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## 1 FIGURE 1 – BACCALA ET AL. (2016) DTF: UNIFIED ASYMPTOTIC THEORY

DESCRIPTION:

Routine **figure1\_example1\_pdc\_dtf\_ns50.m** publish

Linear bivariate VAR(2) model

LA Baccala, DY Takahashi, K Sameshima (2016) Directed Transfer Function: Unified Asymptotic Theory and Some of its Implications. *IEEE Transactions on Biomedical Engineering* **PP**.

<http://dx.doi.org/10.1109/TBME.2016.2550199>

Example 1: Simple example VAR(2) without feedback

$x1 ==> x2$

## Contents

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- Some remarks:

## Generating data set for analysis

```
clear; clc; format compact
flgPlotStyle = 'Print'; % or 'Screen'
flgRandomize = 0; % Generate the specific data set used in Fig. 1.
ns = 50; % number of sample points
nDiscard = 2000; % number of points discarded at beginning of simulation
p = 2; % model order

if (exist('figure1_example1_pdc_dtf_ns50.mat') == 2) & is_octave & ~
    flgRandomize,
    load figure1_example1_pdc_dtf_ns50
else
    [u] = fbaccala2016_example1( ns, nDiscard, flgRandomize );
    if ~is_octave & ~flgRandomize,
        save figure1_example1_pdc_dtf_ns50 u
```

```

    end;
end;

chLabels = [];    % Using default labeling schema for channel identification

```

```

=====
"DTF Unified Asymptotic Theory" Simple Example 1, ns =50
      x1 ==> x2
=====

```

## Equation

$$\begin{aligned}
 x_1(n) &= 0.95\sqrt{2}x_1(n-1) - 0.9025x_1(n-2) + w_1(n) \\
 x_2(n) &= -0.5x_1(n-1) + 0.5x_2(n-1) + w_2(n)
 \end{aligned}$$

Equation (8) from Baccala et al. *IEEE Trans Biomed Engin.*, 2016.

Data pre-processing: detrending and normalization options

```

flgDetrend = 1;    % Detrending the data set
flgStandardize = 0; % No standardization
[nChannels,nSegLength] = size(u);
if nChannels > nSegLength, u = u.';
    [nChannels,nSegLength] = size(u);
end;
if flgDetrend,
    for i=1:nChannels, u(i,:) = detrend(u(i,:)); end;
    disp('Time series were detrended.');
```

```

Time series were detrended.

```

## MVAR model estimation

```

maxIP = 30;    % maximum model order to consider.
alg = 1;    % 1: Nutall-Strand MVAR estimation algorithm
criterion = 1;    % 1: AIC, Akaike Information Criteria
disp('Running MVAR estimation and GCT analysis routines.')
[IP,pf,A,pb,B,ef,eb,vaic,Vaicv] = mvar(u,maxIP,alg,criterion);
disp(['Number of channels = ' int2str(nChannels) ' with ' ...
    int2str(nSegLength) ' data points; MAR model order = ' int2str(IP) ' .']);

```

```

Running MVAR estimation and GCT analysis routines.
maxOrder limited to 30
IP=1 vaic=467.864503
IP=2 vaic=403.041777
IP=3 vaic=408.042486

```

```

Number of channels = 2 with 50 data points; MAR model order = 2.

```

## Testing for adequacy of MAR model fitting through Portmanteau test

```
h = 20; % testing lag
MVARadequacy_signif = 0.05; % VAR model estimation adequacy significance
                        % level
aValueMVAR = 1 - MVARadequacy_signif; % Confidence value for the testing
flgPrintResults = 1;
```

Granger causality test (GCT) and instantaneous GCT

```
gct_signif = 0.01; % Granger causality test significance level
igct_signif = 0.01; % Instantaneous GCT significance level
metric = 'euc'; % euc = original PDC or DTF;
           % diag = generalized PDC (gPDC) or directed coherence (DC);
           % info = information PDC (iPDC) or iDTF.
flgPrintResults = 1;
[Tr_gct, pValue_gct, Tr_igct, pValue_igct] = gct_alg(u,A,pf,gct_signif, ...
                                                    igct_signif,flgPrintResults);
```

```
-----
                        GRANGER CAUSALITY TEST
=====
Connectivity matrix:
    NaN    0
     1    NaN
Granger causality test p-values:
    NaN    0.2550
     0     NaN
-----
                        INSTANTANEOUS GRANGER CAUSALITY TEST
=====
Instantaneous connectivity matrix:
    NaN    0
     0    NaN
Instantaneous Granger causality test p-values:
    NaN    0.9098
    0.9098    NaN
>>>> Instantaneous Granger causality NOT detected.
```

## DTF estimation

DTF analysis results are saved in **c** structure. See `asympt_dtf.m` or issue `>> help asympt_dtf` command for more detail.

```
metric = 'euc'; % euc = Euclidian = original DTF;
%           % diag = diagonal = DC;
%           % info = information = iDTF.
nFreqs = 128;
alpha = 0.01;

c = asympt_dtf(u,A,pf,nFreqs,metric,alpha);
```

```
* Original DTF and asymptotic statistics
```

$|DTF(\lambda)|^2$  Matrix Layout Plotting

```

switch lower(flagPlotStyle)
case 'print'
    flgColor = [0];          % white background
    flgMax = 'TCI';
    flgSignifColor = 1; % black + gray
    flgScale = 3;           % [0 max(flagMax)]
otherwise % 'screen'
    flgColor = [1];          % Colored background
    flgMax = 'TCI';
    flgSignifColor = 3; % red + green
    flgScale = 2;           % [0 1]/[0 .1]/[0 .01]
end;

% -----Plotting options flag setting-----
%           [1 2 3 4 5 6 7]
flagPrinting=[1 1 1 2 2 0 1];
%           | | | | | 7 Spectra(0: w/o SS; 1: Linear; 2: log-scale)
%           | | | | | 6 Coherence
%           | | | | 5 Plot lower confidence limit (legacy)
%           | | | 4 Plot upper confidence limit
%           | | 3 Significant DTF(w) in red line (legacy)
%           | 2 Patnaik threshold level in black dashed-line
%           1 plot DTF
%-----

fs = 1;          % sampling frequency
% axis_scale = [0 0.50 -0.02 1.05];
w = fs*(0:(nFreqs-1))/2/nFreqs;
w_max = fs/2;

h = figure;
set(h,'NumberTitle','off','MenuBar','none', ...
    'Name','[Asymptotic DTF] Fig 1. Example 1 DTF, ns = 50')
[hxlabel hylabel] = xplot(c,...
    flagPrinting,fs,w_max,chLabels,flagColor,flagScale,flagMax,flagSignifColor);

[ax,hT] = suplabel(['Example 1 - simple example; sigma2 = 100,' ...
    ' sigma1 = sigma3 = 1; ns = ' ...
    int2str(ns) '; p = ' int2str(p) ...
    ' \alpha = ' int2str(100*alpha) '%.'], 't');
set(hT,'FontSize',10); % Subtitle font size
drawnow

```

Uncomment the command line bellow to generate an eps output file

```
% print -depsc2 -painters Fig1_example1_dtf_ns50.eps
```

## PDC estimation

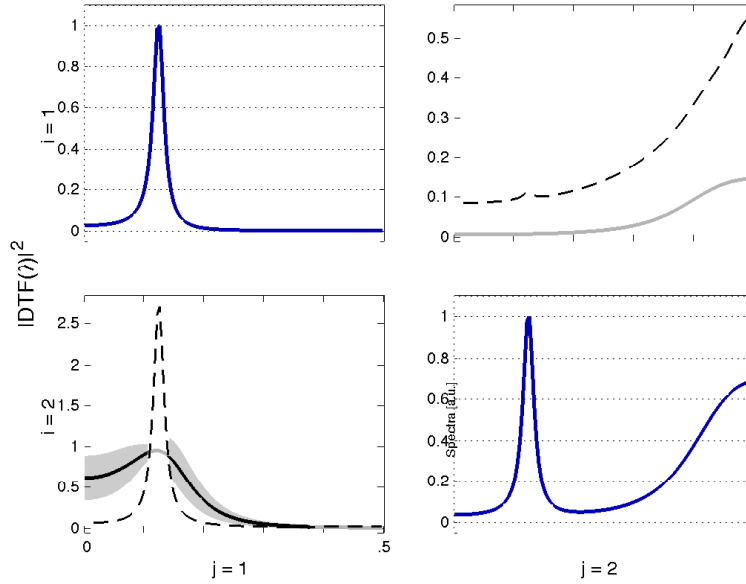
PDC analysis results are saved in **d** structure. See `asympt_dtf.m` or issue `>> help asympt_pdc` command for more detail.

```

nFreqs = 128;
metric = 'euc'; % euc = Euclidian = original PDC;
%           % diag = diagonal = generalized PDC or gPDC;
%           % info = information = iPDC

```

Example 1 - simple example; sigma2 = 100, sigma1 = sigma3 = 1; ns = 50; p = 2 alpha = 1%.



```
alpha = 0.01;
d = asymp_pdc(u,A,pf,nFreqs,metric,alpha);
```

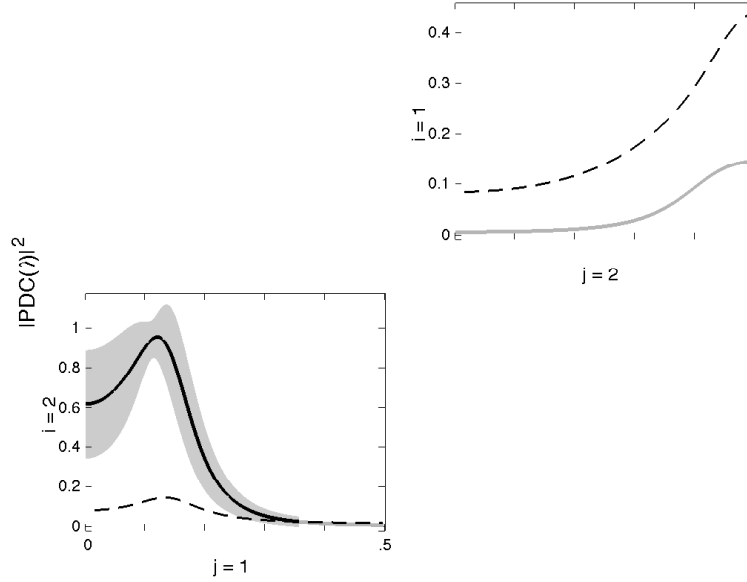
\* Original PDC and asymptotic statistics

## $|PDC(\lambda)|^2$ Matrix Layout Plotting

```
% -----Plotting options flag setting-----
%           [1 2 3 4 5 6 7]
flgPrinting=[1 1 1 2 2 0 0];
%           | | | | | 7 Spectra(0: w/o SS; 1: Linear; 2: log-scale)
%           | | | | | 6 Coherence
%           | | | | 5 Plot lower confidence limit (legacy)
%           | | | 4 Plot upper confidence limit
%           | | 3 Significant DTF(w) in red line (legacy)
%           | 2 Patnaik threshold level in black dashed-line
%           1 plot DTF
% -----
fs = 1;           % sampling frequency
w_max = fs/2; % x-axis maximum value
h = figure;
set(h,'NumberTitle','off','MenuBar','none', ...
    'Name','[Asymptotic DTF] Fig 1. Example 1 PDC, ns = 50')
[hxlabel hylabel] = xplot(d,...
    flgPrinting,fs,w_max,chLabels,flgColor,flgScale,flgMax,flgSignifColor);

[ax,hT] = suplabel(['Example 1 - simple example; sigma2 = 100,' ...
    ' sigma1 = sigma3 = 1; ns = ' ...
    int2str(ns) '; p = ' int2str(p) ...
    ' \alpha = ' int2str(100*alpha) '%.'], 't');
set(hT,'FontSize',10); % Subtitle font size
drawnow
```

Example 1 - simple example;  $\sigma_2 = 100$ ,  $\sigma_1 = \sigma_3 = 1$ ;  $n_s = 50$ ;  $p = 2$   $\alpha = 1\%$ .



Uncomment the command line bellow to generate an eps output file

```
%print -depsc2 -painters Fig1_example1_pdc_ns50.eps
```

## Result from the original article, Baccala et al (2016)

Figure 1, from article.

### Some remarks:

1. For the final figure 1, we did some cosmetic edit with AI
2. Erratum: **If we were not able to correct this typo.** In the caption of Figure 1, it states that "Gray shades describe 95% confidence intervals when above threshold.", however as we have used  $\alpha = 1\%$ , conversely the confidence interval is actually 99%, instead of 95%.

This completes the Figure 1 generation (Baccala et al, 2016)'

