

A Probabilistic Distance Clustering Algorithm Using Gaussian and Student-*t* Multivariate Density Distributions

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Abstract

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8 A new dissimilarity measure for cluster analysis is presented and used in the context of probabilistic distance (PD) cluster-9 ing. The basic assumption of PD-clustering is that for each unit, the product between the probability of the unit belonging to 10 a cluster and the distance between the unit and the cluster is constant. This constant is a measure of the classifiability of the 11 point, and the sum of the constant over units is called joint distance function (JDF). The parameters that minimize the JDF 12 maximize the classifiability of the units. The new dissimilarity measure is based on the use of symmetric density functions 13 and allows the method to find clusters characterized by different variances and correlation among variables. The multivariate 14 Gaussian and the multivariate Student-t distributions have been used, outperforming classical PD clustering, and its variation 15 PD clustering adjusted for cluster size, on simulated and real datasets.

¹⁶ Keywords Cluster analysis · PD-clustering · Multivariate distributions · Dissimilarity measures

¹⁷ Introduction

18 Cluster analysis refers to a wide range of numerical meth-19 ods aiming to find distinct groups of homogeneous units. 20 Clustering in two or three dimensions is a natural task that 21 humans can often do visually; however, machine approaches 22 are needed for all but such low dimensions. We focus on 23 partitioning clustering methods; given a number of clus-24 ters K, partitioning methods assign units to the K clusters 25 optimizing a given criterion. These methods are generally 26 divided into not model-based and model-based, according 27 to the distributional assumptions. Model-based clustering or 28 finite mixture model clustering assumes that the population 29 probability density function is a convex linear combination 30 of a finite number of density functions; accordingly, they are 31 very well suited to clustering problems. A variety of meth-32 ods and algorithms have been proposed for finite mixture

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model parameter estimation. The most widely used strategy is to find the parameters that maximize the complete-data likelihood function using the expectation-maximization (EM) algorithm, which was proposed in 1977 [10] building on prior work (e.g., [5, 7, 23, 24]). The non-model-based methods generally optimize a criterion based on distance or dissimilarity measures. Different dissimilarity measures can be used based on the type of data, in this paper we focus on continuous data.

Formally, let us consider an $n \times J$ data matrix **X**, with generic row vector $\mathbf{x}_i = (x_{i1}, \dots, x_{iJ})$. Partitioning algorithms aim to find a set of *K* clusters, \mathcal{C}_k , with $k = 1, \dots, K$, such that the elements inside a cluster are homogeneous and $\mathcal{C}_1 \cup \mathcal{C}_2 \cup \dots \cup \mathcal{C}_K = X$. If, for any pair $\{k, k'\} \in 1, \dots, K$, $\mathcal{C}_k \cap \mathcal{C}_{k'} = \emptyset$, then the clustering technique is called hard or crisp, otherwise it is called fuzzy or soft. In the latter case, each unit can belong to more than one cluster with certain membership degree.

The most frequently used non-model-based methods for continuous data are k-means [20] and its fuzzy analogue c-means [4], which minimize the sum of the within groups sum of squares over all variables. In spite of their simplicity, the optimal solution can only be found applying an iterative intuitively reasonable procedure. More recently, [3] proposed probabilistic distance (PD) clustering, a distribution free fuzzy clustering technique (i.e., non-model-based),

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where the membership degree is defined as heuristic proba-59 bility. PD clustering optimization problems represents a spe-60 cial case of the Weber-Fermat's problem, when the number 61 62 of the 'attraction points' is greater or equal to three, see [16] among others. In this framework, PD clustering assumes 63 that the product of the probability of a point belonging to 64 a cluster and the distance of the point from the center of 65 the cluster is constant, and this constant is a measure of the 66 classificability of the point. The method obtains the centers 67 that maximize the classificability of all the points. A newer 68 version of the algorithm that considers clusters of different 69 size, PDQ-clustering, was proposed by [14] and an extension 70 for high-dimensional data was proposed by [35, 36]. 71

Generally, non-model-based clustering techniques are only 72 based on the distances between the points and the centers: 73 therefore, they do not take into account the shape and the size of 74 the clusters. Accordingly, these techniques may fail when clus-75 ters are either non-spherical or spherical with different radii. To 76 77 overcome this issue we propose a new dissimilarity measure based on symmetric density functions that have the advantage 78 of considering the variability and the correlation among the 79 80 variables. We use two different density functions, the multivariate Gaussian and the multivariate Student-t, but it could 81 be extended to other symmetric densities. We then integrate 82 this measure with PD-clustering and obtain new more flexible 83 clustering techniques. Preliminary results can be found in [29]. 84

After a background section on PD-clustering and PDQclustering, Sect. "Background", we introduce the new dissimilarity measure and the new techniques, Sect. "Flexible Extensions of PD-Clustering". We then compare them with some model-based and distance-based algorithms on simulated and real datasets, Sect. Empirical Evidence from Simulated and Real Data.

92 Background

In this section we briefly introduce PD-clustering [3], a
distance-based soft clustering algorithm, and its extension,
PD-clustering adjusted for cluster size [14].

96 Probabilistic Distance Clustering

Ben-Israel and Iyigun [3] proposed a non-hierarchical distance-based clustering method, called probabilistic distance
(PD) clustering. They then extended the method to account for clusters of different size, i.e., PDQ [14]. Tortora et al.
[35] proposed a factor version of the method to deal with high-dimensional data. Recently, [29] further extended the method to include more flexibility.

In PD-clustering, the number of clusters *K* is assumed to be a priori known, and a wide review on how to choose *K* can be found in [8]. Given some random centers, the probability of any point belonging to a cluster is assumed to be inversely 107 proportional to the distance from the center of that cluster [13]. 108 Suppose we have a data matrix **X** with *N* units and *J* variables, 109 and consider K (non-empty) clusters. PD-clustering is based 110 on two quantities: the distance of each data point \boldsymbol{x} , from each 111 cluster centre \mathbf{c}_k , denoted $d(\mathbf{x}_i, \mathbf{c}_k)$, and the probability of each 112 point belonging to a cluster, i.e., $p(\mathbf{x}_i, \mathbf{c}_k)$, for $k = 1, \dots, K$ and 113 i = 1, ..., N. 114

For convenience, define $p_{ik} := p(\mathbf{x}_i, \mathbf{c}_k)$ and $d_{ik} := d(\mathbf{x}_i, \mathbf{c}_k)$. 115 PD-clustering is based on the principle that the product of 116 the distances and the probabilities is a constant depending 117 only on \mathbf{x}_i [3]. Denoting this constant as $F(\mathbf{x}_i)$, we can write 118 this principle as 119

$$p_{ik}d_{ik} = F(\boldsymbol{x}_i),\tag{1}$$

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where $F(\mathbf{x}_i)$ depends only on \mathbf{x}_i , i.e., $F(\mathbf{x}_i)$ does not depend 122 on the cluster k. As the distance from the cluster centre 123 decreases, the probability of the point belonging to the 124 cluster increases. The quantity $F(\mathbf{x}_i)$ is a measure of the 125 closeness of x_i to the cluster centres, and it determines the 126 classificability of the point x_i with respect to the centres c_k , 127 for k = 1, ..., K. The smaller the $F(\mathbf{x}_i)$ value, the higher the 128 probability of the point belonging to one cluster. If all of the 129 distances between the point x_i and the centers of the clusters 130 are equal to d_i , then $F(\mathbf{x}_i) = d_i/K$ and all of the probabilities 131 of belonging to each cluster are equal, i.e., $p_{ik} = 1/K$. The 132 sum of $F(\mathbf{x}_i)$ over *i* is called joint distance function (JDF). 133 Starting from (1), it is possible to compute p_{ik} , i.e., 134

$$p_{ik} = \frac{\prod_{m \neq k} d_{im}}{\sum_{m=1}^{K} \prod_{r \neq m} d_{ir}},$$
(2)
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for k = 1, ..., K, and i = 1, ..., N. The whole clustering problem consists in the identification of the centers that minimize the JDF:

$$JDF = \sum_{i=1}^{n} \sum_{k=1}^{K} d_{ik} p_{ik}.$$
 (3)

Extensive details on PD clustering are given in [3], who suggest using p^2 in (3) because it is a smoothed version of the problem. It follows that the optimized functions become

$$JDF = \sum_{i=1}^{n} \sum_{k=1}^{K} d_{ik} p_{ik}^{2},$$
(4)

and the centers can be computed as

$$c_k = \frac{\sum_{i=1}^N u_{ik} \boldsymbol{x}_i}{\sum_{i=1}^N},\tag{5}$$

$$\sum_{j=1}^{n} u_{jk}$$

with
$$u_{ik} = p_{ik}^2 / d_{ik}$$
.

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It is worth noting that the function p_{ik} respects all neces-151 sary conditions to be a probability and yet no assumptions 152 are made on the distribution of this function; further, p_{ik} can 153 only be computed given x_i and for every c_k [28]. Following 154 [13], we refer to p_{ik} as subjective probabilities, which are 155 based on degree of belief (see [2]). 156

PD Clustering Adjusted for Cluster Size 157

The probabilities obtained using 1 do not consider the clus-158 ter size, and the algorithm tends to fail when clusters are 159 unbalanced. Moreover, the resulting clusters have similar 160 variance and covariance matrices. To overcome these issues 161 [14] proposed PD-clustering adjusted for cluster size (PDQ). 162 They assume that 163

$$\frac{p_{ik}^2 d_{ik}}{q_k} = F(\boldsymbol{x}_i),\tag{6}$$

where q_k is the cluster size, with the constraint that 166 $\sum_{k=1}^{K} q_k = N$. The p_{ik} can then be computed via 167

$$p_{ik} = \frac{\prod_{m \neq k} d_{im}/q_m}{\sum_{m=1}^{K} \prod_{r \neq m} d_{ir}/q_r}.$$
(7)

The cluster size is considered a variable, the value of q_k that 170 minimizes (6) is 171

¹⁷²
$$q_{k} = N \frac{\left(\sum_{i=1}^{N} d_{ik} p_{ik}^{2}\right)^{1/2}}{\sum_{k=1}^{K} \left(\sum_{i=1}^{N} d_{ik} p_{ik}^{2}\right)^{1/2}},$$
(8)

for k = 1, ..., K - 1, and 174

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 $q_K = N - \sum_{i=1}^{K-1} q_k.$ 176

Flexible Extensions of PD-Clustering 177

Gaussian PD-Clustering 178

The PDQ algorithm can detect clusters of different size and 179 with different within-cluster variability; however, it can still 180 fail at detecting the clustering partition when variables are 181 correlated or when there are outliers in the data. To over-182 come these issues we proposed a new dissimilarity measure 183 based on a density function. Let $M_k = \max\{f(\mathbf{x}_i; \boldsymbol{\mu}_k, \boldsymbol{\theta}_k)\}$ and 184 define the quantity 185

$$\delta_{ik} = \log \left(M_k f(\boldsymbol{x}_i; \boldsymbol{\mu}_k, \boldsymbol{\theta}_k)^{-1} \right),$$
(9)
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which is a dissimilarity measure where $f(\mathbf{x}_i; \boldsymbol{\mu}_k, \boldsymbol{\theta}_k)$ is a sym-188 metric unimodal density function with location parameter μ_k 189 and parameter vector $\boldsymbol{\theta}_k$. See appendix for the proof. 190

Recall that the density of a multivariate Gaussian distribution is

$$\phi(\mathbf{x}_{i};\boldsymbol{\mu},\boldsymbol{\Sigma}) = \frac{1}{(2\pi)^{\frac{J}{2}}} |\boldsymbol{\Sigma}|^{-\frac{1}{2}} \exp\left\{-\frac{1}{2}(\mathbf{x}-\boldsymbol{\mu})'\boldsymbol{\Sigma}^{-1}(\mathbf{x}-\boldsymbol{\mu})\right\}, \qquad ^{193}$$

194 (10)

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and define $\phi_{ik} := \phi(\mathbf{x}_i; \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k)$, where $k = 1, \dots, K$. Using 195 (10) in (9) and the result in (6), the JDF becomes 196

$$JDF = \sum_{i=1}^{n} \sum_{k=1}^{K} \frac{p_{ik}^{2}}{q_{k}} \log(M_{k}) + \sum_{i=1}^{n} \sum_{k=1}^{K} \frac{1}{2} \frac{p_{ik}^{2}}{q_{k}} \log((2\pi)^{J} |\boldsymbol{\Sigma}_{k}|) + \sum_{i=1}^{n} \sum_{k=1}^{K} \frac{1}{2} \frac{p_{ik}^{2}}{q_{k}} (\boldsymbol{x}_{i} - \boldsymbol{\mu}_{k})' \boldsymbol{\Sigma}_{k}^{-1} (\boldsymbol{x}_{i} - \boldsymbol{\mu}_{k}).$$
(11) 198

This technique is called Gaussian PD-clustering, GPDC, and 199 it offers many advantages when compared to PD-clustering. 200 The new dissimilarity measure already takes into account 201 the impact of different within cluster variances and the cor-202 relation among variables. 203

The clustering problem now consists in the estimation of μ_{μ} 204 and Σ_k , with k = 1, ..., K, that minimize (11). A differentia-205 tion procedure leads to these estimates. An iterative algorithm 206 is then used to compute the belonging probabilities and update 207 the parameter estimates. More specifically, differentiating (11)208 with respect to $\boldsymbol{\mu}_k$ gives 209

$$\frac{\partial \text{JDF}}{\partial \boldsymbol{\mu}_k} = -\frac{1}{2} \sum_{i=1}^n \frac{p_{ik}^2}{q_k} \boldsymbol{\Sigma}_k^{-1} (\boldsymbol{x}_i - \boldsymbol{\mu}_k).$$
(12)

Setting (12) equal to zero and solving for μ_k gives

$$\boldsymbol{\mu}_{k} = \frac{\sum_{i=1}^{n} p_{ik}^{2} \boldsymbol{x}_{i}}{\sum_{i=1}^{n} p_{ik}^{2}}$$
(13)

Now, differentiating (11) with respect to Σ_k gives

$$\frac{\partial \text{JDF}}{\partial \boldsymbol{\Sigma}_{k}} = \sum_{i=1}^{n} \frac{1}{2} \frac{p_{ik}^{2}}{q_{k}} \boldsymbol{\Sigma}_{k}^{-1} - \boldsymbol{\Sigma}_{k}^{-1} \sum_{i=1}^{n} \frac{1}{2} (\boldsymbol{x}_{i} - \boldsymbol{\mu}_{k}) (\boldsymbol{x}_{i} - \boldsymbol{\mu}_{k})' \frac{p_{ik}^{2}}{q_{k}} \boldsymbol{\Sigma}_{k}^{-1} = \frac{1}{2} \boldsymbol{\Sigma}_{k}^{-1} \left[\sum_{i=1}^{n} \frac{p_{ik}^{2}}{q_{k}} - \sum_{i=1}^{n} (\boldsymbol{x}_{i} - \boldsymbol{\mu}_{k}) (\boldsymbol{x}_{i} - \boldsymbol{\mu}_{k})' p_{ik}^{2} \boldsymbol{\Sigma}_{k}^{-1} \right].$$
(14) 217

Setting (14) equal to zero and solving for Σ_k gives

$$\Sigma_{k} = \frac{\sum_{i=1}^{n} (\mathbf{x}_{i} - \boldsymbol{\mu}_{k}) (\mathbf{x}_{i} - \boldsymbol{\mu}_{k})' p_{ik}^{2}}{\sum_{i=1}^{n} p_{ik}^{2}}.$$
(15)

It follows that, at generic iteration (t + 1), the parameters that 221 minimize the (11) are: 222

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$$\boldsymbol{\mu}_{k}^{(t+1)} = \frac{\sum_{i=1}^{n} p_{ik}^{2} \boldsymbol{x}_{i}}{\sum_{i=1}^{n} p_{ik}^{2}},$$
(16)

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Author Proof

Our iterative procedure for Gaussian mixture model-based clustering parameter estimation can be summarized as follows:

 $\boldsymbol{\Sigma}_{k}^{(t+1)} = \frac{\sum_{i=1}^{n} (\boldsymbol{x}_{i} - \boldsymbol{\mu}_{k}^{(t+1)}) (\boldsymbol{x}_{i} - \boldsymbol{\mu}_{k}^{(t+1)})' \boldsymbol{p}_{ik}^{2}}{\sum_{i=1}^{n} \boldsymbol{p}_{ik}^{2}}.$

Algo	rithm 1 GPDC	
1: p	rocedure GPDC(X , <i>K</i>)	
2:	for $k = 1, \ldots, K$ do	▷ Initialization
3:	Random initialization of $\boldsymbol{\mu}_k$	
4:	set $\boldsymbol{\Sigma}_k$ as identity matrix	
5:	while $\boldsymbol{\mu}_k^{(t)} eq \boldsymbol{\mu}_k^{(t+1)}$ do	⊳ Core
6:	for $k = 1, \ldots, K$ do	
7:	update q_k according to (8)	
8:	update p_{ik} according to (7)	
9:	update $\boldsymbol{\mu}_k$ according to (16)	
10:	update $\boldsymbol{\Sigma}_k$ according to (17)	
11:	return $p_{ik}, \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k$	

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Generalization to a Multivariate Student-t 231 Distribution 232

The same procedure can be generalized to any symmet-233 ric distribution. In this subsection we use the multivariate 234 Student-t distribution, generating an algorithm identified as 235 Student-t PD-Clustering (TPDC). TPDC can detect clusters 236 characterized by heavy tails; furthermore, the Student-t dis-237 tribution has been often used on datasets characterized by 238 outliers [17]. Now, replace (10) with a multivariate Student-t239 distribution, i.e., 240

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$$f(\boldsymbol{x}, \boldsymbol{\mu}, \boldsymbol{\Sigma}, \boldsymbol{\nu}) = \frac{\Gamma\left(\frac{\nu+J}{2}\right) |\boldsymbol{\Sigma}|^{-\frac{1}{2}}}{(\pi\nu)^{\frac{1}{2}J} \Gamma\left(\frac{\nu}{2}\right) \left\{1 + \frac{\delta(\boldsymbol{x}, \boldsymbol{\mu}, \boldsymbol{\Sigma})}{\nu}\right\}^{\frac{1}{2}(\nu+J)}},$$
(18)

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where $\delta(\mathbf{x}, \boldsymbol{\mu}, \boldsymbol{\Sigma}) = (\mathbf{x} - \boldsymbol{\mu})' \boldsymbol{\Sigma}^{-1} (\mathbf{x} - \boldsymbol{\mu})$, and proceed as in 243 Sect. 3.1 Then, the JDF becomes: 244

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$$JDF = \sum_{i=1}^{n} \sum_{k=1}^{K} \frac{p_{ik}^{2}}{q_{k}} \log(M_{k}) + \sum_{i=1}^{n} \sum_{k=1}^{K} \frac{p_{ik}^{2}}{q_{k}} \left[-\log\left\{\Gamma\left(\frac{v_{k}+J}{2}\right)|\boldsymbol{\Sigma}_{k}|^{-\frac{1}{2}}\right\} \right] + \sum_{i=1}^{n} \sum_{k=1}^{K} \frac{p_{ik}^{2}}{q_{k}} \log\left\{\pi v_{k}\right)^{\frac{j}{2}} \Gamma\left(\frac{v_{k}}{2}\right) \left(1 + \frac{\delta(\boldsymbol{x}_{i}, \boldsymbol{\mu}_{k}, \boldsymbol{\Sigma}_{k})}{v_{k}}\right)^{\frac{v_{k}+J}{2}} \right\}.$$
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(19)

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The parameters that optimize (19) can be found by differ-247 entiating with respect to μ_k , Σ_k , and v_k , respectively. Spe-248 cifically, at a generic iteration (t + 1), the parameters that 249 minimize (19) are: 250

$$\boldsymbol{\mu}_{k}^{(t+1)} = \frac{\sum_{i=1}^{n} w_{ik} \boldsymbol{x}_{i}}{\sum_{i=1}^{n} w_{ik}},$$
(20)

with
$$w_{ik} = p_{ik}^2 / [v_k^{(t)} + \delta(\mathbf{x}_i, \boldsymbol{\mu}_k^{(t)}, \boldsymbol{\Sigma}_k^{(t)})],$$
 253

$$\boldsymbol{\Sigma}_{k}^{(t+1)} = \frac{\sum_{i=1}^{n} p_{ik}^{2} (\boldsymbol{x}_{i} - \boldsymbol{\mu}_{k}^{(t+1)}) (\boldsymbol{x}_{i} - \boldsymbol{\mu}_{k}^{(t+1)})' \boldsymbol{s}_{ik}}{\sum_{i=1}^{n} p_{ik}^{2}}, \qquad (21)$$

with $s_{ik} = (v_k^{(t)} + J) / [v_k^{(t)} + \delta(\mathbf{x}_i, \boldsymbol{\mu}_k^{(t+1)}, \boldsymbol{\Sigma}_k^{(t)})]$, and the degrees of freedom update $v_k^{(t+1)}$ is the solution to the following 256 257 equation: 258

$$\sum_{i=1}^{n} p_{ik}^{2} \left[\psi\left(\frac{v_{k}}{2}\right) - \psi\left(\frac{v_{k}+J}{2}\right) + \frac{J}{2v_{k}} \right]$$

$$+ \sum_{i=1}^{n} p_{ik}^{2} \left[\frac{1}{2} \log \left(1 + \frac{\delta\left(x_{i}, \mu_{k}^{(t+1)}, \Sigma_{k}^{(t+1)}\right)}{v_{k}^{(t)}} \right) \right]$$

$$- \frac{1}{2} \frac{v_{k}+J}{v_{k}} \sum_{i=1}^{n} p_{ik}^{2} \frac{\delta\left(x_{i}, \mu_{k}^{(t+1)}, \Sigma_{k}^{(t+1)}\right)}{v_{k}^{(t)} + \delta\left(x_{i}, \mu_{k}^{(t+1)}, \Sigma_{k}^{(t+1)}\right)} = 0,$$
(22)

where

(17)

$$\psi(v) = \left(\frac{1}{\Gamma(v)}\right) \frac{\delta \Gamma(v)}{\delta v}.$$
²⁶²

Our iterative algorithm can be summarized as follows:

Algo	Algorithm 2 TPDC						
1: p	rocedure TPDC(X , <i>K</i>)						
2:	for $k = 1,, K$ do	▷ Initialization					
3:	Random initialization of $\boldsymbol{\mu}_k$						
4:	set $\boldsymbol{\Sigma}_k$ as identity matrix						
5:	$v_k = 20$						
6:	while $oldsymbol{\mu}_k^{(t)} eq oldsymbol{\mu}_k^{(t+1)}$ do	⊳ Core					
7:	for $\vec{k} = 1, \dots, K$ do						
8:	update q_k according to (8)						
9:	update p_{ik} according to (7)						
10:	update $\boldsymbol{\mu}_k$ according to (20)						
11:	update $\boldsymbol{\Sigma}_k$ according to (21)						
12:	update v_k solving (22)						
13:	return p_{ik} , $\boldsymbol{\mu}_k$, $\boldsymbol{\Sigma}_k$, v_k						

Algorithm Details

All the proposed techniques require a random initialization. 267 Random starts can lead to unstable solutions, to avoid this 268

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problem the algorithms use multiple starts. Moreover, the 269 functions include the option to use PD-clustering or parti-270 tion around medoids (PAM; [15]) to start. As for many other 271 clustering techniques, the optimized function, the JDF in 272 (4), is not convex—not even quasi-convex—and may have 273 other stationary points. For a fixed value of Σ_k , the JDF is a 274 monotonically decreasing function, this guarantees that the function converges to a minimum, not necessarily a global minimum. The proposed techniques, GPDC and TPDC, introduce the estimate of Σ_k , giving much more flexibility, albeit the JDF is no longer monotonically decreasing. Using (9) in (4), we obtain

$$JDF = \sum_{i=1}^{n} \sum_{k=1}^{K} p_{ik}^{2} (\log M_{k} - \log \phi(\boldsymbol{x}_{i}; \boldsymbol{\mu}_{k}, \boldsymbol{\Sigma}_{k}))$$

with $M_k \ge \phi(\mathbf{x}_i; \boldsymbol{\mu}_k, \boldsymbol{\Sigma}_k)$. Therefore, for every $k = 1, \dots, K$, the function is upper-bounded for non-degenerate density functions. The convergence of the algorithm cannot depend on the JDF but is based on μ_k . The time complexity of the algorithm is comparable to the EM algorithm, both algorithms require the inversion and the determinant of a $J \times J$ 288 matrix, therefore, the time complexity is of $O(n^3 JK)$, where 289 *n* is the number of observations, *J* the number of variables, 290 and K the number of clusters. 291

Empirical Evidence from Simulated and Real 292 Data 293

The proposed algorithm has been evaluated on real and sim-294 ulated datasets. The simulated datasets have been used to 295 illustrate the ability of the algorithms to recover the param-296 eters of the distributions and to compare the new techniques 297 with some existing methods. In the following sessions we 298 used the software R [26], the functions for both GPDC and 299 TPDC are included in the R package FPDclustering 300 [37]. 301

Simulation Study 302

The same design was used twice, the first time each clus-303 ter was generated from a multivariate Gaussian distribution 304 with three variables and K = 3 clusters. The second time, 305 using a multivariate Student-t distribution with five degrees 306 of freedom, same number of variables and clusters. We set 307 the parameter using a four factor full factorial design. There 308 are two factors per each level, where the levels are 309

- Overlapping and not overlapping clusters 310
- Different number of elements per clusters 311
- Unitary variance and variance bigger than 1 312
- Uncorrelated and correlated variables _ 313

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Table 1 Model parameters used to generate the simulated datasets

Not overlapping	Overlapping
$\mu_1 = (0, 0, 0)'$	$\mu_1 = (0, 0, 0)'$
$\mu_2 = (-7, 7, 0)'$	$\mu_2 = (-4, 4, 0)'$
$\mu_3 = (-7, 0, 7)'$	$\mu_3 = (-4, 0, 4)'$
Option 1	Option 2
$n_1 = 200$	$n_1 = 200$
$n_2 = 300$	$n_2 = 100$
$n_3 = 100$	$n_3 = 300$
Unitary variance	bigger than 1
$\operatorname{diag}(\boldsymbol{\Sigma}_1) = 1$	$\operatorname{diag}(\boldsymbol{\Sigma}_1) = 1$
$\operatorname{diag}(\boldsymbol{\Sigma}_2) = 1$	$\operatorname{diag}(\boldsymbol{\Sigma}_2) = 16$
$\operatorname{diag}(\boldsymbol{\Sigma}_3) = 1$	$\operatorname{diag}(\boldsymbol{\Sigma}_3) = 2.25$
Not correlated	Correlated
$\boldsymbol{\Sigma}_{1} = \begin{pmatrix} 1 & 0 & 0 \\ 0 & 1 & 0 \\ 0 & 0 & 1 \end{pmatrix}$	$\boldsymbol{\Sigma}_{1} = \begin{pmatrix} 1 & 0 & 0 \\ 0 & 1 & 0 \\ 0 & 0 & 1 \end{pmatrix}$
$\boldsymbol{\Sigma}_2 = \begin{pmatrix} 1 & 0 & 0 \\ 0 & 1 & 0 \\ 0 & 0 & 1 \end{pmatrix}$	$\boldsymbol{\Sigma}_2 = \begin{pmatrix} 1 & -0.5 & -0.5 \\ -0.5 & 1 & 0.5 \\ -0.5 & 0.5 & 1 \end{pmatrix}$
$\boldsymbol{\Sigma}_{3} = \begin{pmatrix} 1 & 0 & 0 \\ 0 & 1 & 0 \\ 0 & 0 & 1 \end{pmatrix}$	$\boldsymbol{\Sigma}_3 = \begin{pmatrix} 1 & 0.7 & 0.7 \\ 0.7 & 1 & 0.7 \\ 0.7 & 0.7 & 1 \end{pmatrix}$

Table 1 shows the parameters used in the simulation study.

The datasets have been generated using the R package 316 mvtnorm [11]. Tables 5, 6, 7, 8, 9, 10, 11, 12 in Appen-317 dix B.2 show the true and the average estimated values of 318 the parameters obtained from 50 runs of the GPDC and 319 TPDC algorithms. For sake of space, comments are limited 320 to groups of scenarios. The factors that affect the estimates 321 the most are the change in variances and the amount of 322 overlap. Specifically, when data are simulated using mul-323 tivariate Gaussian distributions, in cases 5-8 and 13-16, 324 the variances are not homogeneous and the GPDC tends 325 to underestimate the bigger variances and overestimate the 326 smaller ones. The TPDC is less affected by this issue, i.e., 327 it underestimates some of the variances but the degrees of 328 freedom recover; however, in the two extreme scenarios, 329 8 and 16, it cannot recover the cluster structures. Similar 330 outcomes occur when data are simulated using a multivari-331 ate Student-t distribution; moreover, as expected on those 332 datasets, the GPDC tends to overestimate the variances and 333 TPDC tends to underestimate the variances and compensate 334 with the degrees of freedom. 335

On the same datasets we used the functions gpcm, option 336 VVV, of the R package mixture [6] for the Gaussian mix-337 ture models (GMM) and the function teigen, option 338 UUUU, of the homonymous R package [1] for the mixtures 339 of multivariate Student-t distributions (TMM). The k-means 340

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Table 2 Average ARI and standard deviation on 50 datasets per scenario

	Correlated	Unitary	Over-lapping	n_k	GPDC TPDC		2	PDQ		k-mea	ins	GMM	[TMM		
		Variance			ARI	SD	ARI	SD	ARI	SD	ARI	SD	ARI	SD	ARI	SD
Mul	tivariate Gauss	sian distribu	tion													
1	No	Yes	No	Op 1	1.00	0.00	1.00	0.00	1.00	0.00	0.91	0.20	0.99	0.06	1.00	0.00
2	No	Yes	no	Op 2	1.00	0.00	1.00	0.00	1.00	0.00	0.92	0.18	0.96	0.12	1.00	0.00
3	No	Yes	Yes	Op 1	0.97	0.01	0.98	0.01	0.98	0.01	0.99	0.01	0.99	0.01	0.99	0.01
4	No	Yes	Yes	Op 2	0.97	0.01	0.98	0.01	0.98	0.01	0.98	0.01	0.99	0.01	0.99	0.01
5	No	No	No	Op 1	0.80	0.03	0.84	0.03	0.74	0.04	0.61	0.08	0.93	0.07	0.93	0.02
6	No	No	No	Op 2	0.94	0.02	0.93	0.02	0.74	0.05	0.89	0.09	0.97	0.01	0.97	0.01
7	No	No	Yes	Op 1	0.59	0.04	0.60	0.05	0.30	0.04	0.35	0.06	0.76	0.08	0.76	0.04
8	No	No	Yes	Op 2	0.56	0.12	0.50	0.09	0.60	0.06	0.75	0.08	0.87	0.02	0.87	0.02
9	Yes	Yes	No	Op 1	1.00	0.00	1.00	0.00	1.00	0.00	0.86	0.22	0.95	0.14	1.00	0.00
10	Yes	Yes	No	Op 2	1.00	0.00	1.00	0.00	1.00	0.00	0.91	0.19	0.91	0.18	1.00	0.00
11	Yes	Yes	Yes	Op 1	0.97	0.02	0.98	0.02	0.98	0.01	0.93	0.15	0.96	0.09	0.99	0.01
12	Yes	yes	Yes	Op 2	0.98	0.02	0.99	0.01	0.98	0.01	0.98	0.07	0.98	0.08	1.00	0.00
13	Yes	No	No	Op 1	0.76	0.08	0.90	0.03	0.67	0.05	0.52	0.08	0.95	0.07	0.96	0.01
14	Yes	No	No	Op 2	0.97	0.02	0.96	0.02	0.77	0.06	0.91	0.02	0.99	0.01	0.99	0.01
15	Yes	No	Yes	Op 1	0.44	0.08	0.45	0.08	0.33	0.05	0.32	0.04	0.82	0.09	0.83	0.03
16	Yes	No	Yes	Op 2	0.54	0.07	0.45	0.08	0.54	0.08	0.80	0.10	0.94	0.01	0.94	0.01
Mult	tivariate Stude	nt-t distribut	ion, 5 df													
1	no	Yes	No	Op 1	0.98	0.01	0.99	0.01	0.99	0.01	0.98	0.07	0.97	0.06	0.99	0.01
2	No	Yes	No	Op 2	0.99	0.01	0.99	0.01	0.99	0.01	0.93	0.17	0.95	0.11	0.99	0.01
3	No	Yes	Yes	Op 1	0.88	0.02	0.89	0.02	0.91	0.02	0.90	0.02	0.90	0.02	0.91	0.02
4	No	Yes	Yes	Op 2	0.88	0.03	0.89	0.02	0.90	0.02	0.89	0.07	0.89	0.05	0.90	0.02
5	No	No	No	Op 1	0.92	0.02	0.94	0.02	0.94	0.02	0.89	0.07	0.93	0.07	0.95	0.01
6	No	No	No	Op 2	0.96	0.02	0.96	0.01	0.93	0.02	0.96	0.01	0.91	0.06	0.96	0.01
7	No	No	Yes	Op 1	0.69	0.04	0.71	0.04	0.70	0.04	0.63	0.04	0.56	0.11	0.76	0.03
8	No	No	Yes	Op 2	0.76	0.03	0.76	0.03	0.62	0.05	0.81	0.03	0.66	0.06	0.78	0.05
9	Yes	Yes	No	Op 1	0.99	0.01	0.99	0.01	0.99	0.01	0.90	0.19	0.92	0.17	0.99	0.01
10	Yes	Yes	No	Op 2	0.99	0.01	0.99	0.01	0.99	0.01	0.87	0.22	0.93	0.14	0.99	0.01
11	Yes	Yes	Yes	Op 1	0.87	0.03	0.88	0.03	0.90	0.02	0.83	0.16	0.90	0.07	0.93	0.02
12	Yes	Yes	Yes	Op 2	0.87	0.05	0.89	0.04	0.93	0.02	0.92	0.11	0.94	0.06	0.96	0.01
13	Yes	No	No	Op 1	0.93	0.02	0.95	0.02	0.92	0.02	0.79	0.17	0.93	0.10	0.96	0.01
14	Yes	No	No	Op 2	0.97	0.01	0.98	0.01	0.95	0.02	0.97	0.01	0.96	0.03	0.98	0.01
15	Yes	No	Yes	Op 1	0.56	0.10	0.59	0.08	0.65	0.04	0.55	0.08	0.72	0.13	0.82	0.03
16	Yes	No	Yes	Op 2	0.49	0.29	0.51	0.32	0.56	0.31	0.67	0.37	0.64	0.35	0.71	0.38

algorithm is part of the stats package [27], and the PDQclust function for PDQ clustering is part of the FPDclustering package [37].

To compare the clustering performance of the methods 344 we used the adjusted Rand index (ARI) [12]. It compares 345 predicted classifications with true classes. The ARI corrects 346 the Rand index [30] for chance, its expected value under ran-347 dom classification is 0, and it takes a value of 1 when there 348 is perfect class agreement. Steinley [31] gives guidelines for 349 350 interpreting ARI values. Table 2 shows the average ARI and the standard deviation on 50 runs for each algorithm. 351

As pointed out in the previous sections, GPDC and TPDC are framed in a non-parametric view; however, to evaluate the performance we compare them with the GMM 354 and TMM. The performance is not expected to be better 355 than those techniques, although in most scenarios GPDC 356 and TPDC perform as well as finite mixture models. As 357 expected, k-means results are impacted by correlations and 358 not homogeneous variances. PDQ cannot recover the correct 359 clustering partition in case of overlapping and not homoge-360 neous variance. It is not affected by changes in group size 361 or correlation. The proposed techniques GPDC and TPDC 362 outperform k-means and PDQ in most scenarios, they show 363 weakness in the two most extreme situations, i.e., scenarios 364 8 and 16. Specifically, when clusters have different variances 365 and the biggest variance is associated with the smallest 366

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Table 3 Number of units, variables, and clusters for the three real datasets

	n	J	K	
Seed	210	4	3	UCI machine learning repository
HSCT	9702	3	4	Terry Fox Lab
AIS	202	4	2	EMMIXuskew R package

Table 4 Adjusted Rand index for the real datasets

	GPDC	TPDC	PDQ	k-means	GMM	TMM
Seed	0.41	0.33	0.17	0.16	0.44	0.53
HSCT	0.99	0.98	0.98	0.72	0.88	0.99
AIS	0.88	0.90	0.16	0.06	0.72	0.81

cluster, they fail detecting the clustering partitions. Fig-367 ures 5, 6, 7, 8, 9, 10, 11, 12 in Appendix B.2 show examples 368 369 of simulated datasets for each scenario.

Real Data Analysis 370

371 We performed a real data analysis on three datasets that differ in size and number of clusters (details in Table 3). 372 We performed variable selection prior to cluster analysis 373 (details in Appendix B.1). The seed dataset¹ contains infor-374 mation about kernels belonging to three different varieties of 375 wheat-Kama, Rosa and Canadian-with 70 observations 376 per variety (see Fig. 1). We used the variables: compactness, 377 length of kernel, width of kernel, and asymmetry coefficient. 378 The hematopoietic steam cell transplant (HSCT) data were 379 collected in the Terry Fox Lab at the British Columbia Can-380 cer Agency. It contains information about 9780 cells, each 381 stained with four fluorescent dyes. Experts identified four 382 clusters; moreover, 78 cells were deemed "dead", leaving a 383 total of 9702 observation, we selected the three most inform-384 ative variables. Figure 2 shows the partitions defined by the 385 experts. The Australian Institute of sport dataset². contains 386 data on 102 male and 100 female athletes for the Australian 387 institute of sports. We selected the variables: height in cm, 388 hematocrit, plasma ferritin concentration, and percent body 389 fat, see Fig. 3. 390

391 Table 4 shows the ARI on the three datasets. On the seed dataset, GPDC and TPDC perform better than PDQ and 392 k-means. The improvement from PDQ is noticeable, PDQ 393 gives an ARI of 0.17, while GPDC gives an ARI of 0.41. 394 On this dataset, TMM gives the best performance. On the 395 HSCT dataset, GPDC, TPDC, PDQ, and TMM have a very 396

² GLMsData R package. 2EL 01

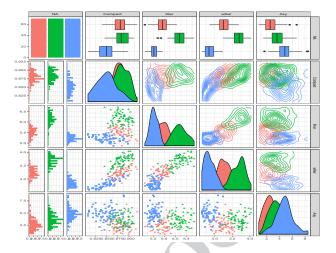


Fig. 1 Seed dataset, each color and symbol representing a different variety of wheat

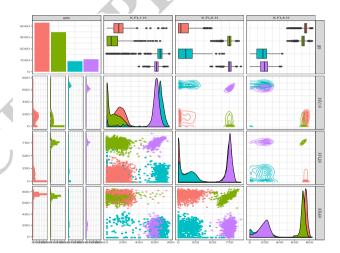


Fig. 2 HSCT dataset, each color and symbol representing a partition defined by the experts

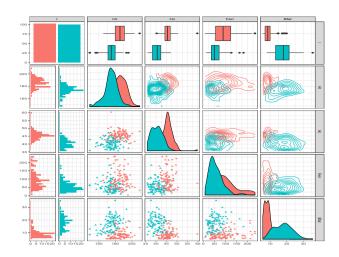
high ARI. On the AIS dataset, GPDC and TPDC give the 397 best performance. 398

Conclusion

A new distance measure based on density functions is intro-400 duced and used in the context of probabilistic distance clus-401 tering adjusted for cluster size (PDQ). PDQ assumes that, 402 for a generic unit, the product between the probability of 403 belonging to a cluster and its distance from the cluster is 404 constant. The minimization of the sum of these constants 405 over the units leads to clusters that maximize the classifi-406 ability of the data. We introduce two algorithms based on 407 PDQ that use distance measures based on the multivariate 408

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 $\ensuremath{\mbox{Fig. 3}}$ AIS dataset, each color and symbol representing male and female athletes

Gaussian distribution and on the multivariate Student-*t* distribution. Using simulated and real datasets we show how
the new algorithms over-perform PDQ and the well known
k-means algorithm.

The algorithm could be extended using different distributions. Further to this point, we mentioned outliers as a possible motivation for the PDQ approach with the multivariate Student-*t* distribution (Sect. 3). However, if the objective is dealing with outliers, it will be better to consider the PDQ approach with the multivariate contaminated normal dis-

tribution [25] and this will be a topic of future work. Other

approaches for handling cluster concentration will also be
considered (e.g., [9]) as will methods that accommodate
asymmetric, or skewed, clusters (e.g., [18, 19, 21, 22, 32,
34]).420
421
422

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Compliance with Ethical Standards

Conflict of interestOn behalf of all authors, the corresponding author433states that there is no conflict of interest.434

Dissimilarity Measure

435

424

432

A general measure $d(\mathbf{x}, \mathbf{y})$ is a dissimilarity measure if the following conditions are verified [33, p.404]: 437

- $1. \quad \mathbf{d}(\mathbf{x}, \mathbf{y}) \ge 0 \tag{438}$
- 2. $d(x, y) = 0 \Leftrightarrow x = y$ 439
- 3. d(x, y) = d(x, y). 440

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Let $f(\mathbf{x}_i; \boldsymbol{\mu}_k, \boldsymbol{\theta}_k)$ be the generic symmetric unimodal multivari-441 ate density function of the random variable \mathbf{X} with parameter 442 $\boldsymbol{\theta}_k$ and location parameter $\boldsymbol{\mu}_k$ then 443

⁴⁴⁴
$$d(\boldsymbol{x}_i, \boldsymbol{\mu}_k) = \log\left(\frac{M_k}{f(\boldsymbol{x}_i; \boldsymbol{\mu}_k, \boldsymbol{\theta}_k)}\right),$$
⁴⁴⁵ (23)

satisfies all the three properties and it is a dissimilarity meas-446 ure for $k = 1, \ldots, K$. 447

448 1. d(
$$\boldsymbol{x}_i, \boldsymbol{\mu}_k$$
) > 0, $\forall \boldsymbol{x}_i$.
449 Proof

449

4

⁴⁵⁰
$$0 < \frac{f(\boldsymbol{x}_i; \boldsymbol{\mu}_k, \boldsymbol{\theta}_k)}{M_k} \le 1 \Rightarrow \frac{M_k}{f(\boldsymbol{x}_i; \boldsymbol{\mu}_k, \boldsymbol{\theta}_k)} \ge 1 \Rightarrow$$

 $\Rightarrow \log\left(\frac{M_k}{f(\boldsymbol{x}_i; \boldsymbol{\mu}_k, \boldsymbol{\theta}_k)}\right) \ge 0.$

451

454

Author Proof

2. d($\mathbf{x}_i, \mathbf{\mu}_k$) = 0 $\Leftrightarrow \mathbf{x}_i = \mathbf{\mu}_k$. 452 453 2a. $\mathbf{x}_i = \mathbf{\mu}_k \Rightarrow d(\mathbf{x}_i, \mathbf{\mu}_k) = 0 \ \forall \mathbf{x}_i$. Proof

$$\begin{aligned} \mathbf{x}_i &= \mathbf{\mu}_k \Rightarrow f(\mathbf{x}_i; \mathbf{\mu}_k, \mathbf{\theta}_k) = f(\mathbf{\mu}_k; \mathbf{\mu}_k, \mathbf{\theta}_k) &= M_k \Rightarrow \\ &\Rightarrow \frac{M_k}{M_k} = 1 \Rightarrow \log(1) = 0, \end{aligned}$$

 $1 \Rightarrow$

 $x_i = \mu_k, \ \forall x_i$ Proof 457

$$\log\left(\frac{M_k}{f(\boldsymbol{x}_i;\boldsymbol{\mu}_k,\boldsymbol{\theta}_k)}\right) = 0 \Rightarrow \frac{M_k}{f(\boldsymbol{x}_i;\boldsymbol{\mu}_k,\boldsymbol{\theta}_k)} = 1 \Rightarrow$$
$$\Rightarrow f(\boldsymbol{x}_i;\boldsymbol{\mu}_k,\boldsymbol{\theta}_k) = M_k$$
$$= f(\boldsymbol{\mu}_k;\boldsymbol{\mu}_k,\boldsymbol{\theta}_k) \Rightarrow \boldsymbol{x}_i = \boldsymbol{\mu}_k.$$

459

458

3. $d(\mathbf{x}_i, \boldsymbol{\mu}_k) = d(\boldsymbol{\mu}_k, \mathbf{x}_i), \ \forall \mathbf{x}_i \text{ Proof Given } \boldsymbol{\theta}_k$ 460

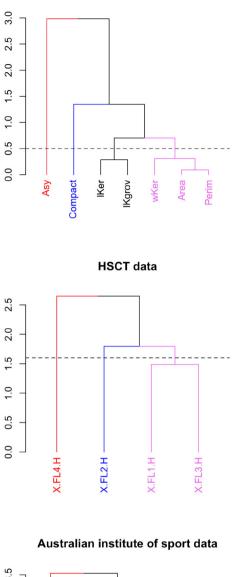
461
$$f(\mathbf{x}_i; \boldsymbol{\mu}_k, \boldsymbol{\theta}_k) = f(\boldsymbol{\mu}_k; \mathbf{x}_i, \boldsymbol{\theta}_k), \Rightarrow \log\left(\frac{M_k}{f(\mathbf{x}_i; \boldsymbol{\mu}_k, \boldsymbol{\theta}_k)}\right) = \log\left(\frac{M_k}{f(\boldsymbol{\mu}_k; \mathbf{x}_i, \boldsymbol{\theta}_k)}\right)$$

463



Variable Selection 465

On each dataset we selected one variable per group using 466 hierarchical clustering (Fig. 4).



Seed data

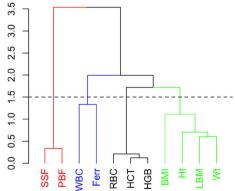


Fig. 4 Variable selection

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455	
456	2b. d($\mathbf{x}_i, \boldsymbol{\mu}_k$) = 0 \Rightarrow

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Simulated Datasets

 Table 5
 Simulated datasets
 using multivariate Gaussian distributions Scenarios 1-4

5 Simulated datasets g multivariate Gaussian		True			GPDC			TPDC		
butions Scenarios 1-4	μ_1	0.00	0.00	0.00	- 0.02	0.03	-0.01	0.01	-0.00	-0.01
	μ_2	- 7.00	7.00	0.00	- 7.00	7.00	0.01	-7.01	7.01	0.01
	μ_3	-7.00	0.00	7.00	-6.98	0.05	6.97	-6.99	0.02	7.02
	$\boldsymbol{\Sigma}_1$	1.00	0.00	0.00	1.29	-0.29	-0.02	0.97	-0.06	0.01
	-	0.00	1.00	0.00	-0.29	1.23	0.00	-0.06	0.95	0.00
		0.00	0.00	1.00	-0.02	0.00	0.93	0.01	0.00	0.84
	$\boldsymbol{\Sigma}_2$	1.00	0.00	0.00	1.04	-0.11	0.00	0.90	-0.02	-0.00
	_	0.00	1.00	0.00	-0.11	1.09	-0.04	-0.02	0.93	-0.01
		0.00	0.00	1.00	0.00	-0.04	0.90	-0.00	-0.01	0.84
	$\boldsymbol{\Sigma}_3$	1.00	0.00	0.00	1.04	-0.02	-0.11	0.87	-0.01	-0.01
	-	0.00	1.00	0.00	-0.02	1.18	-0.28	-0.01	0.89	-0.06
		0.00	0.00	1.00	-0.11	-0.28	1.35	-0.01	-0.06	0.95
	μ_1	0.00	0.00	0.00	-0.02	-0.00	0.03	0.02	-0.00	-0.01
	μ_2	-7.00	7.00	0.00	-6.99	6.94	0.07	-7.00	6.99	0.04
	μ_3	-7.00	0.00	7.00	-6.99	0.01	7.00	-7.00	0.01	7.01
	$\boldsymbol{\Sigma}_1$	1.00	0.00	0.00	1.28	-0.03	-0.27	0.97	-0.00	-0.04
		0.00	1.00	0.00	-0.03	0.91	0.00	-0.00	0.82	0.00
		0.00	0.00	1.00	-0.27	0.00	1.25	-0.04	0.00	0.97
	$\boldsymbol{\Sigma}_2$	1.00	0.00	0.00	1.03	-0.12	-0.03	0.85	-0.02	-0.02
		0.00	1.00	0.00	-0.12	1.36	-0.30	-0.02	0.95	-0.07
		0.00	0.00	1.00	-0.03	-0.30	1.23	-0.02	-0.07	0.92
	$\boldsymbol{\Sigma}_3$	1.00	0.00	0.00	1.03	0.00	-0.12	0.90	0.00	-0.02
		0.00	1.00	0.00	0.00	0.91	-0.03	0.00	0.84	-0.01
		0.00	0.00	1.00	-0.12	-0.03	1.07	-0.02	-0.01	0.90
	μ_1	0.00	0.00	0.00	-0.08	0.12	-0.02	0.01	0.03	-0.03
	μ_2	-4.00	4.00	0.00	-3.99	4.04	-0.01	-4.02	4.05	-0.01
	μ_3	-4.00	0.00	4.00	-3.92	0.22	3.79	-3.95	0.10	3.97
	$\boldsymbol{\Sigma}_1$	1.00	0.00	0.00	1.51	-0.52	-0.05	1.00	-0.20	-0.01
		0.00	1.00	0.00	-0.52	1.38	-0.00	-0.20	0.94	0.00
		0.00	0.00	1.00	-0.05	-0.00	0.87	-0.01	0.00	0.68
	Σ_2	1.00	0.00	0.00	1.01	-0.19	-0.00	0.80	-0.07	-0.00
		0.00	1.00	0.00	-0.19	1.13	-0.05	-0.07	0.87	-0.02
C		0.00	0.00	1.00	-0.00	-0.05	0.81	-0.00	-0.02	0.69
	$\boldsymbol{\Sigma}_3$	1.00	0.00	0.00	1.16	-0.05	-0.26	0.71	-0.02	-0.09
		0.00	1.00	0.00	-0.05	1.50	-0.70	-0.02	0.85	-0.27
		0.00	0.00	1.00	-0.26	-0.70	1.90	-0.09	-0.27	0.99
	μ_1	0.00	0.00	0.00	-0.08	-0.01	0.11	0.01	-0.01	0.02
	μ_2	-4.00	4.00	0.00	-3.93	3.77	0.24	-3.97	3.94	0.13
	μ_3	-4.00	0.00	4.00	-3.98	-0.01	4.03	-4.01	-0.01	4.05
	$\boldsymbol{\Sigma}_1$	1.00	0.00	0.00	1.51	-0.06	-0.51	0.99	-0.02	-0.19
		0.00	1.00	0.00	-0.06	0.86	-0.01	-0.02	0.67	-0.01
		0.00	0.00	1.00	-0.51	-0.01	1.40	-0.19	-0.01	0.94
	$\boldsymbol{\Sigma}_2$	1.00	0.00	0.00	1.14	-0.28	-0.05	0.70	-0.11	-0.02
		0.00	1.00	0.00	-0.28	1.91	-0.72	-0.11	1.01	-0.29
		0.00	0.00	1.00	-0.05	-0.72	1.54	-0.02	-0.29	0.88
	$\boldsymbol{\Sigma}_3$	1.00	0.00	0.00	1.02	-0.00	-0.20	0.78	-0.00	-0.07
		0.00	1.00	0.00	-0.00	0.82	-0.05	-0.00	0.67	-0.02
		0.00	0.00	1.00	-0.20	-0.05	1.12	-0.07	-0.02	0.82
		0								

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 Table 6
 Simulated datasets
 using multivariate Gaussian distributions Scenarios 5-8

6 Simulated datasets		True			GPDC			TPDC		
multivariate Gaussian		IIuc			OIDC		1	IIDC		
outions Scenarios 5-8	μ_1	0.00	0.00	0.00	-0.20	0.23	-0.03	-0.02	0.04	-0.02
	μ_2	-7.00	7.00	0.00	-7.33	8.08	-0.46	-7.28	7.85	-0.40
	μ_3	-7.00	0.00	7.00	-7.09	0.86	6.33	-7.03	0.40	6.78
	$\boldsymbol{\Sigma}_1$	1.00	0.00	0.00	2.68	-0.93	0.09	0.90	-0.09	0.02
		0.00	1.00	0.00	-0.93	2.60	-0.06	-0.09	0.89	-0.00
		0.00	0.00	1.00	0.09	-0.06	1.69	0.02	-0.00	0.74
	$\boldsymbol{\Sigma}_2$	16.00	0.00	0.00	12.84	0.95	0.09	11.36	0.46	-0.00
		0.00	16.00	0.00	0.95	13.43	1.24	0.46	12.01	0.80
		0.00	0.00	16.00	0.09	1.24	11.82	-0.00	0.80	10.36
	$\boldsymbol{\Sigma}_3$	2.25	0.00	0.00	4.40	-0.23	0.39	2.14	-0.09	0.13
		0.00	2.25	0.00	-0.23	6.37	-2.79	-0.09	2.79	-0.85
		0.00	0.00	2.25	0.39	-2.79	7.28	0.13	-0.85	2.94
	μ_1	0.00	0.00	0.00	-0.12	0.06	0.01	-0.00	0.01	-0.02
	μ_2	-7.00	7.00	0.00	-7.10	7.23	0.80	-7.07	7.11	0.87
	μ_3	-7.00	0.00	7.00	-7.02	0.06	7.00	-7.01	0.01	7.06
	$\boldsymbol{\Sigma}_1$	1.00	0.00	0.00	2.09	-0.29	-0.27	0.91	-0.04	-0.02
		0.00	1.00	0.00	-0.29	1.31	-0.11	-0.04	0.73	-0.02
		0.00	0.00	1.00	-0.27	-0.11	1.76	-0.02	-0.02	0.87
	$\boldsymbol{\Sigma}_2$	16.00	0.00	0.00	11.06	-0.17	-0.08	9.14	-0.44	-0.06
		0.00	16.00	0.00	-0.17	17.36	-4.90	-0.44	14.91	-4.73
	F	0.00	0.00	16.00	-0.08	-4.90	14.27	-0.06	-4.73	11.95
	Σ_3	2.25	0.00	0.00	2.42	7 0.03	-0.02	1.86	-0.01	-0.02
		0.00	2.25	0.00	-0.03	2.29	-0.44	-0.01	1.70	-0.09
		0.00	0.00	2.25	-0.02	-0.44	3.02	-0.02	-0.09	2.01
	μ_1	0.00	0.00	0.00	-0.07	0.17	-0.03	0.02	0.04	-0.03
	μ ₂	-4.00	4.00 0.00	0.00 4.00	-4.65 -4.12	5.39 0.66	-0.79 3.84	-4.55 -4.07	5.17 0.39	-0.65 4.00
	μ_3	-4.00	0.00	0.00	4.12 2.67	-0.07	0.01	1.04	0.39	4.00 0.00
	$\boldsymbol{\Sigma}_1$	1.00 0.00	1.00	0.00	-0.07	2.53	-0.01	0.01	1.02	-0.01
		0.00	0.00	1.00	0.07	-0.04	2.18	0.01	-0.01	0.01
	$\boldsymbol{\Sigma}_1$	16.00	0.00	0.00	12.16	1.15	-0.07	9.85	0.67	-0.06
	21	0.00	16.00	0.00	1.15	13.16	1.16	0.67	11.00	0.70
		0.00	0.00	16.00	-0.07	1.16	11.51	-0.06	0.70	9.30
	Σ_3	2.25	0.00	0.00	5.01	0.07	0.18	2.59	-0.01	0.04
	-3	0.00	2.25	0.00	0.07	5.12	-0.24	-0.01	2.60	-0.05
		0.00	0.00	2.25	0.18	-0.24	6.08	0.04	-0.05	2.92
	μ_1	0.00	0.00	0.00	-0.31	0.16	-0.01	-0.18	0.10	0.03
	μ_2	-4.00	4.00	0.00	-3.84	2.77	2.35	-1.30	0.33	1.09
	μ_3	-4.00	0.00	4.00	-4.18	-0.23	3.82	-4.21	-0.25	3.86
	Σ_1	1.00	0.00	0.00	3.00	-0.40	-0.15	1.75	-0.20	-0.20
	1	0.00	1.00	0.00	-0.40	2.29	-0.32	-0.20	1.40	-0.11
		0.00	0.00	1.00	-0.15	-0.32	2.80	-0.20	-0.11	1.61
	$\boldsymbol{\Sigma}_2$	16.00	0.00	0.00	5.58	-0.02	0.27	1.99	-0.17	-0.12
	2	0.00	16.00	0.00	-0.02	9.10	-3.30	-0.17	1.88	-0.38
		0.00	0.00	16.00	0.27	-3.30	6.61	-0.12	-0.38	2.05
	Σ_3	2.25	0.00	0.00	2.72	-0.01	0.18	2.01	-0.05	0.13
	2	0.00	2.25	0.00	-0.01	2.83	-0.61	-0.05	2.04	-0.47
		0.00	0.00	2.25	0.18	-0.61	3.37	0.13	-0.47	2.34

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 Table 7
 Simulated datasets
 using multivariate Gaussian distributions Scenarios 9-12

le 7 Simulated datasets g multivariate Gaussian		True			GPDC			TPDC		
ributions Scenarios 9-12	μ_1	0.00	0.00	0.00	-0.04	0.04	0.02	0.01	0.00	0.02
	μ_2	-7.00	7.00	0.00	-7.01	7.00	0.03	-7.02	7.01	0.02
	μ_3	-7.00	0.00	7.00	-6.98	0.03	7.02	-6.97	0.03	7.03
	Σ_1	1.00	0.00	0.00	1.36	-0.34	-0.01	1.00	-0.07	0.00
	1	0.00	1.00	0.00	-0.34	1.30	-0.01	-0.07	0.98	-0.00
		0.00	0.00	1.00	-0.01	-0.01	0.96	0.00	-0.00	0.86
	$\boldsymbol{\Sigma}_2$	1.00	-0.50	-0.50	1.07	-0.60	-0.47	0.92	-0.47	-0.43
		-0.50	1.00	0.50	-0.60	1.11	0.46	-0.47	0.94	0.43
		-0.50	0.50	1.00	-0.47	0.46	0.91	-0.43	0.43	0.85
	Σ_3	1.00	0.70	0.70	0.93	0.63	0.64	0.86	0.59	0.61
		0.70	1.00	0.70	0.63	0.93	0.62	0.59	0.85	0.60
		0.70	0.70	1.00	0.64	0.62	0.98	0.61	0.60	0.89
	μ_1	0.00	0.00	0.00	-0.04	-0.00	0.05	0.00	-0.00	0.01
	μ_2	-7.00	7.00	0.00	-7.00	6.97	0.05	-7.02	7.00	0.04
	μ_3	-7.00	0.00	7.00	-6.99	0.02	7.01	-6.99	0.02	7.01
	$\boldsymbol{\Sigma}_1$	1.00	0.00	0.00	1.40	-0.01	-0.35	1.01	-0.00	-0.06
		0.00	1.00	0.00	-0.01	0.97	0.01	-0.00	0.88	0.00
		0.00	0.00	1.00	-0.35	0.01	1.36	-0.06	0.00	1.01
	$\boldsymbol{\Sigma}_2$	1.00	-0.50	-0.50	1.06	-0.59	-0.50	0.90	-0.46	-0.46
		-0.50	1.00	0.50	-0.59	1.19	0.39	-0.46	0.93	0.43
		-0.50	0.50	1.00	-0.50	0.39	1.06	-0.46	0.43	0.91
	$\boldsymbol{\Sigma}_3$	1.00	0.70	0.70	0.94	0.63	0.64	0.88	0.60	0.61
		0.70	1.00	0.70	0.63	0.88	0.63	0.60	0.84	0.60
		0.70	0.70	1.00	0.64	0.63	0.94	0.61	0.60	0.88
	μ_1	0.00	0.00	0.00	-0.19	0.22	0.04	-0.09	0.11	0.05
	μ_2	-4.00	4.00	0.00	-4.05	4.04	0.08	-4.06	4.06	0.07
	μ_3	-4.00	0.00	4.00	-3.95	0.05	4.04	-3.94	0.05	4.08
	$\boldsymbol{\Sigma}_1$	1.00	0.00	0.00	1.78	-0.76	-0.00	1.32	-0.41	-0.01
		0.00	1.00	0.00	-0.76	1.62	-0.08	-0.41	1.20	-0.05
		0.00	0.00	1.00	-0.00	-0.08	0.97	-0.01	-0.05	0.83
	Σ_2	1.00	-0.50	-0.50	1.05	-0.64	-0.44	0.84	-0.47	-0.38
		-0.50	1.00	0.50	-0.64	1.17	0.43	-0.47	0.91	0.37
		-0.50	0.50	1.00	-0.44	0.43	0.83	-0.38	0.37	0.72
	$\boldsymbol{\Sigma}_3$	1.00	0.70	0.70	0.86	0.57	0.57	0.57	0.40	0.42
		0.70	1.00	0.70	0.57	0.90	0.55	0.40	0.59	0.42
		0.70	0.70	1.00	0.57	0.55	0.99	0.42	0.42	0.65
	μ_1	0.00	0.00	0.00	-0.17	-0.00	0.24	-0.08	0.01	0.12
	μ_2	-4.00	4.00	0.00	-3.94	3.85	0.12	-4.03	4.02	0.08
	μ_3	-4.00	0.00	4.00	-3.99	0.03	4.05	-3.99	0.03	4.04
	$\boldsymbol{\Sigma}_1$	1.00	0.00	0.00	1.89	0.04	-0.79	1.34	-0.02	-0.40
		0.00	1.00	0.00	0.04	0.97	0.02	-0.02	0.85	0.00
		0.00	0.00	1.00	-0.79	0.02	1.70	-0.40	0.00	1.25
	$\boldsymbol{\Sigma}_2$	1.00	-0.50	-0.50	1.19	-0.79	-0.46	0.68	-0.40	-0.32
		-0.50	1.00	0.50	-0.79	1.66	0.22	-0.40	0.81	0.28
	v	-0.50	0.50	1.00	-0.46	0.22	1.23	-0.32	0.28	0.70
	Σ_3	1.00	0.70	0.70	0.88	0.59	0.62	0.78	0.53	0.56
		0.70	1.00	0.70	0.59	0.77	0.59	0.53	0.70	0.53
		0.70	0.70	1.00	0.62	0.59	0.91	0.56	0.53	0.81

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Table 8Simulated datasetsusing multivariate GaussiandistributionsScenarios13–16

	True			GPDC			TPDC		
μ_1	0.00	0.00	0.00	-0.44	0.48	-0.24	-0.09	0.11	-0.06
μ_2	-7.00	7.00	0.00	-7.94	8.25	0.64	-7.73	7.75	0.58
μ_3	-7.00	0.00	7.00	-6.98	0.63	6.52	-6.92	0.09	7.09
$\boldsymbol{\Sigma}_1$	1.00	0.00	0.00	4.11	-2.42	0.32	1.24	-0.36	0.10
I	0.00	1.00	0.00	-2.42	3.92	-0.39	-0.36	1.20	-0.11
	0.00	0.00	1.00	0.32	-0.39	2.86	0.10	-0.11	1.06
$\boldsymbol{\Sigma}_2$	16.00	-8.00	-8.00	13.01	-6.32	-6.60	12.04	-5.73	-5.83
2	-8.00	16.00	8.00	-6.32	13.59	6.70	-5.73	12.79	5.60
	-8.00	8.00	16.00	-6.60	6.70	11.52	-5.83	5.60	11.29
$\boldsymbol{\Sigma}_3$	2.25	1.57	1.57	3.11	0.73	0.83	1.29	0.88	0.93
-3	1.57	2.25	1.57	0.73	5.46	-0.75	0.88	1.37	0.90
	1.57	1.57	2.25	0.83	-0.75	6.25	0.93	0.90	1.46
μ_1	0.00	0.00	0.00	-0.16	0.10	0.00	-0.01	0.02	-0.01
μ_2	-7.00	7.00	0.00	-7.72	7.58	0.98	-7.61	7.51	0.92
μ_3	-7.00	0.00	7.00	-6.96	0.05	7.05	-6.96	0.05	7.06
Σ_1	1.00	0.00	0.00	2.40	-0.40	-0.55	0.94	-0.05	-0.04
1	0.00	1.00	0.00	-0.40	1.46	-0.08	-0.05	0.76	-0.03
	0.00	0.00	1.00	-0.55	-0.08	2.40	-0.04	-0.03	0.96
$\boldsymbol{\Sigma}_2$	16.00	-8.00	-8.00	12.21	-5.92	-5.73	10.56	-5.29	-4.72
2	-8.00	16.00	8.00	-5.92	15.67	3.66	-5.29	13.38	2.96
	-8.00	8.00	16.00	-5.73	3.66	13.43	-4.72	2.96	11.09
$\boldsymbol{\Sigma}_3$	2.25	1.57	1.57	2.00	1.32	1.37	1.80	1.21	1.27
3	1.57	2.25	1.57	1.32	1.87	1.29	1.21	1.66	1.20
	1.57	1.57	2.25	1.37	1.29	2.11	1.27	1.20	1.84
μ_1	0.00	0.00	0.00	-0.25	0.41	-0.21	-0.21	0.17	0.02
μ_2	-4.00	4.00	0.00	-4.99	5.11	0.35	-5.06	4.94	0.45
μ_3	-4.00	0.00	4.00	-4.12	1.09	3.35	-3.52	0.39	3.62
$\boldsymbol{\Sigma}_1$	1.00	0.00	0.00	3.91	-1.63	-1.06	2.06	-0.51	-0.61
-	0.00	1.00	0.00	-1.63	3.78	0.90	-0.51	1.79	0.34
	0.00	0.00	1.00	-1.06	0.90	3.57	-0.61	0.34	2.05
Σ_2	16.00	-8.00	-8.00	11.09	-5.47	-5.53	10.45	-5.07	-5.30
	-8.00	16.00	8.00	-5.47	12.68	6.01	-5.07	11.75	5.11
	-8.00	8.00	16.00	-5.53	6.01	10.45	-5.30	5.11	10.05
Σ_3	2.25	1.57	1.57	5.54	-1.45	-0.85	1.76	0.22	0.27
	1.57	2.25	1.57	-1.45	6.76	1.81	0.22	1.96	0.84
	1.57	1.57	2.25	-0.85	1.81	6.94	0.27	0.84	2.03
μ_1	0.00	0.00	0.00	-0.37	0.29	-0.14	-0.32	0.32	-0.02
μ_2	-4.00	4.00	0.00	-2.03	1.59	1.47	-0.32	0.32	-0.02
μ_3	-4.00	0.00	4.00	-4.66	-0.55	3.40	-4.41	-0.34	3.66
$\boldsymbol{\Sigma}_1$	1.00	0.00	0.00	3.80	-1.26	-1.06	2.70	-1.22	-0.64
	0.00	1.00	0.00	-1.26	2.89	0.49	-1.22	2.39	0.36
	0.00	0.00	1.00	-1.06	0.49	3.82	-0.64	0.36	2.35
$\boldsymbol{\Sigma}_2$	16.00	-8.00	-8.00	4.97	-2.68	-0.61	2.70	-1.22	-0.64
	-8.00	16.00	8.00	-2.68	6.30	-0.13	-1.22	2.39	0.36
	-8.00	8.00	16.00	-0.61	-0.13	5.50	-0.64	0.36	2.35
Σ_3	2.25	1.57	1.57	1.92	0.60	0.88	1.20	0.45	0.62
	1.57	2.25	1.57	0.60	2.02	0.78	0.45	1.18	0.50
	1.57	1.57	2.25	0.88	0.78	2.02	0.62	0.50	1.27

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 Table 9
 Simulated datasets
 using multivariate Student-t distributions Scenarios 1-4

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le 9 Simulated datasets ng multivariate Student- <i>t</i>		True			GPDC			TPDC		
ributions Scenarios 1–4	μ_1	0.00	0.00	0.00	-0.05	0.06	-0.01	0.01	-0.00	-0.01
	μ_2	-7.00	7.00	0.00	-6.98	7.02	-0.00	-7.00	7.02	0.00
	μ_3	-7.00	0.00	7.00	-6.96	0.11	6.90	-6.99	0.01	7.02
	$\boldsymbol{\Sigma}_1$	1.00	0.00	0.00	2.02	-0.60	-0.08	0.96	-0.09	-0.03
		0.00	1.00	0.00	-0.60	1.94	-0.01	-0.09	0.94	-0.01
		0.00	0.00	1.00	-0.08	-0.01	1.27	-0.03	-0.01	0.76
	$\boldsymbol{\Sigma}_2$	1.00	0.00	0.00	1.47	-0.22	-0.01	1.01	-0.04	-0.01
		0.00	1.00	0.00	-0.22	1.62	-0.04	-0.04	1.06	-0.00
		0.00	0.00	1.00	-0.01	-0.04	1.24	-0.01	-0.00	0.90
	$\boldsymbol{\Sigma}_3$	1.00	0.00	0.00	1.65	-0.06	-0.32	0.81	-0.03	-0.04
		0.00	1.00	0.00	-0.06	2.13	-0.72	-0.03	0.88	-0.08
		0.00	0.00	1.00	-0.32	-0.72	2.59	-0.04	-0.08	0.97
	μ_1	0.00	0.00	0.00	-0.06	-0.01	0.07	0.01	-0.01	-0.00
	μ_2	-7.00	7.00	0.00	-6.96	6.87	0.11	-7.02	7.00	0.01
	μ_3	-7.00	0.00	7.00	-7.00	-0.02	7.02	-7.02	-0.01	7.03
	$\boldsymbol{\Sigma}_1$	1.00	0.00	0.00	2.05	-0.06	-0.63	0.97	-0.02	-0.11
		0.00	1.00	0.00	-0.06	1.28	-0.01	-0.02	0.78	-0.00
		0.00	0.00	1.00	-0.63	-0.01	1.92	-0.11	-0.00	0.95
	$\boldsymbol{\Sigma}_2$	1.00	0.00	0.00	1.67	-0.34	-0.01	0.80	-0.05	0.00
		0.00	1.00	0.00	-0.34	2.51	-0.73	-0.05	0.95	-0.10
		0.00	0.00	1.00	-0.01	-0.73	2.07	0.00	-0.10	0.90
	$\boldsymbol{\Sigma}_3$	1.00	0.00	0.00	1.52	-0.04	-0.25	1.03	-0.03	-0.06
		0.00	1.00	0.00	-0.04	1.24	-0.05	-0.03	0.92	-0.02
		0.00	0.00	1.00	-0.25	-0.05	1.60	-0.06	-0.02	1.05
	μ_1	0.00	0.00	0.00	-0.09	0.13	-0.03	-0.01	0.05	-0.02
	μ_2	-4.00	4.00	0.00	-3.99	4.06	-0.03	-4.02	4.07	-0.02
	μ_3	-4.00	0.00	4.00	-3.88	0.29	3.71	-3.94	0.14	3.89
	$\boldsymbol{\Sigma}_1$	1.00	0.00	0.00	1.98	-0.61	-0.08	0.95	-0.21	-0.03
		0.00	1.00	0.00	-0.61	1.86	-0.01	-0.21	0.91	-0.01
		0.00	0.00	1.00	-0.08	-0.01	1.19	-0.03	-0.01	0.64
	Σ_2	1.00	0.00	0.00	1.33	-0.20	-0.01	0.75	-0.05	-0.01
		0.00	1.00 0.00	0.00	-0.20	1.54	-0.03	-0.05	0.81	0.00
	F	0.00 1.00	0.00	1.00	-0.01	-0.03	1.11	-0.01	0.00	0.64
	Σ_3	0.00	1.00	0.00 0.00	1.70 0.09	-0.09 2.22	-0.40 -0.88	0.86 0.05	-0.05 1.09	-0.16 -0.37
		0.00	0.00	1.00	-0.40	-0.88	2.85	-0.16	-0.37	1.32
		0.00	0.00	0.00	-0.10	-0.03	0.13	-0.01	-0.02	0.04
	μ_1 μ_2	-4.00	4.00	0.00	-3.90	3.67	0.13	-3.97	3.87	0.14
	μ_2 μ_3	-4.00	0.00	4.00	-4.00	-0.05	4.06	-4.03	-0.03	4.08
	Σ_1	1.00	0.00	0.00	2.01	-0.05	-0.64	0.96	-0.02	-0.23
	-1	0.00	1.00	0.00	-0.05	1.20	-0.02	-0.02	0.64	-0.01
		0.00	0.00	1.00	-0.64	-0.02	1.82	-0.23	-0.01	0.91
	$\boldsymbol{\Sigma}_2$	1.00	0.00	0.00	1.72	-0.40	-0.06	0.83	-0.15	-0.02
	-2	0.00	1.00	0.00	-0.40	2.75	-0.90	-0.15	1.25	-0.39
		0.00	0.00	1.00	-0.06	-0.90	2.18	-0.02	-0.39	1.07
	$\boldsymbol{\Sigma}_3$	1.00	0.00	0.00	1.38	-0.03	-0.23	0.78	-0.02	-0.08
	- 5	0.00	1.00	0.00	-0.03	1.12	-0.03	-0.02	0.66	-0.01
		0.00	0.00	1.00	-0.23	-0.03	1.53	-0.08	-0.01	0.82
			0.00	1.00	0.20	0.05	1.00	0.00	0.01	

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Table 10Simulated datasetsusing multivariateStudent-tdistributionsScenarios5-8

	True			GPDC			TPDC		
μ_1	0.00	0.00	0.00	-0.16	0.17	- 0.02	- 0.01	0.02	- 0.01
μ_2	-7.00	7.00	0.00	-7.04	7.27	- 0.07	- 7.04	7.18	-0.05
μ_3	-7.00	0.00	7.00	- 6.99	0.47	6.59	- 7.00	0.14	6.93
$\boldsymbol{\Sigma}_1$	1.00	0.00	0.00	2.80	- 1.12	- 0.04	0.97	-0.15	-0.02
-	0.00	1.00	0.00	-1.12	2.72	- 0.05	-0.15	0.96	-0.01
	0.00	0.00	1.00	-0.04	- 0.05	1.60	-0.02	-0.01	0.74
$\boldsymbol{\Sigma}_2$	16.00	0.00	0.00	4.83	- 0.20	0.02	3.58	-0.14	-0.02
2	0.00	16.00	0.00	-0.20	5.39	0.09	-0.14	3.84	0.06
	0.00	0.00	16.00	0.02	0.09	4.25	- 0.02	0.06	3.14
$\boldsymbol{\Sigma}_3$	2.25	0.00	0.00	2.90	-0.14	-0.21	1.32	-0.07	-0.05
	0.00	2.25	0.00	-0.14	4.69	-2.42	- 0.07	1.77	- 0.56
	0.00	0.00	2.25	-0.21	-2.42	5.80	- 0.05	- 0.56	1.97
μ_1	0.00	0.00	0.00	-0.10	0.00	0.08	0.00	- 0.01	-0.00
μ_2	-7.00	7.00	0.00	- 6.96	6.76	0.36	- 7.01	6.93	0.18
μ_3	- 7.00	0.00	7.00	-7.02	- 0.02	7.04	- 7.03	-0.02	7.06
Σ_1	1.00	0.00	0.00	2.32	-0.18	- 0.65	0.91	-0.03	-0.10
	0.00	1.00	0.00	-0.18	1.42	-0.03	- 0.03	0.71	-0.01
	0.00	0.00	1.00	-0.65	- 0.03	2.06	-0.10	-0.01	0.87
$\boldsymbol{\Sigma}_2$	16.00	0.00	0.00	5.13	- 0.65	-0.00	2.94	-0.34	-0.02
	0.00	16.00	0.00	- 0.65	8.04	- 2.46	- 0.34	4.20	- 1.14
	0.00	0.00	16.00	- 0.00	- 2.46	6.76	-0.02	-1.14	3.78
$\boldsymbol{\Sigma}_3$	2.25	0.00	0.00	2.13	- 0.05	-0.22	1.48	-0.04	- 0.08
	0.00	2.25	0.00	-0.05	1.83	- 0.20	- 0.04	1.32	-0.04
	0.00	0.00	2.25	-0.22	- 0.20	2.40	-0.08	-0.04	1.56
μ_1	0.00	0.00	0.00	-0.16	0.21	-0.05	- 0.05	0.09	-0.03
μ_2	-4.00	4.00	0.00	-4.12	4.46	- 0.19	-4.12	4.38	-0.15
μ_3	-4.00	0.00	4.00	- 3.99	0.62	3.53	- 3.98	0.40	3.72
$\boldsymbol{\Sigma}_1$	1.00	0.00	0.00	2.57	- 0.62	- 0.02	1.11	-0.20	-0.02
	0.00	1.00	0.00	-0.62	2.48	-0.07	- 0.20	1.08	-0.02
	0.00	0.00	1.00	-0.02	-0.07	1.80	- 0.02	-0.02	0.82
$\boldsymbol{\Sigma}_2$	16.00	0.00	0.00	4.42	- 0.02	0.03	2.61	- 0.05	-0.01
	0.00	16.00	0.00	-0.02	5.15	0.19	- 0.05	2.91	0.10
	0.00	0.00	16.00	0.03	0.19	4.06	- 0.01	0.10	2.34
Σ_3	2.25	0.00	0.00	3.18	-0.14	-0.17	1.56	-0.10	-0.10
	0.00	2.25	0.00	-0.14	3.89	- 1.26	-0.10	1.96	- 0.63
	0.00	0.00	2.25	-0.17	- 1.26	4.89	-0.10	- 0.63	2.30
μ_1	0.00	0.00	0.00	-0.16	0.01	0.10	- 0.05	- 0.00	0.03
μ_2	-4.00	4.00	0.00	- 3.95	3.53	0.75	- 3.95	3.45	0.75
μ_3	-4.00	0.00	4.00	-4.06	- 0.05	4.11	-4.06	-0.05	4.14
$\boldsymbol{\Sigma}_1$	1.00	0.00	0.00	2.40	-0.17	- 0.49	1.08	-0.05	-0.21
	0.00	1.00	0.00	-0.17	1.57	-0.12	- 0.05	0.76	-0.04
	0.00	0.00	1.00	-0.49	-0.12	2.15	-0.21	-0.04	1.02
$\boldsymbol{\Sigma}_2$	16.00	0.00	0.00	4.25	-0.42	- 0.04	2.83	-0.38	-0.09
	0.00	16.00	0.00	-0.42	6.72	-2.02	- 0.38	4.79	- 1.79
	0.00	0.00	16.00	- 0.04	- 2.02	5.15	- 0.09	- 1.79	3.74
$\boldsymbol{\Sigma}_3$	2.25	0.00	0.00	2.04	- 0.05	-0.13	1.10	- 0.03	-0.07
	0.00	2.25	0.00	-0.05	1.81	-0.16	- 0.03	0.97	-0.02
	0.00	0.00	2.25	-0.13	-0.16	2.41	- 0.07	- 0.02	1.21

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Table 11Simulated datasetsusing multivariate Student-tdistributions Scenarios 9–12

	True			GPDC			TPDC		
μ_1	0.00	0.00	0.00	-0.11	0.11	0.04	- 0.01	0.01	0.03
μ_2	- 7.00	7.00	0.00	- 7.01	7.02	0.04	- 7.02	7.03	0.03
μ_3	- 7.00	0.00	7.00	- 6.97	0.03	7.03	- 6.98	0.02	7.03
Σ_1	1.00	0.00	0.00	2.37	- 0.83	- 0.08	1.16	-0.14	-0.04
1	0.00	1.00	0.00	- 0.83	2.26	- 0.06	-0.14	1.12	- 0.02
	0.00	0.00	1.00	- 0.08	- 0.06	1.43	- 0.04	-0.02	0.91
$\boldsymbol{\Sigma}_2$	1.00	-0.50	-0.50	1.55	-0.92	-0.69	1.07	-0.57	-0.51
2	- 0.50	1.00	0.50	- 0.92	1.69	0.68	-0.57	1.12	0.51
	-0.50	0.50	1.00	- 0.69	0.68	1.31	-0.51	0.51	0.97
Σ_3	1.00	0.70	0.70	1.32	0.89	0.89	0.77	0.53	0.56
	0.70	1.00	0.70	0.89	1.44	0.89	0.53	0.80	0.57
	0.70	0.70	1.00	0.89	0.89	1.54	0.56	0.57	0.87
μ_1	0.00	0.00	0.00	-0.10	- 0.01	0.14	- 0.01	- 0.01	0.02
μ_2	- 7.00	7.00	0.00	- 6.98	6.92	0.06	- 7.03	7.01	0.02
μ_3	- 7.00	0.00	7.00	- 7.01	- 0.00	7.01	- 7.01	- 0.00	7.01
Σ_1	1.00	0.00	0.00	2.46	- 0.00	- 0.87	1.11	- 0.02	-0.14
	0.00	1.00	0.00	- 0.00	1.47	0.03	- 0.02	0.91	-0.00
	0.00	0.00	1.00	- 0.87	0.03	2.26	-0.14	- 0.00	1.07
$\boldsymbol{\Sigma}_2$	1.00	-0.50	-0.50	1.71	- 1.04	- 0.66	0.83	- 0.45	-0.40
	- 0.50	1.00	0.50	- 1.04	2.24	0.43	-0.45	0.92	0.39
	- 0.50	0.50	1.00	- 0.66	0.43	1.72	-0.40	0.39	0.86
Σ_3	1.00	0.70	0.70	1.32	0.86	0.89	1.04	0.69	0.72
	0.70	1.00	0.70	0.86	1.18	0.86	0.69	0.97	0.70
	0.70	0.70	1.00	0.89	0.86	1.34	0.72	0.70	1.06
μ_1	0.00	0.00	0.00	-0.26	0.29	0.02	-0.13	0.14	0.03
μ_2	- 4.00	4.00	0.00	- 4.08	4.09	0.09	-4.07	4.08	0.07
μ_3	-4.00	0.00	4.00	- 3.93	0.07	4.01	-3.96	0.02	4.06
$\boldsymbol{\Sigma}_1$	1.00	0.00	0.00	2.49	- 1.03	- 0.08	1.27	-0.44	- 0.04
	0.00	1.00	0.00	- 1.03	2.30	- 0.07	- 0.44	1.19	- 0.06
	0.00	0.00	1.00	- 0.08	-0.07	1.47	- 0.04	- 0.06	0.84
Σ_2	1.00	- 0.50	-0.50	1.36	-0.80	- 0.62	0.89	- 0.49	-0.41
	- 0.50	1.00	0.50	- 0.80	1.55	0.61	- 0.49	0.98	0.42
	- 0.50	0.50	1.00	- 0.62	0.61	1.14	-0.41	0.42	0.77
Σ_3	1.00	0.70	0.70	1.25	0.77	0.73	0.60	0.42	0.44
	0.70	1.00	0.70	0.77	1.43	0.76	0.42	0.65	0.46
	0.70	0.70	1.00	0.73	0.76	1.62	0.44	0.46	0.73
μ_1	0.00	0.00	0.00	- 0.28	- 0.04	0.36	-0.11	-0.01	0.18
μ_2	-4.00	4.00	0.00	- 3.89	3.71	0.18	- 4.01	3.97	0.04
μ_3	-4.00	0.00	4.00	- 4.01	- 0.00	4.06	- 4.01	0.01	4.05
$\boldsymbol{\Sigma}_1$	1.00	0.00	0.00	2.81	0.18	- 1.16	1.36	0.03	- 0.50
	0.00	1.00	0.00	0.18	1.43	0.11	0.03	0.82	0.03
	0.00	0.00	1.00	-1.16	0.11	2.44	- 0.50	0.03	1.25
$\boldsymbol{\Sigma}_2$	1.00	- 0.50	- 0.50	1.84	- 1.10	-0.55	0.81	- 0.49	- 0.35
	-0.50	1.00	0.50	- 1.10	2.72	0.13	- 0.49	1.02	0.30
_	- 0.50	0.50	1.00	- 0.55	0.13	2.07	- 0.35	0.30	0.84
Σ_3	1.00	0.70	0.70	1.19	0.79	0.87	0.86	0.57	0.63
	0.70	1.00	0.70	0.79	1.00	0.81	0.57	0.76	0.59
	0.70	0.70	1.00	0.87	0.81	1.25	0.63	0.59	0.91

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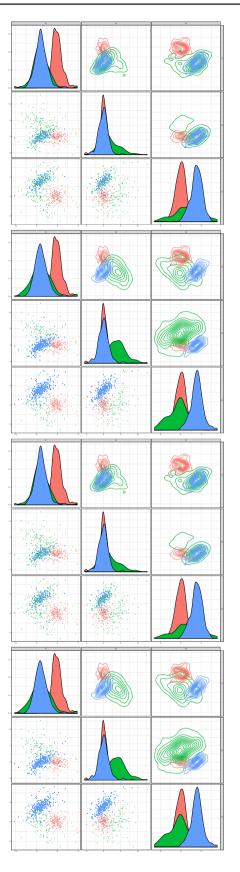
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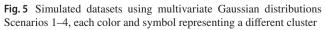
 Table 12
 Simulated datasets
 using multivariate Student-t distributions Scenarios 13-16

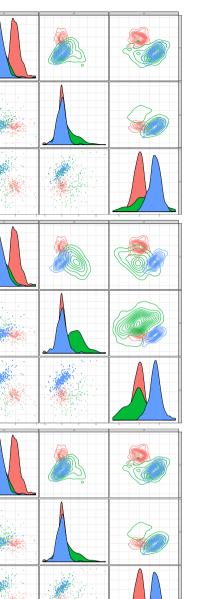
12 Simulated datasets multivariate Student- <i>t</i>		True			GPDC			TPDC		
outions Scenarios 13-16	μ_1	0.00	0.00	0.00	-0.35	0.35	-0.04	- 0.06	0.06	0.01
	μ_2	- 7.00	7.00	0.00	-7.22	7.28	0.20	-7.15	7.17	0.15
	μ_3	-7.00	0.00	7.00	- 6.97	0.12	6.95	-6.97	0.02	7.05
	Σ_1	1.00	0.00	0.00	3.87	-2.17	- 0.03	1.22	-0.33	- 0.00
		0.00	1.00	0.00	-2.17	3.67	-0.13	-0.33	1.18	- 0.06
		0.00	0.00	1.00	-0.03	-0.13	2.15	-0.00	-0.06	0.90
	$\boldsymbol{\Sigma}_2$	16.00	- 8.00	- 8.00	5.07	- 2.80	- 2.48	3.94	- 2.09	- 1.86
		- 8.00	16.00	8.00	-2.80	5.56	2.46	- 2.09	4.22	1.84
		- 8.00	8.00	16.00	- 2.48	2.46	4.51	- 1.86	1.84	3.55
	Σ_3	2.25	1.57	1.57	2.05	1.22	1.19	1.01	0.70	0.74
		1.57	2.25	1.57	1.22	2.70	0.89	0.70	1.09	0.76
		1.57	1.57	2.25	1.19	0.89	2.98	0.74	0.76	1.18
	μ_1	0.00	0.00	0.00	-0.15	0.02	0.15	-0.01	0.00	0.02
	μ_2	- 7.00	7.00	0.00	- 7.09	6.93	0.29	- 7.12	7.09	0.13
	μ_3	- 7.00	0.00	7.00	- 7.01	- 0.01	7.03	- 7.01	- 0.00	7.02
	$\boldsymbol{\Sigma}_1$	1.00	0.00	0.00	2.83	-0.15	- 0.99	1.03	-0.04	-0.14
		0.00	1.00	0.00	-0.15	1.62	0.04	-0.04	0.82	-0.01
		0.00	0.00	1.00	- 0.99	0.04	2.56	-0.14	-0.01	0.99
	$\boldsymbol{\Sigma}_2$	16.00	- 8.00	-8.00	5.42	- 3.00	- 2.19	2.84	- 1.57	- 1.29
		- 8.00	16.00	8.00	- 3.00	7.28	1.35	- 1.57	3.40	1.13
		- 8.00	8.00	16.00	- 2.19	1.35	5.99	- 1.29	1.13	3.04
	$\boldsymbol{\Sigma}_3$	2.25	1.57	1.57	1.89	1.24	1.31	1.43	0.94	1.01
		1.57	2.25	1.57	1.24	1.69	1.24	0.94	1.31	0.96
		1.57	1.57	2.25	1.31	1.24	1.96	1.01	0.96	1.48
	μ_1	0.00	0.00	0.00	- 0.42	0.51	- 0.16	-0.30	0.31	- 0.08
	μ_2	-4.00	4.00	0.00	- 4.50	4.61	0.31	- 4.37	4.41	0.34
	μ_3	-4.00	0.00	4.00	- 3.83	0.40	3.57	- 3.96	0.09	4.05
	$\boldsymbol{\Sigma}_1$	1.00	0.00	0.00	3.44	- 1.68	- 0.32	1.95	-0.79	-0.07
		0.00	1.00	0.00	- 1.68	3.25	0.21	-0.79	1.84	- 0.06
		0.00	0.00	1.00	-0.32	0.21	2.63	-0.07	- 0.06	1.50
	$\boldsymbol{\Sigma}_2$	16.00	- 8.00	- 8.00	4.40	- 2.35	- 2.20	2.98	- 1.53	- 1.41
		- 8.00	16.00	8.00	-2.35	5.01	2.21	-1.53	3.31	1.41
		- 8.00	8.00	16.00	-2.20	2.21	3.83	- 1.41	1.41	2.68
	Σ_3	2.25	1.57	1.57	2.70	0.49	0.35	1.02	0.62	0.66
		1.57	2.25	1.57	0.49	3.49	0.71	0.62	1.19	0.64
	1	1.57	1.57	2.25	0.35	0.71	4.04	0.66	0.64	1.35
	μ_1	0.00	0.00	0.00	- 0.44	0.26	0.17	- 0.27	0.16	0.13
	μ_2	- 4.00	4.00	0.00	- 2.50	2.03	0.77	-2.42	2.15	0.36
	μ_3	-4.00	0.00	4.00	- 3.11	0.13	3.09	- 3.10	0.09	3.20
× ×	$\boldsymbol{\Sigma}_1$	1.00	0.00	0.00	3.12	- 0.84	- 0.85	1.78	-0.48	- 0.50
		0.00	1.00	0.00	-0.84	2.19	0.23	-0.48	1.29	0.08
-	-	0.00	0.00	1.00	-0.85	0.23	2.47	- 0.50	0.08	1.41
	$\boldsymbol{\Sigma}_2$	16.00	-8.00	- 8.00	3.66	- 2.07	-0.63	2.46	- 1.40	- 0.64
		- 8.00	16.00	8.00	- 2.07	4.99	0.16	- 1.40	3.27	0.14
	F	- 8.00	8.00	16.00	- 0.63	0.16	3.83	- 0.64	0.14	2.48
	Σ_3	2.25	1.57	1.57	1.49	0.66	0.82	0.90	0.52	0.63
		1.57	2.25	1.57	0.66	1.53	0.57	0.52	0.83	0.50
		1.57	1.57	2.25	0.82	0.57	1.82	0.63	0.50	1.03

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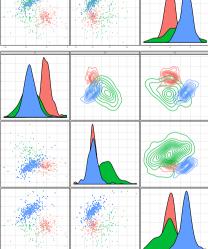
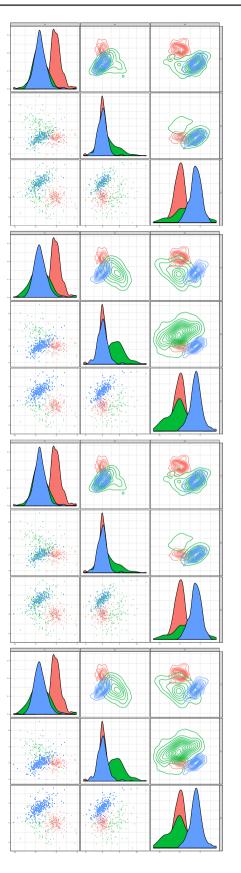
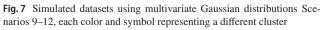


Fig. 6 Simulated datasets using multivariate Gaussian distributions Scenarios 5–8, each color and symbol representing a different cluster

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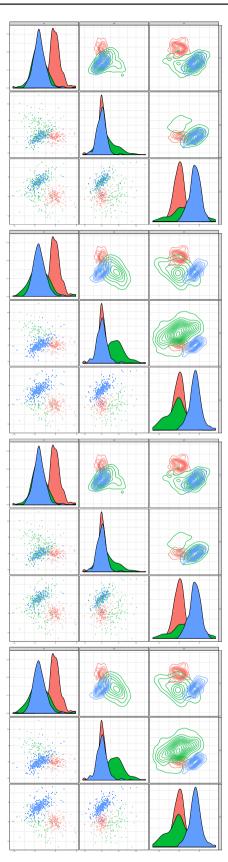
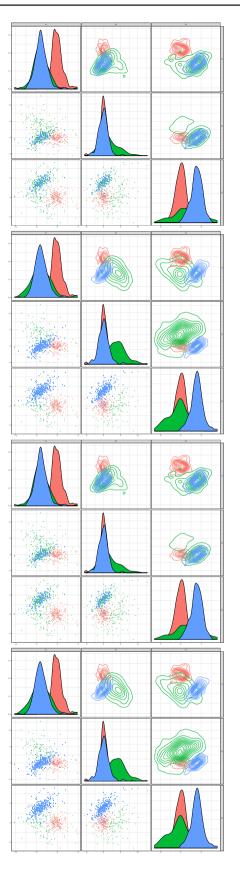


Fig. 8 Simulated datasets using multivariate Gaussian distributions Scenarios 13–16, each color and symbol representing a different cluster

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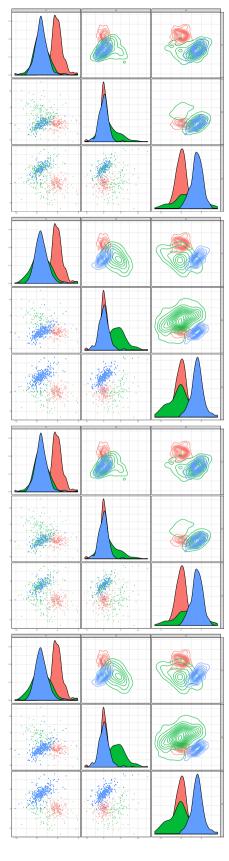
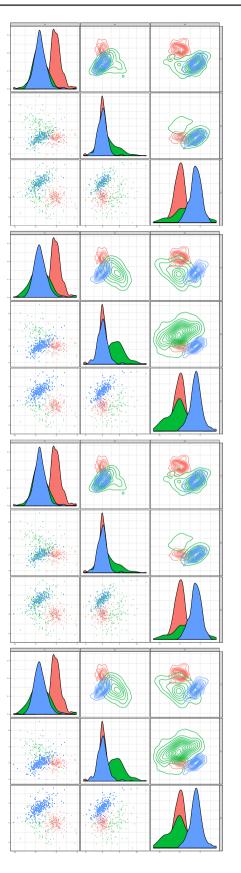


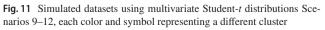
Fig.9 Simulated datasets using multivariate Student-*t* distributions Scenarios 1–4, each color and symbol representing a different cluster

Fig. 10 Simulated datasets using multivariate Student-*t* distributions Scenarios 5–8, each color and symbol representing a different cluster

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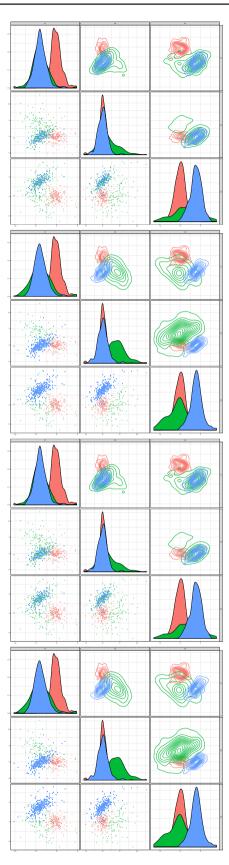


Fig. 12 Simulated datasets using multivariate Student-*t* distributions Scenarios 13–16, each color and symbol representing a different cluster

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