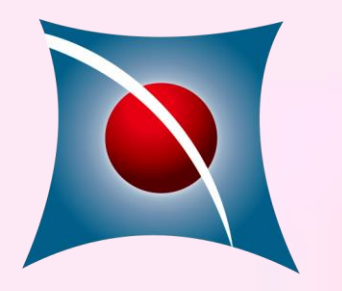


Mixtures of unrestricted skew normal factor analyzers with missing information



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Abstract

Mixtures of factor analyzers (**MFA**) based on the restricted skew normal distribution (rMSN) has been shown to be a flexible tool for modeling **asymmetrical high-dimensional data with heterogeneity**. However, the rMSN distribution is oft-criticized a lack of sufficient ability to accommodating skewness arising from more than one feature space. This thesis presents an alternative extension of MFA by assuming the unrestricted skew normal (uMSN) distribution for the component factors. In particular, the proposed mixtures of uMSN factor analyzers (**MuSNFA**) can simultaneously accommodate multiple directions of skewness and the occurrence of missing values or nonresponses. Under the missing at random mechanism, we develop a computationally feasible **expectation conditional maximization (ECM) algorithm** for computing the maximum likelihood estimates of model parameters. Practical aspects related to model-based clustering, prediction of factor scores and missing values are also discussed. The utility of the proposed methodology is illustrated with the analysis of simulated data and an application to the Pima Indian women diabetes data containing genuine **missing values**.

Methodology

MuSNFA model

$Y_j = \mu_i + B_i U_{ij} + \varepsilon_{ij}$, with probability π_i ($i = 1, \dots, g$)

$U_{ij} \sim uSN_q(-c\Delta_i^{-1/2}\Lambda_i\mathbf{1}_q, \Delta_i^{-1}, \Delta_i^{-1/2}\Lambda_i)$, $\varepsilon_{ij} \sim N_p(\mathbf{0}, D_i)$, $U_{ij} \perp \varepsilon_{ij}$

where Λ_i be a $q \times q$ **skewness parameter** matrix, and $\Delta_i = I_q + (1 - c^2)\Lambda_i\Lambda_i^T$ with $c = \sqrt{2/\pi}$.

Introduce two permutation matrices $O_j(p_j^o \times p)$ and $M_j((p - p_j^o) \times p)$ such that

$Y_j = O_j^T Y_j^o + M_j^T Y_j^m$. The hierarchical representation of **MuSNFA** is

$Y_j^o | (Z_{ij} = 1) \sim CFUSN_{p_j^o, q}(\mu_j^o - c\alpha_{ij}^o \mathbf{1}_q, \Sigma_{ij}^o, \alpha_{ij}^o)$

$Z_j \sim \mathcal{M}(1; \pi_1, \dots, \pi_g)$,

where $Y_j^o = O_j Y_j$, $Y_j^m = M_j Y_j$, $\mu_j^o = O_j \mu_j$, $\Sigma_{ij}^o = O_j \Sigma_i O_j^T$, $\alpha_{ij}^o = O_j \alpha_i$, $\Sigma_i = B\Delta_i^{-1}B^T + D_i$,

$\alpha_i = B\Delta_i^{-1/2}\Lambda_i$, and Z_j is a set of binary allocation indicators.

The component pdf of $Y_j^o | (Z_{ij} = 1)$ is given by

$\phi(y_j^o; \theta_i) = 2^q \phi_{p_j^o}(y_j^o; \mu_j^o - c\alpha_{ij}^o \mathbf{1}_q, \Omega_{ij}^o) \times \Phi_q(\alpha_{ij}^{oT} \Omega_{ij}^{oo-1}(y_j^o - \mu_j^o + c\alpha_{ij}^o \mathbf{1}_q); I_q - \alpha_{ij}^{oT} \Omega_{ij}^{oo-1} \alpha_{ij}^o)$,

where $\Omega_{ij}^{oo} = O_j \Omega_i O_j^T$ with $\Omega_i = \Sigma_i + \alpha_i \alpha_i^T$.

For ease of estimation, we reparameterize

$\tilde{B}_i \triangleq B_i \Delta_i^{-1/2}$ and $\tilde{U}_{ij} \triangleq \Delta_i^{1/2} U_{ij}$

It follows that $Y_j = \mu_i + \tilde{B}_i \tilde{U}_{ij} + \varepsilon_{ij}$, where $\tilde{U}_{ij} \sim uSN_q(-c\Lambda_i \mathbf{1}_q, I_q, \Lambda_i)$.

Let $\Theta = \{\pi_1, \dots, \pi_{g-1}, \theta_1, \dots, \theta_g\}$ represent all the unknown parameters of the mixture model,

where $\theta_i = (\mu_i, B_i, D_i, \Lambda_i)$.

The complete data likelihood function of Θ given $Y_c = (y^o, y^m, \tilde{U}, \gamma, Z)$ is

$$L_c(\Theta | Y_c) = \prod_{j=1}^n f(y_j^o | Z_{ij} = 1) f(Z_j) = \prod_{j=1}^n \prod_{i=1}^g \{\pi_i \phi(y_j^o | \theta_i)\}^{Z_{ij}} \\ = \prod_{j=1}^n \prod_{i=1}^g \left\{ \pi_i \phi_{p_j^o}(y_j^o; \mu_j^o; \tilde{B}_{ij}^o \tilde{U}_{ij} + D_{ij}^o) \phi_q(\tilde{U}_{ij}; -c\Lambda_i \mathbf{1}_q + \Lambda_i \gamma_j, I_q) \right\}^{Z_{ij}}$$

Proposition:

For notational simplicity, the symbol " $| \dots$ " stands for conditioning on y_j^o and $Z_{ij} = 1$.

Given the hierarchical representation, we have the following conditional expectations:

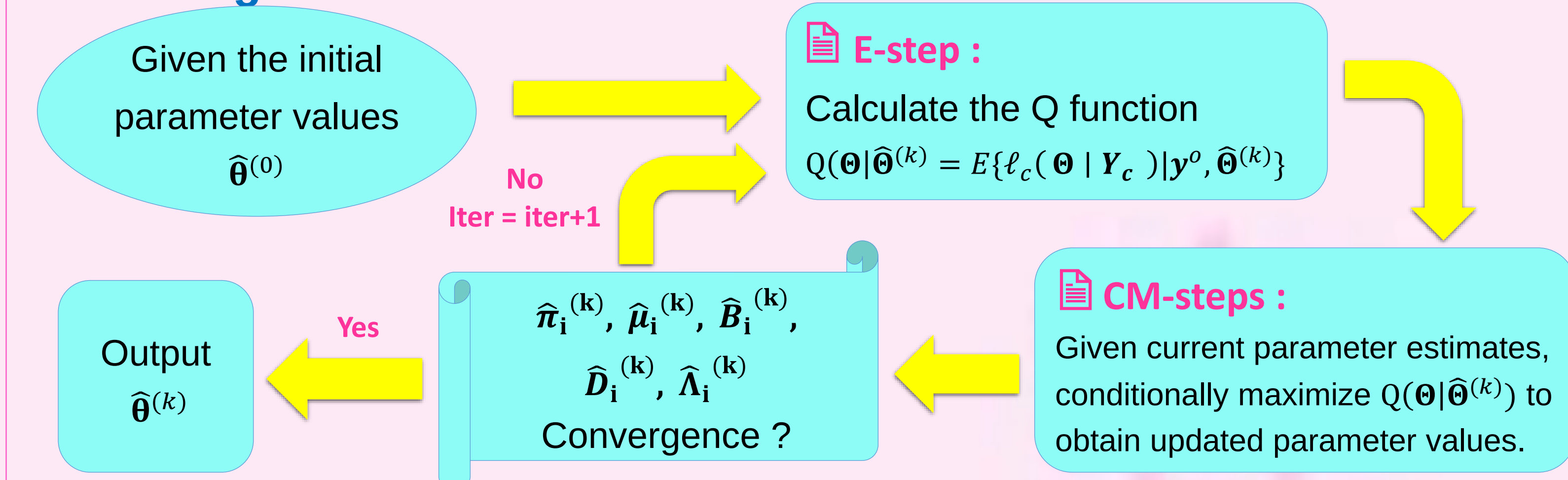
$E(\tilde{U}_{ij} | \dots) = W_{ij}^{oo} \{v_{ij}^o + \Lambda_i [E(Y_j - c\mathbf{1}_q | \dots)]\}$, $W_{ij}^{oo} = I_q + \tilde{B}_i^T C_{ij}^{oo} \tilde{B}_i^{-1}$, $v_{ij}^o = \tilde{B}_i^T C_{ij}^{oo} (y_j - \mu_i)$,

$E(\tilde{U}_{ij} y_j^T | \dots) = W_{ij}^{oo} \{v_{ij}^o E(y_j^T | \dots) + \Lambda_i [E(y_j y_j^T | \dots) - c\mathbf{1}_q E(y_j^T | \dots)]\}$ and

$E(\tilde{U}_{ij} \tilde{U}_{ij}^T | \dots) = \{I_q + E(\tilde{U}_{ij} | \dots) v_{ij}^{oT} + [E(\tilde{U}_{ij} y_j^T | \dots) - cE(\tilde{U}_{ij} | \dots) \mathbf{1}_q^T \Lambda_i] W_{ij}^{ooT}\}$,

where $E(y_j^T | \dots)$ and $E(y_j y_j^T | \dots)$ are the first-two moments of the TMVN distribution.

ECM algorithm



Conclusions and future work

The proposed MuSNFA model is a new model-based clustering tool for handling asymmetric high-dimensionality data with possibly missing values.

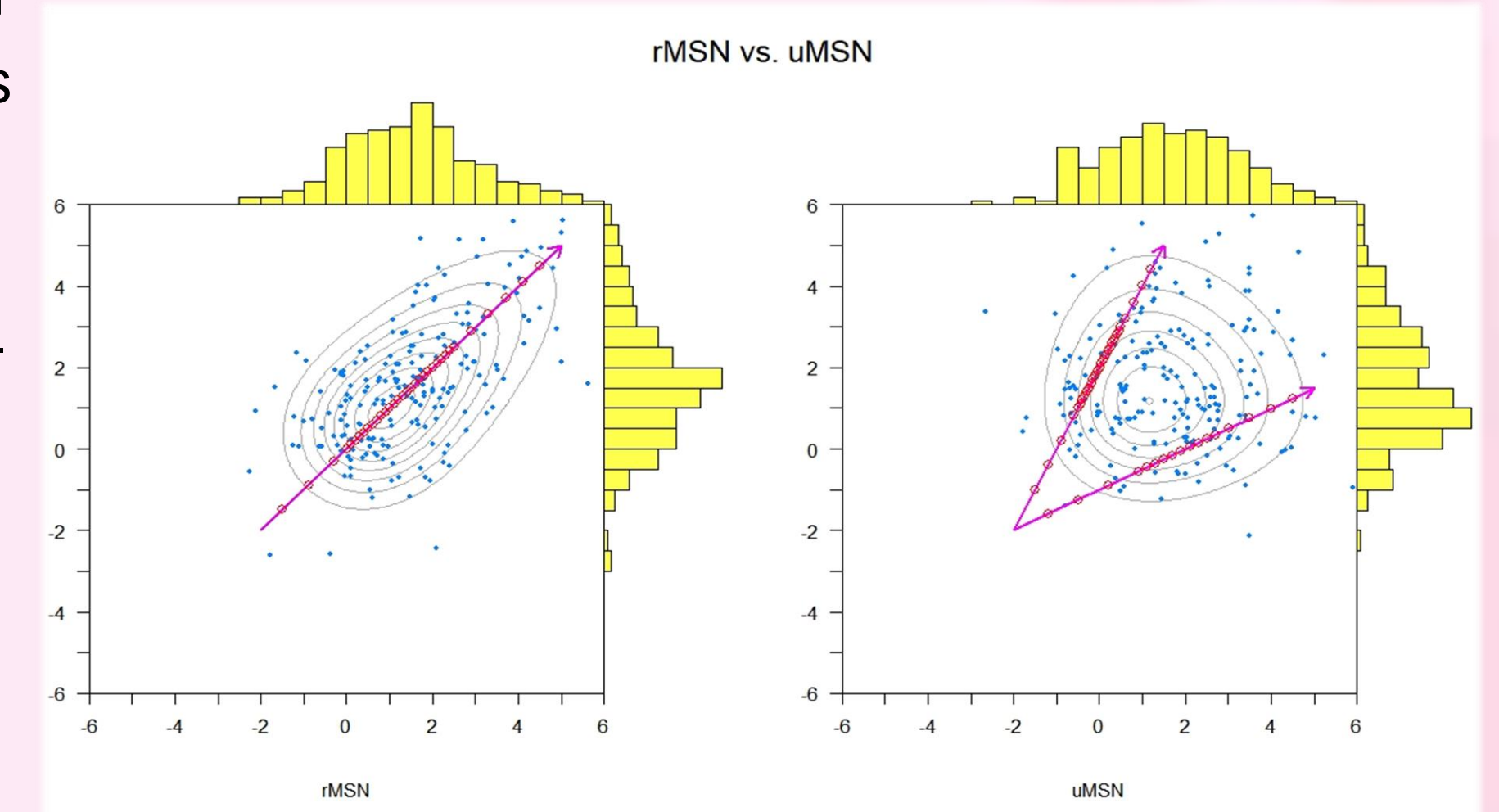
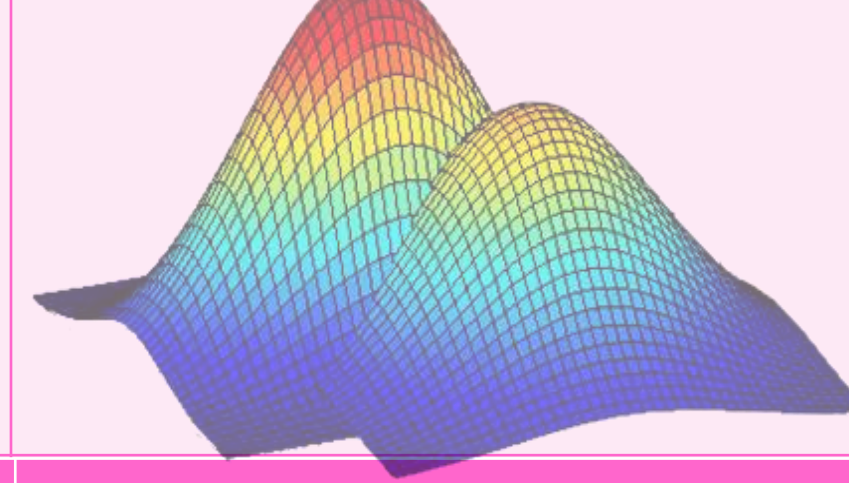
Experimental results demonstrate that the proposed MuSNFA model may outperform MFA and MrSNFA on providing a better fit, improved classification performance, and more accurate prediction for missing values.

Future research can expand the capabilities of the current method to analyze the data with multimodality, multiple skewness, missing values and censored responses simultaneously, see Lin et al. (2018) and Lin and Wang (2020) in the context of multivariate linear mixed models.

Motivation

Unlike the rMSN distribution, the uMSN distribution is able to capture skewness in more than one direction.

Aim: propose a skew extension of MFA based on the uMSN distribution and allows the handling of missing values.



Simulation & Real data

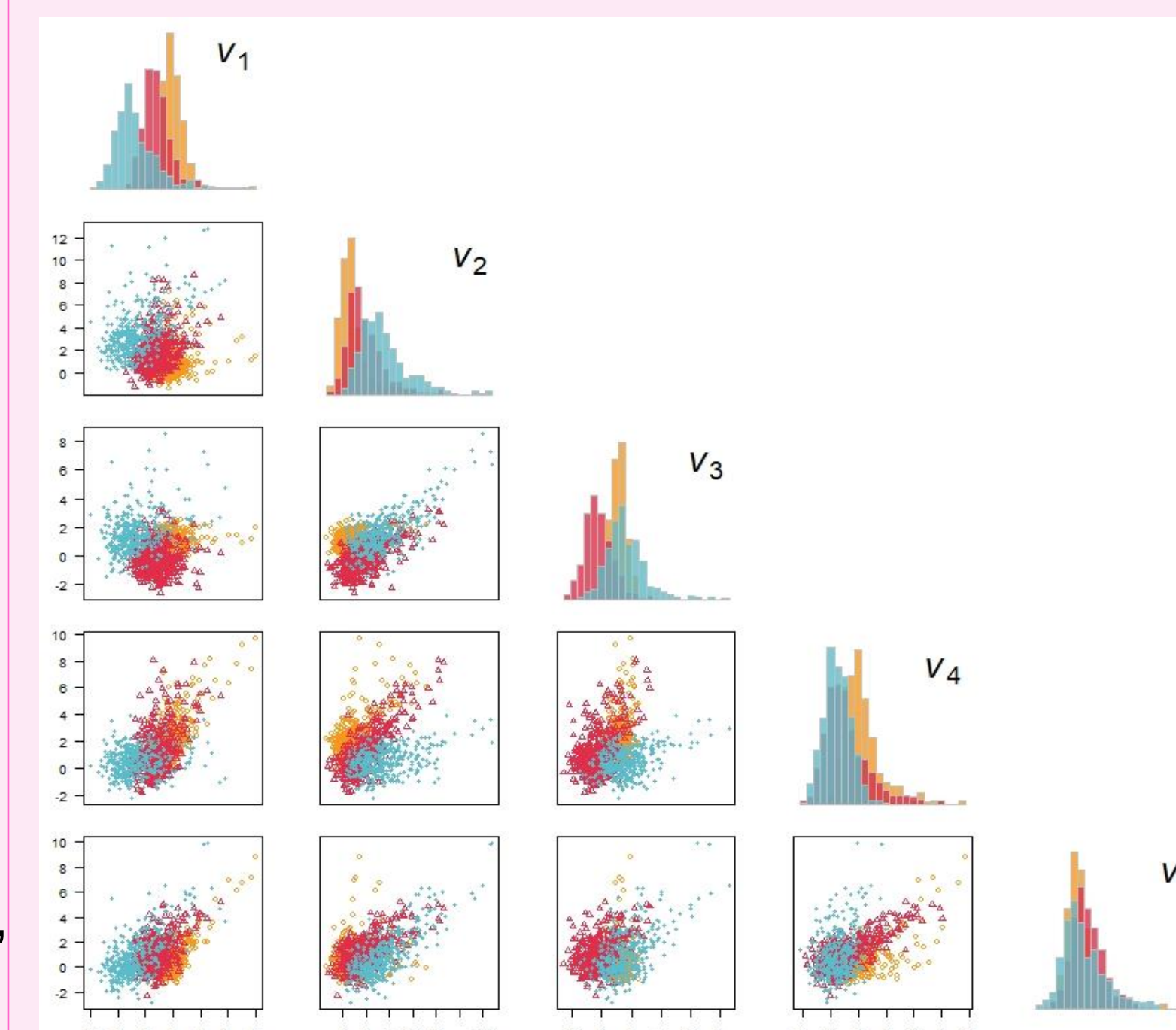
Simulation

A simulation study is undertaken to examine the performance of the proposed MuSNFA model in comparison with the MFA and MrSNFA approaches under varying proportions of **synthetic missing values**.

100 Monte Carlo samples are created from the **3-component MFA model with p=5 and q=2 factors**, while the latent factors are independently generated from the chi-square distribution with one degree of freedom χ_1^2 for yielding strong effects of skewness and kurtosis.

MuSNFA model outperforms the other two competing approaches in almost all trials.

The MFA model has the worst performance due to a lack of sufficient robustness against serious departure of normality assumption.



Missing Criterion rate	n = 300			n = 600			n = 1200		
	MFA	MrSNFA	MuSNFA	MFA	MrSNFA	MuSNFA	MFA	MrSNFA	MuSNFA
BIC	Mean 4790.81	4756.70	4735.59	8958.33	8828.85	8782.34	17664.59	17352.88	17243.15
	Freq 4	14	82	0	4	96	0	6	94
	Mean 4902.15	4864.91	4834.12	9246.50	9112.59	9022.93	18265.53	17944.62	17717.26
ICL	Freq 6	11	83	0	6	94	0	2	98
	Mean	0.834	0.842	0.870	0.834	0.858	0.886	0.832	0.873
	Freq	25	19	56	12	25	63	8	5
CCR	Mean	0.623	0.634	0.689	0.617	0.646	0.710	0.617	0.675
	Freq	23	17	60	12	24	64	8	3
ARI	Mean	1.184	1.139	1.112	1.183	1.151	1.130	1.142	1.097
	Freq	14	35	51	6	28	66	2	16
MSE	Mean	3757.54	3702.84	3693.55	7281.53	7134.67	7113.46	14331.74	13997.08
	Freq	0	16	84	0	12	88	0	9
BIC	Mean	3972.24	3882.21	3871.68	7747.61	7552.67	7513.12	15312.47	14892.26
	Freq	0	24	76	0	20	80	0	7
ICL	Mean	0.781	0.745	0.766	0.803	0.759	0.803	0.813	0.782
	Freq	49	16	35	34	5	61	28	12
CCR	Mean	0.475	0.439	0.473	0.508	0.451	0.523	0.526	0.486
	Freq	47	16	37	32	5	63	25	14
ARI	Mean	1.529	1.485	1.457	1.432	1.393	1.379	1.419	1.353
	Freq	16	29	55	15	26	59	1	28
MSE	Mean	1.529	1.485	1.457	1.432	1.393	1.379	1.419	1.353
	Freq	16	29	55	15	26	59	1	28
	MSE	Mean 1.529	1.485	1.457	1.432	1.393	1.379	1.419	1.353

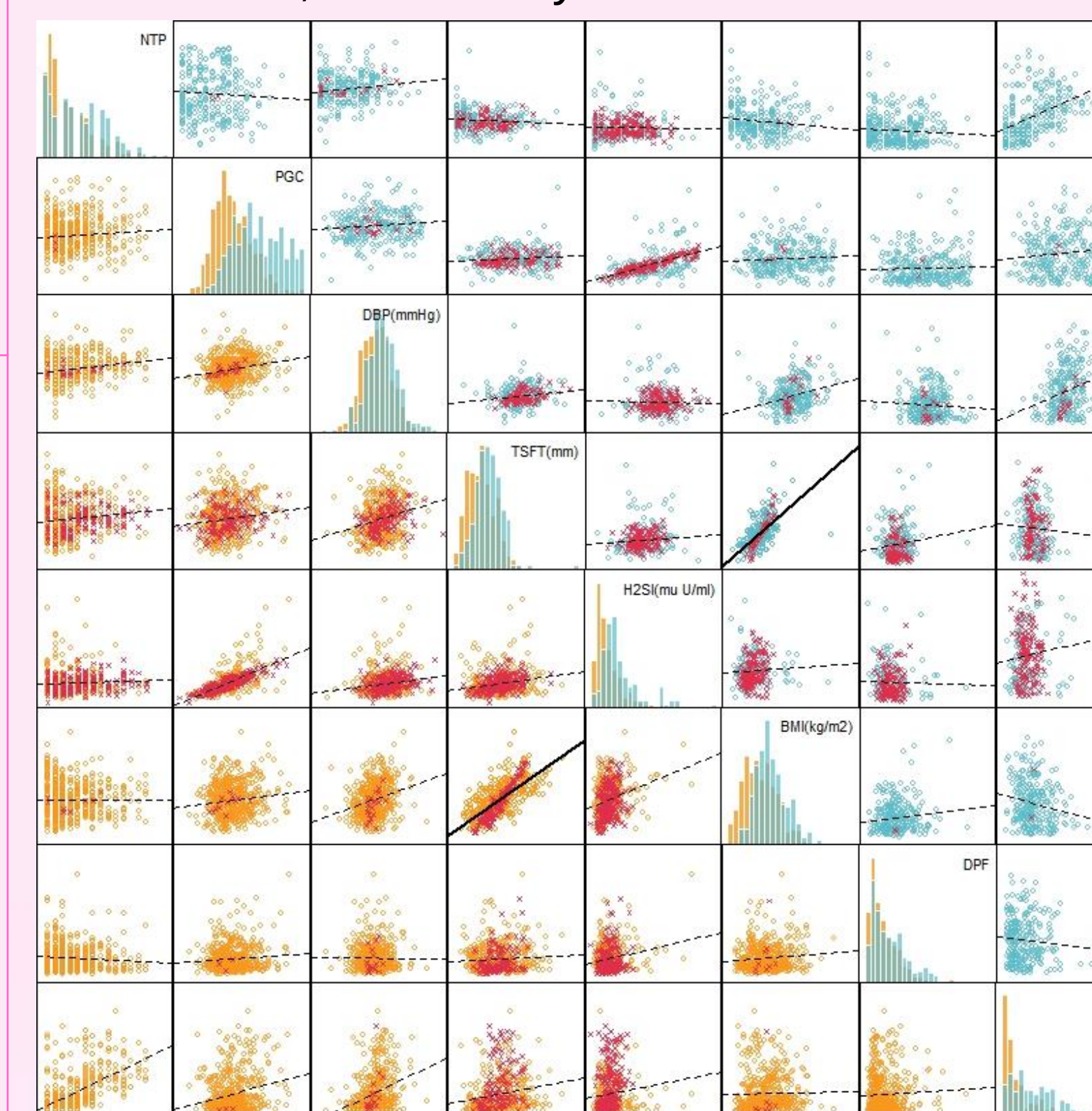
Pima Indians Diabetes Data Set (PIMA data)

There are ($p=8$) attributes measured on 500 **non-diabetes** and 278 **diabetes** female patients.

We consider the fitting of 2-component MFA, MrSNFA and MuSNFA models to this dataset with q ranging from 1-4.

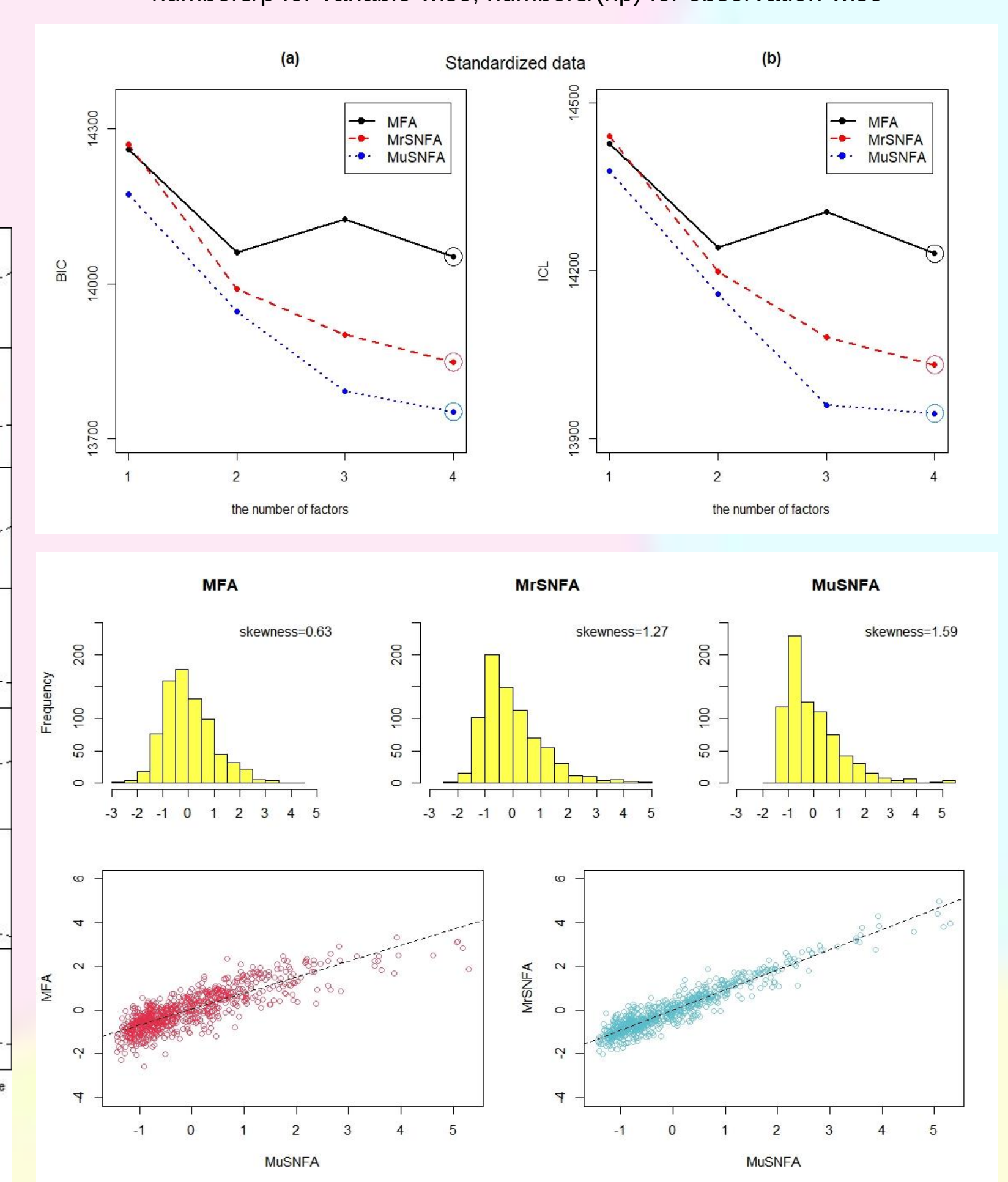
The **factor scores** predicted by the three approaches are all positively skewed, but the scattering patterns somewhat different.

The **MuSNFA** offer the strongest degree of skewness, followed by MrSNFA and MFA.



Number of missing items	Patients					NA-patient	NA-var	NA-obs
	0	1	2	3	4			
Non-diabetes	262	95	124	13	6	500	238	5
	(52.40)	(19.00)	(24.80)	(2.60)	(1.20)	(0.00)	(65.10)	(30.99)
Diabetes	130	47	75	15	1	0	268	138
	(48.50)	(17.54)	(27.99)	(5.60)	(0.37)	(0.00)	(34.90)	(17.97)
Total	392	142	199	28	7	500	506	193
	(51.04)	(18.49)	(25.91)	(3.65)	(0.91)	(0.00)	(48.96)	(10.61)

The percentages (%) are listed in parentheses (numbers/n for patient-wise; numbers/p for variable-wise; numbers/(np) for observation-wise



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