

Quasi-random sampling for multivariate distributions via generative neural networks

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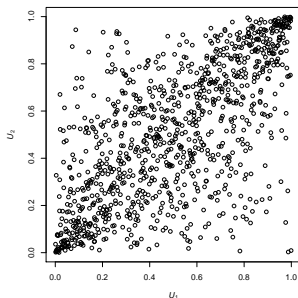
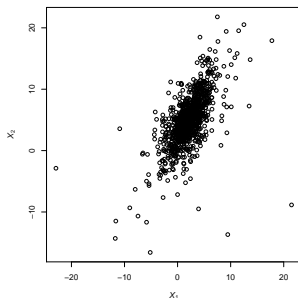
1 Classical copula modeling

- **General goal:** Modeling a df H or $\mathbf{X} \sim H$ with continuous margins F_1, \dots, F_d (for possibly high-dimensional, computational applications).
- **Examples:**
 - ▶ **Static:** \mathbf{X} models a joint loss in a risk management context; e.g., $\mu = \mathbb{P}(\mathbf{X} > \mathbf{x})$ or $\text{ES}_\alpha(S) = \mathbb{E}(S \mid S > F_S^{-1}(\alpha))$ for $S = X_1 + \dots + X_d$.
 - ▶ **Dynamic:** \mathbf{X} models joint innovations of an ARMA–GARCH process or increments of dependent geometric Brownian motions
- **Sklar's Theorem:**
 - ▶ **Analytical:** $H(\mathbf{x}) = C(F_1(x_1), \dots, F_d(x_d))$ for a copula C , a distribution function with $U(0, 1)$ margins.
 - ▶ **Stochastic:** $\mathbf{X} = (F_1^{-1}(U_1), \dots, F_d^{-1}(U_d)) \sim H$ (quantile transformation) and $\mathbf{U} = (F_1(X_1), \dots, F_d(X_d)) \sim C$ (probability transf.).
- **Why:** Useful under **asymmetric information** (margins known, dependence unknown) or from a **computational point of view** (e.g., estimation).

- Statistics:** Instead of \mathbf{X} , we now have $X = (\mathbf{X}_1^\top, \dots, \mathbf{X}_n^\top)^\top \in \mathbb{R}^{n \times d}$.
 - Given X , compute the *pseudo-observations* $U = (\mathbf{U}_1^\top, \dots, \mathbf{U}_n^\top)^\top \in \mathbb{R}^{n \times d}$ with

$$U_{ij} = \hat{F}_{n,j}(X_{ij}) = \frac{1}{n+1} \sum_{k=1}^n \mathbb{1}_{\{X_{kj} \leq X_{ij}\}} = \frac{R_{ij}}{n+1},$$

where $R_{ij} = \text{rank}(X_{ij})$ among the component sample X_{1j}, \dots, X_{nj} .



- Based on U , one needs to **fit, test and select** an **adequate copula C** .
- Simulate $\mathbf{U}_1, \dots, \mathbf{U}_{n_{\text{gen}}}$** from the fitted C and **estimate**, for example:

- ▶ $F_S^{-1}(u)$ where $S = X_1 + \dots + X_d = F_1^{-1}(U_1) + \dots + F_d^{-1}(U_d)$

Examples (QRM):

- Value-at-risk $\text{VaR}_\alpha(S) = F_S^{-1}(\alpha)$ of the total loss S
- Expected shortfall $\text{ES}_\alpha(S) = \mathbb{E}(S | S > F_S^{-1}(\alpha))$
- ▶ Expectations $\mu = \mathbb{E}(\Psi_0(\mathbf{X})) = \mathbb{E}(\Psi(\mathbf{U}))$ where $\Psi(\mathbf{u})$ is given by $\Psi_0(F_1^{-1}(u_1), \dots, F_d^{-1}(u_d))$ and so $\mu \approx \frac{1}{n_{\text{gen}}} \sum_{i=1}^{n_{\text{gen}}} \Psi(\mathbf{U}_i)$.

Examples (MC):

- With $\Psi(\mathbf{u}) = \mathbb{1}_{\{u_1 > u, \dots, u_d > u\}}$ we obtain that

$$\mu = \mathbb{E}(\Psi(\mathbf{U})) = \mathbb{P}(U_1 > u, \dots, U_d > u),$$

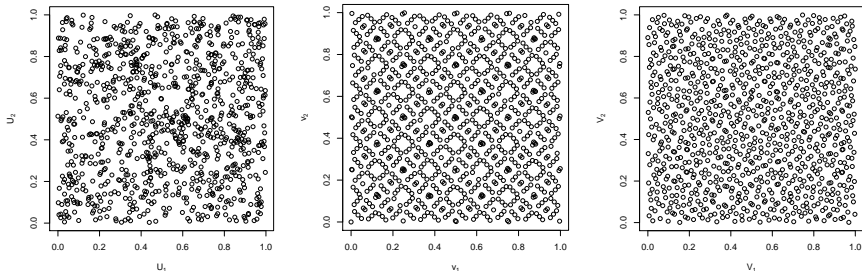
so exceedance probabilities, e.g., the probability of a flood over multiple dikes or joint losses in a stock portfolio.

- With $\Psi(\mathbf{u}) = \max\{(F_1^{-1}(u_1) + \dots + F_d^{-1}(u_d))/d - K, 0\}$ we obtain that $\mu = \mathbb{E}(\Psi(\mathbf{U}))$ is the expected payoff of a European basket call option.

- **Problems (with this classical modeling approach):**
 - 1) Step 2) above: Finding an adequate copula model C for the pseudo-observations, especially in more than two dimensions.
 - 2) Large variance $\text{var}\left(\frac{1}{n_{\text{gen}}}\sum_{i=1}^{n_{\text{gen}}}\Psi(U_i)\right)$ of the Monte Carlo estimator under rare-event simulation.
- **Ideas:**
 - 1) Use (specific) neural networks (NNs) (\Rightarrow flexible dependence).
 - 2) Use quasi-random sampling for such NNs (\Rightarrow variance reduction).
- **Outline:**
 - ▶ We first address 2), under dependence.
 - ▶ We then study how NNs can be random number generators (RNGs) from copulas, so 1).
 - ▶ We then address how NNs can be quasi-RNGs (QRNGs).
 - ▶ For simplicity, we focus on the case where $F_1 = \dots = F_d$ are $U(0, 1)$ in what follows (except for an ES_α example).

2 Quasi-random numbers for copulas

- If $C = \Pi$ (independence), pseudo-random numbers (PRNs) can be replaced by **quasi-random numbers (QRNs)** to reduce the variance.
- QRN sequences are **low-discrepancy sequences** $P_n = \{\mathbf{v}_i\}_{i=1}^n$ (middle) with $D^*(P_n) = \sup_{z \in (0,1]^d} \left| \frac{\#\{i: \mathbf{v}_i \in [0, z]\}}{n} - \lambda([0, z]) \right| \in O(n^{-1} \log^d n)$.
- **Example:** $U_1, \dots, U_{n_{\text{gen}}}$ (left), **randomized Sobol'** $V_1, \dots, V_{n_{\text{gen}}}$ (right):



- The **RQMC estimator** $\frac{1}{n_{\text{gen}}} \sum_{i=1}^{n_{\text{gen}}} \Psi(\mathbf{V}_i)$ is **unbiased**, **fast** and its **variance** can be estimated (from repeated randomizations).

- **Question:** How can we obtain QRNs from a general copula C ?
- **Idea:** Could define $D_C^*(P_n) = \sup_{z \in (0,1]^d} \left| \frac{\#\{i: v_i \in [\mathbf{0}, z]\}}{n} - \mathbb{P}(U \in [\mathbf{0}, z]) \right|$, but this **does not lead to a construction principle**. However, one can transform $V_1, \dots, V_{n_{\text{gen}}}$ to samples from C ; see Cambou et al. (2017).
- **Inverse Rosenblatt transform \mathcal{R}^{-1} :** Bijection to transform $U' \sim U(0,1)^d$ to $U \sim C$ (known as *conditional distribution method (CDM)* for sampling, a generalization of the *inversion method* to $d > 1$):

$$U_1 = U'_1,$$

$$U_2 = C_{2|1}^{-1}(U'_2 | U_1), \quad (\text{compare with } X_2 = F_2^{-1}(U'_2) \sim F_2)$$

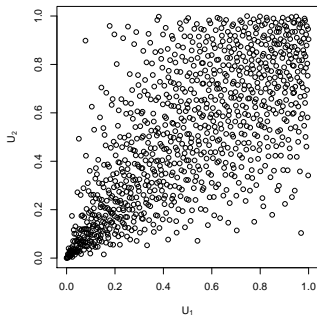
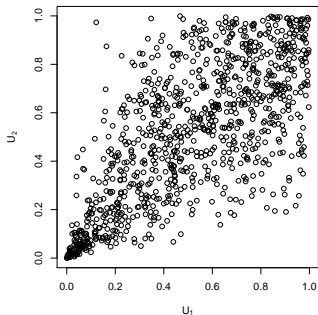
$$U_j = C_{j|1, \dots, j-1}^{-1}(U'_j | U_1, \dots, U_{j-1}), \quad j \in \{2, \dots, d\}.$$

- **Formula** for the implementation (which **needs to be inverted!**):

$$C_{j|1, \dots, j-1}(u_j | u_1, \dots, u_{j-1}) = \frac{D_{j-1, \dots, 1} C_{1, \dots, j}(u_1, \dots, u_j)}{D_{j-1, \dots, 1} C_{1, \dots, j-1}(u_1, \dots, u_{j-1})}.$$

\Rightarrow Known analytically tractable $C_{j|1, \dots, j-1}^{-1}$: **Normal, t , Clayton.**

- **Example:** 1000 PRNs (left) and QRNs (right) from a Clayton copula.



- **Disclaimer:** Even if not visible, there may be a variance reduction effect.
- For large j , evaluating $C_{j|1,\dots,j-1}^{-1}$ is time-consuming even for normal, t , Clayton.
- For Gumbel copulas, $C_{j|1,\dots,j-1}^{-1}$ is not tractable.
- See Cambou et al. (2017) for a different approach, but conceptually (first component \Rightarrow frailty) and numerically challenging (α -stable qf).

3 PRNs for copulas via GMMNs

- **Idea:** To address Problem 1), we consider

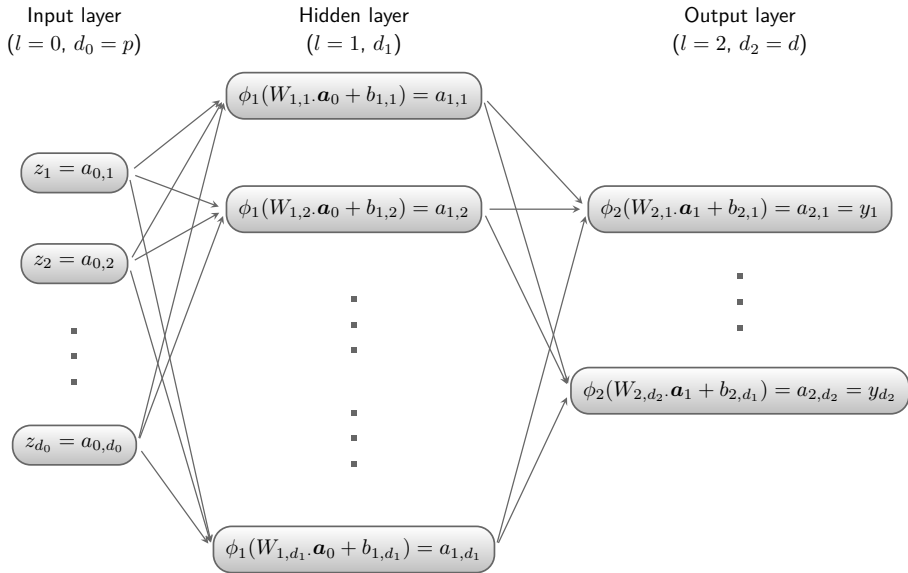
$$\mathbb{E}(\Psi(\mathbf{U})) \approx \frac{1}{n_{\text{gen}}} \sum_{i=1}^{n_{\text{gen}}} \Psi(\mathbf{U}_i), \quad (1)$$

$$\mathbf{U}_i = f_{\hat{\theta}}(F_{\mathbf{Z}}^{-1}(\mathbf{U}'_i)), \quad i = 1, \dots, n_{\text{gen}}, \quad (2)$$

where

- ▶ $\mathbf{U}'_1, \dots, \mathbf{U}'_{n_{\text{gen}}} \stackrel{\text{ind.}}{\sim} \text{U}(0, 1)^p$ (later: $\mathbf{V}_1, \dots, \mathbf{V}_{n_{\text{gen}}}$; randomized Sobol');
 - ▶ $F_{\mathbf{Z}}^{-1}(\mathbf{u}) = (F_{Z_1}^{-1}(u_1), \dots, F_{Z_p}^{-1}(u_p))$ maps to a *prior distribution* (Φ);
 - ▶ $f_{\hat{\theta}}$ is a trained generative neural network f_{θ} (GMMNs).
- Training of f_{θ} learns a map from pseudo-random numbers from \mathbf{Z} (here: $\text{N}_p(\mathbf{0}, I_p)$) to pseudo-random numbers from $\mathbf{U} \sim \mathcal{C}$.
 - This is similar to \mathcal{R}^{-1} , but computationally simpler to evaluate.
 - **Errors:**
 - 1) Monte Carlo error (1) (can be made arbitrarily small by the SLLN)
 - 2) Neural network (NN) “bottleneck” (2) (small if NN trained correctly)

3.1 What are NNs?



- **Hyperparameters** (fixed before training):
 - ▶ The ϕ_l 's are the *activation functions* (e.g., *ReLU* $\phi_l(x) = \max\{0, x\}$, *sigmoid* $\phi_l(x) = 1/(1 + e^{-x})$).
 - ▶ Number of layers, number of neurons per layer, number of epochs in training and batch size (more later).
- **Parameter vector:**

$\theta = (W_1, \dots, W_{L+1}, \mathbf{b}_1, \dots, \mathbf{b}_{L+1})$ consists of *weight matrices* and *biases* (initialized randomly and with zeros, resp.).
- θ is fitted (or: the NN is *trained*) by *stochastic gradient descent* based on a *cost function* $E(U, Y)$ computed between (a subset of)
 - ▶ the *training data* $U = (\mathbf{U}_1^\top, \dots, \mathbf{U}_{n_{\text{trn}}}^\top)^\top$ from \mathcal{C} (= target) and
 - ▶ the *outputs* $Y = f_\theta(Z)$ of the NN from the prior sample $Z = (\mathbf{Z}_1^\top, \dots, \mathbf{Z}_{n_{\text{trn}}}^\top)^\top$.
- Often E is taken as the scaled **MSE** $E(U, Y) = \frac{1}{2n_{\text{trn}}} \sum_{i=1}^{n_{\text{trn}}} \|\mathbf{u}_i - \mathbf{y}_i\|_2^2$
 \Rightarrow **Fails to learn the map from Z to U properly.**

3.2 What are GMMNs?

- *Generative moment matching networks (GMMNs)* use as $E(U, Y)$ the sample *maximum mean discrepancy* $MMD(U, Y)$

$$\sqrt{\frac{1}{n_{\text{trn}}^2} \sum_{i_1=1}^{n_{\text{trn}}} \sum_{i_2=1}^{n_{\text{trn}}} (K(U_{i_1}, U_{i_2}) - 2K(U_{i_1}, Y_{i_2}) + K(Y_{i_1}, Y_{i_2}))}$$

where K is a mixture of Gaussian kernels of different bandwidths. After experimentation, we chose

$$K(\mathbf{u}, \mathbf{y}) = \sum_{i=1}^6 e^{-\frac{\|\mathbf{u}-\mathbf{y}\|_2^2}{2\sigma_i^2}} \quad \boldsymbol{\sigma} = (0.001, 0.01, 0.15, 0.25, 0.50, 0.75).$$

- Intuitively, **MMD** takes into account all pairs of observations between U_{i_1} and Y_{i_2} (desirable, but costly \Rightarrow mini-batch optimization).
- For the population version, one can show that $MMD(U, Y) = 0$ if and only if $U \stackrel{d}{=} Y$.

3.3 How are GMMNs trained?

- **Training algorithm** (*mini-batch optimization*):
 - 1) Initialize θ (weights W_1, W_2 : uniform entries; biases $\mathbf{b}_1, \mathbf{b}_2$: $\mathbf{0}$)
 - 2) Partition the (n_{trn}, d) -data matrix U and prior (n_{trn}, d) -matrix Z into *batches* ($\approx n_{\text{trn}}/n_{\text{bat}}$ blocks of n_{bat} (batch size) rows each).
 - 3) For each batch, update θ by a stochastic gradient step (Adam).
 - 4) After $\approx n_{\text{trn}}/n_{\text{bat}}$ gradient steps, U and Z are exhausted and one epoch is completed. Shuffle their rows and go to Step 2).
 - 5) Training finishes after n_{epo} epochs.
- **Setup** in our experiments:
 - ▶ $p = d$ (inspired from \mathcal{R}^{-1} in the CDM);
 - ▶ single hidden layer (*universal approximation theorem*: given suitable activation functions, single hidden layer NNs with $d_1 < \infty$ can approximate any continuous function on a compact subset of \mathbb{R}^d);
 - ▶ $d_1 = 300$ neurons in the hidden layer;

- ▶ ϕ_1 to be ReLU (fast) and ϕ_2 to be sigmoid (maps to $(0, 1)$);
- ▶ batch size $n_{\text{bat}} = 5000$ (trade-off: large enough for “bottleneck” error to be small; small enough to not be memory-prohibitive);
- ▶ number of epochs $n_{\text{epo}} = 300$; and
- ▶ $n_{\text{trn}} = 60\,000$ pseudo-samples from C (available for all models).

3.4 How can GMMNs generate QRNs from C ?

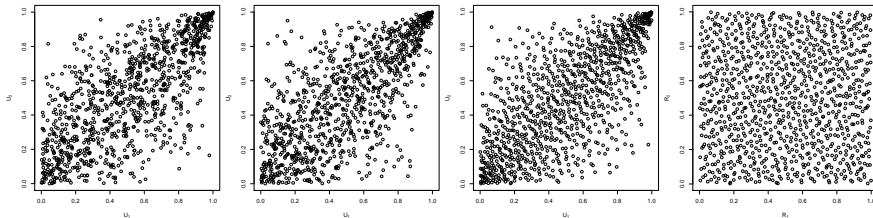
» Algorithm 3.1 (GMMN quasi-random sampling)

- 1) Compute the RQMC points $\mathbf{V}_1, \dots, \mathbf{V}_{n_{\text{gen}}}$ (e.g., randomized Sobol’).
- 2) Compute the prior samples $\mathbf{Z}_i = F_{\mathbf{Z}}^{-1}(\mathbf{V}_i)$, $i = 1, \dots, n_{\text{gen}}$.
- 3) Return the pseudo-observations of $\mathbf{Y}_i = f_{\hat{\theta}}(\mathbf{Z}_i)$, $i = 1, \dots, n_{\text{gen}}$.

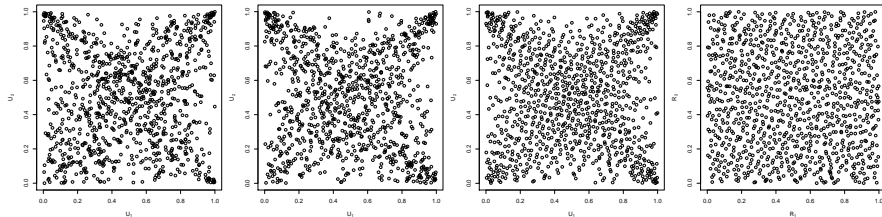
GMMNs are fast to evaluate and sufficiently smooth to preserve low discrepancy. If all the mixed partial derivatives of $h = \Psi \circ f_{\hat{\theta}} \circ F_{\mathbf{Z}}^{-1}$ exist a.e. and are continuous, then under Owen-type scrambling, $\text{var}\left(\frac{1}{n_{\text{gen}}} \sum_{i=1}^{n_{\text{gen}}} h(\mathbf{V}_i)\right) = O(n_{\text{gen}}^{-3} (\log n_{\text{gen}})^{p-1})$.

3.5 Do GMMNs generate samples from \mathcal{C} well?

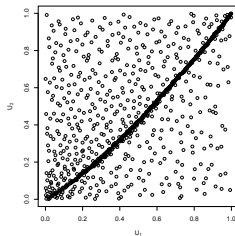
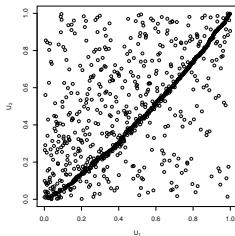
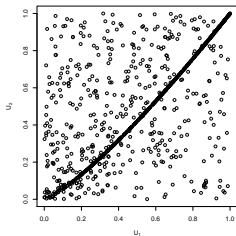
Gumbel PRNs, GMMN PRNs, GMMN QRNs, \mathcal{R} -transformed GMMN QRNs:



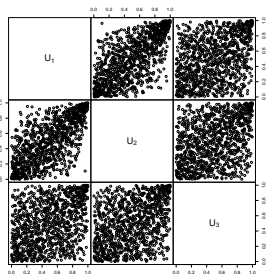
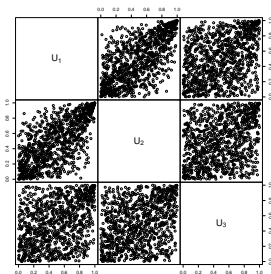
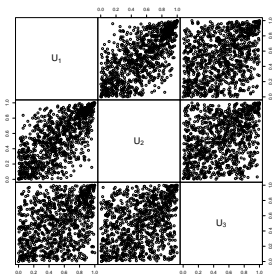
Gumbel-rotated t_4 mixture:



Marshall–Olkin copula $C(u_1, u_2) = \min\{u_1^{1-\alpha_1}u_2, u_1u_2^{1-\alpha_2}\}$ (without \mathcal{R}):



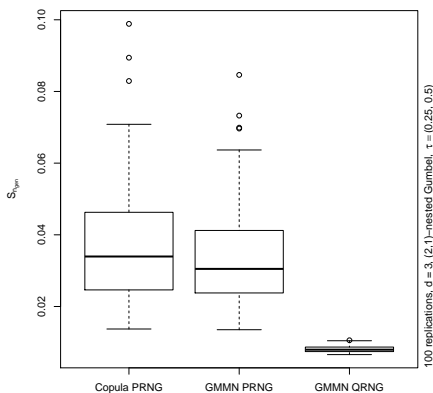
Nested Gumbel copula $C(u) = C_0(C_1(u_1, u_2), u_3)$ ($\tau \in \{0.25, 0.5\}$):



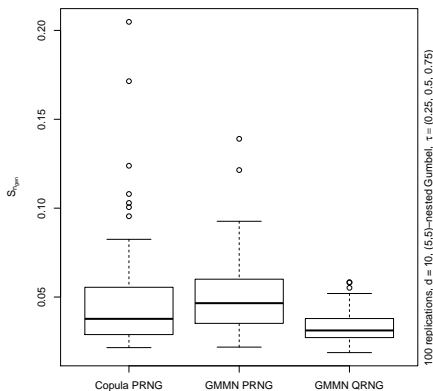
One can consider **box plots** of realization of the *Cramér–von Mises statistic*

$$S_{n_{\text{gen}}} = \int_{[0,1]^d} n_{\text{gen}} (C_{n_{\text{gen}}}(\mathbf{u}) - C(\mathbf{u}))^2 dC_{n_{\text{gen}}}(\mathbf{u}),$$

where $C_{n_{\text{gen}}}$ is the *empirical copula* (ecdf of the pseudo-obs.) of $n_{\text{gen}} = 1000$ PRNs, GMMN PRNs and GMMN QRNs ($d \in \{3, 10\}$):



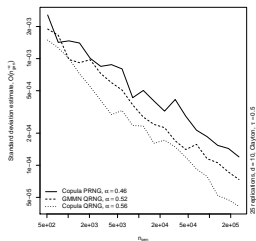
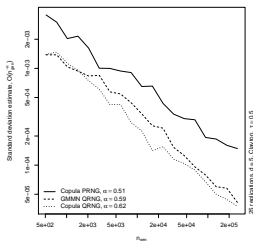
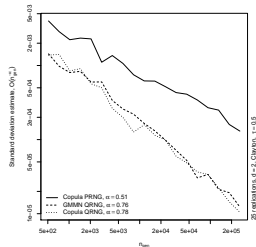
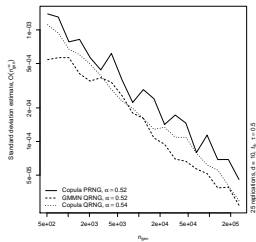
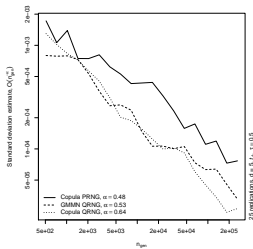
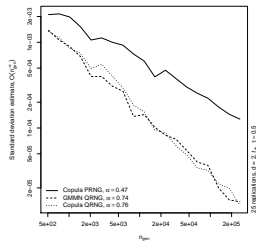
100 replications, $d = 3$, (2,1)-nested Gumbel, $\tau = (0.25, 0.5)$



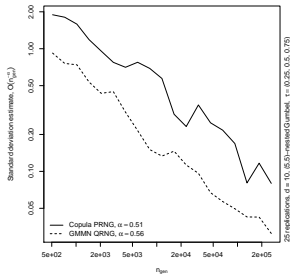
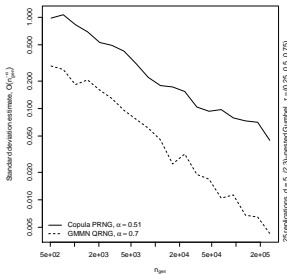
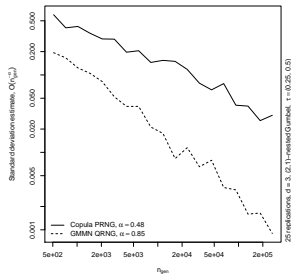
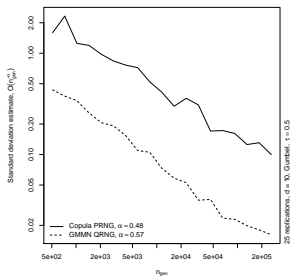
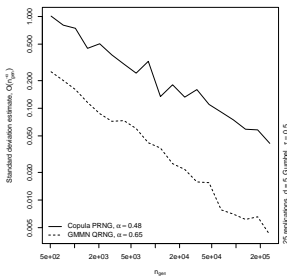
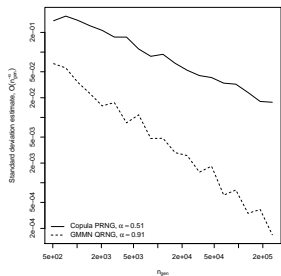
100 replications, $d = 10$, (5,5)-nested Gumbel, $\tau = (0.25, 0.5, 0.75)$

3.6 Variance-reduction effect?

Sample standard deviation of the estimator of $\mathbb{P}(U > 0.99)$ computed from PRNs, GMMN QRNs and QRNs:



Sample standard deviation of the estimator of $\mathbb{E}(S | S > F_S^{-1}(0.99))$ (N(0,1) margins) computed from PRNs and GMMN QRNs (no QRNs):



Summary

- We can learn a QRNG from any joint model based on a PRNG for C .
- If generated from a known C , this is easy. If only a sample is available, n_{trn} needs to be sufficiently large (depending on the application).
- **Gain:** **Universality** (all models), **computability** (robustness, run time), especially useful for **real data** (where the true model is unknown).
- **Challenges:**
 - 1) Kendall's tau near 1;
 - 2) training needs a GPU server (evaluation "needs" TensorFlow);
 - 3) **joint tail behavior**;
 - 4) $d \gg 10$.
- **Open problem:** For $d \gg 10$, must training be improved (as distributions become harder to learn) or do RQMC point sets generally deteriorate? (GMMNs are still smooth). One could try (t, m, s) -nets other than Sobol' (but no related R package or standalone C code available).

Outlook: Application to time series

- **Copula–GARCH model:** $X_{t,j} = \mu_{t,j} + \sigma_{t,j}Z_{t,j}$, $j = 1, \dots, d$, and $Z_t = (Z_{t,1}, \dots, Z_{t,d}) \stackrel{\text{ind.}}{\sim} H$ with **copula** C .
- **GMMN–GARCH model:** $X_{t,j} = \mu_{t,j} + \sigma_{t,j}Z_{t,j}$, $j = 1, \dots, d$, and $Z_t = (Z_{t,1}, \dots, Z_{t,d}) \stackrel{\text{ind.}}{\sim} H$ with **GMMN** for C .
- **Goal:** Improving empirical predictive distributions (here: 1-step ahead).
- **Applications:**
 - 1) **ZCB yield curves** (term structure of interest rates)
 - ▶ **US** ($d = 30$ annual times to maturity) and **CA** ($d = 120$ quarterly times to maturity) ZCB yield curves (**training: 1995–2015**)
 - ▶ **Marginal deARMA–GARCHing** \Rightarrow standardized residuals $(\hat{Z}_t)_t$
 - ▶ **PCA** $\Rightarrow d = 3$ (US data) and $d = 4$ (CA data) account for $\geq 95\%$ of the total variance
 - ▶ **Performance** evaluation on **1y** (CA) and **1/2y** (US) **test data:**

- **Dependence:** $AMMD = \frac{1}{100} \sum_{i=1}^{100} MMD(U^{(i)}, \hat{U}; K_{\text{tst}})$, where $U^{(i)}$ is the i th matrix of generated data and \hat{U} are the pseudo-observations of the test data.
- **Better one-day ahead empirical predictive distributions:**

$$AVS^{1/4} = \frac{1}{T-\tau} \sum_{t=\tau+1}^T \sum_{j_1=1}^d \sum_{j_2=1}^d \left(|X_{t,j_1} - X_{t,j_2}|^{1/4} - \frac{1}{1000} \sum_{i=1}^{1000} |\hat{X}_{t,j_1}^{(i)} - \hat{X}_{t,j_2}^{(i)}|^{1/4} \right)^2$$

For both datasets, the **GMMN–GARCH models better matched the cross-sectional dependence** on the test datasets.

2) Exchange rates w.r.t. USD and w.r.t. GBP

- ▶ CAD, GBP, EUR, CHF, JPY w.r.t. USD and of CAD, USD, EUR, CHF, JPY, CNY w.r.t. GBP (training: 2000–2015; test: 2015)
- ▶ AMMD and $AVS^{1/4}$ as before.
- ▶ $VEAR_{\alpha} = \left| \alpha - \frac{1}{T-\tau} \sum_{t=\tau+1}^T \mathbb{1}_{\{S_t < \widehat{\text{VaR}}_{\alpha}(\hat{S}_t)\}} \right|$ (VaR ex. abs. err.)
- ▶ For both datasets, the **GMMN–GARCH models produced better daily $\text{VaR}_{0.05}(S_t)$ forecasts.**

References

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