

NON-NESTED HYPOTHESIS TESTS FOR VINE COPULAS AND STATISTICAL  
LEARNING TECHNIQUES IN PROCESS MONITORING

By

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A dissertation submitted in partial fulfillment of  
the requirements for the degree of

DOCTOR OF PHILOSOPHY

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To the Faculty of Washington State University:

The members of the Committee appointed to examine the dissertation of  
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NON-NESTED HYPOTHESIS TESTS FOR VINE COPULAS AND STATISTICAL  
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Abstract

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In this dissertation, we first introduce three bootstrap-based non-nested hypothesis tests for regular vine-copulas. These test statistics are derived from log-likelihood ratio test statistics and Cox test statistics. This study presents the power study comparing the proposed tests with existing vine-copula non-nested hypothesis tests. Across models with varying structures of regular copulas, our hypothesis tests consistently achieve higher power. Deriving from statistical algorithms, we also propose two different control charts that can be applied to multivariate statistical process control (MSPC). The first one is based on support vector data description (SVDD). We propose a SVDD control chart using the Mahalanobis distance kernel (Mahalanobis k-chart). Mathematical illustrations and statistical comparisons are presented on the

basis of both simulations and a real example of electricity consumption. The results show that Mahalanobis k-chart can achieve lower Phase II average run length (ARL) in most shifted-process scenarios. Another control chart we propose is random oversampling gradient boosting real-time contrasts (ROGB-RTC) chart. Real-time contrasts (RTC) control charts convert the statistical process monitoring problem into a dynamic binary classification problem. But only a limited number of RTC studies have scrutinized the imbalance problem between the sample size of the reference data and that of the sliding window data. Our control chart handles the imbalance problem in terms of both classifier and monitoring statistics. Experiments show that the proposed method achieves better performance than that of the original real-time contrasts chart.

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PREVIEW

# CHAPTER 1. NON-NESTED HYPOTHESIS TEST FOR VINE COPULAS

## Abstract

We introduce three bootstrap-based non-nested hypothesis tests for regular vine-copulas. These test statistics are derived from log-likelihood ratio test statistics and Cox test statistics. This study presents a power study that compares the proposed tests with existing vine-copula non-nested hypothesis tests. Across models with varying structures of regular copulas, our hypothesis test procedures consistently achieve higher power.

### 1.1 Introduction

A copula is a multivariate distribution with a uniform marginal probability distribution. It is useful to model the dependence structure among variables since it allows to model dependencies and marginal distributions separately. Copula based models are widely used in risk management and option pricing. Model selection on copula-based models is an essential part of empirical work. However, there is a limited

number of studies working on non-nested hypothesis testing on copulas, vine copulas in particular.

Suppose there are two models, say  $f$  and  $g$ , which may or may not be different from each other. If one model  $f$  cannot be derived from the other by parametric restriction or process limiting. then these two models are called “non-nested”. To compare two non-nested models, one approach is to conduct a likelihood ratio test which tests the null hypothesis  $H_0$  (model  $f$ ) is significantly better than the alternative hypothesis  $H_1$  (model  $g$ ). For non-nested models, the log-likelihood ratio statistic is  $\lambda = l_f - l_g$ , where  $l_f$  and  $l_g$  are the log-likelihood under null and alternative models. Nevertheless, the non-nested condition violates the asymptotical Chi-square distribution when doubling log-likelihood difference. In this scenario, one can still conduct a hypothesis test based on the asymptotic distribution of test statistics. In a seminal study, Cox [15] develops an adjusted log-likelihood ratio statistic for non-nested test. In line with Cox’s work, some researchers use simulation approaches to compute which still assumes an asymptotic distribution (Pesaran and Pesaran [32], [62], and Weeks [80]). Moreover, when various asymptotic ways are used to estimate the variance of Cox test statistic, they give rise to variants of Cox test statistics. In contrast to previous approaches based on Cox test statistic, Vuong [78] proposes a test based on the Kullback-Leibler information criterion  $KLIC = E_0(l_f - l_g)$ , where  $E_0$  is the expected value under the true model. He proves under the regular conditions,

the likelihood ratio statistic with maximum likelihood estimators of parameters is asymptotically normally distributed. Following Vuong's work, Clarke [11] proposes a distribution-free paired sign test with an asymptotic Binomial distribution under the null hypothesis.

Apart from the above approaches to find an asymptotic distribution, another manner is to use bootstrap (Efron[20]). Beran [3] uses a parametric bootstrap procedure to approximate the distribution of test statistics. In this case, the significance of the test is evaluated by the observed test statistic and the empirical reference distribution.

Some researchers have also combined a variety procedures to improve the non-nested hypothesis. For example, Kapetanios and Weeks [44] combine both simulation and bootstrap procedures in their non-nested hypothesis test to adjust both the likelihood ratio and Cox test statistics. Comparing several bootstrap procedures with test statistics on asymptotic approximation, they conclude that a simple bootstrap procedure has a significantly better performance than that of complicated bootstrap procedures and asymptotic procedures.

This paper is outlined as follows. Section 1.2 introduces copula and vine copula models. Section 1.3 describes the Cox test statistic and the bootstrap procedure. In Section 1.3, we perform a simulation study with various 5-dimensional vine copulas and compared the results of different tests in terms of power. Finally, Section 1.5 summarizes the results and discusses some potential future work.

## 1.2 Copula and Regular Vine Copula

### 1.2.1 Definitions

**Definition 1.1.** A  $d$ -dimensional copula,  $C : [0, 1]^d \rightarrow [0, 1]$  is a cumulative distribution function (CDF) with uniform marginals.

We use the notation  $C(\mathbf{u}) = C(u_1, u_2, \dots, u_d)$  to denote a copula. The distribution condition on  $C$  has the following properties:

- $C(u_1, u_2, \dots, u_d)$  is always non-decreasing in each component  $u_i$ .
- Since  $C$  is uniformly distributed, the  $i^{th}$  marginal distribution is obtained by setting  $u_j = 1$  for all  $j \neq i$ ,

$$C(1, \dots, 1, u_i, 1, \dots, 1) = u_i.$$

- For  $a_i \leq b_i$ ,  $P(U_1 \in [a_1, b_1], \dots, U_d \in [a_d, b_d])$  must be non-negative. This implies the rectangle inequality

$$\sum_{i_1=1}^2 \cdots \sum_{i_d=1}^2 (-1)^{i_1+\dots+i_d} C(u_{1,i_1}, \dots, u_{d,i_d}) \geq 0,$$

where  $u_{j,1} = a_j$  and  $u_{j,2} = b_j$ , for  $j = 1, \dots, d$

Any function that satisfies those 3 properties above is a copula.

### 1.2.2 Sklar's theorem

Let  $\mathbf{X} = (X_1, \dots, X_d)$  be a multivariate random vector with joint cdf  $F_X$  with continuous and increasing marginals. It can be easily verified that the joint distribution of  $F_1(X_1), \dots, F_d(X_d)$  is a copula. With this result, Sklar's theorem provides a theoretical foundation for obtaining multivariate distributions with a copula and marginal distributions.

**Theorem 1.1** (Sklar's Theorem). *Consider a  $d$ -dimensional cdf  $F$  with marginal cdf's  $F_1, \dots, F_d$ . Then there exists a copula  $C$  such that*

$$F(x_1, \dots, x_d) = C(F_1(x_1), \dots, F_d(x_d)) \quad (1.1)$$

for all  $x_i \in (-\infty, \infty)$  and  $i = 1, \dots, d$ .

If  $F_i$  is continuous for all  $i = 1, \dots, d$ , then  $C$  is unique. Otherwise  $C$  is uniquely determined on  $Ran(F_1) \times \dots \times Ran(F_d)$ , where  $Ran(F_i)$  is the range of cdf  $F_i$ .

Consider a copula  $C$  and univariate cdf's  $F_1, \dots, F_d$ . Then  $F$  as defined in (1.1) is a multivariate cdf with marginal cdf's  $F_1, \dots, F_d$

If all marginal cdf's  $F_1, \dots, F_n$  are continuous, it can be obtained that

$$F_i(F_i^\leftarrow(y)) = y.$$

where  $F^\leftarrow(x) := \inf\{v : f(v) \geq x\}$ . Apply above equation to (1.1), we obtain

$$C(\mathbf{u}) = F(F_1^\leftarrow(u_1), \dots, F_d^\leftarrow(u_d)).$$

### 1.2.3 Copula density and conditional distribution

The copula density can be obtained as below:

$$c(\mathbf{u}) := \frac{\partial^d C(\mu_1, \dots, \mu_d)}{\partial \mu_1 \dots \partial \mu_d},$$

if it exists. When the marginal cdf  $F_i$  is differentiable, we have  $F_i^\leftarrow = F_i^{-1}$ . If the copula is in the form of (1.2) and has a density, the copula density can be computed as

$$c(\mathbf{u}) = \frac{f(F_1^{-1}(u_1), \dots, F_d^{-1}(u_d))}{f_1(F_1^{-1}(u_1)) \dots f_d(F_d^{-1}(u_d))},$$

where  $f$  is the joint density and  $f_i$ ,  $i = 1, \dots, d$ , are the marginal densities. Assuming sufficient regularity, the copula conditional cdf can be obtained by

$$P(U_2 \leq u_2 | U_1 = u_1) = \frac{\partial}{\partial u_1} C(u_1, u_2)$$

Thus, the conditional cdf of a copula can be directly derived from itself.

### 1.2.4 Pair-copula constructions (PCC)

Copula models are powerful tools for modeling the dependence structure. However, traditional multivariate copulas such as Archimedean copulas cannot model different dependencies between pairs of variables in high-dimension. A flexible way

to construct high-dimentional copulas is to construct multivariate densities with  $d(d - 1)/2$  bivariate copulas (Joe [41] , Bedford and Cooke [1],[2]).

Let  $X_1, \dots, X_d$  be a random variables with a joint distribution  $F$  and a density  $f$ . We can decompose the joint density  $f$  by

$$f(x_1, \dots, x_d) = f(x_d|x_1, \dots, x_{d-1})f(x_1, \dots, x_{d-1}) \quad (1.2)$$

$$= \prod_{i=2}^d f(x_i|x_1, \dots, x_{i-1}) \times f(x_1) \quad (1.3)$$

where  $F(\cdot|\cdot)$  and  $f(\cdot|\cdot)$  are the conditional cdf and the density function, respectively.

If we apply Sklar's theorem with  $d = 2$ , we can derive

$$f(x_1, x_2) = c_{12}(F_1(x_1), F_2(x_2)) \cdot f_1(x_1) \cdot f_2(x_2) \quad (1.4)$$

where  $c_{12}$  is a bivariate copula density. With (1.4), the conditional density of  $X_2$  given  $X_1$  is computed as

$$f(x_1|x_2) = c_{12}(F_1(x_1), F_2(x_2)) \cdot f_1(x_1) \quad (1.5)$$

Using Equation (1.3), (1.4) and (1.5), the pair-copula construction in  $d$  dimensions is listed as follows:

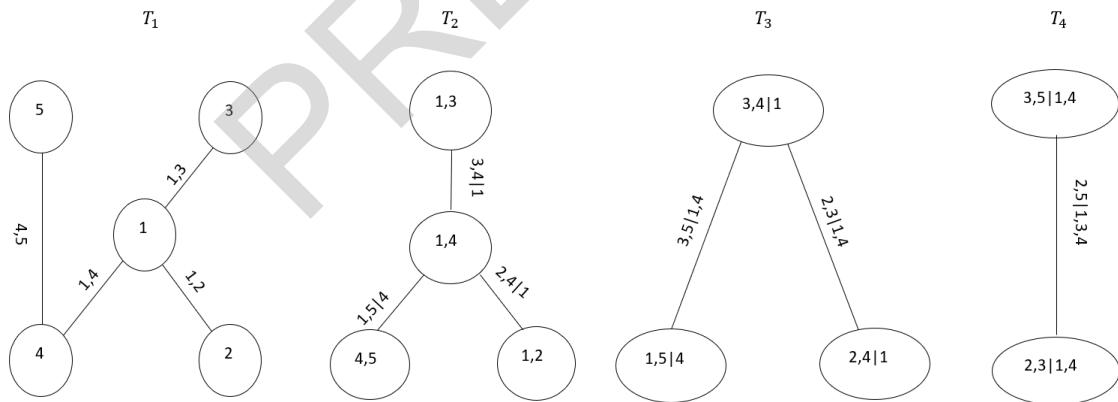
$$f(x_1, \dots, x_d) = \prod_{j=1}^{d-1} \prod_{i=1}^{d-j} c_{i,(i+j)|(i+1),\dots,(i+j-1)} \cdot \prod_{k=1}^d f_k(x_k) \quad (1.6)$$

where  $c_{i,j|i_1,\dots,i_k} := c_{i,j|i_1,\dots,i_k}(F(x_i|x_{i_1}, \dots, x_{i_k}), F(x_j|x_{i_1}, \dots, x_{i_k}))$ , for  $i, j, i_1, \dots, i_k = 1, \dots, d$  with  $i < j$  and  $i_1 < \dots < i_k$ .

### 1.2.5 Regular vine copula

Bedford and Cooke[1],[2] use trees to represent the graphic structure of multivariate copulas. Denote trees by  $T_i = (V_i, E_i)$ ,  $i = 1, \dots, d-1$ , where  $V_i$  is the set of nodes and  $E_i$  denotes the set of edges. A vine is a sequence of trees  $\nu = (T_1, \dots, T_{d-1})$  that satisfies the following conditions:

- $T_i$  is a tree with nodes  $V_i = 1, \dots, d$  and edges  $E_i$ .
- For  $i \geq 2$ ,  $T_i$  is a tree with nodes  $V_i = E_{i-1}$  and edges  $E_i$ .
- Two nodes in the tree  $T_{i+1}$  are joined by an edge only if their corresponding edges in  $T_i$  share a common node.

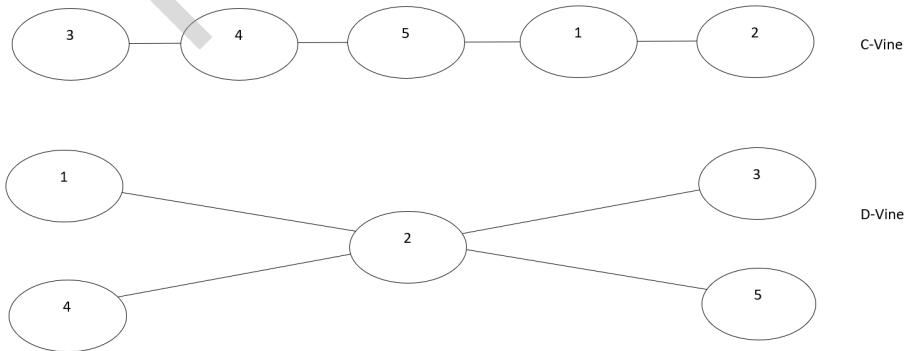


**Figure 1.1:** The tree structure of a 5 dimensional R-vine copula

Figure 1.1 is an example of a 5-dimensional R-vine. Tree  $T_1$  has nodes  $V_1 = 1, \dots, 5$

and edges  $E_1 = \{\{1, 2\}, \{1, 3\}, \{1, 4\}, \{4, 5\}\}$  with unconditional pair-copulas. For trees  $T_2, \dots, T_4$ , their edges  $E_2, \dots, E_4$  form conditional pair-copulas. The joint copula density of the 5-dimensional R-vine copula in Figure 1.1 can be decomposed as

$$\begin{aligned}
& c_{12345}(u_1, \dots, u_5) \\
&= c_{1,2}(u_1, u_2) \cdot c_{1,3}(u_1, u_3) \cdot c_{1,4}(u_1, u_4) \cdot c_{4,5}(u_4, u_5) \\
&\quad \cdot c_{1,5:4} (C_{1|4}(u_1|u_4), C_{5|4}(u_5|u_4)) \cdot c_{2,4;1} (C_{2|1}(u_2|u_1), C_{4|1}(u_4|u_1)) \\
&\quad \cdot c_{3,4;1} (C_{3|1}(u_3|u_1), C_{4|1}(u_4|u_1)) \\
&\quad \cdot c_{2,3;1,4} (C_{2|1,4}(C_{2|1}(u_2|u_1), C_{4|1}(u_4|u_1)), C_{3|1,4}(C_{3|1}(u_3|u_1), C_{4|1}(u_4|u_1))) \\
&\quad \cdot c_{3,5;1,4} (C_{3|1,4}(C_{3|1}(u_2|u_1), C_{4|1}(u_4|u_1)), C_{5|1,4}(C_{5|1}(u_5|u_1), C_{4|1}(u_4|u_1))) \\
&\quad \cdot c_{2,5;1,3,4} (C_{2|1,3,4}(C_{2|1,4}(C_{2|1}(u_2|u_1), C_{4|1}(u_4|u_1)), C_{3|1,4}(C_{3|1}(u_3|u_1), C_{4|1}(u_4|u_1))), \\
&\quad C_{5|1,3,4}(C_{3|1,4}(C_{3|1}(u_3|u_1), C_{4|1}(u_4|u_1)), C_{5|1,4}(C_{1|4}(u_1|u_4), C_{5|4}(u_5|u_4)))) .
\end{aligned}$$



**Figure 1.2:** The first tree structures of C-vine and D-vine Copulas

Figure 1.2 is an example of two special R-vine tree structures, namely, C-vine and D-vine. C-vine has one root node with degree  $d - 1$ , that connects to another  $d - 1$  nodes. All other nodes only have degree 1. In D-vine, two nodes have degree 1 and others have degree 2.

### 1.3 Cox Test Statistic and Bootstrap Procedure

#### 1.3.1 Cox Test statistic

Consider two models,  $F_\theta = f(\mathbf{y}_i|\mathbf{x}_i, \theta)$  and  $G_\lambda = g(\mathbf{y}_i|\mathbf{x}_i, \lambda)$ , where  $\theta$  and  $\lambda$  are two vectors of unknown parameters belonging to two respective parameter spaces  $\Theta$  and  $\Lambda$ . We write the null and alternative hypotheses of a test as

$$\mathbf{H}_0 : F = F_\theta$$

$$\mathbf{H}_a : F = G_\lambda$$

Here  $F$  is the true population distribution. To simplify notations, we use  $f_i(\theta)$  and  $g_i(\lambda)$  to represent  $f(\mathbf{y}_i|\mathbf{x}_i, \theta)$  and  $g(\mathbf{y}_i|\mathbf{x}_i, \lambda)$ . The Cox test statistic is obtained by

$$T_f = l_f(\hat{\theta}) - l_g(\hat{\lambda}) - C_{fg}(\hat{\theta}, \hat{\lambda})$$

where  $l_f(\hat{\theta})$  and  $l_g(\hat{\lambda})$  are the average log-likelihood functions

$$l_f(\hat{\theta}) = \frac{1}{n} \sum_{i=1}^n \log f_i(\hat{\theta}), \quad l_g(\hat{\lambda}) = \frac{1}{n} \sum_{i=1}^n \log g_i(\hat{\lambda}),$$