

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 → pattern models
 - Ranking → pattern models
 - Rating → pattern models

The Basic Bradley-Terry Model (BT)

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

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

$$N_{(jk)} = n_{(j\succ k)} + n_{(k\succ j)}$$

total number of responses
to comparison (jk)

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

$$P(j \succ k) = \frac{\pi_j}{\pi_j + \pi_k}$$

π 's are called *worth parameters*
and are non-negative numbers
describing the location of the objects

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 $m_{(j \succ k)}$ of $n_{(j \succ k)}$ is

$$m_{(j \succ k)} = N_{(jk)} p_{(j \succ k)}$$

$$P(j \succ k) = \frac{\pi_j}{\pi_j + \pi_k} = c_{(jk)} \frac{\sqrt{\pi_j}}{\sqrt{\pi_k}}$$

where $c_{(jk)}$ is constant for a given comparison

then our basic paired comparison model for one comparison is

$$\ln m_{(j \succ k)} = \mu_{(jk)} + \lambda_j - \lambda_k$$

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

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

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

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	$n_{(1\succ 2)}$	1	1	-1	0
(12)	O_2	$n_{(2\succ 1)}$	1	-1	1	0
(13)	O_1	$n_{(1\succ 3)}$	2	1	0	-1
(13)	O_3	$n_{(3\succ 1)}$	2	-1	0	1
(23)	O_2	$n_{(2\succ 3)}$	3	0	1	-1
(23)	O_3	$n_{(3\succ 2)}$	3	0	-1	1

the **design structure** consists of counts (dependent variable) and the **design matrix \mathbf{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 (V_1, \dots, V_6) is

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

First respondent gave the following responses:

V_1	V_2	V_3	V_4	V_5	V_6
(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) \Rightarrow 0 means first object preferred

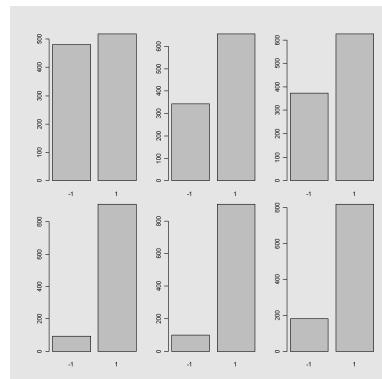
Wine data

- 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
```

prefmod > LLBT Wine Example - Result

```
> 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 modell $n_{ij} = m_{ij}$

- 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)
```

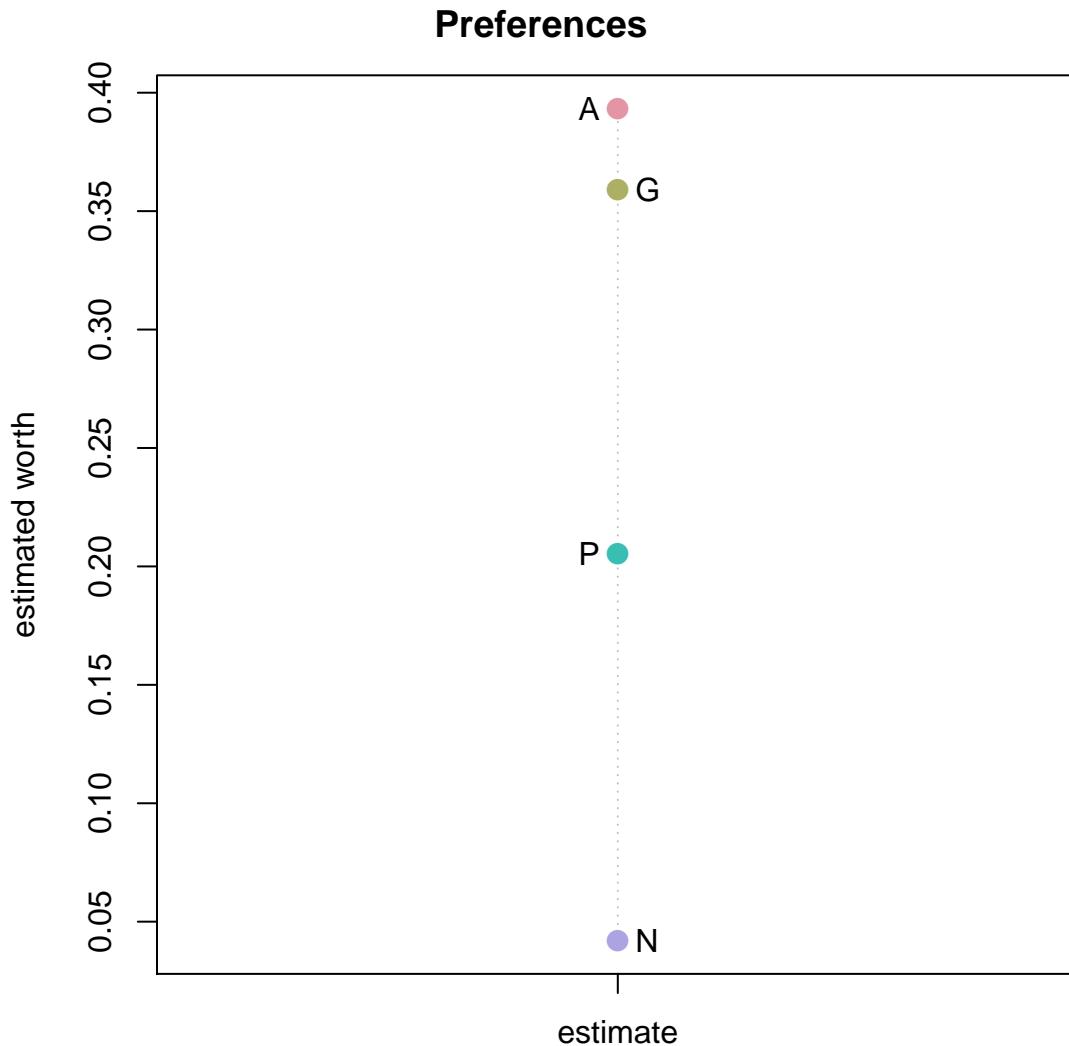
Remember:
$$\pi_j = \frac{\exp(2\lambda_j)}{\sum_j \exp(2\lambda_j)}$$
 $\lambda_A = 1.118, \lambda_G = 1.073, \lambda_P = 0.794, \lambda_N = 0$

- 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 ofInterest 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))
```

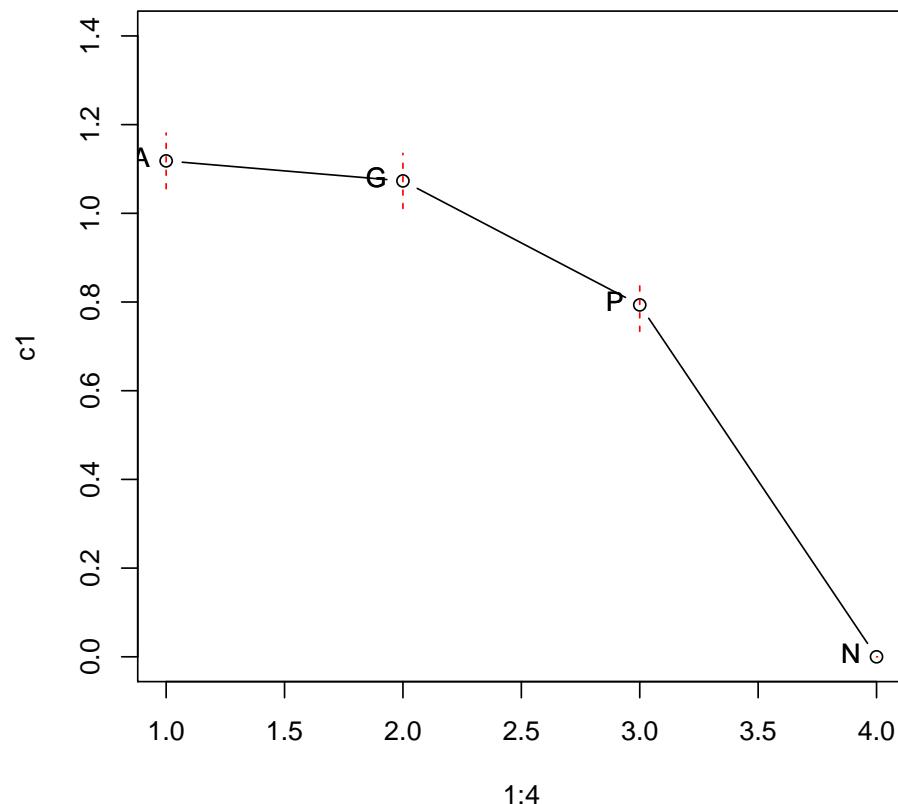
- How to make a confidence interval plot

```
> 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 estimates and confidence intervals

```
> 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**

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

insert equations

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

- 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 $m_{(A \succ G)}$ and for $m_{(G \succ A)}$?
- 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

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

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

- The observed preferences (counts) $n_{(i \succ j)}$ 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 $m_{(i \succ j)}$ 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:

$$\begin{aligned} j &\succ k \\ k &\succ j \\ j &= k \end{aligned}$$

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

$$\begin{aligned} LO &\succ PA \\ PA &\succ LO \\ LO &= PA \end{aligned}$$

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:

$$\ln m_{(j\succ k)} = \mu_{(jk)} + \lambda_j - \lambda_k$$

$$\ln m_{(k\succ j)} = \mu_{(jk)} - \lambda_j + \lambda_k$$

$$\ln m_{(j=k)} = \mu_{(jk)} + \gamma$$

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 $(j \succ k)$ to "no decision" $(j = k)$

$$\ln \left(\frac{P(j \succ k)}{P(j = k)} \right) = \ln m_{(j \succ k)} - \ln m_{(j = k)}$$

insert equations

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

for $\lambda_j = \lambda_k$

- log odds preferring $(j \succ k)$ 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	<i>decision</i>	<i>counts</i>	μ	γ	λ_1	λ_2	λ_3
(12)	O_1	$n_{(1\succ 2)}$	1	0	1	-1	0
(12)	<i>no</i>	$n_{(1=2)}$	1	1	0	0	0
(12)	O_2	$n_{(2\succ 1)}$	1	0	-1	1	0
(13)	O_1	$n_{(1\succ 3)}$	2	0	1	0	-1
(13)	<i>no</i>	$n_{(1=3)}$	2	1	0	0	0
(13)	O_3	$n_{(3\succ 1)}$	2	0	-1	0	1
(23)	O_2	$n_{(2\succ 3)}$	3	0	0	1	-1
(23)	<i>no</i>	$n_{(2=3)}$	3	1	0	0	0
(23)	O_3	$n_{(3\succ 2)}$	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

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

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)	O_1	$n_{(1\succ 2) 1}$	1	0	0	1	-1	0	0	0	0
(12)	no	$n_{(1=2) 1}$	1	0	1	0	0	0	0	0	0
(12)	O_2	$n_{(2\succ 1) 1}$	1	0	0	-1	1	0	0	0	0
:	:	:	:					:			
(12)	O_1	$n_{(1\succ 2) 2}$	4	1	0	1	-1	0	1	-1	0
(12)	no	$n_{(1=2) 2}$	4	1	1	0	0	0	0	0	0
(12)	O_2	$n_{(2\succ 1) 2}$	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)
```

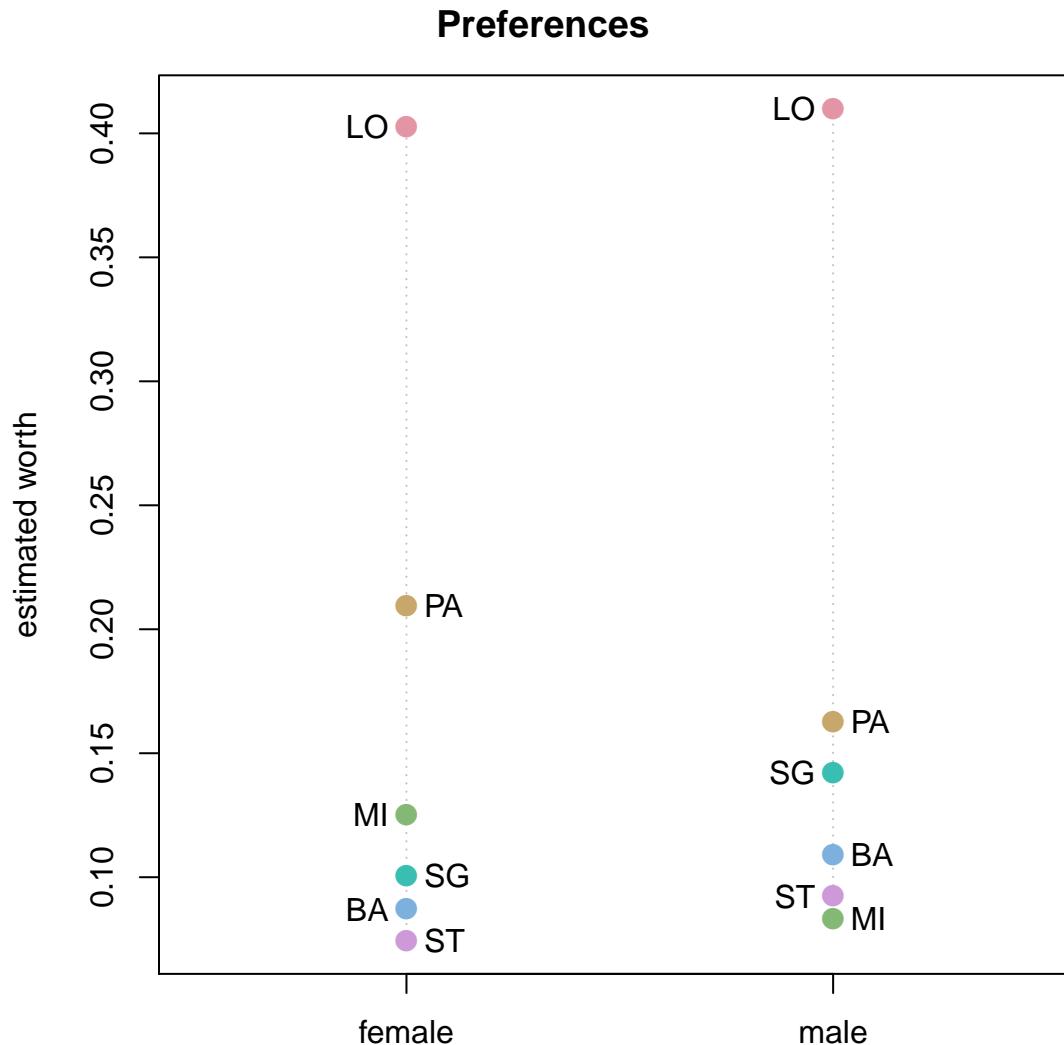
fit model with different preference scales for SEX

```
> 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 of Interest

- $\lambda_j^{O_j}$ 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	$\lambda^{PA} + \lambda^{PA:SEX2}$

Interpretation of parameters of Interest

```
> 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	$\lambda_{male}^{PA} = 0.284$