# SAS macro to generate Network Balanced Type II Designs (NetBD2) for multi-location agroforestry trial 

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Sometimes a single continuous piece of land may not be available at one location to the researcher for implementation of the designs. In such situations, it will be preferable to have suitable designs which can be implemented in multiple locations despite the limitations associated with the available land. A new class of designs called Network Balanced Designs Type II (NetBD2) is proposed for use in such experimental situations.

Network Balanced Design (NetBD): NetBD refers to a circular design based on the network model in which each tree species on a plot has every other tree species appearing as left, right, top and bottom neighbours equal (constant) number of times. Here, balance indicates combinatorial balance only.

Experimental setup and model: The linear network effects model of Parker et al. (2017) can be considered, which should effectively account for main and nondirectional interference effects of trees in an agroforestry trial as stated by Birteeb (2021).

Consider an agroforestry experiment where same crop but $(v)$ different tree species are planted on $n$ plots, each plot has only one tree species. Assuming that the response $Y_{i}$ (measured from the crop) is a result of "tree effect" $\left(\tau_{j, i}\right)$ from plot $i$ having tree species $j$, and "tree network effect" $\left(\delta_{l, k}\right)$ if tree species $l$ is planted on an adjacent connected plot $k$. Each of the $n$ plots is connected by 4 other plots surrounding it. Let $\boldsymbol{A}_{n \times n}$ is a symmetric adjacency matrix for this experiment, and let $\boldsymbol{u}_{j}$ be the indicator vector with $j^{\text {th }}$ element equal to 1 when tree species $j$ is planted on plot $i$ and 0 otherwise, then the network effects model is expressed in matrix notation as:

$$
\boldsymbol{Y}=\mu \mathbf{1}+\boldsymbol{U} \boldsymbol{\tau}+\boldsymbol{A} \boldsymbol{U} \boldsymbol{\delta}+\boldsymbol{\varepsilon}
$$

where, $\boldsymbol{Y}$ is $n \times 1$ vector of observations, $\mu$ is grand mean, $\mathbf{1}$ is a $n \times 1$ vector of unities, $\boldsymbol{U}$ is a $n \times v$ design matrix of observations versus direct tree species, $\boldsymbol{\tau}$ is a $v \times 1$ vector of direct tree effects, $\boldsymbol{N}=\boldsymbol{A} \boldsymbol{U}$ is a $n \times v$ design matrix of observations versus all adjacent treatments, $\boldsymbol{\delta}$ is a $v \times 1$ vector of all network effects, $\boldsymbol{X}$ is a $n \times(2 v+1)$ design matrix, $\boldsymbol{\theta}$ is a $(2 v+1) \times 1$ vector of parameters, $\boldsymbol{\varepsilon}$ is a $n \times 1$ vector of errors, with $E(\boldsymbol{\varepsilon})=\mathbf{0}$ and $D(\boldsymbol{\varepsilon})=\sigma^{2} \boldsymbol{I}_{n}$.

Example: Let there be $v=7$ tree species. Then the number of arrays required is $\frac{v-1}{2}=3$. The initial arrays (for $g=1,2,3$ ) are developed as:

| Array I |  |  |  |  |  |  | Array II |  |  |  |  |  |  | Array III |  |  |  |  |  |  |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 0 | 1 | 2 | 3 | 4 | 5 | 6 | 0 | 2 | 4 | 6 | 1 | 3 | 5 | 0 | 3 | 6 | 2 | 5 | 1 | 4 |
| 1 | 2 | 3 | 4 | 5 | 6 | 0 | 2 | 4 | 6 | 1 | 3 | 5 | 0 | 3 | 6 | 2 | 5 | 1 | 4 | 0 |
| 2 | 3 | 4 | 5 | 6 | 0 | 1 | 4 | 6 | 1 | 3 | 5 | 0 | 2 | 6 | 2 | 5 | 1 | 4 | 0 | 3 |
| 3 | 4 | 5 | 6 | 0 | 1 | 2 | 6 | 1 | 3 | 5 | 0 | 2 | 4 | 2 | 5 | 1 | 4 | 0 | 3 | 6 |
| 4 | 5 | 6 | 0 | 1 | 2 | 3 | 1 | 3 | 5 | 0 | 2 | 4 | 6 | 5 | 1 | 4 | 0 | 3 | 6 | 2 |
| 5 | 6 | 0 | 1 | 2 | 3 | 4 | 3 | 5 | 0 | 2 | 4 | 6 | 1 | 1 | 4 | 0 | 3 | 6 | 2 | 5 |
| 6 | 0 | 1 | 2 | 3 | 4 | 5 | 5 | 0 | 2 | 4 | 6 | 1 | 3 | 4 | 0 | 3 | 6 | 2 | 5 | 1 |

After adding 1 to every element and making each array circular, the final NetBD2 is:


|  |  | 5 |  |  |  |  |  |  |
| :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- | :--- |
| 5 | 1 | 4 | 7 | 3 | 6 | 2 |  |  |
|  | 1 | 4 | 7 | 3 | 6 | 2 | 5 | 1 |
| 1 | 4 | 7 | 3 | 6 | 2 | 5 | 1 | 4 |
| 4 | 7 | 3 | 6 | 2 | 5 | 1 | 4 | 7 |
| 7 | 3 | 6 | 2 | 5 | 1 | 4 | 7 | 3 |
| 3 | 6 | 2 | 5 | 1 | 4 | 7 | 3 | 6 |
| 6 | 2 | 5 | 1 | 4 | 7 | 3 | 6 | 2 |
| 2 | 5 | 1 | 4 | 7 | 3 | 6 | 2 | 5 |
|  | 1 | 4 | 7 | 3 | 6 | 2 | 5 |  |

The layout of the design can be obtained as generated using the developed SAS macro by just entering the number of lines.
/*Developed by- Peter T. Birteeb, Eldho Varghese, Cini Varghese, Seema Jaggi and Mohd Harun*/
/*Date: 10-07-2022*/
/*VERSION 1.0: 10-07-2022*/
/*It provides generation of Network Balanced Designs Type II (NetBD2) */
OPTIONS NODATE NOSTIMER LS=78 PS=60;
\%let $\mathrm{v}=7$; /*Enter the number of treaments (v must be prime number) */
ods rtf file="NETWORK BALANCED DESIGN.rtf";
title 'NETWORK BALANCED DESIGN';

```
proc iml;
pp1=1;
do i=2 to &v-1;
pp=mod(&v,i);
if pp=0 then ppl=0;
end;
if pp1=0 then do;
print 'Entered number is not a prime number';
end;
if pp1^=0 then do;
print "NETWORK BALANCED DESIGN for v = &v";
Square=j(&v,&v,0);
do i=1 to &v;
do j=1 to &v;
Square[i,j]=mod((i-1)+(j-1),&v)+1;
end;
end;
do k=1 to (&v-1)/2;
if k=1 then do;
NBD1=Square;
end;
else do;
do i=1 to nrow(square);
do j=1 to ncol(square);
NBD1[i,j]=mod(Square[i,j]+(j-1)+(i-1),&v);
if NBD1[i,j]=0 then NBD1[i,j]=&v;
end;
end;
end;
Square=NBD1;
a0=j(1,1,'Border_Plots');
a1_l=NBD1[ ,ncol(NBD1)];
a2_l=char(a1_1,4,0);
*a3_l=a0//a2_l//a0;
a1_r=NBD1[ ,1];
a2_r=char(a1_r,4,0);
*a3_r=a0//a2_r//a0;
a1_t=NBD1[nrow(NBD1), ];
a2_t=char(a1_t,4,0);
a3_t=a0||a2_t|a0;
a1_b=NBD1[1, ];
a2_b=char(a1_b,4,0);
a3_b=a0|a2_b||a0;
a4=char(NBD1,4,0);
NBD=a3_t//(a2_l|a4||a2_r)//a3_b;
print 'Square Number=' K;
print NBD;
```

print 'Note: Circular border plots has been considered for all the four sides viz., left, right, top and bottom';
end;
end;
run;
ods rtf close;
quit;

## SAS Output

| NETWORK BALANCED DESIGN |  |  |  |  |  |  |  |  |  |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| NETWORK BALANCED DESIGN for $v=7$ |  |  |  |  |  |  |  |  |  |
|  |  |  |  |  |  | k |  |  |  |
|  |  |  |  | Square Nu | umber= | 1 |  |  |  |
| NBD |  |  |  |  |  |  |  |  |  |
|  | COL1 | COL2 | COL3 | COL4 | COL5 | COL6 | COL7 | COL8 | COL9 |
| ROW1 | Border_Plots | 7 | 1 | 2 | 3 | 4 | 5 | 6 | Border_Plots |
| ROW2 | 7 | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 1 |
| ROW3 | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 1 | 2 |
| ROW4 | 2 | 3 | 4 | 5 | 6 | 7 | 1 | 2 | 3 |
| ROW5 | 3 | 4 | 5 | 6 | 7 | 1 | 2 | 3 | 4 |
| ROW6 | 4 | 5 | 6 | 7 | 1 | 2 | 3 | 4 | 5 |
| ROW7 | 5 | 6 | 7 | 1 | 2 | 3 | 4 | 5 | 6 |
| ROW8 | 6 | 7 | 1 | 2 | 3 | 4 | 5 | 6 | 7 |
| ROW9 | Border_Plots | 1 | 2 | 3 | 4 | 5 | 6 | 7 | Border_Plots |
| Note: Circular border plots has been considered for all the four sides viz., left, right, top and bottom |  |  |  |  |  |  |  |  |  |



## References

Birteeb, P. T. (2021). Designing agroforestry systems for sustainable livelihood. Unpublished Ph.D. Thesis, ICAR-Indian Agricultural Research Institute, New Delhi.

Parker, B. M., Gilmour, S. G., Schormans, J. (2017). Optimal design of experiments on connected units with application to social networks. Journal of the Royal Statistical Society, Series C, 66(3), 455-480.

