Bayesian Nonparametric Conditional Copula Estimation of Twin Data*

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Abstract

Several studies on heritability in twins aim at understanding the different contribution of environmental and genetic factors to specific traits. Considering the National Merit Twin Study, our purpose is to correctly analyse the influence of the socioeconomic status on the relationship between twins’ cognitive abilities. Our methodology is based on conditional copulas, which allow us to model the effect of a covariate driving the strength of dependence between the main variables. We propose a flexible Bayesian nonparametric approach for the estimation of conditional copulas, which can model any conditional copula density. Our methodology extends the work of Wu et al. (2015) by introducing dependence from a covariate in an infinite mixture model. Our results suggest that environmental factors are more influential in families with lower socio-economic position.

Keywords: Bayesian nonparametrics, Conditional Copula models, Slice sampling, National Merit Twin Study, Social Science.

1 Introduction

The literature on heritability of traits in children often focusses on twins, due to the shared environmental factors and the association of genetical characteristics. Among studies on the heritability of diseases, Wang et al. (2011) applied an efficient estimation method to mixed-effect models to analyze disease inheritance in twins.

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One of the main purposes of studies on heritability is to estimate the different contribution of genetic and environmental factors to traits or outcomes (see, for example, the latent class twin method of Baker (2016)). Bates et al. (2013) studied the interactions between environmental and genetic effects to intelligence in twins, showing that higher socioeconomic status is associated with higher intelligence scores. Bioecological theory states that environmental factors may significantly influence the heritability of certain characteristics, such as cognitive ability, which is the readiness for future intellectual or educational pursuits. Several studies have found that cognitive ability is more pronounced and evident among children raised in higher socioeconomic status families. Such families can offer greater opportunities to children, due to their socioeconomic wealth status, and represent stimulating environments where children’s inherited capabilities may become more manifest.

The aim of this paper is to correctly analyse the effect of socioeconomic factors on the relationship between twins’ cognitive abilities. From a sample of 839 US adolescent twin pairs who completed the National Merit Scholarship Qualifying Test, we consider each twin’s overall school performance (measured by a total score including English, Mathematics, Social Science, Natural Science and Word Usage), the mother’s and father’s education level and the family income. The data are plotted in Figure 1, which shows the scatterplots of the twins’ school performances, on each axis, against the socioeconomic variables, whose values are in different colours (dark brown denotes low values, while light brown denotes high values). Figure 1 indicates that the twins’ school performances are positively correlated and their dependence is influenced by the values of the socioeconomic variables (the mother’s (panel (a)), the father’s level of education (panel (b)) and the family income (panel (c))). Indeed, most of the light brown dots (denoting high values of the covariates) are grouped in the upper right corner, while the dark brown dots (denoting low values of the covariates) lie in the bottom left corner of each plot. Hence, the higher the parents’ education or family income, the higher the twins’ school performance. This means that the twins’ performance scores are functions of each covariate and they vary according to the values of the covariates.

In Figure 2 we discretized the covariates into low, medium and high values and we produced scatterplots of the twins’ school performance scores. The top plots refer to the mother’s level of education, the central plots refer to the father’s level of education and the bottom plots refer to the family income. From left to right, the plots correspond to low [0,2), medium [2,4) and high [4,6] ([4,7] for the family income) levels of the covariates. As we move from low to high levels of the covariates, the point clouds tend to change shape around the diagonal and move to the upper right corner. Therefore, in all three cases, as already pointed out, low values of covariates correspond to low performance scores, while high values of covariates correspond to high performance scores. In addition, the different shapes of the scatterplot points corresponding to low, medium and high level of covariates suggest that the dependence structure between the twins’ school outcomes changes according to the levels of the covariates.
Therefore, a flexible model, able to capture the effect of a covariate on the dependence between the kids’ performance scores is necessary.

In order to model the dependence structure between the twins’ school performances, we use copulas, which are popular modeling approaches in multivariate statistics allowing the separation of the marginal components of a joint distribution from its dependence structure. More precisely, Sklar (1959) proved that a $d$-dimensional distribution $H$ of the random variables $Y_1, \ldots, Y_d$ can be fully described by its marginal distributions and a function $C : [0, 1]^d \rightarrow [0, 1]$, called copula, through the relation $H(y_1, \ldots, y_d) = C(F_1(y_1), \ldots, F_d(y_d))$. In the literature, copulas have been applied to model the dependence between variables in a wide variety of fields (see Kolev et al. (2006) and Cherubini et al. (2004)). In particular, applications of copula models involved lifetime data analysis (Andersen (2005)), survival analysis of Atlantic halibut (Brackers and Veraverbeke (2005)) and transfusion-related AIDS and cancer analysis (Emura and Wang (2012), Huang and Zhang (2008) and Owzar et al (2007)).
Figure 2: Scatterplots of the twins overall scores with respect to the mother’s (top panels) and father’s level of education (middle panels) and family income (bottom panels). From left to right, the plots correspond to low [0,2), medium [2,4) and high [4,6] ([4,7] for the family income) levels of the covariates. The black dashed line corresponds to the 45 degrees diagonal.

The introduction of covariate adjustments to copulas has attracted an increased interest in recent years. [Craiu and Sabeti (2012)] propose a conditional copula approach in regression settings where the bivariate outcome can be continuous or mixed. [Patton (2006)] introduces time-variation in the dependence structure of ARMA models (see also [Jondeau and Rockinger (2006)] and [Bartram et al. (2007)] for other applications of time-series analysis to dependence modelling). The paper of [Acar et al. (2010)] provides a nonparametric procedure to estimate the functional relationship between copula parameters and covariates, showing that the gestational age drives the strength of dependence between the birth weights of twins. [Abegaz et al. (2012)] and [Gijbels et al. (2012)] propose semiparametric and nonparametric methodologies for the estimation of conditional copulas, establishing consistency and asymptotic normality results for the estimators. The methodology is then applied to examine the influence of the gross domestic product (GDP), in USD per capita, on the life expectancy of males and females at birth.

In a similar vein, parametric models such as Bayesian regression copulas allow the
specification of Bayesian marginal regressions for a set of outcomes, linking the marginals to covariates, and combining them via a copula to form a joint model. The general framework of Bayesian Gaussian regression copulas with discrete, continuous or mixed outcomes is presented by Pitt et al. (2006) and allows to handle a multivariate regression with Gaussian and non-Gaussian marginal distributions. Yin and Yuan (2009) adopt a Bayesian regression copula model in cancer clinical trials for dose finding to account for the synergistic effect of combinations of multiple drugs. A copula constructed from the skew t distribution is employed by Smith et al. (2012) to capture asymmetric and extreme dependence between variables modelled via Bayesian marginal regressions. While most Bayesian regression copula models focus on covariate adjustments for the marginals, recently Klein and Kneiss (2016) proposed simultaneous Bayesian inference for both the marginal distributions and the copula. Other contributions along the same lines are Taglioni et al. (2016), Stander et al. (2015a) and Stander et al. (2015b). However, the authors selected the copula family by using the deviance information criterion, which may suffer from limitations, as discussed for example by Plummer (2008). Indeed, the choice of the copula family may be controversial and it is still an open problem (see Joe (2014)). The literature offers a rich range of copula families, such as elliptical copulas (e.g. Gaussian and Student's t) and archimedean copulas (e.g. Frank, Gumbel, Clayton and Joe) to accommodate various dependence structures. In this paper, we adopt a Bayesian nonparametric approach which allows us to overcome the issue of the choice of copula and we adopt a conditional copula approach to model the effect of a covariate on the dependence between variables. Our methodology builds on Wu et al. (2015), who propose a Bayesian nonparametric procedure to estimate any unconditional copula density function. The authors combine the well-known Gaussian copula density with the modeling flexibility of the Bayesian nonparametric approach, proposing to use an infinite mixture of Gaussian copulas. Burda and Prokhorov (2014) propose to use nonparametric univariate Gaussian mixtures for the marginals and a multivariate random Bernstein polynomial copula for the link function under the Dirichlet process prior. Our paper extends the work of Wu et al. (2015) to the conditional copula setting, by proposing a novel methodology which combines the advantages of a conditional copula approach with the modeling flexibility of Bayesian nonparametrics. In particular, we include a conditional covariate component to explain the variable dependence structure, allowing us further flexibility to the copula density modelling. Up to our knowledge, this is the first Bayesian nonparametric proposal in the conditional copulas literature.

The outline of the paper is the following. In Section 2 we briefly review the literature about conditional copulas and Bayesian nonparametric copula estimation. In Section 3 we introduce our novel Bayesian nonparametric conditional copula setting. Section 4 provides an algorithm for estimating the posterior parameters and Section 5 illustrates the performance of the methodology. Section 6 is devoted to the application of our methodology to the analysis of the National Merit Twin Study. Concluding remarks are given in Section 7.
2 Preliminaries

In this Section, we review some preliminary notions about conditional copulas and illustrate the Bayesian nonparametric copula density estimation introduced in Wu et al. (2015). In what follows, we focus on the bivariate case for simplicity, however the arguments can be easily extended to more than two dimensions.

2.1 The conditional copula

Let $Y_1$ and $Y_2$ be continuous variables of interest and $X$ be a covariate that may affect the dependence between $Y_1$ and $Y_2$. Following Gijbels et al. (2012), Abegaz et al. (2012) and Acar et al. (2010), we suppose that the conditional distribution of $(Y_1, Y_2)$ given $X = x$ exists and we denote the corresponding conditional joint distribution function by

$$H_x(y_1, y_2) = P(Y_1 \leq y_1, Y_2 \leq y_2 | X = x).$$

If the marginals of $H_x$, denoted as $F_{1x}(y_1) = P(Y_1 \leq y_1 | X = x), F_{2x}(y_2) = P(Y_2 \leq y_2 | X = x)$, are continuous, then according to Sklar’s theorem there exists a unique copula $C_x$ which equals

$$C_x(u, v) = H_x(F_{1x}^{-1}(u), F_{2x}^{-1}(v))$$

where $F_{1x}^{-1}(u) = \inf\{y_1 : F_{1x} \geq u\}$ and $F_{2x}^{-1}(v) = \inf\{y_2 : F_{2x} \geq v\}$ are the conditional quantile functions and $u = F_{1x}(y_1)$ and $v = F_{2x}(y_2)$ are called pseudo-observations. The conditional copula $C_x$ fully describes the conditional dependence structure of $(Y_1, Y_2)$ given $X = x$. An alternative expression for (1) is

$$H_x(y_1, y_2) = C_x(F_{1x}(y_1), F_{2x}(y_2)).$$

2.2 Bayesian nonparametric copula density estimation

Let $\Phi_{\rho}(y_1, y_2)$ denote the standard bivariate normal distribution function with correlation coefficient $\rho$. Then, $C_\rho$ is the copula corresponding to $\Phi_{\rho}$, taking the form:

$$C_\rho(u, v) = \Phi_\rho(\Phi^{-1}(u), \Phi^{-1}(v))$$

where $\Phi$ is the univariate standard normal distribution function. The Gaussian copula density is:

$$c_\rho(u, v) = |\Sigma|^{-\frac{1}{2}} \exp\left\{-\frac{1}{2}(\Phi^{-1}(u), \Phi^{-1}(v))(\Sigma^{-1} - I)(\Phi^{-1}(u), \Phi^{-1}(v))\right\}$$
where the correlation matrix is:
\[ \Sigma = \begin{bmatrix} 1 & \rho \\ \rho & 1 \end{bmatrix}. \]

Wu et al. (2015) proposed to use an infinite mixture of Gaussian copulas for the estimation of a copula density, as follows
\[ c(u, v) = \sum_{j=1}^{\infty} w_j c_{\rho_j}(u, v) \] (5)

where the weights \( w_j \)'s sum up to 1 and the \( \rho_j \)'s vary in \((-1, 1)\). Given a set of \( n \) observations \((u_1, v_1), \ldots, (u_n, v_n)\), their model can be described through a hierarchical specification, i.e.
\[ (u_i, v_i) \mid \rho_i \overset{\text{ind}}{\sim} c_{\rho_i}(u_i, v_i), \quad i = 1, \ldots, n, \]
\[ \rho_i \mid G \overset{\text{ind}}{\sim} G, \]
\[ G \sim DP(\lambda, G_0), \] (6)

where \( G \) is a Dirichlet Process prior with total mass \( \lambda \) and base measure \( G_0 \). This proposal is motivated by the fact that bivariate density functions on the real plain can be arbitrarily well approximated by a mixture of a countably infinite number of bivariate normal distributions of the form
\[ f(y_1, y_2) = \sum_{j=1}^{\infty} w_j N((y_1, y_2) | (\mu_{1j}, \mu_{2j}), \Sigma_j) \]

where \( N((y_1, y_2) | (\mu_{1j}, \mu_{2j}), \Sigma_j) \) is the joint bivariate normal density with mean vector \((\mu_{1j}, \mu_{2j})\) and correlation matrix \( \Sigma_j \) (see Lo (1984) and Ferguson (1983)). Roughly speaking, the authors are mimicking the Dirichlet process mixture model in the copula setting (see Escobar (1994) and Escobar and West (1995)). The sampling strategy follows the slice sampler of Walker (2007) and Kalli et al. (2011). The authors show that the Gaussian mixture is flexible enough to accurately approximate any bivariate copula density.

3 Conditional copula estimation with Dirichlet process priors

The data object of study requires a model which can take into account the effect of a covariate. We build on the model introduced by Wu et al. (2015) and illustrated in the previous section. The idea is to replace the Gaussian copula with a conditional version where the correlation is a function of the covariate, i.e.
\[ c_{\rho}(u, v | x) = c_{\rho(x)}(u, v). \]
The function \( \rho(x) \) can be modelled as preferred, for instance, with a generalized linear model or with a non-linear function. In any case, we have that \( \rho(x) \) will depend on a vector of parameters \( \beta \), so that

\[
c_{\rho(x)}(u, v) = c_{\rho(x|\beta)}(u, v).
\]

We assume a Dirichlet process prior on the vector of parameters \( \beta = (\beta_1, \ldots, \beta_d) \). Following the model description provided in equation (6), we can summarize our model as follows,

\[
(u_i, v_i) \mid \rho(x_i|\beta_i) \overset{\text{iid}}{\sim} c_{\rho(x_i|\beta_i)}(u_i, v_i), \quad i = 1, \ldots, n,
\]

\[
\beta_i \mid G \overset{\text{iid}}{\sim} G, \\
G \sim DP(\lambda, G_0),
\]

(7)

where \( G \) is a Dirichlet process prior with total mass \( \lambda \) and base measure \( G_0 \). As in Wu et al. (2015), our model can be described as an infinite mixture of Normal distributions,

\[
c_{\rho}(u, v|x) = \sum_{j=1}^{\infty} w_j c_{\rho(x|\beta_j)}(u, v),
\]

(8)

and hence suitable for implementing a slice sampling algorithm, as explained in the next section.

In order to model the function \( \rho(x|\beta) \), we would like to follow some standard approaches in the literature. Abegaz et al. (2012) model the dependence of the parameter of interest, with respect to the covariate, through a calibration function \( \theta(x|\beta) \). It is important to highlight that in many copula families the parameter space is restricted. In contrast, the calibration function \( \theta(x|\beta) \) can assume any value on the real line. In our case, the parameter is restricted to the interval \((-1, 1)\) and we need a transformation which can link the calibration function \( \theta(x|\beta) \) to \( \rho(x|\beta) \). In this paper, we adopt the following transformation,

\[
\rho(x|\beta) = \frac{2}{|\theta(x|\beta)| + 1} - 1.
\]

In our simulated and real data examples we focus on two particular calibration functions studied in the literature, which are

\[
\theta(x|\beta) = \beta_1 + \beta_2 x^2
\]

\[
\theta(x|\beta) = \beta_1 + \beta_2 x + \beta_3 \exp(-\beta_4 x^2)
\]

respectively, such that \( \theta(x|\beta) \in (-\infty, +\infty) \) and, consequently, \( \rho(x|\beta) \in (-1, 1) \).
4 Posterior sampling algorithm

The observations \((y_1, y_2, \ldots, y_n)\), for \(i = 1, \ldots, n\), are transformed into the corresponding pseudo-observations \((u_i, v_i)\) using a nonparametric estimation approach, as in Gijbels et al. (2011). The pseudo-observations are then plugged into the copula. Following equation \((8)\), given \((u_i, v_i)\) for \(i = 1, \ldots, n\), and the conditional variable \(x_i\), the conditional copula density function for each pair \((u_i, v_i)\) can be written as an infinite mixture of conditional Gaussian copulas, such that:

\[
c(u_i, v_i | x_i) = \sum_{j=1}^{\infty} w_j c_{\rho(x_i|\beta_j)}(u_i, v_i)
\]  

(9)

where \(w_j\)'s are the stick-breaking weights, i.e.

\[
w_j = \pi_j \prod_{l=1}^{j-1} (1 - \pi_l)
\]

where the \(\pi_j\) are distributed as a \(\text{Be}(1, \lambda)\), \(\lambda > 0\). In order to sample from the infinite mixture displayed in equation \((9)\), we use the slice sampling algorithm for mixture models proposed by Walker (2007) and Kalli et al. (2011). To reduce the dimensionality of the problem, the authors introduce a latent variable \(z_i\) for each \(i\) which allows us to write the infinite mixture model as follows:

\[
c(u_i, v_i, z_i | x_i) = \sum_{j=1}^{\infty} I(z_i < w_j) c_{\rho(x_i|\beta_j)}(u_i, v_i).
\]

(10)

The introduction of the slice variable \(z_i\) reduces the sampling complexity to the analogous of a finite mixture model. In particular, letting

\[
A_w = \{j : z_i < w_j\},
\]

(11)

then it can be proved that the cardinality of the set \(A_w\) is almost surely finite. Consequently, there is a finite number of parameters to be estimated. By iterating the data augmentation principle further, we introduce another latent variable \(d_i\), which is called allocation variable, allowing us to allocate each observation to one component of the mixture model. Then, the conditional copula density \(c(u_i, v_i, z_i, d_i | x_i)\) takes the form:

\[
c(u_i, v_i, z_i, d_i | x_i) = I(z_i < w_{d_i}) c_{\rho(x_i|\beta_{d_i})}(u_i, v_i)
\]

(12)

where \(d_i \in \{1, 2, \ldots\}\). Hence, the full likelihood function of the conditional copula model is:

\[
\prod_{i=1}^{n} c(u_i, v_i, z_i, d_i | x_i) = \prod_{i=1}^{n} I(z_i < w_{d_i}) c_{\rho(x_i|\beta_{d_i})}(u_i, v_i).
\]

(13)
We use the notation \((U, V) = \{i = 1, \ldots, n : (u_i, v_i)\}, X = \{x_1, \ldots, x_n\}\) to describe the pseudo-observations and the covariate values, respectively. We denote with \(\beta = \{\beta_1, \beta_2, \ldots\}\) the vector of parameters and \(D = \{d_1, \ldots, d_n\}, Z = \{z_1, \ldots, z_n\}\) and \(\pi = \{\pi_1, \pi_2, \ldots\}\) the new variables introduced so far.

![Scatter plots and histograms](image1)

(a) Simulated, 1st cal. fun.  (b) Simulated, 1st cal. fun.  (c) Predictive, 1st cal. fun.  (d) Predictive, 1st cal. fun.

![Scatter plots and histograms](image2)

(e) Simulated, 2nd cal. fun.  (f) Simulated, 2nd cal. fun.  (g) Predictive, 2nd cal. fun.  (h) Predictive, 2nd cal. fun.

Figure 3: Gaussian copula with sample size \(n = 500\). Panels (a), (b), (c) and (d) depict the scatter plots and histograms, obtained with the first calibration function, of the simulated and predictive samples, respectively; panels (e), (f), (g) and (h) depict the scatter plots and histograms, obtained with the second calibration function, of the simulated and predictive sample, respectively.

Therefore, we used a Gibbs sampler to simulate iteratively from the posterior distribution function, according to the following steps:

1. The stick-breaking components \(\pi\) are updated given \([Z, D, \beta, (U, V), X]\);
2. The latent slice variables \(Z\) are updated given \([\pi, D, \beta, (U, V), X]\);
3. The allocation variables \(D\) are updated given \([\pi, Z, \beta, (U, V), X]\);
4. The vector of parameters \(\beta\) is updated given \([\pi, Z, D, (U, V), X]\).

The Gibbs sampling details are explained in Appendix A.
Figure 4: Frank copula with sample size $n = 500$. Panels (a), (b), (c) and (d) depict the scatter plots and histograms, obtained with the first calibration function, of the simulated and predictive samples, respectively; panels (e), (f), (g) and (h) depict the scatter plots and histograms, obtained with the second calibration function, of the simulated and predictive sample, respectively.

5 Simulation experiments

This section illustrates the performance of the Bayesian nonparametric conditional copula model with simulated data. We generate datasets $(U, V)$ of sizes $n = 250, 500$ and $1000$ from different copula families, such as the Gaussian and Frank copulas. The copula dependence parameter is considered as a function of the exogenous variable $X$, which is simulated from an Uniform distribution in the interval $[-2, 2]$.

For the Dirichlet process prior, we fix the total mass $\lambda = 1$ and, for the base measure $G_0$, we adopt a bivariate normal distribution with zero mean vector and covariance matrix $\sigma^2 I$, where $\sigma^2 = 100$. The following calibration functions are selected for $\theta(x|\beta)$,

$\theta(x|\beta) = \beta_1 + \beta_2 x^2$
\[
\theta(x|\beta) = \beta_1 + \beta_2 x + \beta_3 \exp (-\beta_4 x^2).
\]

As highlighted in Section 3, we link the calibration functions $\theta(x|\beta)$ with $\rho(x|\beta)$ through the following transformation:

$\rho(x|\beta) = \frac{2}{\theta(x|\beta)} - 1.$
This ensures that $\rho(x|\beta)$ assumes values between $(-1, 1)$.

We run the Gibbs sampler algorithm described in Section 4 for 4000 iterations with (i) 500 burn-in iterations and (ii) 3500 burn-in iterations. Aiming at a parsimonious representation of the results, we focused on 3500 burn-in iterations, since 500 burn-in iterations gave very similar results.

![Gaussian Copula](a) ![Frank Copula](b) Figure 5: Number of components ($y$-axis) generated at each MCMC iteration ($x$-axis) for the Gaussian (left) and for the Frank copula (right) with sample size $n = 500$.

Figure 5 illustrates the results of the application of the Bayesian nonparametric conditional copula model to data simulated from a Gaussian copula, with sample size $n = 500$. Figure 4 illustrates similar results for the Frank copula. Since the performances of the model with sample sizes $n = 250$ and 1000 for both copula families were analogous, here we omit the results. In Figures 3 and 4, panels (a), (b), (c) and (d) show the scatter plots and histograms of the simulated data and the predictive samples obtained using the first calibration function; while panels (e), (f), (g) and (h) show the scatter plots and histograms of the simulated data and the predictive sample obtained using the second calibration function. The comparison between the simulated and predictive outputs highlights the excellent fit of the Bayesian nonparametric conditional copula model using either calibration function and for different copula families. The model performance appears to be consistent across both copula families, demonstrating that the approach is suitable to model different dependence patterns and tail structures. Figure 5 shows the plot of the number of components generated at each MCMC iteration for both the Gaussian and the Frank copula. In Figure 5, we focus on the first calibration function, since the second calibration function gave similar results. Table 1 shows the summary statistics of the number of components generated at each MCMC iteration for both copulas, indicating that the posterior median of the number of components is equal to 2. For the two most significant
components, we estimated the weights generated at each MCMC iteration for both copulas. In Figure 6, we show the trace plots of the last 500 iterations of the first two weights, as defined in Section 4. Figure 6 suggests that the first weight is much more important than the second weight, since the first weight tends to take values close to 1, while the second weight takes values close to 0. For each of the two components we also estimated the posterior mean of the copula correlation coefficient $\rho_j$ (defined in eq. 5), obtaining, for the Gaussian copula, a value of 0.6701 for the first component and $-0.9860$ for the second component. On the other hand, for the Frank copula we obtained a posterior mean of 0.7941 for the first component and $-0.9722$ for the second component, respectively.

|          | Min. | 1st Quant. | Median | Mean  | 3rd Quant. | Max. |
|----------|------|------------|--------|-------|------------|------|
| Gaussian | 1    | 1          | 2      | 2.102 | 3          | 7    |
| Frank    | 1    | 2          | 2      | 2.546 | 3          | 7    |

Table 1: Summary statistics of the number of components generated at each MCMC iterations for the first calibration function.
6 Real Data application

We now apply the proposed Bayesian nonparametric conditional copula method to a sample of 839 adolescent twin pairs, which is a subset of the National Merit Twin Study (Loehlin and Nichols, 2009, 2014). The dataset contains questionnaire data from 17 years old twins and their parents, where the twins were identified among 600,000 US high school juniors who took part to the National Merit Scholarship Qualifying Test (NMSQT).

The NMSQT was designed to measure cognitive aptitude, that is students’ readiness for future intellectual or educational pursuits. The participants to the test include identical twins and same-sex fraternal twins who were asked to fill in a complete questionnaire in order to understand their school performance and attitude. Our purpose is to examine whether the relationship between twins’ cognitive ability, measured by the NMSQT, is influenced by their socioeconomic status, measured by parent education and parental income. The variables we considered from this study are the overall measures of each twin’s performance at school (obtained as the sum of individual scores in English Usage, Mathematics Usage, Social Science Reading, Natural Science Reading and Word Usage/Vocabulary), the mother’s and father’s level of education and the family income. The overall scores range from 30 to 160, the education covariates range from 0 to 6, while the family income covariate ranges from 0 to 7. The levels of the education covariates correspond to: less than 8-th grade, 8-th grade, part high school, high

![Figure 7: Panels (a) and (b): scatterplots of the twins’ overall scores for the real and pseudo-observations with respect to the mother’s level of education; panels (c) and (d): scatterplots of the predictive and transformed predictive samples; panels (e) and (f): histograms of the real data and the predictive sample.](image)
school graduate, part college or junior college, college graduate, and graduate or professional degree beyond the bachelor’s degree. The levels of the income covariate correspond to values going from less than $5000 per year to over than $25000 per year.

Figure 8: Panels (a) and (b): scatterplots of the twins’ overall scores for the real and pseudo-observations with respect to the father’s level of education; panels (c) and (d): scatterplots of the predictive and transformed predictive samples; panels (e) and (f): histograms of the real data and the predictive sample.

As discussed in Section 1, the scatterplots in Figure 1 clearly show that there is a positive correlation between the twins’ school performance and the strength of dependence varies according to the values of a covariate, which is the mother’s (panel (a)) or father’s level of education (panel (b)) or the family income (panel (c)). In Figure 1 the effect of the covariates is illustrated by dots of different colours, where we notice that most of the light brown dots are grouped in the upper right corner, while the dark brown dots lie in the bottom left corner. Therefore, the higher the parents’ education or family income, the higher the twins’ school performance. In order to model the effect of a covariate, such as the mother’s and father’s education and family income, on the dependence between the overall scores of the twins, we implement the Bayesian nonparametric conditional copula model.

Figure 9 shows the relationship between the covariates of the twins dataset, where the lower-triangular panels represent pairwise scatterplots, the upper-triangular panels show pairwise Pearson’s correlation coefficients, and the diagonal panels represent the histograms of each covariate. The scatterplots and Pearson’s correlation coefficients in Figure 9 indicate a rather strong positive correlation between each pair of covariates, especially between the mother’s and father’s level of education. The high correlations indicate that the data do not contain much
information on the independent effects of each covariate, and suggest the inclusion of only one of them in the model. For this reason we decided to include only one of the redundant covariates at a time. Note that, with a different dataset, the methodology may be extended to include more than one covariate. However, model specification issues and increased computational costs must be carefully considered.

Figure 9: Relationship between the covariates of the twins dataset. The lower-triangular panels represent pairwise scatterplots, the upper-triangular panels show pairwise Pearson’s correlation coefficients, and the diagonal panels represent the histograms of each covariate. (Note that jittering was used in the scatterplots to prevent overplotting).

Adopting the same priors of the simulation studies, we run the Gibbs sampling algorithm described in Section 4 for 4000 iterations. Figures 7, 8 and 10 show, for the mother’s and father’s education and family income, respectively, the scatterplots of the twins’ overall scores using the real and transformed pseudo-observations (panels (a) and (b)), the scatterplots of the predictive and transformed predictive samples (panels (c) and (d)) and the histograms of the real and the predictive samples (panels (e) and (f)). Note that the pseudo-observations are obtained using the nonparametric estimation approach described in Section 4. From the comparison between the scatterplots and histograms of the real and predictive samples obtained with the three different covariates, it emerges that the Bayesian nonparametrics conditional copula model accurately captures the tail structures and the dependence patterns between the twins’ overall scores. Moreover, the posterior means of the number of mixture components
Figure 10: Panels (a) and (b): scatterplots of the twins’ overall scores for the real and pseudo-observations with respect to the family income; panels (c) and (d): scatterplots of the predictive and transformed predictive samples; panels (e) and (f): histograms of the real data and the predictive sample.

for the conditional copula are 26.82, 24.49 and 27.11, for the mother’s and father’s level of education and the family income, respectively, supporting the need for non-Gaussian copulas. We note that the good performance of this approach in tail modelling makes it suitable to various applications focussing on extremes. To quantify the degree of dependence between the twins’ scores, we use the conditional Kendall’s tau, which is a nonparametric measures of correlation, known as concordance, between two ranked variables \((Y_1, Y_2)\) with respect to a covariate \(X = x\). The conditional Kendall’s tau takes the following form:

\[
\tau(x) = 4 \int \int C_x(u_1, u_2) dC_x(u_1, u_2) - 1
\]

where \(C_x\) is the appropriate conditional copula. Figure 11 shows Kendall’s tau estimated from the model against the mother’s (top panel) and father’s level of education (middle panel) and the family income (bottom panel), together with 95% credible intervals. The plots clearly illustrate the negative effect of all three covariates on the dependence between the twins’ overall scores. The effect is greater for the family income, where the Kendall’s tau decreases from approximately 0.83 to 0.45, while for the parents’ education levels the Kendall’s tau decreases from approximately 0.8 to 0.6. Therefore, the higher the parents’ education and family income, the better the socioeconomic status and the higher the differences between the twins’ school performances. The cognitive aptitudes of twins from less advantaged families are
Figure 11: Estimated Kendall’s tau against the mother’s (top panel) and father’s level of education (middle panel) and the family income (bottom panel) and an approximate 95% credible intervals (dotted lines)

more similar to each other than those from high income, highly educated families. Families of high socioeconomic status provide supportive and challenging environments, able to offer a wide range of opportunities and choices to their children, and allowing them to express themselves freely. Hence, twins raised in wealthy families are encouraged to develop differences in their traits, and may show rather dissimilar cognitive abilities, albeit high on average. On the contrary, families of low socioeconomic status offer scarce opportunities to their children and may provide limiting and restrictive environments. In less advantaged families, twins cannot develop their full potential and individuality, hence both tend to show low cognitive abilities.

This might suggest, as in [Loehlin et al. (2009)], an interaction between genetic and environmental factors. Genes multiply environmental inputs that support intellectual growth such that an increased socioeconomic status raises the average cognitive ability but also magnifies individual differences in cognitive ability (see [Bates et al. (2013)].)

7 Conclusion

In this paper we proposed a Bayesian nonparametric conditional copula approach to model the strength and type of dependence between two variables of interest and we applied the methodology to the National Merit Twin Study. In order to capture the dependence structure between two variables, we introduced two different calibration functions expressing the functional form of a covariate variable. The statistical inference was obtained implementing a slice sampling algorithm, assuming an infinite mixture model for the copula. The methodology combines the advantages of the conditional copula approach with the modeling flexibility of Bayesian nonparametrics.

The simulation studies illustrated the excellent performance of our model with three distinct
copula families and different sample sizes. The application to the twins data revealed the importance of the environment in the development of twins’ cognitive abilities and suggests that environmental factors are more influential in families with higher socioeconomic position. On the contrary, other factors, such as genetic causes, may be more dominant in families with lower socioeconomic position.

Although this paper focuses on bivariate copula models, the methodology can be extended to multivariate copulas including more than one covariate. However, the inclusion of multiple covariates needs special attention regarding the choice of variables prior to estimate the calibration functions. Moreover, the increasing computational cost due to the additional covariates should be taken carefully into consideration.

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A Gibbs sampling details

Let $\mathcal{D}_j = \{i = 1, \ldots, n : d_i = j\}$ be the set of indexes of the observations allocated to the $j$-th component of the mixture, while $\mathcal{D} = \{j : \mathcal{D}_j \neq \emptyset\}$ is the set of indexes of non-empty mixtures components. Let $D^* = \sup \{\mathcal{D}\}$ be the number of stick-breaking components used in the mixture. As in Kalli et al. (2011), the sampling of infinite elements of $\pi$ and $\beta$ is not necessary, since only the elements of the full conditional probability density functions of $D$ are need.

The maximum number of stick-breaking components to be sampled is:

$$N^* = \max \{i = 1, \ldots, n | N^*_i\},$$

where $N^*_i$ is the smallest integer such that $\sum_{j=1}^{N^*_i} w_j > 1 - z_i$.

A.1 Update of $\pi$

We update the stick-breaking components and consequently the weights $w_j$ based on the equation $w_j = \pi_j \prod_{k<j}(1 - \pi_k)$. Assuming that $\pi_j$ is distributed as a Beta ($Be(1, \lambda)$), the full conditional distribution of $\pi_j$ is:

$$\begin{aligned}
\pi_j | \cdots & \sim Be(1 + \#\{d_i = j\}, \lambda + \#\{d_i > j\}),
\end{aligned}$$ (14)

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where \( \#\{d_i = j\} \) are the number of \( d_i \) equal to \( j \) and \( \#\{d_i > j\} \) is the number of \( d_i \) greater than \( j \) for \( j < D^* \).

On the other hand, if \( j = D^* + 1, \ldots, N^* \) we have that
\[
\pi_j | \cdots \sim \text{Be}(1, \lambda).
\]

### A.2 Update of \( Z \)

From the full likelihood function (13), \( z_i \) follows a uniform distribution
\[
z_i | \cdots \sim \mathcal{U}(0, w_{d_i})
\]
and it is sampled accordingly.

### A.3 Update of \( D \)

The allocation variable \( d_i \) values lie between 0 and \( N_i \) and the density of \( d_i \) satisfies
\[
P(d_i = j | \cdots) \propto I(z_i < w_{d_i})c_{\rho(x_i|\beta_{d_i})}(u_i, v_i).
\]

### A.4 Update of \( \beta \)

The full conditional of the vector of parameters \( \beta_k \), for \( k \geq 1 \) is:
\[
f(\beta_k | \cdots) \propto \pi(\beta_k) \prod_{d_i = k} c_{\rho(x_i|\beta_k)}(u_i, v_i),
\]
where \( \pi(\beta_k) \) is the prior on \( \beta \). Since the (17) is not a standard distribution, we used a Random Walk Metropolis Hastings.