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## 1 FIGURE 2 – BACCALA ET AL. (2016) DTF: UNIFIED ASYMP-TOTIC THEORY

DESCRIPTION:

Routine `figure2_example1_pdc_dtf_ns500.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`

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### Generating data set for analysis

```
clear; clc; format compact
flgPlotStyle = 'Print'; % or 'Screen' mode
flgRandomize = 0; % Generate the specific data set used in Fig. 1.
ns = 500; % number of sample points
nDiscard = 2000; % number of points discarded at beginning of simulation
p = 2; % model order

if (exist('figure2_example1_pdc_dtf_ns500.mat') == 2) & is_octave & ~
    flgRandomize,
    load figure2_example1_pdc_dtf_ns500
else
    [u] = fbaccala2016_example1(ns, nDiscard, flgRandomize);
    if ~is_octave & ~flgRandomize,
        save figure2_example1_pdc_dtf_ns500 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.');
end;
if flgStandardize,
for i=1:nChannels, u(i,:) = u(i,:)/std(u(i,:)); end;
disp('Time series were scale-standardized.');
end;
```

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=7020.488123
IP=2  vaic=6181.594313
IP=3  vaic=6189.212782
```

Number of channels = 2 with 500 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      0.4494
      0     NaN
-----
                               INSTANTANEOUS GRANGER CAUSALITY TEST
=====
Instantaneous connectivity matrix:
    NaN      0
      0     NaN
Instantaneous Granger causality test p-values:
    NaN      0.0366
  0.0366      NaN
>>> Instantaneous Granger causality NOT detected.

```

## DTF estimation

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

```

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

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

* Original DTF and asymptotic statistics

```

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

```

switch lower(flgPlotStyle)
    case 'print'
        flgColor =[0];          % white background
        flgMax = 'TCI';
        flgSignifColor = 1; % black + gray
        flgScale = 2;          % [0 max(flgMax)]
    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]
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 = fs*(0:(nFreqs-1))/2/nFreqs;
w_max = fs/2;

h = figure;
set(h, 'NumberTitle','off','MenuBar','none', ...
    'Name','[Asymptotic DTF] Fig 2. Example 1 DTF, ns = 500')
[h xlabel ylabel] = xplot(c, ...
    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

```

Uncomment the command line bellow to generate an eps output file

```
% print -depsc2 -painters Fig2_example1_dtf_ns500.eps
```

## PDC estimation

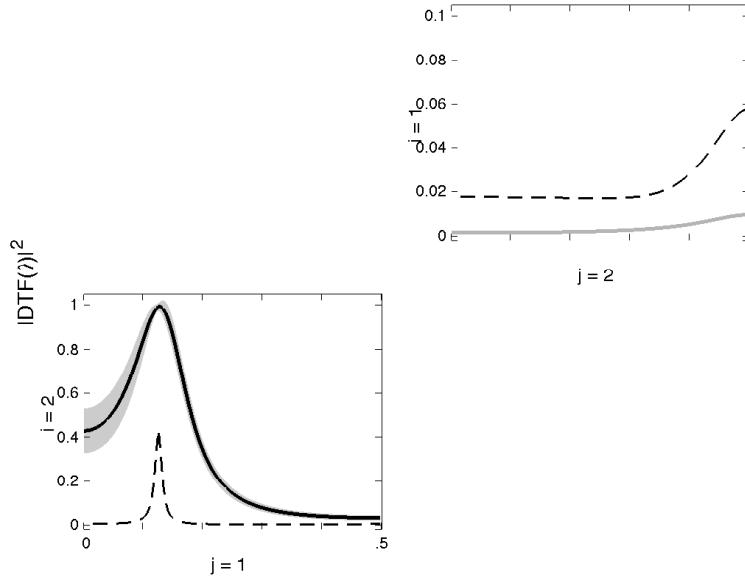
PDC analysis results are saved in **d** structure. See asymp\_dtf.m or issue >> help asymp\_pdc command for more detail.

```

nFreqs = 128;
metric = 'euc'; % euc = Euclidian = original PDC;
%                 % diag = diagonal = generalized PDC or gPDC;
%                 % info = information = iPDC
alpha = 0.01;

```

Example 1 - simple example; sigma2 = 100, sigma1 = sigma3 = 1; ns = 500; p = 2  $\alpha$  = 1%.



```
d = asymp_pdc(u,A,PF,nFREQS,metric,alpha);
* Original PDC and asymptotic statistics
```

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

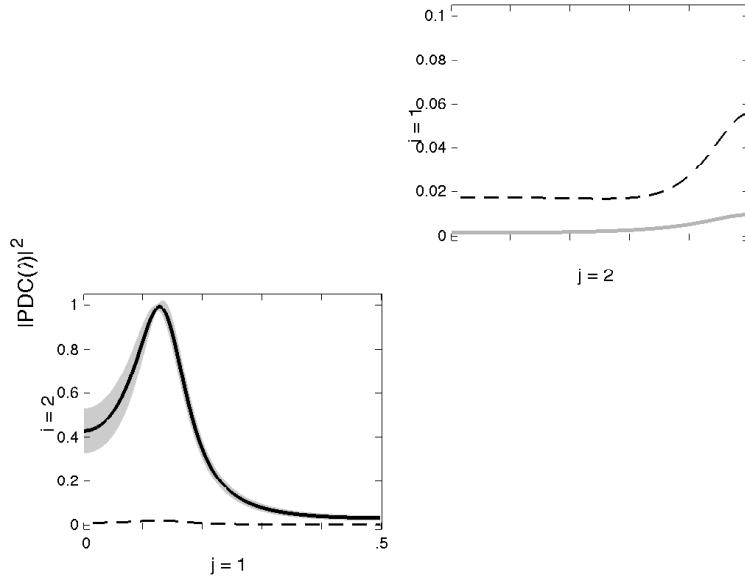
```
% -----Plotting options flag setting-----
% [1 2 3 4 5 6 7]
flgPrinting=[1 1 1 2 2 0 0];
% | | | | | 7 Spectra(0: w/o SS; 1: Linear; 2: log-scale)
% | | | | 6 Coherence
% | | | | 5 Plot lower confidence limit (legacy)
% | | | | 4 Plot upper confidence limit
% | | | 3 Significant DTF(w) in red line (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 2. Example 1 PDC, ns = 500')
[h xlabel ylabel] = xplot(d,...
    flgPrinting,fs,w_max,chLabels,flgColor,flgScale,flgMax,flgSignifColor);

xplot_title(alpha,metric, measure(d));

[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
```

Uncomment the command line bellow to generate an eps output file

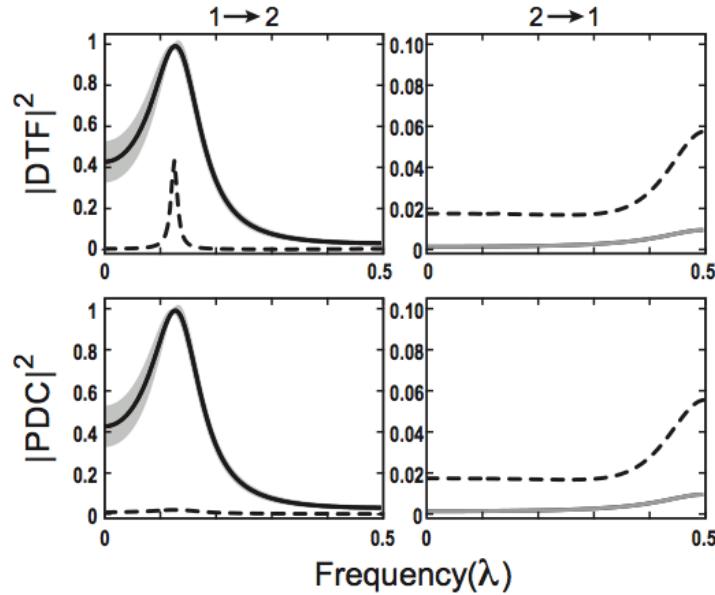
Example 1 - simple example; sigma2 = 100, sigma1 = sigma3 = 1; ns = 500; p = 2  $\alpha$  = 1%.



```
%print -depsc2 -painters Fig2_example1_pdc_ns500.eps
```

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

Figure 1, from article.



### Some remarks:

1. For the final figure 2, 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 2 generation (Baccala et al, 2016)'