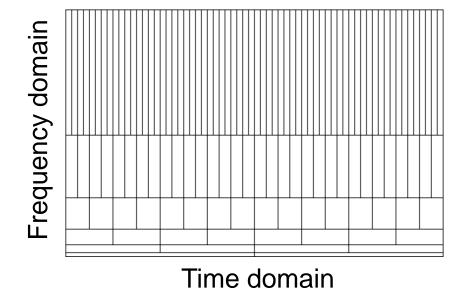


Dyadic Wavelet Transform

In the dyadic wavelet transform, we discretize the wavelet coefficient as

$$W_{x}[j,n] = \sum_{m=0}^{N-1} x[m] \frac{1}{\sqrt{2^{j}}} \psi^{*} \left(\frac{T_{s} \cdot m - 2^{j} \cdot T_{s} \cdot n}{2^{j}} \right), \qquad n, j \in \mathbb{Z}^{+}$$

- At any scale j, the signal is segmented into frames of 2^j samples, with a bandwidth proportional to $\approx 1/(T_{\psi} \cdot 2^j)$
- At the following scale j+1, we obtain two sub-frames of 2^{j+1} samples, with bandwidth proportional to $\approx 1/(T_{\psi} \cdot 2^{j+1})$



Dyadic Wavelet Transform

In the laboratory, we considered an Haar wavelet with support equal to $T_s = T_{\psi}$

This implies that the <u>dyadic</u> wavelet atom associated with scale j=1 is associated with a support of $T_{\psi} \cdot 2^j = 2 \cdot T_s$ seconds and $T_{\psi} \cdot \frac{2^j}{T_s} = 2$ samples

The Wavelet transform $W_x[j,n]$ of $x[\cdot]$ at sample n and scale j=1 is defined as

$$W_{x}[1,n] = \sum_{m=0}^{N-1} x[m] \frac{1}{\sqrt{2}} \psi^{*} \left(\frac{T_{s} \cdot m - 2 \cdot T_{s} \cdot n}{2} \right), \qquad n \in \mathbb{Z}^{+}$$

where $\psi^*\left(\frac{T_S \cdot m - 2 \cdot T_S \cdot n}{2}\right)$ has non-zero values only when $\frac{T_S \cdot m - 2 \cdot T_S \cdot n}{2} \in \left[-\frac{T_S}{2}\right]$, $+\frac{T_S}{2}$

 \blacktriangleright This implies that m must be equal to 2n or 2n-1

Dyadic Wavelet Transform

In the laboratory, we considered an Haar wavelet with support equal to $T_s = T_{\psi}$

This implies that the <u>dyadic</u> wavelet atom associated with a generic scale j is associated with a support of $T_{\psi} \cdot 2^j = 2^j \cdot T_s$ seconds and $T_{\psi} \cdot \frac{2^j}{T_s} = 2^j$ samples

The Wavelet transform $W_x[j,n]$ of $x[\cdot]$ at sample n and scale j is defined as

$$W_{x}[j,n] = \sum_{m=0}^{N-1} x[m] \frac{1}{\sqrt{2^{j}}} \psi^{*} \left(\frac{T_{s} \cdot m - 2^{j} \cdot T_{s} \cdot n}{2^{j}} \right), \qquad n \in \mathbb{Z}^{+}$$

where $\psi^*\left(\frac{T_S \cdot m - 2^j \cdot T_S \cdot n}{2^j}\right)$ has non-zero values only when $\frac{T_S \cdot m - 2^j \cdot T_S \cdot n}{2^j} \in \left[-\frac{T_S}{2}\right]$

> This implies that m must be in $\{2^{j}n - 2^{j-1}, ..., 2^{j}n, ..., 2^{j}n + 2^{j-1} - 1\}$

Given the dyadic WT of $x[\cdot]$ for the scale $j \in \{1, 2, 3, ..., J\}$, we can approximate x[n] as

$$x(n) \approx \frac{\log_e 2}{C_{\psi}} \sum_{j=1}^{J} \frac{1}{2^j} \sum_{m=\lfloor n/2^j \rfloor} W_x[j,m] \psi_{j,m}[n] + \frac{1}{2^J} \sum_{m=\lfloor n/2^J \rfloor} \mathcal{L}_x[J,m] \phi_{J,m}[n]$$

In the case of Haar wavelet, $C_{\psi} = \log_e 2$

This equation is implemented by the method get_reconstructed_signal_v1

The first part of the method implements the first part of the equation, i.e., add to each signal sample x[n] the details associated with all the scales from j = 1 to j = J

```
for sample_index in range(signal_length):
 for j_index, j in enumerate(j_values):
   m = np.floor((sample_index - 2 ** (j-1)) / (2**j)) + 1 # Select the time shift associated with the sample
   atom_domain, atom_wavelet = get_haar_dyadic_wavelet(j, m, sampling_period) # Get the wavelet atom associated with (j,m)
   if atom_domain[0] > 0:
     # We fill with zeros the portion of the signal domain that is not covered by the wavelet
     atom_wavelet = np.concatenate((np.zeros(int(round(sampling_frequency * atom_domain[0]))), atom_wavelet))
     # We do not consider the portion of the wavelet domain that is lower than 0
     atom_wavelet = atom_wavelet[int(np.arqmin(np.abs(atom_domain))):]
   m_values = np.arange(int(signal_length / (2**j))) # Time shifts associated with scale j
   if m in m_values:
     m_index = np.argwhere(m_values == m)
     # We add the detail coefficient, weighted by the wavelet functions, to reconstruct the signal
     reconstructed_signal[sample_index] += detail_coefficients[j_index, m_index] * atom_wavelet[sample_index] / (2 ** j)
     coefficient_masks[j_index, 0, m_index] = 1
```

The first part of the method implements the first part of the equation, i.e., add to each signal sample x[n] the details associated with all the scales from j = 1 to j = J

```
for sample_index in range(signal_length):
 for j_index, j in enumerate(j_values):
   atom_domain, atom_wavelet = get_haar_dyadic_wavelet(j, m, sampling_period) # Get the wavelet atom associated with (j,m)
   if atom_domain[0] > 0:
     # We fill with zeros the portion of the signal domain that is not covered by the wavelet
     atom_wavelet = np.concatenate((np.zeros(int(round(sampling_frequency * atom_domain[0]))), atom_wavelet))
     # We do not consider the portion of the wavelet domain that is lower than 0
     atom_wavelet = atom_wavelet[int(np.arqmin(np.abs(atom_domain))):]
                                                                                                                                                                   W_{x}[j,m]\psi_{j,m}
   m_values = np.arange(int(signal_length / (2**j))) # Time shifts associated with scale j
   if m in m_values:
     m_index = np.argwhere(m_values == m)
     # We add the detail coefficient, weighted by the wavelet functions, to reconstruct the signal
     reconstructed_signal[sample_index] += detail_coefficients[j_index, m_index] * atom_wavelet[sample_index] / (2 ** j)
     coefficient_masks[j_index, 0, m_index] = 1
```

The first part of the method implements the first part of the equation, i.e., add to each signal sample x[n] the details associated with all the scales from j = 1 to j = J

```
for sample_index in range(signal_length):
 for j_index, j in enumerate(j_values):
   m = np.floor((sample_index - 2 ** (j-1)) / (2**j)) + 1 # Select the time shift associated with the sample
   atom_domain, atom_wavelet = get_haar_dyadic_wavelet(j, m, sampling_period) # Get the wavelet atom associated with (j,m)
   m_values = np.arange(int(signal_length / (2**j))) # Time shifts associated with scale j
   if m in m_values:
     m_index = np.argwhere(m_values == m)
     # We add the detail coefficient, weighted by the wavelet functions, to reconstruct the signal
     reconstructed_signal[sample_index] += detail_coefficients[j_index, m_index] * atom_wavelet[sample_index] / (2 ** j)
     coefficient_masks[j_index, 0, m_index] = 1
```

The only scope of this is to align the Wavelet domain with the signal domain, i.e., ensure that the first sample of the Wavelet atom is associated with time t=0

The second part of the method implements the second part of the equation, i.e., add to each signal sample x[n] the approximation associated with the maximum scale j = J

```
if m in m_values:
   m_index = np.argwhere(m_values == m)
   atom_domain, atom_scaling_function = get_haar_dyadic_scaling_function(j, m, sampling_period) # Get the scaling function associated with (j,m)
   if atom_domain[0] > 0:
     # We fill with zeros the portion of the signal domain that is not covered by the wavelet
     atom_scaling_function = np.concatenate((np.zeros(int(round(sampling_frequency * atom_domain[0]))), atom_scaling_function))
                                                                                                                                                                         \mathcal{L}_{x}[J,m]\phi_{J,m}[n]
    else:
                                                                                                                                                            m=|n/2J|
     # We do not consider the portion of the wavelet domain that is lower than 0
      atom_scaling_function = atom_scaling_function[int(np.argmin(np.abs(atom_domain))):]
    # We add the approximation coefficient associated with the maximum scale, weighted by the scaling functions, to reconstruct the signal
   reconstructed_signal[sample_index] += approx_coefficients[j_index, m_index] * atom_scaling_function[sample_index] / (2 ** j)
   coefficient_masks[j_index, 1, m_index] = 1
# and, thus, represents the size of the reconstructed signal
return reconstructed_signal, np.sum(coefficient_masks)
```

The second part of the method implements the second part of the equation, i.e., add to each signal sample x[n] the approximation associated with the maximum scale j = J

```
if m in m_values:
   m_index = np.argwhere(m_values == m)
   atom_domain, atom_scaling_function = get_haar_dyadic_scaling_function(j, m, sampling_period) # Get the scaling function associated with (j,m)
   if atom_domain[0] > 0:
     atom_scaling_function = np.concatenate((np.zeros(int(round(sampling_frequency * atom_domain[0]))), atom_scaling_function))
    else:
                                                                                                                                                             m=|n/2\rangle
     # We do not consider the portion of the wavelet domain that is lower than 0
      atom_scaling_function = atom_scaling_function[int(np.argmin(np.abs(atom_domain))):]
    # We add the approximation coefficient associated with the maximum scale, weighted by the scaling functions, to reconstruct the signal
   reconstructed_signal[sample_index] += approx_coefficients[j_index, m_index] * atom_scaling_function[sample_index] / (2 ** j)
   coefficient_masks[j_index, 1, m_index] = 1
# and, thus, represents the size of the reconstructed signal
return reconstructed_signal, np.sum(coefficient_masks)
```

The second part of the method implements the second part of the equation, i.e., add to each signal sample x[n] the approximation associated with the maximum scale j = J

```
if m in m_values:
   m_index = np.argwhere(m_values == m)
   atom_domain, atom_scaling_function = get_haar_dyadic_scaling_function(j, m, sampling_period) # Get the scaling function associated with (j,m)
                                                                                                                        The only scope of this is to align
                                                                                                                           the Wavelet domain with the
                                                                                                                         signal domain, i.e., ensure that
                                                                                                                         the first sample of the Wavelet
   # We add the approximation coefficient associated with the maximum scale, weighted by the scaling functions, to reconstruct the signa
                                                                                                                      atom is associated with time t = 0
   reconstructed_signal[sample_index] += approx_coefficients[j_index, m_index] * atom_scaling_function[sample_index] / (2 ** j)
   coefficient_masks[j_index, 1, m_index] = 1
# and, thus, represents the size of the reconstructed signal
return reconstructed_signal, np.sum(coefficient_masks)
```