Gumbel 2

Note: Unlike the Gaussian, Student-t, or the Skew-Normal copula, the Gumbel copula is evaluated directly on the $[0,1]^2$ grid without the need for an inverse CDF transformation.

Install and load required packages

```
R_packages <- c("bookdown", "copula", "ggplot2", "gridExtra", "ks", "plot3D", "viridis")
options(repos = c(CRAN="http://cran.rstudio.com"))
if (!requireNamespace("librarian", quietly = TRUE)) install.packages("librarian")
librarian::shelf(R_packages)</pre>
```

Set parameters

```
a <- 4 # copula parameter
gsz <- 1000 # grid size
```

1 Gumbel Copula

Define the Gumbel copula

```
gumbel_cop <- gumbelCopula(param = a, dim = 2)</pre>
```

Create grid for u1, u2 in [0, 1]

```
u1_seq <- seq(0.01, 0.99, length.out = gsz)
u2_seq <- seq(0.01, 0.99, length.out = gsz)
grid <- expand.grid(u1 = u1_seq, u2 = u2_seq)
```

Compute copula density on $[0,1]^2$ copula grid

```
copula_dens <- dCopula(as.matrix(grid), gumbel_cop)
z_mat_3d <- matrix(copula_dens, nrow = gsz, byrow = FALSE)
z_mat_3d <- pmin(z_mat_3d, 10) # Truncate to see the countours change</pre>
```

Plot copula density on $[0,1]^2$ copula grid 3D

```
persp3D(
    x = u1_seq,
    y = u2_seq,
    z = z_mat_3d,
    theta = 25,
    phi = 50,
    col = viridis::viridis(100),
    xlab = "u1", ylab = "u2", zlab = "density",
    main = "Bivariate Gumbel Copula Density (a = 4)",
    cex.main = 0.6, cex.lab = 0.6,
    colkey = list(length = 0.6, width = 0.6, cex.axis = 0.6)
)
```

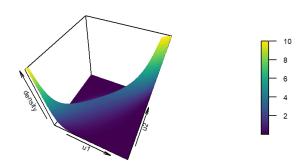


Figure 1.1: Plot copula density on $[0,1]^2$ copula grid contour

```
z_mat_2d <- matrix(copula_dens, nrow = gsz, byrow = TRUE)
#z_mat_2d[is.infinite(z_mat_2d)] <- NaN # RepLace INF entries with NaN
z_mat_2d <- pmin(z_mat_2d, 10) # Truncate to see the contours change

filled.contour(
    x = u1_seq, y = u2_seq, z = z_mat_2d,
    color.palette = viridis::viridis,
    xlab = "", ylab = "",
    plot.axes = {axis(1, at = seq(0.2, 0.8, by = 0.2), line = -0.05); axis(2, line = -3.2)},
    plot.title = title(main = "Copula Density: Contour Plot", cex.main = 0.8),
    key.title = title(main = "Density", cex.main = 0.6), asp = 1, frame.plot = FALSE,
    colkey = list(addlines = FALSE, length = 0.6, width = 0.5, cex.clab = 0.8)
)</pre>
```

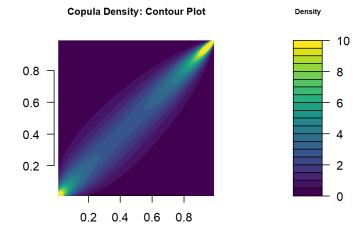


Figure 1.2:

2 Simulated Data

Simulate data

```
n <- 10000 # simulated sample size
set.seed(123456)
cs <- rCopula(n, gumbel_cop)</pre>
```

Gumbel 2

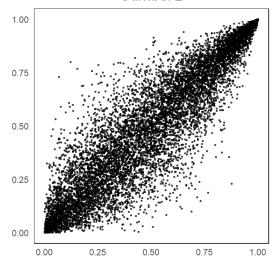


Figure 2.1:
Plot 3D kernel density estimate of the sample

```
# Kernel density estimation
H <- Hpi(cs) # bandwidth matrix via plug-in method
kde_res <- kde(x = cs, H = H, compute.cont = FALSE)
# Extract grid and density matrix
x <- kde_res$eval.points[[1]]</pre>
y <- kde_res$eval.points[[2]]</pre>
z \leftarrow kde\_res\$estimate \# matrix of density values
# 3D interactive plot
persp3D(
 x = x,
 y = y,
 z = z,
 theta = 25,
  phi = 50,
  col = viridis::viridis(100),
 xlab = "u1", ylab = "u2", zlab = "density",
 main = "3D KDE of Gumbel Copula Sample",
 cex.main = 0.6, cex.lab = 0.6,
  colkey = list(length = 0.6, width = 0.6, cex.axis = 0.6)
```

3D KDE of Gumbel Copula Sample

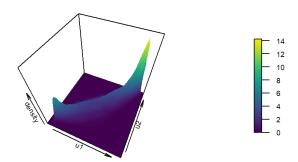


Figure 2.2:

3 Sparse Bernstein Copula

3.1 Set parameters

```
rname <- "Gumbel2"
true_dens <- copula_dens
sample_sizes <- c(100, 200, 300, 400, 500, 1000, 10000)
kmax <- 10 # number of batches for each sample size</pre>
```

3.2 Generate Data of Varying Sample Sizes for Fortran Estimation

Run Fortran estimation of SBP copula

3.3 Plot Simulated Draws from SBP copula

```
n <- 10000 # for sample size
k <- 1 # for data from batch k
csim_SBP <- read.table(file = paste0("../output_sim/",rname,"_csim_",n,"_",k,".out"), header = FALSE)</pre>
```

Plot

```
csim_SBP_df <- as.data.frame(csim_SBP)
colnames(csim_SBP_df) <- c("U1", "U2")

csplot <- ggplot(csim_SBP_df, aes(x=U1, y=U2)) +
    geom_point(alpha=0.7, size=0.5) + coord_fixed(ratio=1) +
    labs(title = "SBP copula", x = "", y = "") +
    theme_minimal() + theme(plot.margin = unit(c(0.1, 0.01, 0, 0), "in")) +
    theme(panel.grid = element_blank(), plot.title = element_text(hjust=0.5, size = 15),
        panel.border = element_rect(colour = "black", fill = NA, linewidth = 0.5))

plot_file = paste0("../graphs_sim/",rname,"_SBP.png")
ggsave(plot_file, plot = csplot, width = 4, height = 4, units = "in", dpi = 300)

csplot</pre>
```

SBP copula

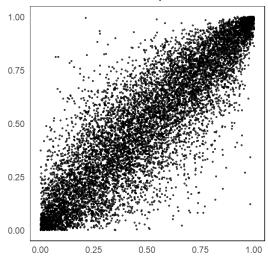


Figure 3.1:

3.4 Evaluate Kullback-Leibler Distance from True Density

```
KL <- numeric()
KLse <- numeric()

for (n in sample_sizes) {

    KL_k <- numeric()
    epsilon <- 1e-10 # Add small epsilon to avoid division by 0

    for (k in 1:kmax) {

        cdens_SBP <- read.table(file = paste0("../output_sim/",rname,"_cdens_",n,"_",k,".out"), header = FALSE)
        cdens_SBP <- as.numeric(unlist(cdens_SBP))
        ratio <- (true_dens + epsilon)/(cdens_SBP + epsilon)
        KL_k_val <- log(ratio)*true_dens # Compute KL integrand
        KL_km <- mean(KL_kval) # approximate integral over the grid
        KL_k <- c(KL_k, KL_km) # add new row
    }

KL <- c(KL, mean(KL_k)) # add new row
    KLse <- c(KLse, sd(KL_k)/sqrt(length(KL_k))) # add new row
}

print(round(KL, 2))</pre>
```

print(round(KLse, 2))

[1] 0.01 0.01 0.01 0.01 0.00 0.00 0.00