

Linear predictive coding

This method combines **linear processing with scalar quantization**.

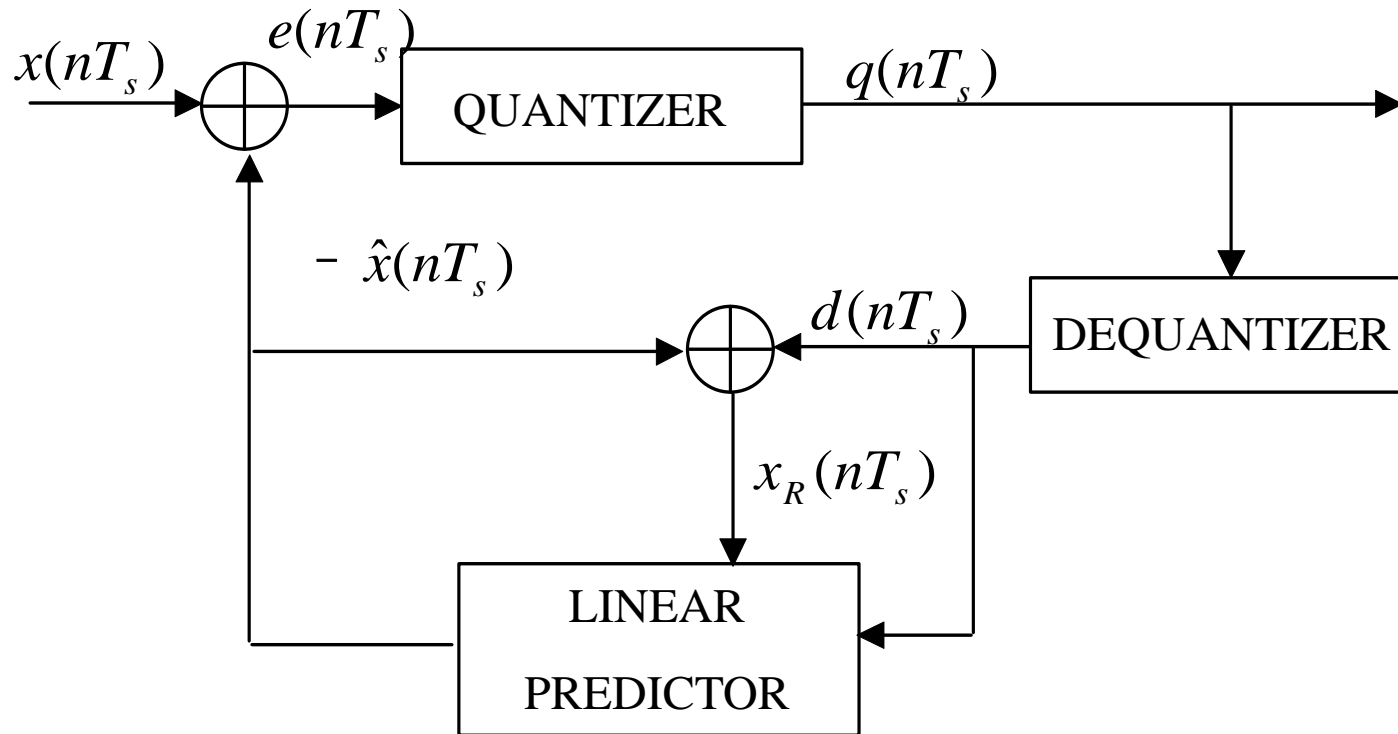
The main idea of the method is **to predict the value of the current sample by a linear combination of previous already reconstructed samples** and then to quantize **the difference** between the actual value and the predicted value.

Linear prediction coefficients are weighting coefficients used in linear combination.

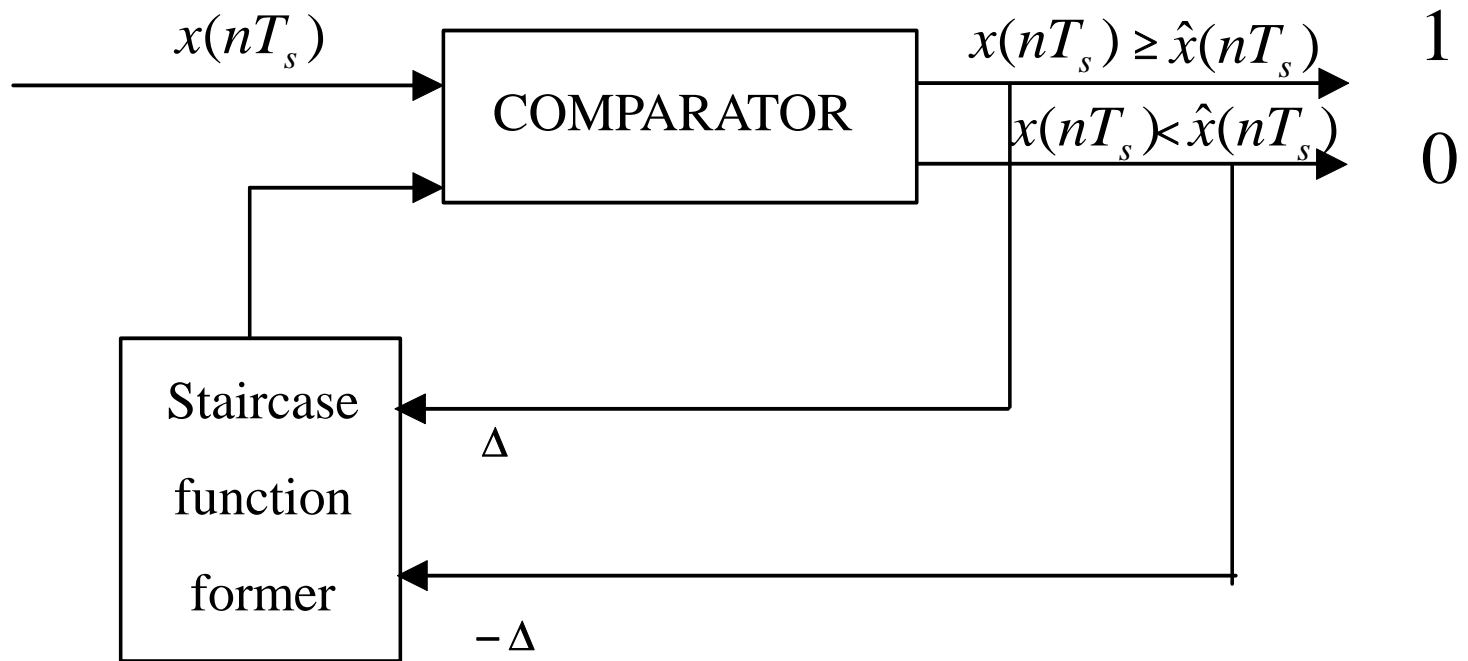
A simple predictive quantizer or **differential pulse-coded modulator** is shown in Fig. 5.1.

If the **predictor** is simply the **last sample** and the quantizer has only one bit, the system becomes a **delta-modulator**. It is shown in Fig. 5.2.

Differential pulse-coded modulator



Delta-modulator



Linear predictive coding

The main feature of the quantizers shown in Figs 5.1, 5.2 is that they exploit not all advantages of predictive coding.

- Prediction coefficients used in these schemes are not optimal
- Prediction is based on past reconstructed samples and not true samples
- Usually coefficients of prediction are chosen by using some empirical rules and are not transmitted

For example quantizer in Fig.5.1 instead of actual value of error e uses reconstructed values d and instead of true sample values x their estimates x_R obtained via d .

Linear predictive coding

The most **advanced quantizer** of linear predictive type represents a basis of the so-called **Code Excited Linear Predictive (CELP) coder**.

- It uses the **optimal set of coefficients** or in other words linear prediction coefficients of this quantizer are determined **by minimizing the MSE** between the current sample and its predicted value.
- It is based on the **original past samples**
- Using the true samples for prediction requires the **“looking-ahead” procedure** in the coder.
- The **predictor coefficients are transmitted**

Linear predictive coding

Assume that prediction coefficients are optimized for each frame and that the original past samples are used for prediction. Let

$x(T_s), x(2T_s), \dots$ be a sequence of samples at the quantizer input. Then each sample $x(nT_s)$ is predicted by the previous samples according to the formula

$$\hat{x}(nT_s) = \sum_{k=1}^m a_k x(nT_s - kT_s), \quad (5.1)$$

where $\hat{x}(nT_s)$ is the predicted value, a_k are prediction coefficients, m denotes the order of prediction. The prediction error is

$$e(nT_s) = x(nT_s) - \hat{x}(nT_s).$$

Linear predictive coding

Prediction coefficients are determined by minimizing the sum of squared errors over a given interval (frame)

$$E = \sum_{n=n_0}^{n_1} e^2(nT_s) \quad (5.2)$$

Inserting (5.1) into (5.2) we obtain

$$\begin{aligned} E &= \sum_{n=n_0}^{n_1} (x(nT_s) - a_1 x(nT_s - T_s) - \dots - a_m x(nT_s - mT_s))^2 = \\ &= \sum_{n=n_0}^{n_1} \{x(nT_s)\}^2 - 2 \sum_{j=1}^m a_j \sum_{n=n_0}^{n_1} x(nT_s) x(nT_s - jT_s) + \\ &+ \sum_{j=1}^m \sum_{k=1}^m a_j a_k \sum_{n=n_0}^{n_1} x(nT_s - jT_s) x(nT_s - kT_s). \quad (5.3) \end{aligned}$$

Linear predictive coding

Differentiating (5.3) over a_k , $k = 1, 2, \dots, m$ yields

$$\partial E / \partial a_k = \sum_{n=n_0}^{n_1} x(nT_s)x(nT_s - kT_s) - \sum_{j=1}^m a_j \sum_{n=n_0}^{n_1} x(nT_s - kT_s)x(nT_s - jT_s) = 0$$

Thus we obtain a system of m linear equations with m unknown quantities a_1, a_2, \dots, a_m

$$\sum_{j=1}^m a_j c_{jk} = c_{ok}, \quad k = 1, 2, \dots, m \quad (5.4)$$

where $c_{jk} = c_{kj} = \sum_{n=n_0}^{n_1} x(nT_s - jT_s)x(nT_s - kT_s).$ (5.5)

The system (5.4) is called **the Yule-Walker equations**.

Linear predictive coding

If a_1, a_2, \dots, a_m are solutions of (5.4) then we can find the minimal achievable prediction error. Insert (5.5) into (5.3).

We obtain that

$$E = c_{00} - 2 \sum_{k=1}^m a_k c_{0k} + \sum_{k=1}^m a_k \sum_{j=1}^m a_j c_{jk}. \quad (5.6)$$

Using (5.3) we reduce (5.6) to

$$E = c_{00} - \sum_{k=1}^m a_k c_{0k}$$

Linear prediction as digital filtering

Eq. (5.1) describes the m th order predictor with transfer function equal to

$$P(z) = \frac{\hat{X}(z)}{X(z)} = \sum_{k=1}^m a_k z^{-k}.$$

z -transform for the prediction error is

$$E(z) = X(z) - \sum_{k=1}^m a_k X(z) z^{-k}.$$

The prediction error is an output signal of the discrete-time filter with transfer function

$$A(z) = \frac{E(z)}{X(z)} = 1 - \sum_{k=1}^m a_k z^{-k}.$$

The problem of finding the optimal set of prediction coefficients = problem of constructing m th order FIR filter.

Linear prediction as digital filtering

Another name of the linear prediction (5.1) is the **autoregressive model (AM)** of signal $x(nT_s)$. It is assumed that the signal $x(nT_s)$ can be obtained as the output of the autoregressive filter with transfer function

$$H(z) = \frac{1}{\left(1 - \sum_{k=1}^m a_k z^{-k}\right)},$$

that is can be obtained as the output of the filter which is **inverse** with respect to the **prediction filter**. This filter is a discrete-time IIR filter. **The Y-W equations determine correspondence between parameters of AM and covariance function of the process.**

Methods of finding coefficients

$$c_{ij}, i = 0, 1, \dots, m, j = 1, 2, \dots, m$$

In order to solve the Yule-Walker eq. (5.4) it is necessary first to evaluate values $c_{ij}, i = 0, 1, \dots, m, j = 1, 2, \dots, m$

There are two approaches to estimating these values:

The autocorrelation method and the covariance method.

The complexity of solving (5.4) is proportional to m^2

The complexity of solving (5.4) is proportional to m^3

Autocorrelation method

The values c_{ij} are computed as

$$c_{ij} = c_{ji} = \sum_{n=i_0}^{i_1} x(nT_s - iT_s)x(nT_s - jT_s). \quad (5.7)$$

We set $i_0 = -\infty, i_1 = \infty$ and $x(nT_s) = 0$ if $n < 0, n \geq N$, where N is called the interval of analysis.

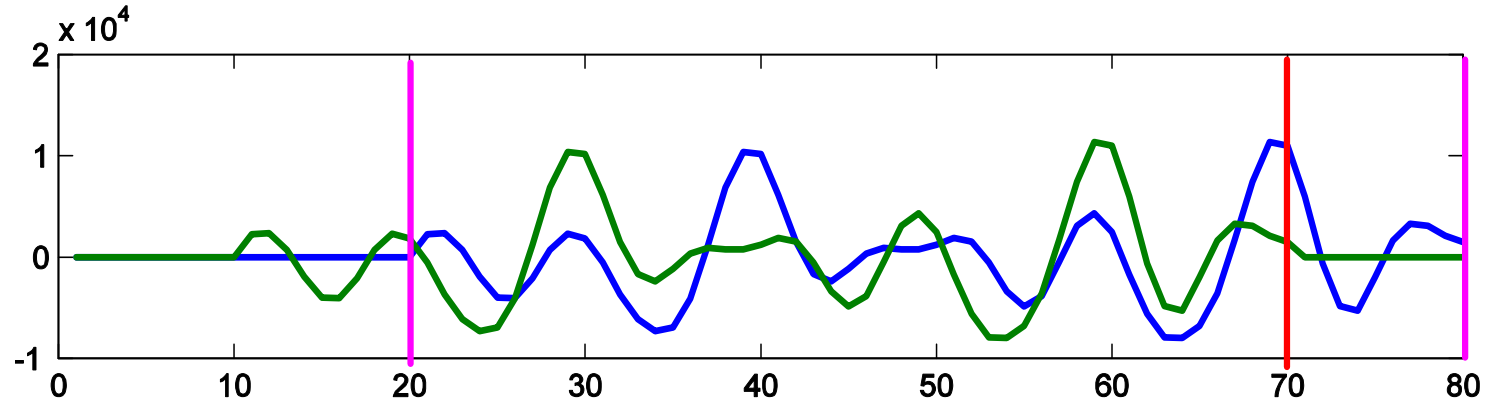
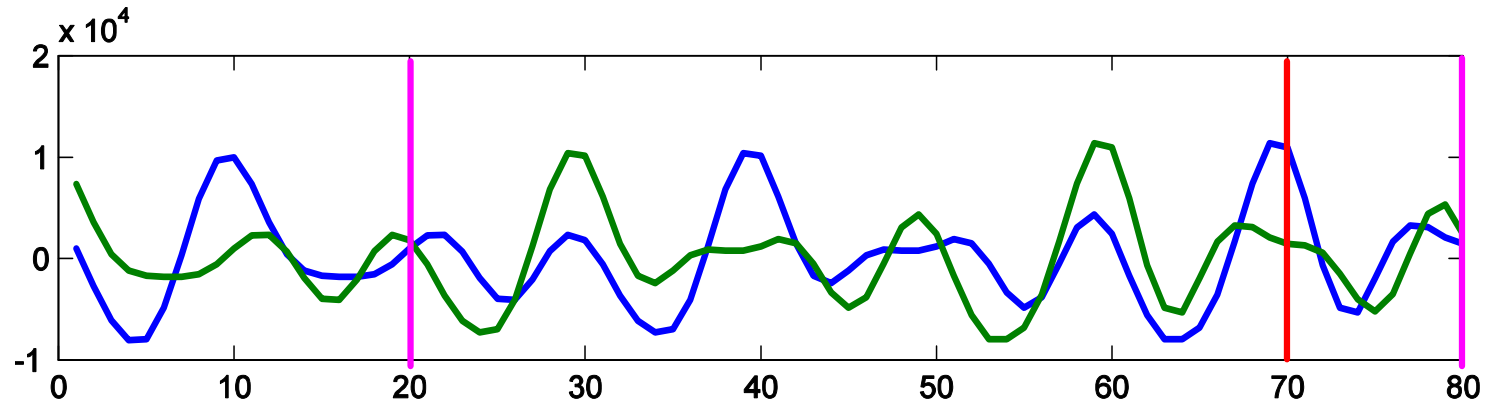
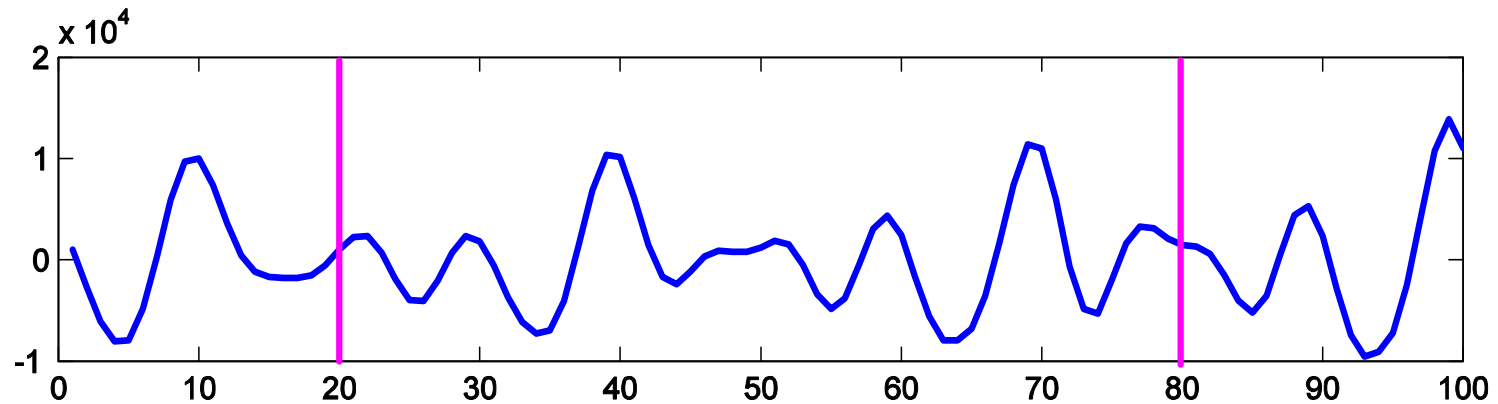
In this case we can simplify (5.7)

$$c_{ij} = \sum_{n=0}^{N-1} x(nT_s - iT_s)x(nT_s - jT_s) = \sum_{n=0}^{N-1-|i-j|} x(nT_s)x(nT_s + |i-j|T_s). \quad (5.8)$$

Normalized by N they coincide with estimates of entries of covariance matrix for $x(nT_s)$

$$\hat{R}(|i-j|) = c_{ij} / N = 1/N \sum_{n=0}^{N-1-|i-j|} x(nT_s)x(nT_s + |i-j|T_s).$$

Autocorrelation method



Autocorrelation method

The Yule-Walker equations for autocorrelation method have the form

$$\sum_{i=1}^m a_i \hat{R}(|i-j|) = \hat{R}(j), j = 1, 2, \dots, m. \quad (5.9)$$

Eq.(5.9) can be given by matrix equation

$$\mathbf{a} \times \mathbf{R} = \mathbf{b},$$

where $\mathbf{a} = (a_1, a_2, \dots, a_m)$, $\mathbf{b} = (\hat{R}(1), \hat{R}(2), \dots, \hat{R}(m))$,

$$\mathbf{R} = \begin{bmatrix} \hat{R}(0) & \hat{R}(1) \dots & \hat{R}(m-1) \\ \hat{R}(1) & \hat{R}(0) \dots & \hat{R}(m-2) \\ \dots & \dots & \dots \\ \hat{R}(m-1) & \hat{R}(m-2) \dots & \hat{R}(0) \end{bmatrix}.$$

Autocorrelation method

It is said that (5.9) relates the parameters of the autoregressive model of m th order with the autocorrelation sequence.

Matrix \mathbf{R} of the autocorrelation method has two important properties.

- It is **symmetric**, that is $\hat{R}(i, j) = \hat{R}(j, i)$
- It has the **Toeplitz** property, that is $\hat{R}(i, j) = \hat{R}(|i - j|)$.

The Toeplitz property of \mathbf{R} makes it possible to reduce the computational complexity of solving (5.4). The fast Levinson-Durbin recursive algorithm requires only m^2 operations.

Covariance method

We choose $i_0 = 0$ and $i_1 = N - 1$ and signal $x(nT_s)$ is not constrained in time. In this case we have

$$c_{ij} = \sum_{n=0}^{N-1} x(nT_s - iT_s)x(nT_s - jT_s). \quad (5.10)$$

Set $k = n - i$ then (5.10) can be rewritten as

$$c_{ij} = \sum_{k=-i}^{N-i-1} x(kT_s)x(kT_s + (i - j)T_s), i = 1, \dots, m, j = 0, \dots, m. \quad (5.11)$$

(5.11) resembles (5.8) but it has other range of definition for k .

- It uses **signal values out of range** $0 \leq k \leq N - 1$,
- The method leads to the **cross-correlation function** between two similar but not exactly the same finite fragments of $x(kT_s)$.

Covariance method

$$\hat{R}(i, j) = c_{ij} / N = 1/N \sum_{n=0}^{N-1} x((n-i)T_s)x((n-j)T_s).$$

The Yule-Walker equations for the covariation method are

$$\sum_{i=1}^m a_i \hat{R}(i, j) = \hat{R}(0, j), j = 1, 2, \dots, m. \quad (5.12)$$

Eq. (5.12) can be given by the matrix equation

$$\mathbf{a} \times \mathbf{P} = \mathbf{c},$$

where $\mathbf{a} = (a_1, a_2, \dots, a_m)$, $\mathbf{c} = (\hat{R}(0,1), \hat{R}(0,2), \dots, \hat{R}(0, m))$,

$$\mathbf{P} = \begin{bmatrix} \hat{R}(1,1)\hat{R}(1,2)\dots\hat{R}(1, m) \\ \hat{R}(2,1)\hat{R}(2,2)\dots\hat{R}(2, m) \\ \dots\dots\dots\dots\dots\dots \\ \hat{R}(m,1)\hat{R}(m,2)\dots\hat{R}(m, m) \end{bmatrix}.$$

Covariance method

Unlike the matrix \mathbf{R} of autocorrelation method the matrix \mathbf{P} is **symmetric** ($\hat{R}(i, j) = \hat{R}(j, i)$) but it is **not Toeplitz**.

Since computational complexity of solving an arbitrary system of linear equations of order m is equal to m^3 then in this case to solve (5.12) it is necessary m^3 operations.

Algorithms for the solution of the Yule-Walker equations

The computational complexity of solving the Yule-Walker equations depends on the method of evaluating values c_{ij} .

Assume that c_{ij} are found by the autocorrelation method.

In this case the Yule-Walker equations has the form (5.9) and the matrix \mathbf{R} is symmetric and the Toeplitz matrix.

These properties make it possible to find the solution of (5.9) by fast methods requiring m^2 operations.

There are a few methods of this type: the Levinson-Durbin algorithm, the Euclidean algorithm and the Berlekamp-Massey algorithm.

The Levinson-Durbin algorithm

It was suggested by Levinson in 1948 and then was improved by Durbin in 1960. Notice that this algorithm works efficiently if matrix of coefficients \mathbf{R} is simultaneously **symmetric and Toeplitz**. The Berlekamp-Massey and the Euclidean algorithms do not require the matrix of coefficients to be symmetric.

We sequentially solve equations (5.9) of order $l = 1, \dots, m$. Let $\mathbf{a}^{(l)} = (a_1^{(l)}, a_2^{(l)}, \dots, a_l^{(l)})$ denote the solution for the system of the l th order. Given $\mathbf{a}^{(l)}$ we find the solution for the $(l + 1)$ th order. At each step of the algorithm we evaluate the prediction error E_l of the l th order system.

The Levinson-Durbin algorithm

Initialization: $l = 0, E_0 = \hat{R}(0), a^{(0)} = 0.$

Recurrent procedure:

For $l = 1, \dots, m$ **compute**

$$a_l^{(l)} = (\hat{R}(l) - \sum_{i=1}^{l-1} a_i^{(l-1)} \hat{R}(l-i)) / E_{l-1},$$

$$a_j^{(l)} = a_j^{(l-1)} - a_l^{(l)} a_{l-j}^{(l-1)}, \quad 1 \leq j \leq l-1,$$

$$E_l = E_{l-1} (1 - (a_l^{(l)})^2).$$

When $l = m$ we obtain the solution

$$\mathbf{a} = (a_1, a_2, \dots, a_m) = \mathbf{a}^{(m)}, E = E_m.$$

Example

$$m = 1 \quad R(0)a_1^{(1)} = R(1), \quad k_1 = -R(1) / R(0),$$

$$a_1^{(1)} = R(1) / R(0),$$

$$a_1^{(1)} = R(1) / R(0),$$

$$E_1 = R(0) - a_1^{(1)} R(1)$$

$$E_1 = (R^2(0) - R^{(2)}(1)) / R(0).$$

$$m = 2 \quad \begin{cases} R(0)a_1^{(2)} + R(1)a_2^{(2)} = R(1) \\ R(1)a_1^{(2)} + R(0)a_2^{(2)} = R(2), \end{cases}$$

$$a_1^{(2)} = \frac{R(1)}{R(0)} (1 - a_2^{(2)}) = a_1^{(1)} - a_2^{(2)} a_1^{(1)}$$

$$a_2^{(2)} = \frac{R(2) - R(1)a_1^{(1)}}{R(0) - a_1^{(1)}R(1)} = \frac{R(2) - R(1)a_1^{(1)}}{E_1}$$

Example

$$a_1^{(2)} = (R(1)R(0) - R(1)R(2)) / (R^2(0) - R^2(1))$$

$$a_2^{(2)} = (R(0)R(2) - R(1)^2) / (R^2(0) - R^{(2)}(1)).$$

$$E_2 = E_1 (1 - (a_2^{(2)})^2)$$

$$E_2 = R(0) - a_1^{(2)} R(1) - a_2^{(2)} R(2) = (R(0) - a_1^{(1)} R(1))(1 - (a_2^{(2)})^2) =$$

$$= E_1 (1 - (a_2^{(2)})^2).$$