

Session 3: Extensions of CCA. The Inter Batteries Analysis

Sessions on Multivariate Modeling.

Course on Kernel Based Learning and Multivariate Modeling

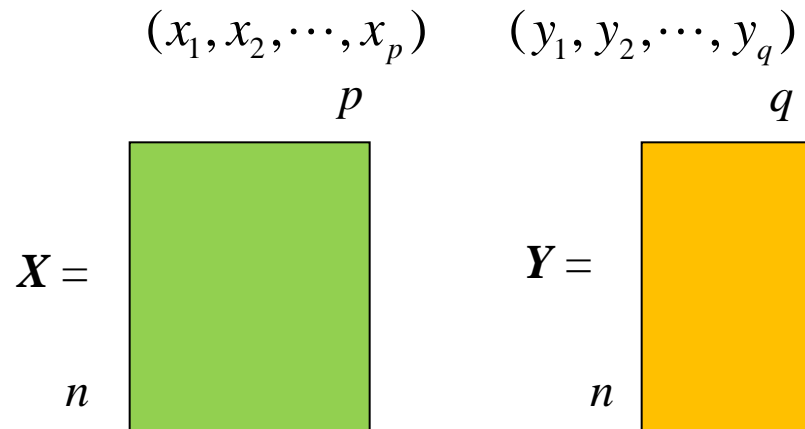
Tomàs Aluja-Banet

tomas.aluja@upc.edu

The problem

We continue having two vectors, measured on n individuals

(Tucker, 1958)



The goal is the same, to measure the relationship between both multivariate vectors by components derived from the original variables

$$t_h = Xa_h \quad u_h = Yb_h$$

Canonical components perform well to explain the other group of variables, but not their own group \rightarrow Hence instead of maximizing the $\text{cor}(t_h, u_h) \dots$

The Inter-batteries analysis

What if we relieve the assumption that components t_h and u_h have to be of unit length.

$$\text{Max } \langle t'_h, u_h \rangle_N = t'_h N u_h = a'_h X' N Y b_h = \gamma_h = \text{cov}(t_h, u_h)$$

$$t_h = X a_h$$

but then we will need to constraint a_h and b_h vectors.

$$u_h = Y b_h$$

$$a'_h a_h = 1; \quad b'_h b_h = 1$$

Restriction on \mathbb{R}^p and \mathbb{R}^q

$$\ell = a'_h X' N Y b_h - \lambda(a'_h a_h - 1) - \mu(b'_h b_h - 1)$$

$$X' N Y b_h - 2\lambda a_h = 0$$

$$a'_h X' N Y b_h = 2\lambda a'_h a_h$$

$$Y' N X a_h - 2\mu b_h = 0$$

$$b'_h Y' N X a_h = 2\mu b'_h b_h$$

$$X' N Y b_h = \gamma_h a_h$$

*

$$Y' N X a_h = \gamma_h b_h$$

$$2\lambda = 2\mu = \gamma_h$$

Transition relations

Here it is enough to consider X and Y centered, but usually they will be scaled, then matrices of covariances become correlations. If X , Y centered, results depend on the scale of variables.

The solution

$$V_{YX}a_h = \gamma_h b_h$$

$$V_{XY}b_h = \gamma_h a_h$$

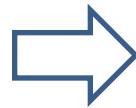
$$\text{rang}(V_{XY}) = \text{rang}(V_{YX}) = s$$

Maximal number of solutions s

Analysis in R^p and R^q

$$V_{XY}V_{YX}a_h = \gamma_h^2 a_h$$

$$V_{YX}V_{XY}b_h = \gamma_h^2 b_h$$



Analysis in R^n

$$XX'NYY'Nt_h = \gamma_h^2 t_h \quad t_h = Xa_h$$

$$YY'NXX'Nu_h = \gamma_h^2 u_h \quad u_h = Yb_h$$

$$A'A = I$$

$$B'B = I$$

$$T = XA$$

$$U = YB$$

$$t_h' N t_h = a_h' V_X a_h$$

$$u_h' N u_h = b_h' V_X b_h$$

Components t_h and u_h neither normalized, nor orthogonal

IBA in practice

usually $q < p$

$$V_{YX}V_{XY}b_h = \gamma_h^2 b_h$$

$$a_h = \frac{1}{\gamma_h} V_{XY} b_h$$

$$t_h = Xa_h$$

$$u_h = Yb_h$$

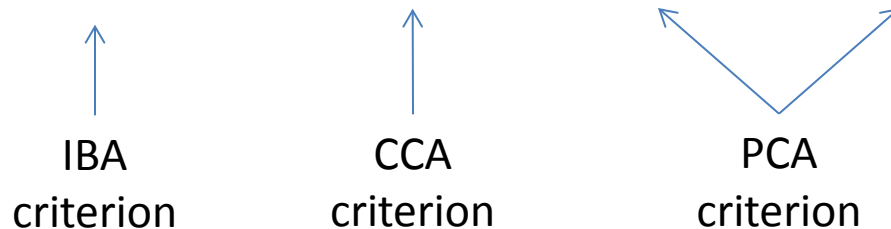
number of solutions = $\text{rank}(V_{XY})$

$$t_h^{\text{test}} = X^{\text{test}} a_h$$

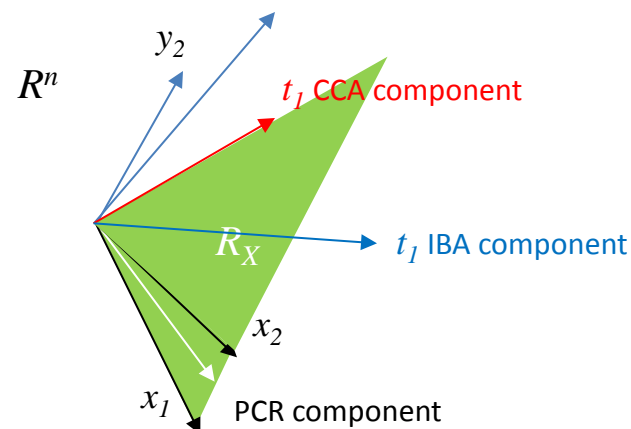
Properties of IBA components

IBA is a compromise between the CCA and the PCA of both blocks

$$\text{cov}(t_h, u_h) = \text{cor}(t_h, u_h) \sqrt{\text{var}(Xa_h)} \sqrt{\text{var}(Yb_h)}$$



The IBA components will compromise the explanation of each own group and the prediction between groups



Properties of IBA components

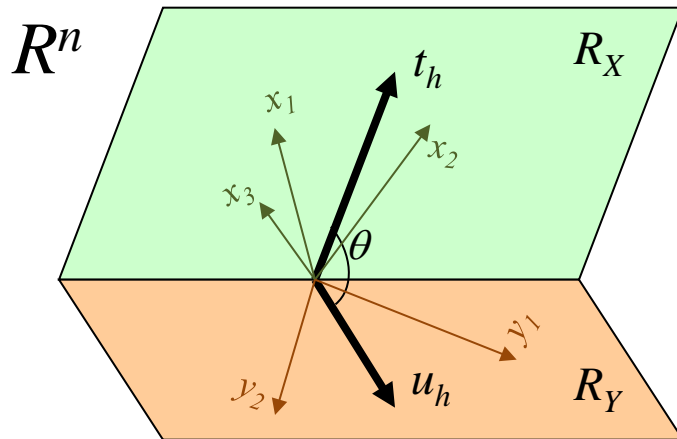
$$\|t_h\|_N^2 = \langle t_h, t_h \rangle_N = a_h' X' N X a_h$$

$$\|u_h\|_N^2 = u_h' N u_h = b_h' Y' N Y b_h$$

$$\text{cov}(t_h, t_l) = a_h' X' N X a_l \neq 0, \quad \text{cov}(u_h, u_l) = b_h' Y' N Y b_l \neq 0$$

***IBA components are
not orthogonal***

Properties of IBA components



$$\begin{aligned}
 * \quad X'NYb_h &= X'Nu_h = \gamma_h a_h \\
 Y'NXa_h &= Y'Nt_h = \gamma_h b_h
 \end{aligned}$$

$$a_h \text{ colinear to } \frac{1}{\gamma_h} \left(X'NYb_h = X'Nu_h = \left(\text{cov}(x_j, u_h) \right) \right)$$

$$b_h \text{ colinear to } \frac{1}{\gamma_h} \left(Y'NXa_h = Y'Nt_h = \left(\text{cov}(y_k, t_h) \right) \right)$$

a_h and b_h are proportional to the covariance of each variable with the IBA component of the other group

Properties of IBA components

$$\gamma_h = \text{cov}(t_h, u_h) = a'_h X' N Y b_h = \langle a_h, X' N Y b_h \rangle = \cos(a_h, X' N u_h) \|X' N u_h\| = \|X' N u_h\|$$

$$\text{Max } \gamma_h \Rightarrow \gamma_h^2 = \|X' N u_h\|^2 = \sum_{j=1}^p \text{cov}^2(x_j, u_h) = \sum_{k=1}^q \text{cov}^2(y_k, t_h)$$

The IBA components maximize the sum of covariances with the variables of the other group

$$\text{cov}(t_h, u_h) = \gamma_h$$

$$\text{cov}(t_h, u_l) = 0 \quad a'_h X' N Y b_l = a'_h V_{XY} b_l = \gamma_l a'_h a_l = 0$$

Covariance with
homonymous
components is
maximized, otherwise is
zero

Singular Value Decomposition

$$\left. \begin{array}{ll} V_{YX} a_h = \gamma_h b_h & V_{YX} A = B \Lambda \\ V_{XY} b_h = \gamma_h a_h & V_{XY} B = A \Lambda \end{array} \right\} V_{XY} = A \Lambda B' \quad \Lambda = \begin{bmatrix} \ddots & & \\ & \gamma_h & \\ & & \ddots \end{bmatrix}$$

$$V_{XY} = \sum_{h=1}^s \gamma_h a_h b_h'$$

$$\text{cov}(x_j, y_k) = \sum_{h=1}^s \frac{1}{\gamma_h} \begin{pmatrix} \vdots \\ \text{cov}(x_j, u_h) \\ \vdots \end{pmatrix} (\cdots \text{cov}(y_k, t_h) \cdots)$$

$$\text{cor}(x_j, y_k) = \sum_{h=1}^s \frac{\text{cor}(x_j, u_h) \times \text{cor}(y_k, t_h)}{\text{cor}(t_h, u_h)}$$

Variable displays

Variables

basis t_1, t_2, \dots
(non orthogonal)

$$\psi_h^X = \begin{pmatrix} \vdots \\ \text{cor}(x_j, t_h) \\ \vdots \end{pmatrix}$$

$$\psi_h^Y = \begin{pmatrix} \vdots \\ \text{cor}(y_k, t_h) \\ \vdots \end{pmatrix}$$

basis u_1, u_2, \dots
(non orthogonal)

$$\varphi_h^X = \begin{pmatrix} \vdots \\ \text{cor}(x_j, u_h) \\ \vdots \end{pmatrix}$$

$$\varphi_h^Y = \begin{pmatrix} \vdots \\ \text{cor}(y_k, u_h) \\ \vdots \end{pmatrix}$$

They are not biplots, and difficult to interpret since dimensions are not orthogonal

Individual displays

Individuals

2 displays

$$Xa_h = t_h$$

$$Yb_h = u_h$$

Display of individuals in the (t_h, u_h) basis.
To reveal the strength of the liaison

$$(t_1, u_1), (t_2, u_2), \dots$$

Relation with the orthogonal procrustean rotation

Orthogonal procrustean rotation

$\text{Min } \|X - YR\|^2$ R rotation matrix

$$X'Y = nV_{XY} = A\Lambda B', \Rightarrow R = BA'$$

We are assuming X and Y of the same dimensions, otherwise we complete with columns of zeroes.

$$\min \left(\|X - YR\|^2 = \text{tr}((X - YR)'(X - YR)) = \text{tr}(X'X + Y'Y - 2X'YR) \right)$$

$$\Rightarrow \max \left(\text{tr}(X'YR) = \text{tr}(A\Lambda B'R) = \text{tr}(\Lambda B'RA) = \text{tr}(\Lambda Q) = \sum \gamma_h q_{hh} \right)$$

$$\begin{aligned} |q_{hh}| \leq 1 &\Rightarrow Q = I \Rightarrow R = BA' \\ &\Rightarrow \text{Max } \text{tr}(X'YR) = \sum \gamma_h \end{aligned}$$

$$\|X - YBA'\|^2 = \text{tr}((X - YBA')(X' - AB'Y')) =$$

$$\text{tr}((X - YBA')AA'(X' - AB'Y')) = \text{tr}(XA - YB)(A'X' - B'Y') = \|T - U\|^2 \quad \begin{aligned} T &= XA \\ U &= YB \end{aligned}$$

IBA components reconstitutes the closest configuration between both blocks

Optimal approximation of V_{XY}

The IBA biplot is not as easy to interpret as the CCA, but it gets optimal approximation of the true covariances

Reconstitution of r rang

$$\sum_{h=1}^s \gamma_h^2 = \text{tr}(V_{XY} V_{YX}) = \|V_{XY}\|^2 = \|\hat{V}_{XY}^r\|^2 + \|V_{XY} - \hat{V}_{XY}^r\|^2 \quad \hat{V}_{XY}^r = \sum_{h=1}^r \gamma_h a_h b_h'$$

$$\|\hat{V}_{XY}^r\|^2 = \sum_{h=1}^r \gamma_h^2$$

$$\|V_{XY} - \hat{V}_{XY}^r\|^2 = \sum_{h=r+1}^s \gamma_h^2$$

Significant components (with the usual assumptions and standardized data)

(Tucker, 1958)

$$\gamma_1 = \gamma_2 = \dots = \gamma_s = 0$$

$$r = 0 \rightarrow n \sum_{h=1}^s \gamma_h^2 \sim \chi_{p \times q}^2$$

$$\gamma_1 > \gamma_r > \gamma_{r+1} = \dots = \gamma_s = 0$$

$$r > 0 \rightarrow \frac{n(p-r)(q-r) \sum_{h=r+1}^s \gamma_h^2}{(p - \sum_{h=1}^r \text{var}(t_h))(q - \sum_{h=1}^r \text{var}(u_h))} \sim \chi_{(p-r)(q-r)}^2$$

IBA Regression

IBA Regression in practice:

We use the t_1, t_2, \dots significant interbattery components as explanatory latent components of the y_j variables.

$$Y = T_{(r)}^{IBA} \tilde{B}_{(r)} + \varepsilon_Y$$

$$\tilde{B}_{(r)} = (T_{(r)}'^{IBA} N T_{(r)}^{IBA})^{-1} T_{(r)}'^{IBA} N Y$$

Expressing the model as function of the original variables

$$Y = X A_{(r)}^{IBA} \tilde{B}_{(r)} + \varepsilon_Y = X B + \varepsilon_Y$$

$$B = A_{(r)}^{IBA} (T_{(r)}'^{IBA} N T_{(r)}^{IBA})^{-1} T_{(r)}'^{IBA} N Y$$

Limitations of IBA

- Number max. of components = $\text{rank}(V_{XY}) \leq \min(p, q, n-1)$
- Solution depends on scale of variables, if data just centered
- Non orthogonal components
- Joint representations are not biplots

Advantages of IBA

- Components are good for prediction and altogether good factors for the own group.
- Deals with collinearity

Running IBA

```
> library(plsdepot)
> iba <- interbat(X, Y, scaled = TRUE)
```

```
> iba$values
[1] 1.272426 0.005657 0.001106
```

max. covariances

```
> iba$x.scores
  t1      t2      t3
1 -0.6429 -0.07471 -0.764316
2 -0.7697 -0.15463 -0.366123
3 -0.9074  0.20078  0.452996
4  0.6884 -0.09726  0.808580
5 -0.4867 -0.24373 -1.363398
6 .....
```

= T

```
> iba$y.scores
  u1      u2      u3
1  0.37145 -0.05444 -0.82290
2  1.34032  0.19638 -0.71715
3  0.08235  0.58493  0.86557
4  0.35497 -0.62863  0.74383
5 -0.46312 -0.39857  0.39749
6 .....
```

= U

```
> iba$x.wgs
```

	t1	t2	t3
Weight	-0.5899	0.7721	-0.2364
Waist	-0.7713	-0.4522	0.4478
Pulse	0.2389	0.4465	0.8623

= A

```
> iba$y.wgs
```

	u1	u2	u3
Tractions	-0.6133	-0.2140	0.76029
Push.ups	-0.7470	-0.1556	-0.64638
Jumps	-0.2567	0.9643	0.06443

= B

```
> iba$cor.xt
```

	t1	t2	t3
Weight	-0.9476	0.4049	-0.19478
Waist	-0.9620	0.1169	-0.07474
Pulse	0.5108	0.6088	0.95034

```
> iba$cor.yu
```

	u1	u2	u3
Tractions	-0.8802	0.1946	0.61617
Push.ups	-0.9397	0.4257	-0.13368
Jumps	-0.7407	0.9420	0.01581

Running IBA

```
> iba$cor.xu
```

	u1	u2	u3
Weight	0.4647	-0.07254	0.01414
Waist	0.6077	0.04249	-0.02679
Pulse	-0.1882	-0.04195	-0.05158


```
> iba$cor.yt
```

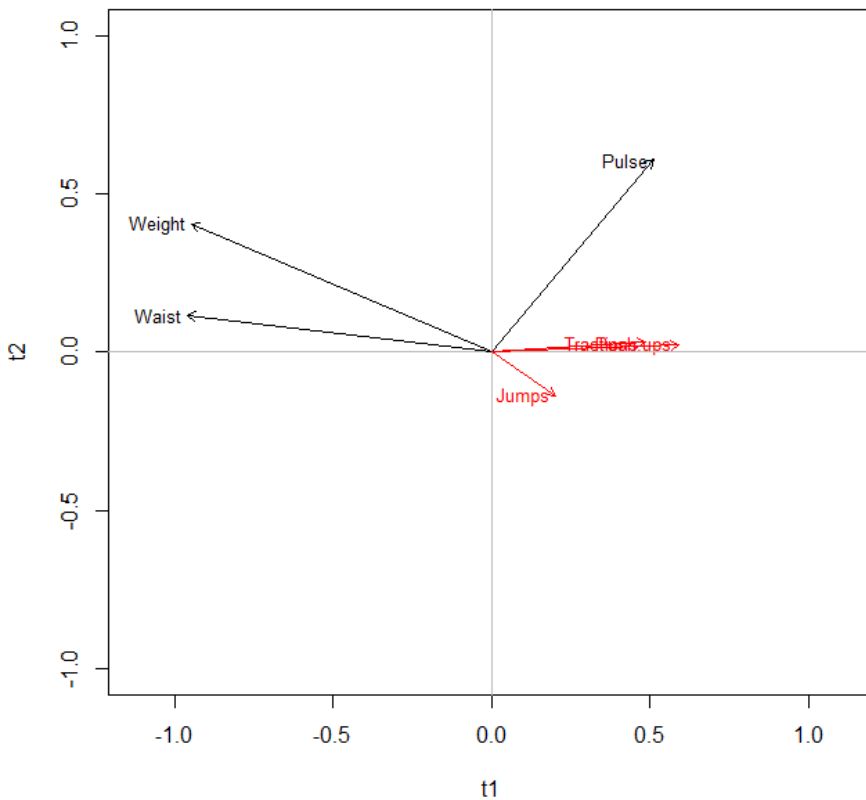
	t1	t2	t3
Tractions	0.4862	0.03028	-0.030384
Push.ups	0.5921	0.02202	0.025832
Jumps	0.2035	-0.13643	-0.002575


```
> iba$cor.tu
```

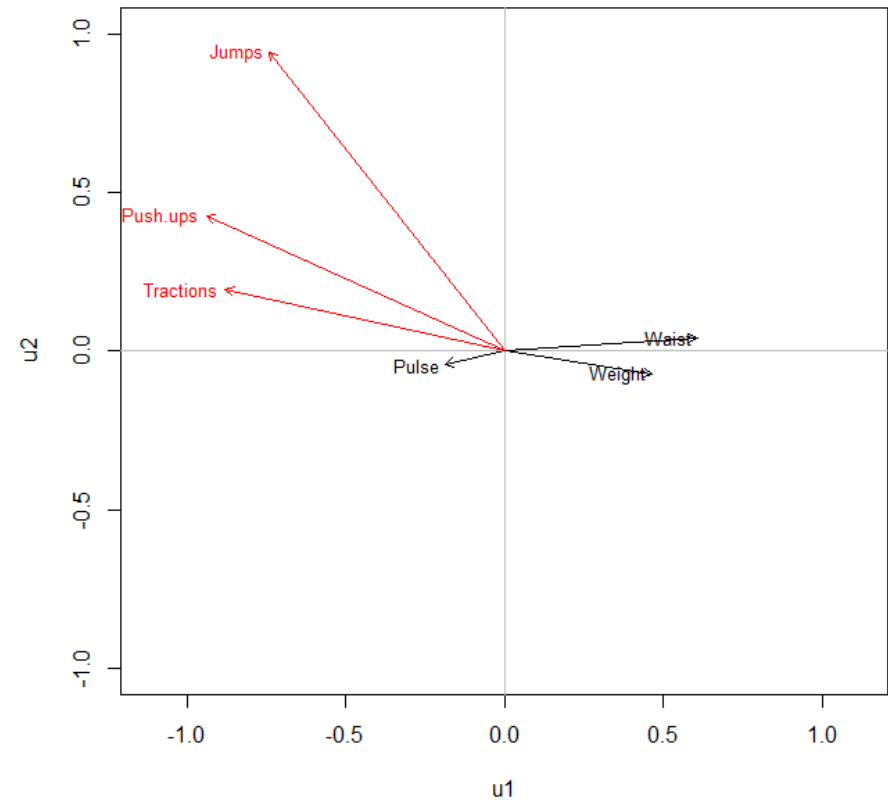
	t1	t2	t3	u1	u2	u3
t1	1.000e+00	-1.290e-01	2.808e-01	-5.536e-01	-6.165e-17	4.072e-16
t2	-1.290e-01	1.000e+00	5.789e-01	1.682e-16	-1.767e-01	-2.827e-15
t3	2.808e-01	5.789e-01	1.000e+00	-1.433e-16	2.595e-15	-7.189e-02
u1	-5.536e-01	1.682e-16	-1.433e-16	1.000e+00	-4.743e-01	-1.970e-01
u2	-6.165e-17	-1.767e-01	2.595e-15	-4.743e-01	1.000e+00	-1.197e-01
u3	4.072e-16	-2.827e-15	-7.189e-02	-1.970e-01	-1.197e-01	1.000e+00

Correlations

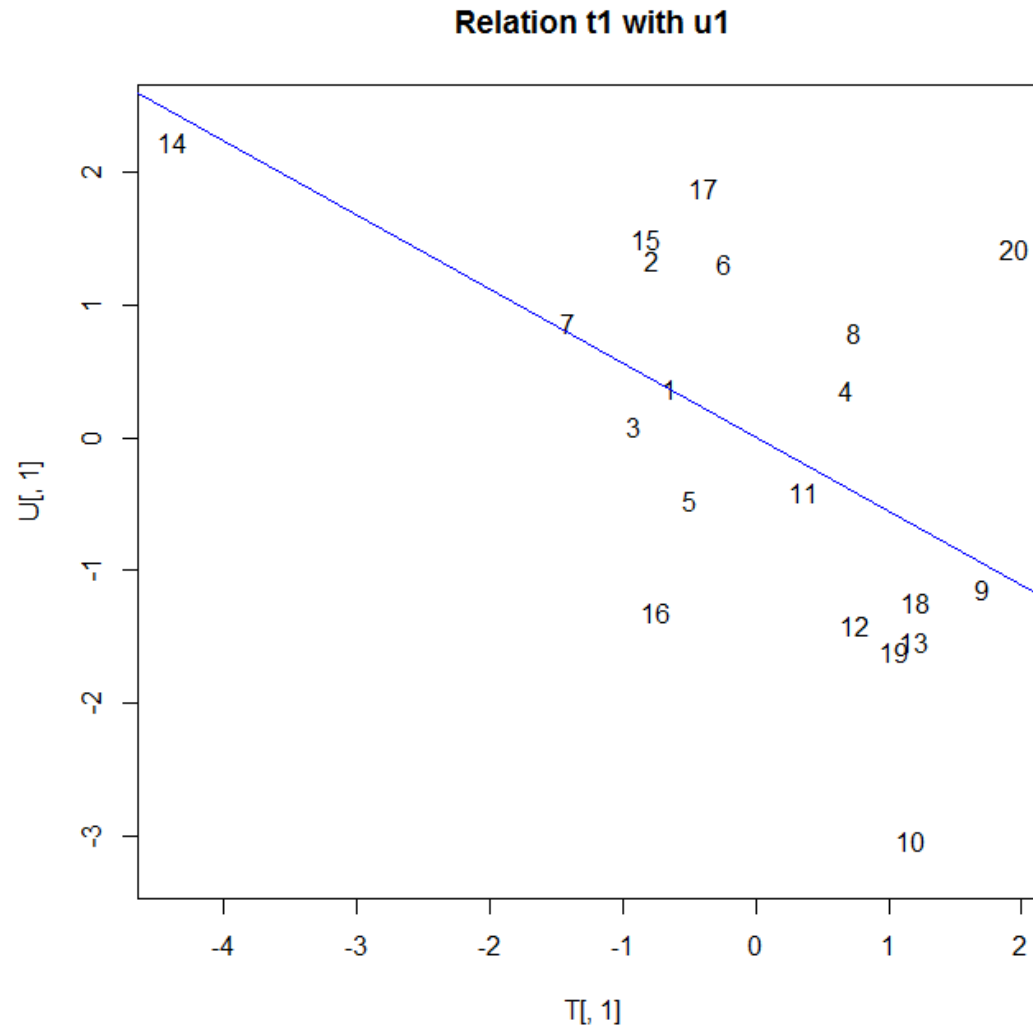
correlations on t1, t2



correlations on u1, u2



Relation t_1 with u_1



Communalities

```
> for (i in 1:p) {for ( j in 1:a) {com.xt[i,j] <- summary(lm(Xs[,i]~T[,1:j]))$r.squared}}
```

```
> rownames(com.xt) <- names(X)
```

```
> print(com.xt)
```

	[,1]	[,2]	[,3]
Weight	0.8980	0.9792	1
Waist	0.9255	0.9255	1
Pulse	0.2609	0.7239	1

```
> for (i in 1:q) {for ( j in 1:a) {com.yu[i,j] <- summary(lm(ys[,i]~U[,1:j]))$r.squared}}
```

```
> rownames(com.yu) <- names(Y)
```

```
> print(com.yu)
```

	[,1]	[,2]	[,3]
Tractions	0.7747	0.8388	1
Push.ups	0.8830	0.8835	1
Jumps	0.5487	0.9988	1

Redundancies

```
> for (i in 1:q) {for ( j in 1:a) {red.yt[i,j] <- summary(lm(Ys[,i]~T[,1:j]))$r.squared}}  
  
> rownames(red.yt) <- names(Y)  
  
> print(red.yt)
```

	[,1]	[,2]	[,3]
Tractions	0.2363	0.24514	0.3396
Push.ups	0.3506	0.36044	0.4365
Jumps	0.0414	0.05374	0.0539

```
> for (i in 1:p) {for ( j in 1:a) {red.xu[i,j] <- summary(lm(Xs[,i]~U[,1:j]))$r.squared}}  
  
> rownames(red.xu) <- names(X)  
  
> print(red.xu)
```

	[,1]	[,2]	[,3]
Weight	0.21597	0.24418	0.26792
Waist	0.36926	0.51037	0.54784
Pulse	0.03542	0.05763	0.07487

Number of significant dimensions

```
> iba$statistic
      phi df  p.value
1 25.5838  9 0.002389
2  0.5842  4 0.964800
3  0.1034  1 0.747824
```

$$H_0 \quad \gamma_{1...h} > 0, \gamma_{h+1...s} = 0$$

```
> print(RMPRESS)
  Traction Push.ups Jumps
1    0.8902    0.8199 0.991
2    0.9046    0.8473 1.054
3    0.9200    0.9124 1.090
>
> print(R2cv)
  Traction Push.ups Jumps
1    0.13655 0.26761 -0.07009
2    0.10841 0.21772 -0.21050
3    0.07789 0.09289 -0.29409
```

Running IBA

```

> lmY <- lm(ys~T[,1]-1)
> summary(lmY)
Response Traction :
Coefficients:
      Estimate Std. Error t value Pr(>|t|)
T[, 1]      0.342      0.141    2.42  0.025 *
Residual standard error: 0.874 on 19 degrees of freedom
Multiple R-squared:  0.236,    Adjusted R-squared:  0.196
F-statistic: 5.88 on 1 and 19 DF,  p-value: 0.0254

Response Push.ups :
Coefficients:
      Estimate Std. Error t value Pr(>|t|)
T[, 1]      0.416      0.130    3.2  0.0047 **
Residual standard error: 0.806 on 19 degrees of freedom
Multiple R-squared:  0.351,    Adjusted R-squared:  0.316
F-statistic: 10.3 on 1 and 19 DF,  p-value: 0.00469

Response Jumps :
Coefficients:
      Estimate Std. Error t value Pr(>|t|)
T[, 1]      0.143      0.158    0.91  0.38
Residual standard error: 0.979 on 19 degrees of freedom
Multiple R-squared:  0.0414,    Adjusted R-squared:  -0.00905
F-statistic: 0.821 on 1 and 19 DF,  p-value: 0.376

> summary(manova(lmY))
      Df Pillai approx F num Df den Df Pr(>F)
T[, 1]   1  0.432    4.31      3    17  0.02 *
Residuals 19

```

Running IBA

```
> # coefficients as functions of the original variables
>
> b.coef <- A[,1:nd] %*% lmY$coefficients
> rownames(b.coef) <- names(X)

> print(b.coef,digits=4)
      Traction Push ups      Jumps
Weight  -0.20152 -0.24545 -0.08434
Waist   -0.26351 -0.32094 -0.11029
Pulse    0.08161  0.09939  0.03415

> cor(X,Y)
      Traction Push ups      Jumps
Weight  -0.3897  -0.4931 -0.22630
Waist   -0.5522  -0.6456 -0.19150
Pulse    0.1506   0.2250  0.03493
```


Summary of IBA results

