



prefmod: modelling preferences using a paired comparison approach

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Response-format	Modell		Designmatrix	Modell-schätzung	Notes
echte Paar-vergleiche	LLBT	Daten	llbt.design()	glm(), gnm()	1,2,(3),4, (5)
		Daten		llbt.fit()	1,3,5
	Pattern	Daten	patt.design()	glm(), gnm()	2,4,(5),6
		Daten	—————>	pattPC.fit()	1,3,(5),6
Rankings	Pattern	Daten	patt.design()	glm(), gnm()	2,4,(5)
		Daten	—————>	pattR.fit()	1,3,5
Ratings (Likert)	Pattern	Daten	patt.design()	glm(), gnm()	2,4,(5)
		Daten	—————>	pattL.fit()	1,3,5,6

- (1) NAs
- (2) R standard Output
- (3) größere Anzahl Vergleiche (Objekte)
- (4) Objektvariablen
- (5) metrische Personenvariablen
- (6) Dependencies



LLBT – Design Matrix

```
> data(cemspc)
> head(cemspc)
  V1 V2 V3 V4 V5 V6 V7 V8 V9 V10 V11 V12 V13 V14 V15 ENG SEX
1  0  0 NA  2  2  2  0  0  1  0  0  0  1  0  1  1  2
2  0  0 NA  0  2  2  0  2  2  2  0  2  2  0  2  1  1
3  1  0 NA  0  0  2  0  0  1  0  0  0  1  0  1  1  2
4  0  0 NA  0  2  0  0  0  0  0  0  0  0  0  2  1  1
5  0  0 NA  2  2  2  2  2  2  0  0  0  0  0  2  2  2
6  2  2 NA  0  0  0  2  2  2  2  0  0  0  0  2  1  2

> des <- llbt.design(cemspc, nitems = 6, cov.sel = "ENG")
> head(des)
  y mu g0 g1 g2 o1 o2 o3 o4 o5 o6 ENG
1 129 1 1 0 0 1 -1 0 0 0 0 0 1
2  20 1 0 1 0 0 0 0 0 0 0 0 1
3  73 1 0 0 1 -1 1 0 0 0 0 0 1
4 167 2 1 0 0 1 0 -1 0 0 0 0 1
5  16 2 0 1 0 0 0 0 0 0 0 0 1
6  39 2 0 0 1 -1 0 1 0 0 0 0 1
```



LLBT – Design Matrix

Usage:

```
llbt.design(obj, nitems = NULL, objnames = "",
            blnCasewise = FALSE, cov.sel = "",
            blnGLIMcmds = FALSE, glimCmdFile = "", outFile = "")
```



LLBT – mit glm()

```
> eng <- factor(des$ENG)
> res <- glm(y ~ o1 + o2 + o3 + o4 + o5 + o6 + eng:(o1 + o2 +
+ o3 + o4 + o5 + o6) + mu * eng, family = poisson, data = des)
> res$coefficients
(Intercept)      o1          o2          o3          o4          o5
  4.2248      1.0696      0.5771      0.1371      0.1762      0.1339
      o6      mu2          mu3          mu4          mu5          mu6
      NA     -0.1919     -0.3055     -0.1709      0.0264      0.0788
      mu7      mu8          mu9          mu10         mu11         mu12
 -0.1936      0.0148      0.0793      0.0787     -0.2708     -0.0288
      mu13      mu14      mu15      eng2      o1:eng2      o2:eng2
  0.0730      0.0689      0.0733     -1.0528     -0.0521     -0.1786
o3:eng2      o4:eng2      o5:eng2      o6:eng2      eng2:mu2      eng2:mu3
  0.0098      0.2452     -0.1001         NA      0.0775     -0.0542
eng2:mu4      eng2:mu5      eng2:mu6      eng2:mu7      eng2:mu8      eng2:mu9
  0.1797      0.0973      0.0202      0.0178      0.0652      0.0403
eng2:mu10      eng2:mu11      eng2:mu12      eng2:mu13      eng2:mu14      eng2:mu15
 -0.0043      0.0755      0.1005      0.0437     -0.0034      0.0502
```

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LLBT – using llbt.fit()

Usage

```
llbt.fit(y, Xmodel, q, ncat, maxiter = 100)
```

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LLBT – using gnm()

```
> eng <- factor(des$ENG)
> res2 <- gnm(formula = y ~ o1 + o2 + o3 + o4 + o5 + o6 + eng:(o1 +
+ o2 + o3 + o4 + o5 + o6), eliminate = eng:mu, family = poisson,
+ data = des)
> res2
Call:
gnm(formula = y ~ o1 + o2 + o3 + o4 + o5 + o6 + eng:(o1 + o2 +
      o3 + o4 + o5 + o6), eliminate = eng:mu, family = poisson,
      data = des)

Coefficients of interest:
      o1      o2      o3      o4      o5      o6      eng2:o1
 1.06960  0.57713  0.13713  0.17625  0.13390      NA  -0.05207
eng2:o2  eng2:o3  eng2:o4  eng2:o5  eng2:o6
-0.17865  0.00984  0.24522  -0.10011         NA

Deviance:      1175
Pearson chi-squared: 1030
Residual df:      50
```

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LLBT – using llbt.fit()

```
> mfr <- llbt.design(cemspc, nitems = 6, objnames = c("lo",
+ "pa", "mi", "sg", "ba", "st"), blnCasewise = TRUE)
> mm <- model.matrix(~lo + pa + mi + sg + ba, data = mfr)
> X <- mm[, -1]
> p <- ncol(X)
> ncat <- 3
> q <- length(levels(mfr$mu)) * length(levels(mfr$CASE))
> llbt.fit(mfr$y, X, q, ncat)
Results of llbt.fit:
```

```
Deviance: 9010
Residual df = 9085
Number of iterations: 22
```

	Estimate	Std. Error	z	P(z)
lo	1.05	0.047	22.6	0.0000
pa	0.53	0.044	12.1	0.0000
mi	0.14	0.043	3.2	0.0006
sg	0.24	0.042	5.7	0.0000
ba	0.11	0.042	2.5	0.0058

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eliminate – treating nuisance parameters

$$\text{IWLS} \quad \hat{\beta}^{(t+1)} = (\tilde{X}^T \hat{W}^{(t)} \tilde{X})^{-1} \tilde{X}^T \hat{W}^{(t)} z^{(t)}$$

partition critical part $\tilde{X} = (X|X_*)$

- X ... covariates for model parameters
- X_* ... covariates for nuisance parameters

$$\tilde{X}^T \hat{W}^{(t)} \tilde{X} = A = \begin{pmatrix} A_{11} & A_{12} \\ A_{21} & A_{22} \end{pmatrix} = \begin{pmatrix} X^T \hat{W}^{(t)} X & X^T \hat{W}^{(t)} X_* \\ X_*^T \hat{W}^{(t)} X & X_*^T \hat{W}^{(t)} X_* \end{pmatrix}$$

for A^{-1} we need inverses of A_{22} and $(A_{11} - A_{12}A_{22}^{-1}A_{21})$

A_{22} is diagonal, even very large models can be fitted



PATTERN MODEL – design matrix

Usage

```
patt.design(obj, nitems = NULL, objnames = "", resptype = "paircomp",
  blnRevert = FALSE, cov.sel = "", blnIntcovs = FALSE,
  blnGLIMcmds = FALSE, glimCmdFile = "", outFile = "", intFile = "")
```



PATTERN MODEL – design matrix

```
> rgbdat <- read.table("RGB_PC.dat", header = TRUE)
> dsgn <- patt.design(rgbdat, 3, objnames = c("R", "G", "B"))

> head(dsgn, 3)
  y R G B
1 10 2 0 -2
2  5 2 -2 0
3  0 0 0 0
> m1 <- glm(y ~ R + G + B, family = poisson, data = dsgn)
> m1
Call:  glm(formula = y ~ R + G + B, family = poisson, data = dsgn)
```

```
Coefficients:
(Intercept)          R          G          B
      1.196      0.510      0.288      NA

Degrees of Freedom: 7 Total (i.e. Null);  5 Residual
Null Deviance:      29
Residual Deviance: 18      AIC: 44
```



PATTERN MODEL – fitting utility for PCs

```
> pattPC.fit(cemspc, nitems = 3)
Results of pattern model for paired comparison
```

```
Call:
pattPC.fit(obj = cemspc, nitems = 3)
```

```
Deviance: 348
log likelihood: -790
```

```
no of iterations: 6 (Code: 1 )
```

	estimate	se	z	p-value
o1	0.93	0.069	13.6	0
o2	0.45	0.070	6.4	0



PATTERN MODEL – fitting utility for PCs

```
> pattPC.fit(cemspc, nitems = 3, NItest = TRUE)
Results of pattern model for paired comparison
```

```
Call:
pattPC.fit(obj = cemspc, nitems = 3, NItest = TRUE)
```

```
Deviance: 348
log likelihood: -790
```

```
no of iterations: 16 (Code: 1 )
```

	estimate	se	z	p-value
o1	0.953	0.080	11.93	0.00
o2	0.457	0.074	6.15	0.00
mis.o1	-0.084	0.172	-0.49	0.63
mis.o2	-0.067	0.218	-0.31	0.76



PATTERN MODEL – fitting utility for PCs

Usage

```
pattPC.fit(obj, nitems, formel = ~1, elim = ~1, resptype = "paircomp",
  obj.names = NULL, undec = FALSE, ia = FALSE, NItest = FALSE,
  NI = FALSE, MISalpha = NULL, MIScommon = FALSE, MISbeta = NULL,
  pr.it = FALSE)
```



PATTERN MODEL – fitting utility for PCs

```
> m1 <- pattPC.fit(cemspc, nitems = 3, formel = ~1, elim = ~SEX)
> m2 <- pattPC.fit(cemspc, nitems = 3, formel = ~SEX, elim = ~SEX)
> m2
Results of pattern model for paired comparison
```

```
Call:
pattPC.fit(obj = cemspc, nitems = 3, formel = ~SEX, elim = ~SEX)
```

```
Deviance: 372
log likelihood: -784
eliminated term(s): ~SEX
```

```
no of iterations: 14 (Code: 1 )
```

	estimate	se	z	p-value
o1	0.71	0.092	7.8	0e+00
o2	0.34	0.096	3.5	4e-04
o1:SEX2	0.47	0.140	3.4	7e-04
o2:SEX2	0.24	0.141	1.7	9e-02



PATTERN MODEL – fitting utility for Rankings

```
> load("conf.Rdata")
> conf[5:8, ]
  PREIS AUSSEN MARKE TECHNIK HERKUNFT INNEN SEX ALTER M.PREF
5  NA      2      4      NA      3      1      1      2      2
6  NA      2      4      3      NA      1      1      1      1
7   3      2      1      NA      NA      NA      2      2      3
8   6      2      4      5      3      1      1      1      2
> mR <- pattR.fit(conf, nitems = 6)
```



PATTERN MODEL – fitting utility for Rankings

```
> mRsex <- pattR.fit(conf, nitens = 6, formel = ~SEX, elim = ~SEX)
```

```
> worthmat <- patt.worth(mRsex)
```

```
> worthmat
```

Worthmatrix:

	SEX1	SEX2
PREIS	0.16	0.15
AUSSEN	0.21	0.18
MARKE	0.18	0.18
TECHNIK	0.17	0.19
HERKUNFT	0.11	0.12
INNEN	0.17	0.18



PATTERN MODEL – fitting utility for Rankings

```
> plotworth(worthmat)
```

