ these are nuisance parameters and they are eliminated by the procedure (fitted, but not displayed)

- Does the model fit?
We can use the Deviance 2.106
degrees of freedom 3

```
> dev1 <- round(m1$deviance, digits = 5)
> df1 <- m1$df.residual
> prob1 <- 1 - pchisq(dev1, df1)
> print(prob1)
[1] 0.55073
```

Model fit is OK (probability is > 0.05)

- Deviance: $\sum o_{ij} \ln\left(\frac{o_{ij}}{e_{ij}}\right) = \sum n_{ij} \ln\left(\frac{n_{ij}}{m_{ij}}\right)$

under full model $n_{ij} = m_{ij}$

- χ^2 - statistic: $\sum \frac{(o_{ij} - e_{ij})^2}{e_{ij}} = \sum \frac{(n_{ij} - m_{ij})^2}{m_{ij}}$

Functions: `llbt.worth()`, `plotworth()`

- calculate worth parameters

```
> worth1 <- llbt.worth(m1)
```

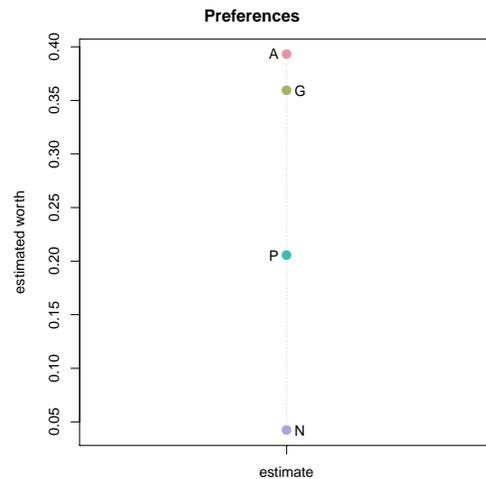
Remember: $\pi_j = \frac{\exp(2\lambda_j)}{\sum_j \exp(2\lambda_j)}$ $\lambda_A = 1.118$, $\lambda_G = 1.073$, $\lambda_P = 0.794$, $\lambda_N = 0$

- or get estimates

```
> est1 <- llbt.worth(m1, outmat = "lambda")
```

- plot worth or estimates

```
> plotworth(worth1, ylab = "estimated worth")
> plotworth(est1)
```



- How to get p.e. from `m1` (which is a `gnm` object)
all parameter estimates $(\mu_1, \mu_2, \dots, \lambda_1, \lambda_2, \dots)$

```
> m1$coefficients
      A      G      P      N
1.1183 1.0731 0.7935      NA
attr(,"eliminated")
[1] 6.2136 6.1628 6.1760 5.6880 5.7241 5.9282
```

- How to get parameter estimates of interest only? $(\lambda_1, \lambda_2, \dots)$

```
> c1 <- coef(m1)
> c1
Coefficients of interest:
      A      G      P      N
1.1183 1.0731 0.7935      NA
```



- Replace last parameter (NA) with 0 and give names

```
> c1 <- c(c1[1:3], 0)
> names(c1) <- c("A", "G", "P", "N")
```

- How to get the μ 's (μ_1, μ_2, \dots)

```
> mu <- attr(m1$coefficients, "eliminated")
> mu[1:6]
[1] 6.2136 6.1628 6.1760 5.6880 5.7241 5.9282
> mu[4]
[1] 5.688
```



- How to get s.e. from m1

```
> cov <- vcov(m1)
> se <- sqrt(diag(cov))
```

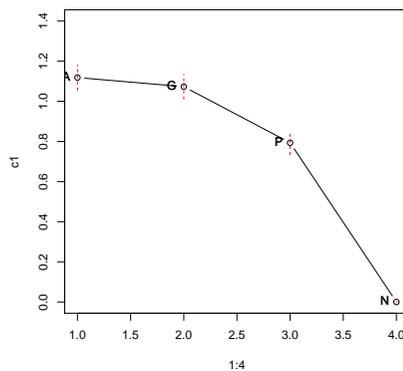
- How to make a confidence interval plot

```
> up1 <- c1 + se * 1.96
> lo1 <- c1 - se * 1.96

> cbind(lo1, c1, up1)
      lo1   c1   up1
A 1.05548 1.1183 1.18114
G 1.01080 1.0731 1.13542
P 0.73392 0.7935 0.85308
N 0.00000 0.0000 0.00000
```



```
> plot(1:4, c1, type = "b", ylim = c(0, 1.4))
> for (i in 1:4) {
+   lines(rep(i, 2), c(lo1[i], up1[i]), col = "red", lty = "dashed")
+   text(rep(i, 2), c1[i], names(c1)[i], pos = 2)
+ }
```



Parameter Interpretation

- log odds for preferring **A** to **N**

$$\ln \left(\frac{P_{(A>N)}}{P_{(N>A)}} \right) = \ln \left(\frac{\frac{m_{(A>N)}}{N_{(A,N)}}}{\frac{m_{(N>B)}}{N_{(A,N)}}} \right) = \ln \left(\frac{m_{(A>N)}}{m_{(N>B)}} \right) = \ln m_{(A>N)} - \ln m_{(N>A)}$$

$$\begin{aligned} \text{insert equations } \ln m_{(A>N)} &= \mu_{(A,N)} + \lambda_A - \lambda_N \\ - \ln m_{(N>A)} &= -\mu_{(A,N)} + \lambda_A - \lambda_N \end{aligned}$$

- log odds preferring **A** to **N** are $2(\lambda_A - \lambda_N)$ is 2.237

parameter estimates are stored in `c2[1]` and `c2[4]`

λ_A is 1.118 and λ_N is 0 (reference object)

- the odds for preferring **A** to **N** are $\exp(2.237) = 9.362$



- How would you calculate the expected preferences for $m_{(A>G)}$ and for $m_{(G>A)}$?

- the λ parameter estimates are:

```
> coef(m1)
Coefficients of interest:
      A      G      P      N
1.1183 1.0731 0.7935   NA
```

- the estimates for the μ s are:

```
> attr(m1$coefficients, "eliminated")
[1] 6.2136 6.1628 6.1760 5.6880 5.7241 5.9282
```

- Hint 1: Use model formula to calculate the logarithm of the expected numbers with

$$\ln m_{(A>G)} = \mu_{(A,G)} + \lambda_A - \lambda_G$$

- Hint 2: the expected numbers are $\exp(\ln m_{(A>G)})$



- The observed preferences (counts) $n_{(i>j)}$ are stored in

```
> m1$y
1.V11 1.V12 1.V21 1.V22 1.V31 1.V32 1.V41 1.V42 1.V51 1.V52 1.V61 1.V62
 519   481   656   344   626   374   908   92   902   98   819   181
```

- The expected preferences $m_{(i>j)}$ are stored in

```
> m1$fitted.values
 1.V11  1.V12  1.V21  1.V22  1.V31  1.V32  1.V41  1.V42  1.V51  1.V52
522.59 477.41 656.92 343.08 636.27 363.73 903.49 96.51 895.32 104.68
 1.V61  1.V62
830.19 169.81
```

- The raw residuals are the difference between observed and expected preferences

```
> m1$y - m1$fitted.values
 1.V11  1.V12  1.V21  1.V22  1.V31  1.V32  1.V41
-3.58549  3.58549 -0.92408  0.92408 -10.27047  10.27047  4.50957
 1.V42  1.V51  1.V52  1.V61  1.V62
-4.50957  6.68499 -6.68499 -11.19455  11.19455
```



Example: CEMS exchange programme

students of the WU can study abroad visiting one of currently 17 CEMS universities

aim of the study:

- preference orderings of students for different locations
- identify reasons for these preferences

data:

- PC-responses about their choices of 6 selected CEMS universities for the semester abroad (London, Paris, Milan, Barcelona, St.Gall, Stockholm)
- answer: *can not decide* was allowed
- several covariates (e.g., gender, working status, language abilities, etc.)



LLBT Extensions Overview

- ▷ [undecided responses \(ties\)](#)
- ▷ [subject covariates](#)
- ▷ [object specific covariates](#)



Undecided Response

- undecided 3 possible responses

for each comparison between two objects j and k the response can be:

$$\begin{aligned} j > k \\ k > j \\ j = k \end{aligned}$$

In our CEMS-example: comparing London (LO) to Paris (PA) the response can be:

$$\begin{aligned} LO > PA \\ PA > LO \\ LO = PA \end{aligned}$$



LLBT with Undecided Response

Using the respecification of the probabilities suggested by Davidson and Beaver (1977):

the LLBT model formulas for the comparison (jk) are now:

$$\begin{aligned} \ln m_{(j>k)} &= \mu_{(jk)} + \lambda_j - \lambda_k \\ \ln m_{(k>j)} &= \mu_{(jk)} - \lambda_j + \lambda_k \\ \ln m_{(j=k)} &= \mu_{(jk)} + \gamma \end{aligned}$$

where γ is the parameter for undecided response (could also be $\gamma_{(jk)}$)

λ 's are the object parameters
 μ 's are nuisance parameters



Interpretation of parameter γ

- log odds for preferring ($j > k$) to "no decision" ($j = k$)

$$\ln \left(\frac{P_{(j>k)}}{P_{(j=k)}} \right) = \ln m_{(j>k)} - \ln m_{(j=k)}$$

insert equations

$$\begin{aligned} \ln m_{(j>k)} &= \mu_{(jk)} + \lambda_j - \lambda_k \\ - \ln m_{(j=k)} &= -\mu_{(jk)} - \gamma \end{aligned}$$

for $\lambda_j = \lambda_k$

- log odds preferring ($j > k$) to "no decision" ($j = k$) is $-\gamma$

if $\gamma < 0$: there is an advantage in favour of a decision



LLBT Design Structure with undecided

PC	decision	counts	μ	γ	λ_1	λ_2	λ_3
(12)	O_1	$n_{(1>2)}$	1	0	1	-1	0
(12)	no	$n_{(1=2)}$	1	1	0	0	0
(12)	O_2	$n_{(2>1)}$	1	0	-1	1	0
(13)	O_1	$n_{(1>3)}$	2	0	1	0	-1
(13)	no	$n_{(1=3)}$	2	1	0	0	0
(13)	O_3	$n_{(3>1)}$	2	0	-1	0	1
(23)	O_2	$n_{(2>3)}$	3	0	0	1	-1
(23)	no	$n_{(2=3)}$	3	1	0	0	0
(23)	O_3	$n_{(3>2)}$	3	0	0	-1	1

- Design matrix consists of $\mu, \lambda_1, \lambda_2, \lambda_3$ and additionally γ where μ is a factor and the n 's are the counts



Function: llbtPC.fit()

preparations

```
> library(prefmod)
> load("cpc.Rdata")
> cities <- c("LO", "PA", "MI", "SG", "BA", "ST")
```

fit simple model

```
> m2 <- llbtPC.fit(cpc, nitems = 6, obj.names = cities)
> summary(m2)
```

fit model including effect for undecided

```
> m3 <- llbtPC.fit(cpc, nitems = 6, undec = TRUE, obj.names = cities)
> summary(m3)
```

compare models

```
> anova(m3, m2)
```



Functions: llbt.worth(), plotworth()

calculate worth parameters

```
> worth2 <- llbt.worth(m2)
> worth3 <- llbt.worth(m3)
> worthmat <- cbind(worth2, worth3)
```

plot them

```
> plotworth(worthmat)
```



Subject Covariates

Are the preference orderings different for different groups of subjects?

For one subject covariate on *s* levels we have now

$$\ln m_{(j>k)|s} = \mu_{(jk)s} + \lambda_s^S + (\lambda_j^{O_j} + \lambda_{js}^{O_j S}) - (\lambda_k^{O_k} + \lambda_{ks}^{O_k S})$$

where

- λ^O object parameters (for subject baseline group)
- λ^{OS} interaction parameter between objects and subject category
- λ_s^S fixing the margin for category *s* of covariate *S* (nuisance)
- μ 's nuisance parameters



LLBT Design structure with 1 subject covariate

PC	decision	counts	μ	λ^S	γ	λ_1^O	λ_2^O	λ_3^O	λ_{12}^{OS}	λ_{22}^{OS}	λ_{32}^{OS}
(12)	<i>O</i> ₁	<i>n</i> _{(1>2) 1}	1	0	0	1	-1	0	0	0	0
(12)	<i>no</i>	<i>n</i> _{(1=2) 1}	1	0	1	0	0	0	0	0	0
(12)	<i>O</i> ₂	<i>n</i> _{(2>1) 1}	1	0	0	-1	1	0	0	0	0
:	:	:	:					:			
(12)	<i>O</i> ₁	<i>n</i> _{(1>2) 2}	4	1	0	1	-1	0	1	-1	0
(12)	<i>no</i>	<i>n</i> _{(1=2) 2}	4	1	1	0	0	0	0	0	0
(12)	<i>O</i> ₂	<i>n</i> _{(2>1) 2}	4	1	0	-1	1	0	-1	1	0
:	:	:	:					:			

- *O* and *OS* are parameter of interest
- *MU* * *S* not related to objects – nuisance parameters



Options for llbtPC.fit(): formel, elim

fit null model (without subject covariates)

```
> mb0 <- llbtPC.fit(cpc, nitems = 6, undec = TRUE, formel = ~1,
+   elim = ~SEX, obj.names = cities)
```

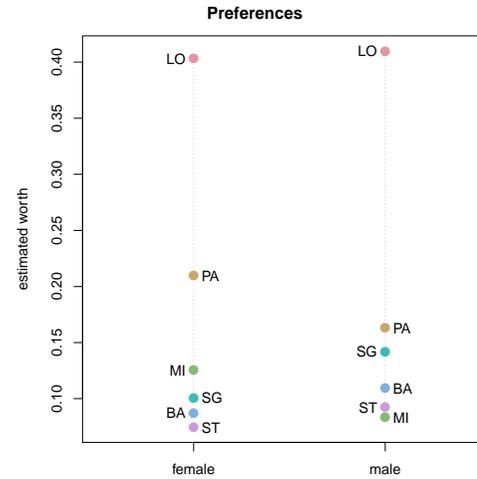
fit model with different preference scales for SEX

```
> mbsex <- llbtPC.fit(cpc, nitems = 6, undec = TRUE, formel = ~SEX,
+   elim = ~SEX, obj.names = cities)
```

Plot for different preference scales for SEX

female=1 and male =2

```
> wbsex <- llbt.worth(mbsex)
> colnames(wbsex) <- c("female", "male")
> plotworth(wbsex, ylab = "estimated worth")
```



Interpretation of parameters ofInterest

- $\lambda_j^{O_j}$ parameter estimate for O_j for the reference group (all subjects covariates on level = 0 in dummy coding)
- $\lambda_{js}^{O_j S}$ change of $\lambda_j^{O_j}$ for group s

model: SEX estimates for object PA

reference group	SEX1	λ^{PA}	
	SEX2	λ^{PA}	$+\lambda^{PA:SEX2}$



Interpretation of parameters ofInterest

```
> cmsw <- coef(mbsex)
> cmsw
Coefficients of interest:
      LO      PA      MI      SG      BA      ST
0.8444125 0.5177037 0.2604805 0.1505150 0.0788050      NA
      u  LO:SEX2  PA:SEX2  MI:SEX2  SG:SEX2  BA:SEX2
-1.3203958 -0.0994717 -0.2341781 -0.3117667 0.0647604 0.0042943
      ST:SEX2
      NA
```

model: SEX estimates for object PA model in **mbsex**

	λ^{PA}	$+\lambda^{PA:SEX2}$	
SEX1	0.518		$\lambda_{female}^{PA} = 0.518$
SEX2	0.518	-0.234	$\lambda_{male}^{PA} = 0.284$