# Dppsampl Package

# 11<sup>th</sup> November 2023

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#### Description:

This package provides algorithms for sampling from the Determinantal Point Processes (DPPs) for SAS.

#### Version:

0.5

#### Modules:

- sample\_from\_beta\_ensemble\_full Function
- sample\_from\_beta\_ensemble\_banded Function

# Overview:

The dppsampl package provides an implementation of the algorithms for sampling from the Determinantal Point Processes (DPPs) written natively in SAS. DPPs are stochastic point processes (so, their result is a subset of that set) which respect the diversity present in the set.

The package offers ways to sample from Finite, Continuous and the so-called Exotic DPPs. For Finite DPPs the set of possible values is finite, for continuous it is infinite and the samples should approximate some continuous distribution. The exotic DPPs however, are a special kind of DPPs which were analyzed for different reasons and one needed a change-of-perspective to deduce that they can be thought of as DPPs.

The available continuous DPPs come from a family of distributions called Beta ensembles. They were initially studied in Physics, as they can be thought of as models of a Coulomb gas.

Before you use the dppsampl package, you must install it by using the package install statement. For example, if the ZIP file is located in the directory C:\Packages, then the following statement installs the package:

```
proc iml;
package install "C:\Packages\dppsampl.zip";
quit;
```

# **Data Sets**

We provide four test datasets which can be used to test the implementation of finite DPPs. All datasets have been constructed from the k.sas7bdat one.

- k.sas7bdat correlation kernel which is a  $10 \times 10$  projection matrix with rank of 8,
- 1.sas7bdat matrix created from k.sas7bdat using the relation  $K = L(I+L)^{-1}$ ,
- eig\_vals.sas7bdat set of eigenvalues of k.sas7bdat,
- $\bullet$  eig\_vals.sas7bdat set of eigenvectors of k.sas7bdat.

# **Available Finite DPP Functions:**

# proj\_dpp\_sampler\_kernel\_Chol Function

#### Syntax:

```
proj_dpp_sampler_kernel_Chol(
    kernel,
    size=.,
    random_state=.
);
```

## Parameters

## Description

For given projection correlation kernel matrix (K), the function returns a sample of given size (if size is not given then the output sample will be of size=rank(kernel). The sample is calculated using Cholesky based method.

```
use data.k;
    read all into kernel;
close;
size=5;
random_state=123;

sample = proj_dpp_sampler_kernel_Chol(
    kernel,
    size,
    random_state
);
```

# proj\_dpp\_sampler\_kernel\_shur Function

## Syntax:

```
proj_dpp_sampler_kernel_shur(
    kernel,
    size=.,
    random_state=.
);
```

#### **Parameters**

# Description

For given projection correlation kernel matrix (K), the function returns a sample of given size (if size is not given then the output sample will be of size=rank(kernel). The sample is calculated using Shur based method.

```
use data.k;
    read all into kernel;
close;
size=5;
random_seed=123;
sample = proj_dpp_sampler_kernel_shur(
    kernel,
    size,
    random_seed
);
```

# proj\_dpp\_sampler\_kernel\_GS Function

## Syntax:

```
proj_dpp_sampler_kernel_GS(
    kernel,
    size=.,
    random_state=.
);
```

#### **Parameters**

kernel Projection correlation matrix K

size Desired size of the output sample. Should be less than or equal to rank(kernel). If none is provided then the sample will be of size equal to rank(kernel)

Seed for the randomness. If provided it should be a positive

random\_state Seed for the randomness. If provided it should be a positive integer

# Description

For given projection correlation kernel matrix (K), the function returns a sample of given size (if size is not given then the output sample will be of size=rank(kernel). The sample is calculated using GS (Gram-Schmidt) based method.

```
use data.k;
    read all into kernel;
close;
size=5;
random_state=123;

sample = proj_dpp_sampler_kernel_GS(
    kernel,
    size,
    random_state
);
```

# proj\_dpp\_sampler\_eig\_GS Function

## Syntax:

```
proj_dpp_sampler_eig_GS(
    eig_vecs,
    size=.,
    random_state=.
);
```

#### **Parameters**

```
eig_vecs Matrix of eigen vectors of a proper correlation kernel matrix.

size Desired size of the output sample. Should be less than or equal to rank(kernel). If none is provided then the sample will be of size equal to rank(kernel)

random_state Seed for the randomness. If provided it should be a positive integer
```

# Description

For given set of eigen vectors of a proper correlation kernel matrix, the function returns a sample of given size (if size is not given then the output sample will be of size=rank(kernel). The sample is calculated using GS (Gram-Schmidt) method.

```
use data.eig_vecs;
    read all into eig_vecs;
close;
size=5;
random_state=123;

sample = proj_dpp_sampler_eig_GS(
    eig_vecs,
    size,
    random_state
);
```

# proj\_dpp\_sampl\_ker\_eig\_KuTa12 Function

## Syntax:

```
proj_dpp_sampl_ker_eig_KuTa12(
    eig_vecs,
    size=.,
    random_state=.
);
```

#### **Parameters**

```
eig_vecs Matrix of eigen vectors of a proper correlation kernel matrix.

size Desired size of the output sample. Should be less than or equal to rank(kernel). If none is provided then the sample will be of size equal to rank(kernel)

random_state Seed for the randomness. If provided it should be a positive integer
```

# Description

For given set of eigen vectors of a proper correlation kernel matrix, the function returns a sample of given size (if size is not given then the output sample will be of size=rank(kernel). The sample is calculated using KuTa12 (Kulesza-Taskar) method.

```
use data.eig_vecs;
    read all into eig_vecs;
close;
size=5;
random_state=123;
sample = proj_dpp_sampl_ker_eig_KuTa12(
    eig_vecs,
    size,
    random_state
);
```

# ${\bf sampler\_generic} \ {\bf Function}$

# Syntax:

```
sampler_generic(
    kernel,
    random_state=.
);
```

#### **Parameters**

kernel Correlation kernel matrix K.

 $\begin{tabular}{ll} {\tt random\_state} & Seed for the randomness. If provided it should be a positive integer \\ \end{tabular}$ 

# Description

For given correlation kernel matrix, the function returns a sample calculated with generic sampling algorithm. It is not possible to determine the size of the sample.

```
use data.k;
    read all into kernel;
close;
random_state=123;

sample = sampler_generic(
    kernel,
    random_state
);
```

# $add\_exchange\_delete\_sampler\ Function$

#### Syntax:

```
add_exchange_delete_sampler(
    kernel,
    s0=.,
    random_state=.,
    nb_iter=10
);
```

#### **Parameters**

kernel Likelihood kernel matrix

so Initial sample from set of values represented by K. If none is provided then the algorithm will generate the initial sample.

random\_state Seed for the randomness. If provided it should be a positive integer

nb\_iter Number of iterations that the algorithm will perform

nb\_trials Number of trials for generating the initial sample

## Description

For given likelihood matrix, the function returns a sample calculated by adding or removing an element (with specific probabilities) from given samples (starting from initial s0 sample) for a number of iterations specified by the nb\_iter parameter. If the initial sample - s0 is not provided, then it is generated by the algorithm . It is not possible to determine the size of the output sample, however it is possible to provide a size of the generated initial sample.

```
use data.1;
    read all into kernel;
close;
sample = add_exchange_delete_sampler(kernel);
```

# $add_delete_sampler\ Function$

#### Syntax:

```
add_delete_sampler(
    kernel,
    s0=.,
    size_s0=.,
    random_state=.,
    nb_iter=100,
    nb_trials=100
);
```

#### **Parameters**

kernel Likelihood kernel matrix

s0 Initial sample from set of values represented by K. If none is provided then the algorithm will generate the initial sample.

size\_s0 Expected size of the initial sample generated by the algorithm (provided the initial sample is not given)

 $\begin{tabular}{ll} \bf random\_state & Seed for the randomness. If provided it should be a positive integer \\ \end{tabular}$ 

nb\_iter Number of iterations that the algorithm will perform.

## Description

For given likelihood matrix, the function returns a sample calculated by adding, exchanging or removing an element (with specific probabilities) from given samples (starting from initial s0 sample) for a number of iterations specified by the nb\_iter parameter. If the initial sample - s0 is not provided, then it is generated by the algorithm. It is not possible to determine the size of the sample.

```
use data.l;
    read all into kernel;
close;
sample = add_delete_sampler(kernel);
```

# basic\_exchange\_sampler Function

#### Syntax:

```
basic_exchange_sampler(
    kernel,
    s0=.,
    size_s0=.,
    random_state=.,
    nb_iter=100,
    nb_trials=100
);
```

#### **Parameters**

kernel Likelihood kernel matrix

so Initial sample from set of values represented by K. If none is provided then the algorithm will generate the initial sample.

size\_s0 Expected size of the initial sample generated by the algorithm (provided the initial sample is not given). In this case this will equal to the size of the output sample.

random\_state Seed for the randomness. If provided it should be a positive integer

nb\_iter Number of iterations that the algorithm will perform.

## Description

For given likelihood matrix, the function returns a sample calculated by exchanging an element (with specific probability) from given samples (starting from initial s0 sample) for a number of iterations specified by the nb\_iter parameter. If the initial sample - s0 is not provided, then it is generated by the algorithm . As the only operation performed during the algorithm is exchanging elements, the output sample will be of the same size as the initial s0 sample, thus it is possible to specify the size of the sample.

```
use data.1;
    read all into kernel;
close;
sample = basic_exchange_sampler(kernel);
```

# **Available Continuous DPP Functions:**

# sample\_from\_beta\_ensemble\_full Function

## Syntax:

```
sample_from_beta_ensemble_full(
    ensemble_version,
    M_1, M_2,
    size=10,
    beta=2,
    normalize=1,
    haar_mode="Hermite",
    heurestic_fix=1,
    random_state=1618
);
```

#### **Parameters**

```
ensemble_version
                    Version of Beta ensemble to use. Available values are
                    "Hermite", "Laguerre", "Jacobi", "Circular", and
                    ""Ginibre"
                    Distribution parameter for the "Laguerre" and "Jacobi"
              M_1
                    ensemble_versions. Should be greater or equal to size.
                    Distribution
                                                                   "Jacobi"
              M_{-}1
                                     parameter
                                                   for
                                                           the
                    ensemble_version. Should be greater or equal to
                    size.
                    Size of the sampled subset.
             size
                    Beta parameter. Should be 1, 2, or 4.
                    Parameter which states whether the sample should be nor-
       normalize
                    malized to fit one of the more known distributions.
                    Which Haar measure mode to use. Can be "Hermite" or
       haar_mode
                    "QR". Influences the result only for the Circular Ensemble.
                    (Should be 1 \text{ or } 0).
                    Whether to apply the heuresis to fix the oversampling pro-
  heurestic_fix
                    blem present in the Circular and Ginibre ensembles. Should
                    be 1 or 0.
                    Seed for the randomness. Should be a positive integer.
    random_state
```

# Description

The function provides the method for sampling from beta ensemble using the full matrix method. There are five versions of Beta ensembles that have been implemented. "Hermite", "Laguerre", "Jacobi", "Circular", and "Ginibre". For the first three, the result is a one-column sample. For the next two, it is a two-column sample.

```
ensemble_version = "Circular";
size=10;
beta=4;
M_1=10; M_2 = 10;
haar_mode="Hermite";
normalize=0;
heurestic_fix=0;
random_state=1618;
sample = sample_from_beta_ensemble_full(
        ensemble_version,
        M_{1}, M_{2},
        size,
        beta,
        normalize,
        haar_mode,
        heurestic_fix,
        random_state
);
run scatter(sample[,1], sample[,2]);
```

## sample\_from\_beta\_ensemble\_banded Function

## **Syntax**

```
sample_from_beta_ensemble_banded(
    ensemble_version,
    size=10,
    beta=2,
    loc=0.0,
    scale=1.0,
    shape = 1.0,
    a = 1.0, b = 1.0,
    normalize=1,
    heurestic_fix=1,
    random_state=1618
);
```

#### **Parameters**

ensemble\_version Version of Beta ensemble to use. Available values are "Hermite", "Laguerre", "Jacobi", and "Ginibre"

size Size of the sampled subset.

beta Beta parameter. Should be positive integer.

- loc Location parameter for the standard deviation for the
  "Hermite" Beta ensemble.
- scale Scale parameter for the expected value for the "Hermite" and "Laguerre" Beta ensembles.

shape Shape parameter for the "Laguerre" Beta ensemble.

- a Parameter for the "Jacobi" Beta ensemble. Related to the Beta distribution.
- b Parameter for the "Jacobi" Beta ensemble. Related to the Beta distribution.

normalize Parameter which states whether the sample should be normalized to fit one of the more known distributions.

heurestic\_fix Whether to apply the heuresis to fix the oversampling problem present in the Circular and Ginibre ensembles. Should be 1 or 0.

random\_state Seed for the randomness. Should be a positive integer.

# Description

The function provides the method for sampling from beta ensemble using the banded matrix method. There are five versions of Beta ensembles that have been implemented. "Hermite", "Laguerre", "Jacobi", and "Ginibre". For the first three, the result is a one-column sample. For the Ginibre ensemble, it is a two-column sample.

```
ensemble_version = "Hermite";
size=1000;
beta=2;
loc=0.0;
scale=1.0;
shape = 1.0;
a = 1.0;
b = 1.0;
normalize=0;
heurestic_fix=0;
random_state=1618;
sample = sample_from_beta_ensemble_banded(
    ensemble_version,
    size,
    beta,
    loc,
    scale,
    shape,
    a, b,
    normalize,
    heurestic_fix,
    random_state
);
run histogram(sample);
```

# **Available Exotic DPP Functions:**

# poiss\_planch\_sample Function

## Syntax

```
poiss_planch_sample(
    theta=10,
    random_state=.
);
```

#### **Parameters**

```
\mbox{ theta} \quad \mbox{parameter of poisson distribution, must be integer} > 1.
```

random\_state Seed for the randomness. Should be a positive integer

## Description

Generates a sample from the Poissonized Plancherel method by using RSK (Robinson-Schensted-Knuth) algorithm on a random permutation on 1, N where N is generated from Poisson distribution with parameter theta. It is not possible to determine the size of the output sample (it will always be  $\leq$  theta).

```
theta = 10;
random_state = 123;

sample = poiss_planch_sample(
    theta,
    random_state);
```

# carries\_sample Function

## Syntax

```
carries_sample(
   base=10,
   size=100,
   random_state=.
);
```

#### **Parameters**

```
base The number by which each element of a sequence is divided to generate the list of rests, must be integer >1.

size Size of the generated list, upper bound of output sample, must be integer >1.

random_state Seed for the randomness. Should be a positive integer
```

## Description

Generates a sample by creating the sequence of Carries. A sequence of i.i.d. digits of a given size is generated and the cumulative sum is computed. The base parameter specifies the number by which each element of a sequence is divided to generate the rests. It is not possible to determine the size of the output sample (it will always be <size).

```
base = 3;
size=20;
random_state = 123;

sample = carries_sample(
    base,
    size,
    random_state
);
```

# $descent\_sample$ Function

# Syntax

```
descent_sample(
    size=100,
    random_state=.
);
```

#### **Parameters**

```
size Size of the generated list, upper bound of output sample, must be integer >1.
```

random\_state Seed for the randomness. Should be a positive integer

# Description

Generates a sample by creating a descent process obtained from a uniformly chosen permutation of 1, ..., size. It is not possible to determine the size of the output sample (it will always be < size).

```
size=20;
random_state = 123;

sample = descent_sample(
    size,
    random_state
);
```

# virtual\_descent\_sample Function

## Syntax

```
virtual_descent_sample(
    size=100,
    x0=0.5,
    random_state=.
);
```

#### **Parameters**

size Size of the generated list, upper bound of output sample, must be integer >1.

x0 The parameter of the binomial distribution that will determine generation of the non-uniform selection of permutation.

random\_state Seed for the randomness. Should be a positive integer

# Description

Generates a sample from a mix of DPPs by obtaining a non-uniformly chosen permutation of  $0, \ldots, size-1$  (using binomial distribution with specified x0 parameter). It is not possible to determine the size of the output sample (it will always be < size).

```
size=20;
x0=0.5;
random_state = 123;
sample = virtual_descent_sample(
    size,
    x0,
    random_state
);
```