

Least-Squares Batch Identification Algorithm - Example

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Identifying an ARX(2, 2) Stochastic Process

Consider the stationary stochastic process \mathcal{S}

$$\mathcal{S} : \quad y(t) = +\frac{6}{5} y(t-1) - \frac{8}{25} y(t-2) + u(t-1) + \frac{1}{2} u(t-2) + \xi(t)$$

$$\xi(\cdot) \sim \text{WN}(0, 1), \quad u(\cdot) \sim \text{WN}(0, 4) \quad e(\cdot), u(\cdot) \text{ uncorrelated}$$

and collect $N_1 = 1500$ samples of both $y(t)$ and $u(t)$

$$Y = \{y(1), y(2), \dots, y(N_1)\} \quad U = \{u(1), u(2), \dots, u(N_1)\}$$

Consider the model \mathcal{M}

$$\mathcal{M}(\vartheta) : y(t) = a_1 y(t-1) + a_2 y(t-2) + b_1 u(t-1) + b_2 u(t-2) + \eta(t) \quad \text{var}[\eta(\cdot)] = \lambda^2$$

and let us identify the model. Let us resort to the **Least-Squares** technique. Then

$$\vartheta = \begin{bmatrix} a_1 \\ a_2 \\ b_1 \\ b_2 \end{bmatrix} \quad \varphi(t) = \begin{bmatrix} y(t-1) \\ y(t-2) \\ u(t-1) \\ u(t-2) \end{bmatrix}$$

and hence

$$\mathcal{M}(\vartheta): y(t) = \varphi(t)^\top \vartheta + \eta(t)$$

$$\widehat{\mathcal{M}}(\vartheta): \widehat{y}(t|t-1) = \varphi(t)^\top \vartheta$$

The **predictor** has a **linear structure** with respect to the **vector ϑ of unknown parameters**.

The prediction error $\varepsilon(t)$ is given by

$$\varepsilon(t) = y(t) - \widehat{y}(t|t-1) = y(t) - \varphi(t)^\top \vartheta$$

Consider the quadratic cost function

$$J(\vartheta) = \frac{1}{N_1} \sum_{t=1}^{N_1} [\varepsilon(t)]^2 = \frac{1}{N_1} \sum_{t=1}^{N_1} [y(t) - \varphi(t)^\top \vartheta]^2$$

and the minimizing vector

$$\vartheta^\circ = \arg \min_{\vartheta} J(\vartheta)$$

If $\sum_{t=1}^N \varphi(t) \varphi(t)^\top$ is **non-singular**, one gets:

$$\widehat{\vartheta}_{N_1} = \left[\sum_{t=1}^{N_1} \varphi(t) \varphi(t)^\top \right]^{-1} \sum_{t=1}^{N_1} \varphi(t) y(t)$$

and the estimate uncertainty is

$$\text{var}[\widehat{\vartheta}_{N_1}] = \widehat{\lambda}_{N_1}^2 \left[\sum_{t=1}^{N_1} \varphi(t) \varphi(t)^\top \right]^{-1}$$

where $\widehat{\lambda}_{N_1}^2$ is an empirical estimate of λ^2

$$\widehat{\lambda}_{N_1}^2 = \frac{1}{N_1} \sum_{t=1}^{N_1} \left[y(t) - \varphi(t)^\top \widehat{\vartheta}_{N_1} \right]^2$$

The condition

$$\det \left[\sum_{t=1}^N \varphi(t) \varphi(t)^\top \right] \neq 0$$

is called **Identifiability Condition**.

ARX Process Simulation

Generation of the Data

```

clear
close all
clc

N1 = 500; % the number of sample data to simulate and collect

var_xi = 1; % the white noise variance
mean_xi = 0; % the white noise mean value

var_u = 4; % the input variable variance
mean_u = 0; % the input variable mean value

Az1 = [1, -6/5, +8/25];
% the coefficients of the polynomial A(z) <-> 1 - a_1 * z^(-1) - a_2 * z^(-2)
%                                     z^2 - a_1 * z^1      - a_2 * z^0

Bz1 = [0, 1, 1/2];
% the coefficients of the polynomial B(z) <-> b_1 * z^(-1) + b_2 * z^(-2)
%                                     b_1 * z^1      + b_2 * z^0
Ts = 1; % sampling time
Gz1 = tf(Bz1, Az1, Ts, 'Variable', 'z^-1')

```

Gz1 =

$$\frac{z^{-1} + 0.5 z^{-2}}{1 - 1.2 z^{-1} + 0.32 z^{-2}}$$

Sample time: 1 seconds
Discrete-time transfer function.

```
Gz = tf(Bz1, Az1, Ts, 'Variable', 'z')
```

Gz =

$$\frac{z + 0.5}{z^2 - 1.2 z + 0.32}$$

Sample time: 1 seconds
Discrete-time transfer function.

Let's estimate how long is the initial transient output of the filter $G(z) = \frac{B(z)}{A(z)}$:

```

p_vector = pole(Gz); % the poles of W(z)
z_vector = zero(Gz); % the zeros of W(z)

slowest_pole = max(abs(p_vector));

zero_threshold = 1e-8;
% let's assume that the value is practically zero if less than or equal to zero_threshold

```

```
Ntransient = 50*(ceil(log10(zero_threshold)/log10(slowest_pole))+10); % a very raw estimate
```

```
N_TOT_data = N1 + Ntransient
```

```
N_TOT_data = 5150
```

```
rng('shuffle');
% seeds the random number generator based on the current
% time so that RAND, RANDI, and RANDN produce a different sequence of
% numbers after each time you call rng.
WN_samples = mean_xi + sqrt(var_xi)*randn(1, N_TOT_data); % the white noise sequence

rng('shuffle');
U_samples = mean_u + sqrt(var_u)*randn(1, N_TOT_data); % the input u sequence
```

Remember: the stochastic process \mathcal{S} is an *ARX* process

$$\mathcal{S}: \quad y(t) = +\frac{6}{5} y(t-1) - \frac{8}{25} y(t-2) + u(t-1) + \frac{1}{2} u(t-2) + \xi(t) \quad (+)$$

Let us rewrite the model \mathcal{S} , pointing in evidence the filters $G(z)$ and $W(z)$:

$$\left[1 - \frac{6}{5}z^{-1} + \frac{8}{25}z^{-2}\right] y(t) = \left[z^{-1} + \frac{1}{2}z^{-2}\right] u(t) + \xi(t)$$

$$A(z) y(t) = B(z) u(t) + \xi(t) \quad A(z) = \left[1 - \frac{6}{5}z^{-1} + \frac{8}{25}z^{-2}\right] \quad B(z) = \left[z^{-1} + \frac{1}{2}z^{-2}\right]$$

$$y(t) = \frac{B(z)}{A(z)} u(t) + \frac{1}{A(z)} \xi(t) \quad G(z) = \frac{B(z)}{A(z)} \quad W(z) = \frac{1}{A(z)}$$

Finally

$$y(t) = G(z) u(t) + W(z) \xi(t) \quad (\star)$$

Now we may generate a set of data of the process \mathcal{S} , using the [Eq. \(\$\star\$ \)](#) or the explicit difference equation in [Eq. \(+\)](#).

```
% let us generate samples using Eq. (*)
% Remember: describe the numerator and denominator of both the transfer
% functions G(z) and W(z) using the notation with ascending powers of z^-1

V_ARX_tot = filter(Bz1, Az1, U_samples) + ...
            filter(1, Az1, WN_samples); % G(z)*u(t) + W(z)*xi(t)

V_ARX = V_ARX_tot(Ntransient+1:end); %#ok<*NASGU>
% let's retrieve the N1 samples when the initial transient part is over
```

Let us implement Eq. (+) to generate the samples.

```
v_y = zeros(1, N_TOT_data); % the array recipient for the r.v. y(t) in Eq.(+)

% --- solving Eq. (+) in a "for" loop ---
for k=1:N_TOT_data

    % ---- checking if the values
    %       at the time instant (t-2) are available ----
    if ((k-2)>0) % if available, then use them
        vy_k2 = v_y(k-2);
        u_k2 = U_samples(k-2);
    else % if not available then set them to zero
        vy_k2 = 0;
        u_k2 = 0;
    end % if ((k-2)>0)

    % ---- checking if the values
    %       at the time instant (t-1) are available ----
    if ((k-1)>0) % if available, then use them
        vy_k1 = v_y(k-1);
        u_k1 = U_samples(k-1);
    else % if not available then set them to zero
        vy_k1 = 0;
        u_k1 = 0;
    end % if ((k-1)>0)

    % ---- evaluate y(t) according to Eq. (+) ----
    v_y(k) = -Az1(2)*vy_k1 -Az1(3)*vy_k2 +...
              +Bz1(1)*U_samples(k) +Bz1(2)*u_k1+Bz1(3)*u_k2 +...
              WN_samples(k);

end
```

Let us compare the resulting sequences:

```
compare_S_sequences = abs(v_y-V_ARX_tot);
V_ARX_tot(1:6)-v_y(1:6)
```

```
ans = 1×6
10-14 ×
    0         0    0.0444         0         0    -0.1776
```

```
fprintf(1, "*** Max abs value of the difference between the two sequences: %e ***\n",
        max(compare_S_sequences));
```

```
*** Max abs value of the difference between the two sequences: 1.421085e-14 ***
```

Both approaches lead to the same sequence.

The Dataset

```
V_ARX = V_ARX_tot(Ntransient+1:end);  
% the N1 samples, collected when the initial transient  
% (due to the initial simulation conditions) has practically disappeared
```

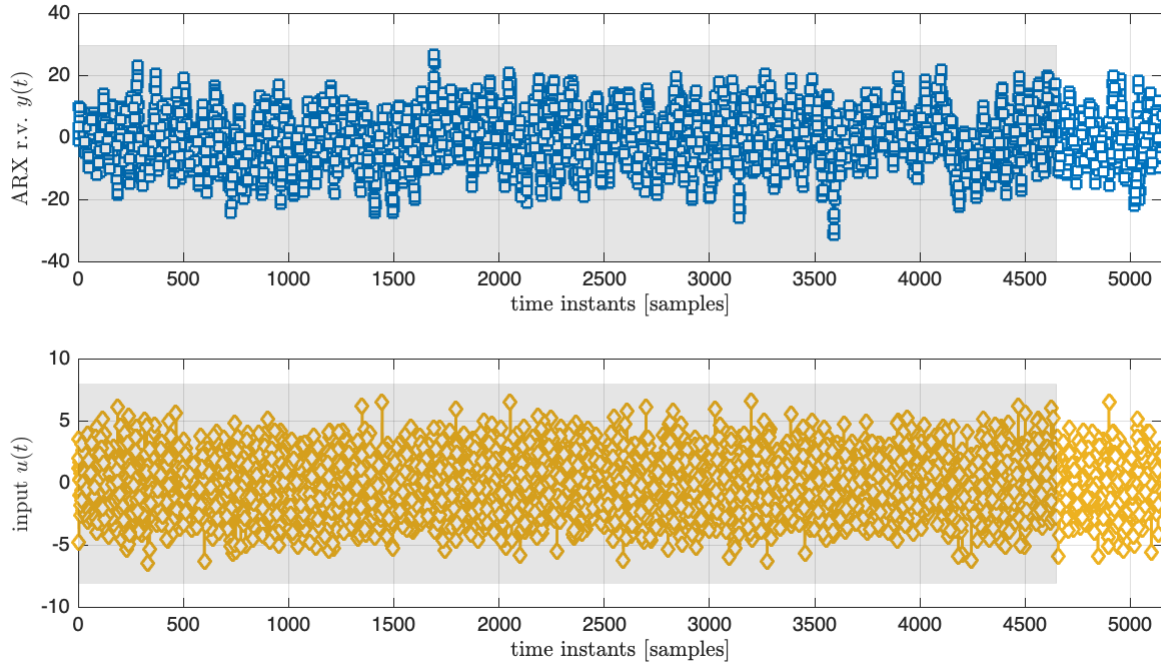
Plot the Process Data

Let us define a set of colors

```
% ---- possible colors ----  
FrenchBlue   = [0 0.4470 0.7410];      % French Blue  
FlameRed     = [0.8500 0.3250 0.0980]; % Flame Red  
Xanthous     = [0.9290 0.6940 0.1250]; % Xanthous  
VividViolet  = [0.4940 0.1840 0.5560]; % Vivid Violet  
AppleGreen   = [0.4660 0.6740 0.1880]; % Apple Green  
BlizzardBlue = [0.3010 0.7450 0.9330]; % Blizzard Blue  
VividBurgundy = [0.6350 0.0780 0.1840]; % Vivid Burgundy  
% ---- possible colors ----
```

and plot the collected data:

```
figure('Units','normalized', 'Position', [0.1, 0.1, 0.9, 0.75]);  
hp1 = subplot(2, 1, 1);  
plot(V_ARX_tot, 'Color', FrenchBlue, 'LineWidth', 1.5, 'LineStyle', '-', 'Marker', 'square',  
      'MarkerSize', 6, 'MarkerEdgeColor', 'auto', 'MarkerFaceColor', 'auto');  
grid on; hold on;  
xlabel('time instants [samples]', 'Interpreter', 'latex');  
ylabel('ARX r.v.  $y(t)$ ', 'Interpreter', 'latex');  
xlim([0, numel(V_ARX_tot)]); % adjusting the x-axis limits  
% the gray part corresponds to the transient part of the whitening filter response  
YL = ylim;  
fill([1, Ntransient(end), Ntransient(end), 1], [YL(1), YL(1), YL(2), YL(2)], ...  
     'k', 'FaceAlpha', 0.10, 'EdgeColor', 'k', 'EdgeAlpha', 0.10);  
  
hp2 = subplot(2, 1, 2);  
plot(U_samples, 'Color', Xanthous, 'LineWidth', 1.5, 'LineStyle', '-', 'Marker', 'diamond',  
      'MarkerSize', 6, 'MarkerEdgeColor', 'auto', 'MarkerFaceColor', 'auto');  
grid on; hold on;  
xlabel('time instants [samples]', 'Interpreter', 'latex');  
ylabel('input  $u(t)$ ', 'Interpreter', 'latex');  
xlim([0, numel(V_ARX_tot)]); % adjusting the x-axis limits  
% the gray part corresponds to the transient part of the whitening filter response  
YL = ylim;  
fill([1, Ntransient(end), Ntransient(end), 1], [YL(1), YL(1), YL(2), YL(2)], ...  
     'k', 'FaceAlpha', 0.10, 'EdgeColor', 'k', 'EdgeAlpha', 0.10);  
linkaxes([hp1, hp2], 'x');
```



Persistency of Excitation and the Identifiability Condition

The regressor $\varphi(t)$ is given by

$$\varphi(t) = \begin{bmatrix} y(t-1) \\ y(t-2) \\ u(t-1) \\ u(t-2) \end{bmatrix}$$

and consequently, the matrices S and R take on the expressions

$$S(N_1) = \begin{bmatrix} \sum_{t=3}^{N_1} [y(t-1)]^2 & \sum_{t=3}^{N_1} y(t-1)y(t-2) & \sum_{t=3}^{N_1} y(t-1)u(t-1) & \sum_{t=3}^{N_1} y(t-1)u(t-2) \\ \sum_{t=3}^{N_1} y(t-2)y(t-1) & \sum_{t=3}^{N_1} [y(t-2)]^2 & \sum_{t=3}^{N_1} y(t-2)u(t-1) & \sum_{t=3}^{N_1} y(t-2)u(t-2) \\ \sum_{t=3}^{N_1} u(t-1)y(t-1) & \sum_{t=3}^{N_1} u(t-1)y(t-2) & \sum_{t=3}^{N_1} [u(t-1)]^2 & \sum_{t=3}^{N_1} u(t-1)u(t-2) \\ \sum_{t=3}^{N_1} u(t-2)y(t-1) & \sum_{t=3}^{N_1} u(t-2)y(t-2) & \sum_{t=3}^{N_1} u(t-2)u(t-1) & \sum_{t=3}^{N_1} [u(t-2)]^2 \end{bmatrix} \quad R(N_1) = \frac{1}{N_1 - 2} S(N_1)$$

Remark: there is a more efficient way to evaluate the matrix S . In fact, given N_1 observations, we build the following matrices and vectors

$$\Phi_{N_1} = \begin{bmatrix} \varphi(1)^\top \\ \vdots \\ \varphi(N_1)^\top \end{bmatrix} \quad \varphi(t) = \begin{bmatrix} y(t-1) \\ y(t-2) \\ u(t-1) \\ u(t-2) \end{bmatrix} \quad \vartheta = \begin{bmatrix} a_1 \\ a_2 \\ b_1 \\ b_2 \end{bmatrix} \quad Y_N = \begin{bmatrix} y(1) \\ \vdots \\ y(N_1) \end{bmatrix}$$

Now rewrite the LS problem in a compact form as

$$\Phi_{N_1} \cdot \vartheta = Y_{N_1}$$

To obtain the *LS normal equations* in compact form, simply multiply both terms of this expression to the left using the transposed matrix $\Phi_{N_1}^\top$:

$$\Phi_{N_1}^\top \Phi_{N_1} \vartheta = \Phi_{N_1}^\top Y_{N_1}$$

Finally

$$S = \Phi_{N_1}^\top \Phi_{N_1}$$

Now we may

- check if $\Phi_{N_1}^\top \Phi_{N_1}$ is a full rank matrix (i.e. if the identifiability condition is fulfilled);
- check if the sub-matrix S_{uu} is full-rank (i.e. if we have persistency of excitation - a necessary condition for the identifiability condition to hold true)

$$S_{uu}(N_1) = \begin{bmatrix} \sum_{t=3}^{N_1} [u(t-1)]^2 & \sum_{t=3}^{N_1} u(t-1)u(t-2) \\ \sum_{t=3}^{N_1} u(t-2)u(t-1) & \sum_{t=3}^{N_1} [u(t-2)]^2 \end{bmatrix}$$

The Identifiability Condition

```
U = U_samples(Ntransient+1:end);
Y = V_ARX;
matPHI = zeros(N1-1, 4); % preallocation of the PHI regressors matrix

for k=3:N1
    matPHI(k,:) = [Y(k-1), Y(k-2), U(k-1), U(k-2)];
end % for k
matPHI(1:2,:) = []; % we can not evaluate the model for t=1 and t=2
```



```

Y_N1 = Y(3:end)';    % we can not evaluate the model for t=1 and t=2

S_N1 = matPHI'*matPHI;
if (rank(S_N1)<4)
    error('Matrix S not invertible! Identifiability condition not fulfilled!')
else
    disp('Matrix S invertible! Identifiability condition fulfilled!')
end

```

Matrix S invertible! Identifiability condition fulfilled!

Identification - the LS Algorithm

$$\hat{\vartheta}_{N_1} = \left[\sum_{t=1}^{N_1} \varphi(t) \varphi(t)^\top \right]^{-1} \sum_{t=1}^{N_1} \varphi(t) y(t)$$

or in compact form (refer to the [previous section](#) of the live script where we recall it)

$$\Phi_{N_1} \cdot \vartheta = Y_{N_1}$$

$$\Phi_{N_1}^\top \Phi_{N_1} \vartheta = \Phi_{N_1}^\top Y_{N_1}$$

Thus we may solve the LS equation using the MATLAB operator "`\`" using the compact formulation

$$\Phi_{N_1} \cdot \vartheta = Y_{N_1}$$

```

hat_theta_N1 = matPHI \ Y_N1

```

```

hat_theta_N1 = 4x1
    1.1776
   -0.2972
    0.9878
    0.4731

```

Alternatively, we may solve the LS normal equations $\Phi_{N_1}^\top \Phi_{N_1} \vartheta = \Phi_{N_1}^\top Y_{N_1}$, exploiting one more time the operator "`\`"

```

hat_theta_N1_ALT = S_N1 \ (matPHI'*Y_N1)

```

```

hat_theta_N1_ALT = 4x1
    1.1776
   -0.2972
    0.9878
    0.4731

```

As can be seen, both approaches lead to the same solution.

The Estimate Uncertainty

What about the estimate uncertainty?

According to the theoretical result, described in the class material (refer to L12 - p. 24), for N sufficiently large the variance of the estimator is $\frac{1}{N} \text{var}[\epsilon(\vartheta^o)] [\bar{R}(\vartheta^o)]^{-1}$

Computing the empirical estimate, one gets

$$\text{var}[\hat{\vartheta}_N] = \hat{\lambda}^2 \left[\frac{1}{N} \sum_{t=1}^N \varphi(t) \varphi(t)^\top \right]^{-1} = \hat{\lambda}^2 S(N)^{-1}$$

where

$$\hat{\lambda}^2 = \frac{1}{N} \sum_{t=1}^N [y(t) - \varphi(t)^\top \hat{\vartheta}_N]^2$$

In this example one gets

```
% the estimate of the prediction error variance
pred_residuals = Y_N1 - matPHI*hat_theta_N1;
hat_lambda2_N1 = mean((pred_residuals).^2)
```

```
hat_lambda2_N1 = 0.9470
```

```
cond(S_N1)
```

```
ans = 106.3693
```

```
var_theta_N1 = hat_lambda2_N1 .* inv(S_N1)
```

```
var_theta_N1 = 4x4
    0.0005    -0.0005     0.0000    -0.0006
   -0.0005     0.0005     0.0000     0.0005
    0.0000     0.0000     0.0005    -0.0000
   -0.0006     0.0005    -0.0000     0.0011
```

Alternatively

```
R_N1 = S_N1 ./ N1;
cond(R_N1)
```

```
ans = 106.3693
```

```
var_theta_N1_ALT = (hat_lambda2_N1/N1) .* inv(R_N1)
```

```
var_theta_N1_ALT = 4x4
    0.0005    -0.0005     0.0000    -0.0006
   -0.0005     0.0005     0.0000     0.0005
    0.0000     0.0000     0.0005    -0.0000
   -0.0006     0.0005    -0.0000     0.0011
```

Remark

The covariance matrix is not a diagonal matrix: there is a weak link between some of the parameters.

The Whiteness Test

What about the prediction residuals? Are they samples of a stationary stochastic white noise process or?

Actually we have the prediction residuals, so let us apply the **Anderson's whiteness test**

```
WTest_result = AndersonWhitenessTest(pred_residuals, 0.01, true)
```

```
AndersonWhitenessTest: confidence level 0.010000  
AndersonWhitenessTest: max lag value M 24  
AndersonWhitenessTest: critical value 0  
AndersonWhitenessTest: accept the hypothesis -->  
the prediction error IS a stationary  
white noise stochastic process  
WTest_result = 0
```