

Lecture 8. Interactions in time series

Outline / overview

- **Section 1.** Source mixing
- **Section 2.** Interactions between sources
- **Section 3.** Covariance and Correlation
- **Section 4.** Regression and Partial correlation
- **Section 5.** Zero-lag interactions
- **Section 6.** Non-zero-lag interactions
- **Section 7.** Causal interactions

Section 1. Source mixing

Linear mixing of sources

What is source mixing?

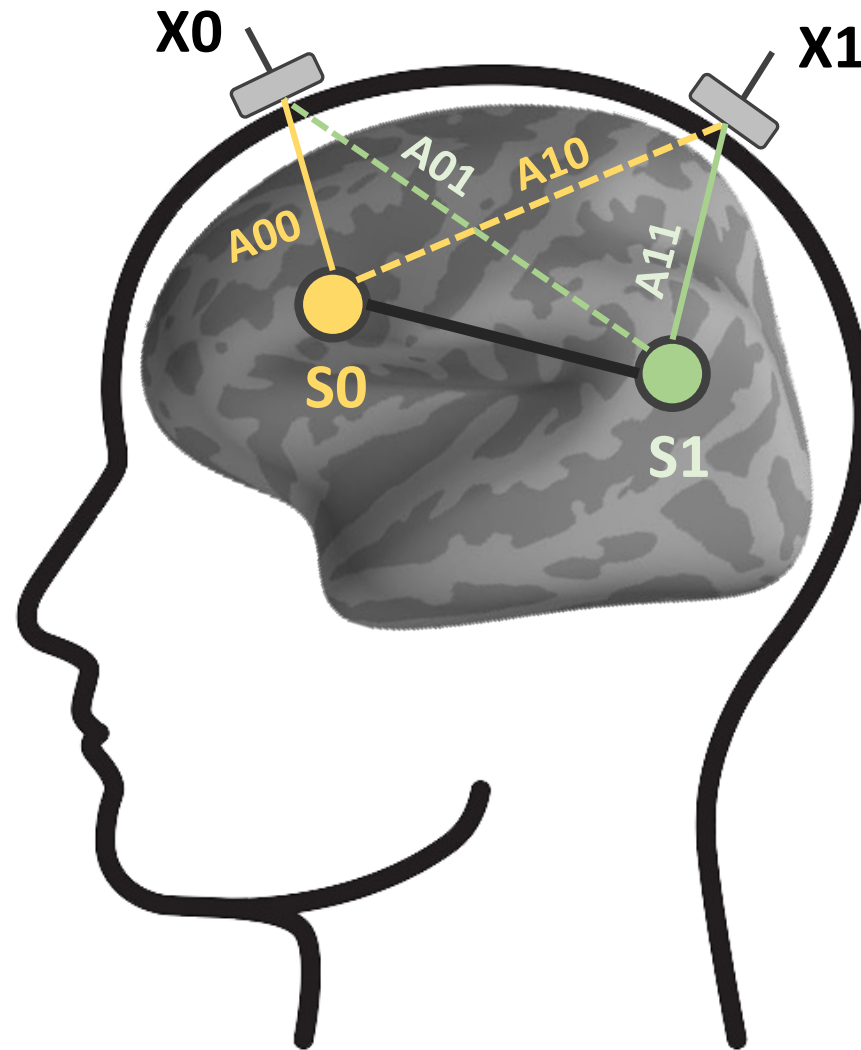
measurements

$$X_0 = A_{00} * S_0 + A_{01} * S_1$$

$$X_1 = A_{10} * S_0 + A_{11} * S_1$$

matrix notation

$$X = \text{np.dot}(A, S)$$



See, “L08_source_mixing.py”

Mixing “strength” (1/2)

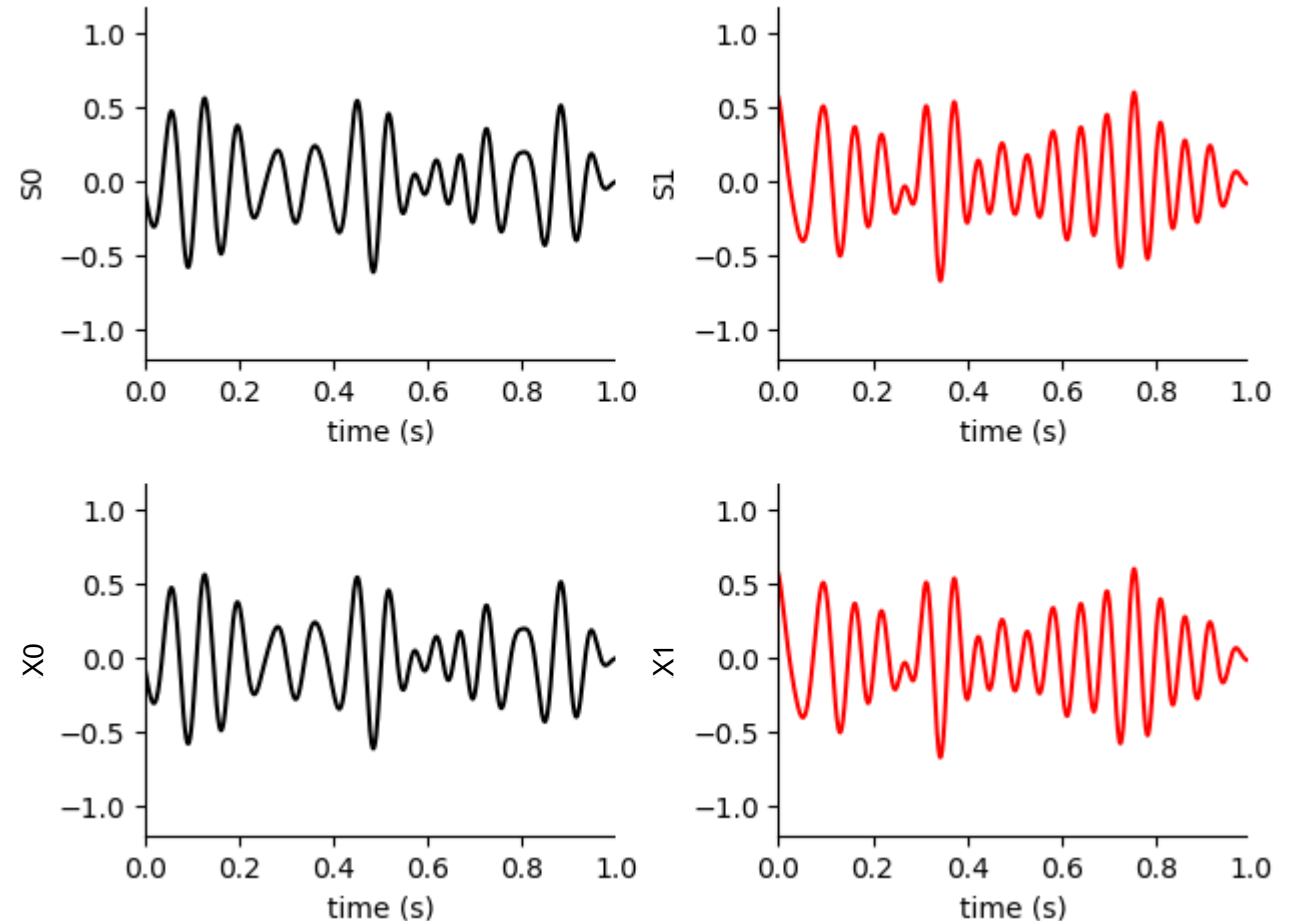
Mixing matrix (A) determines the contribution of different sources.

```
# signal
S0 = np.random.randn(1, N)
S1 = np.random.randn(1, N)
S = np.concatenate((S0, S1))

# mixing matrix
A = np.array([[1.0, 0.0], \
              [0.0, 1.0]])

# mixing
X = np.dot(A, S)
x0 = X[0, :]
x1 = X[1, :]

# what is dot?
X[0, :] = A[0,0]*X[0, :] + A[0,1]*X[1, :]
X[1, :] = A[1,0]*X[0, :] + A[1,1]*X[1, :]
```



See, “L08_source_mixing.py”

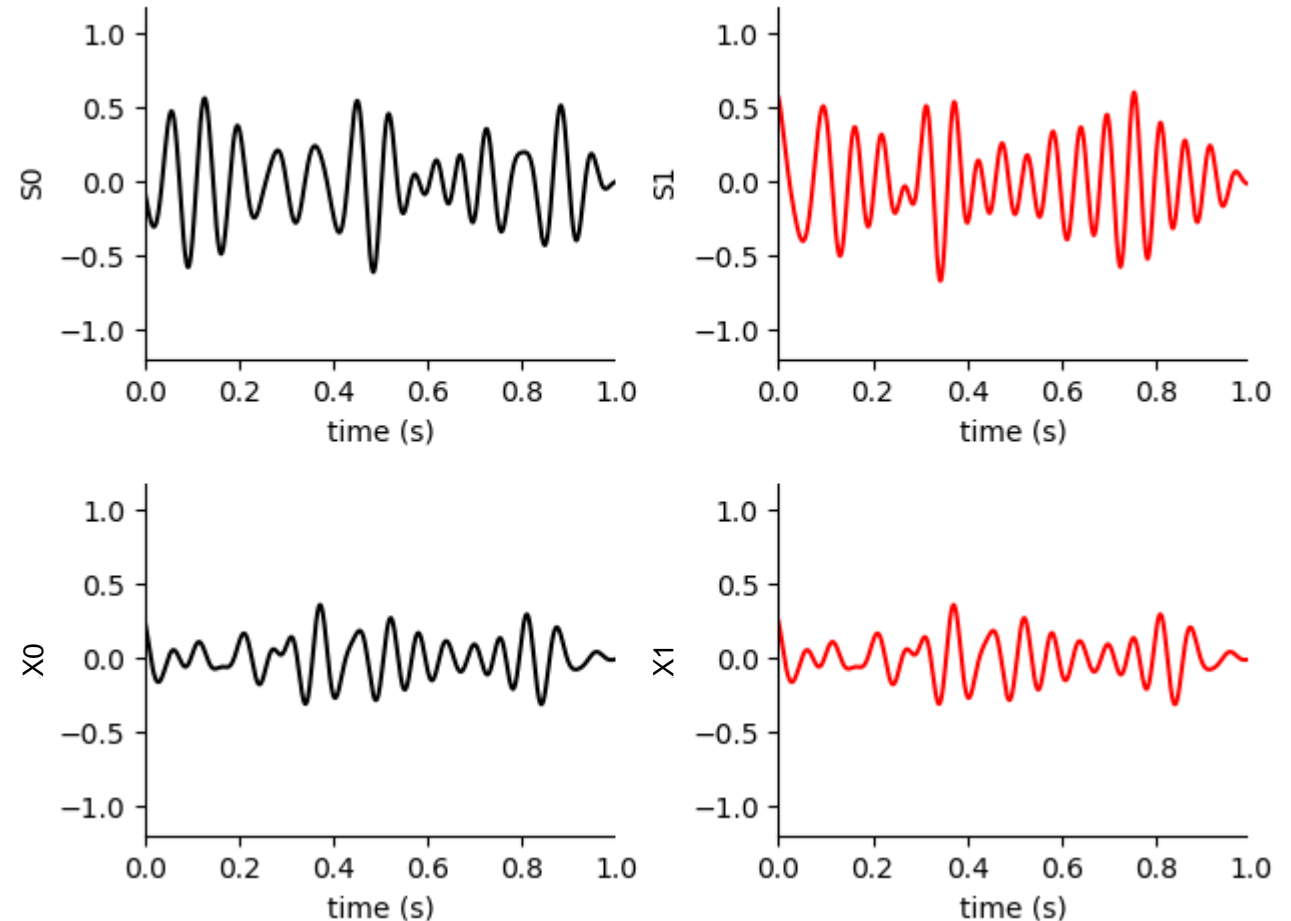
Mixing “strength” (2/2)

Mixing matrix (A) determines the contribution of different sources.

```
# signal
S0 = np.random.randn(1, N)
S1 = np.random.randn(1, N)
S = np.concatenate((S0, S1))

# mixing matrix
A = np.array([[0.5, 0.5], \
              [0.5, 0.5]])

# mixing
X = np.dot(A, S)
X0 = X[0, :]
X1 = X[1, :]
```

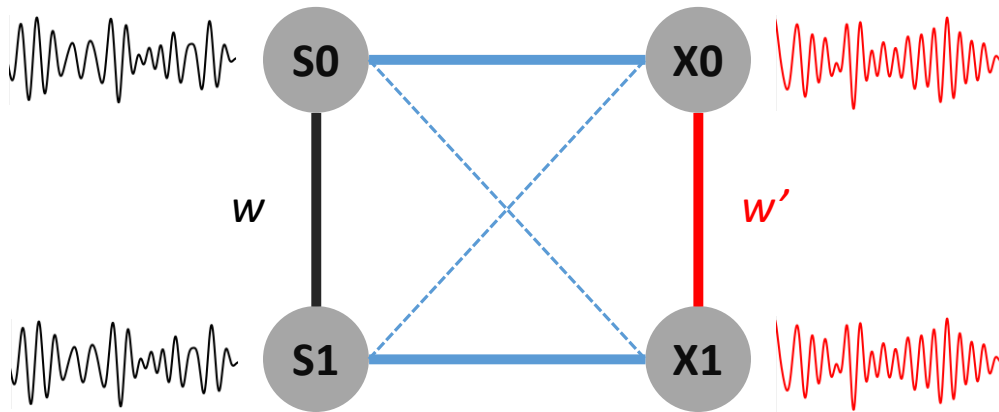


See, “L08_source_mixing.py”

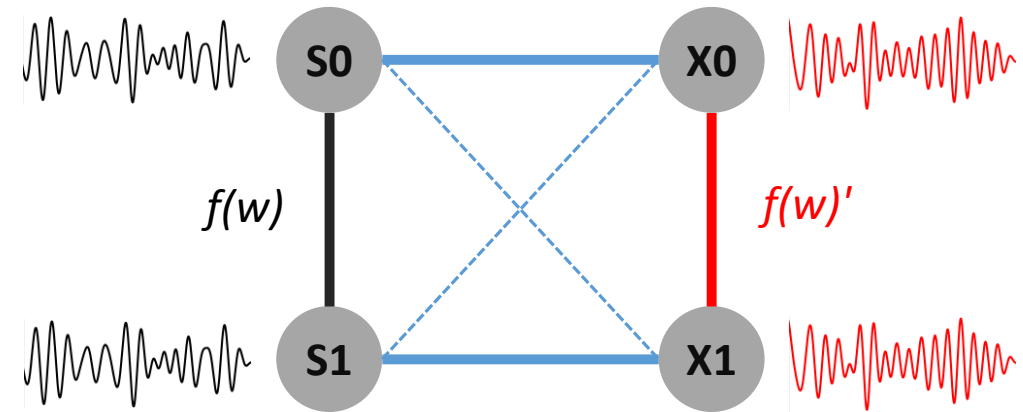
Section 2. Interactions between sources

Linear and Non-linear interactions

Linear interactions

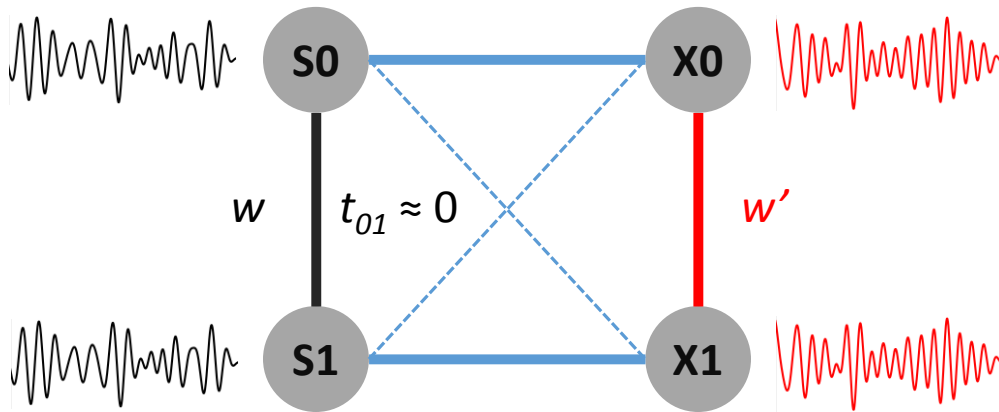


Non-linear interactions

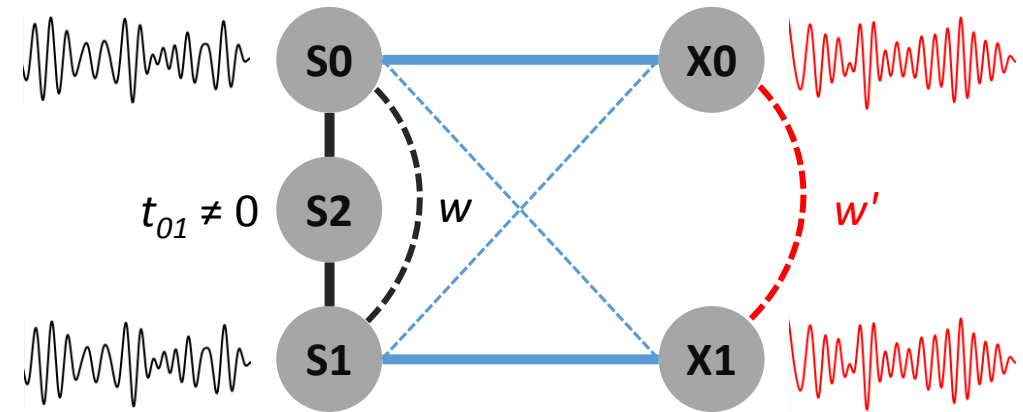


Zero-lag and Non-zero-lag interactions

Zero-lag interactions

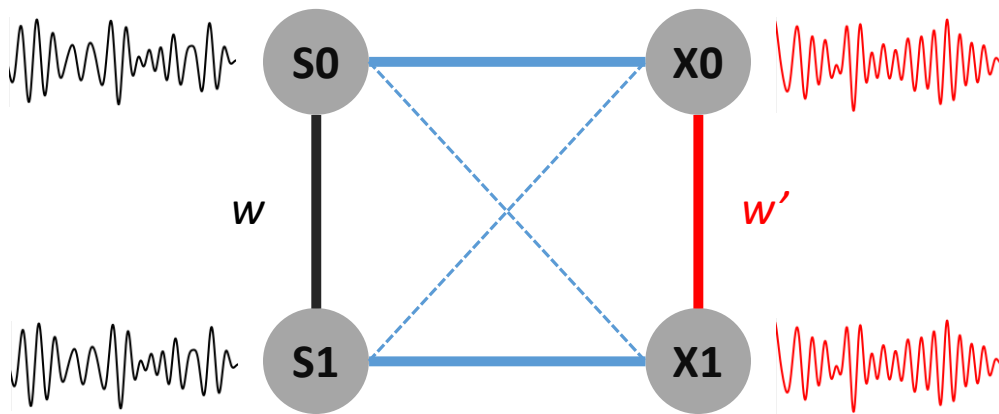


Non-zero-lag interactions

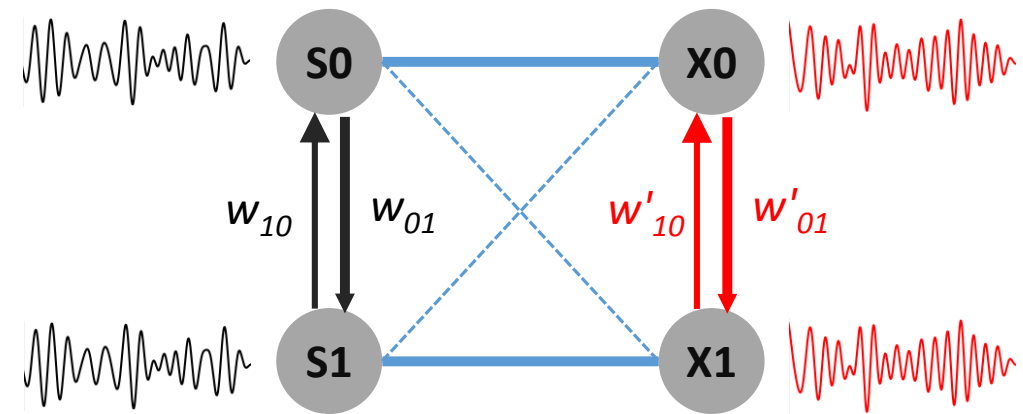


Directed/Causal and Non-directed interactions

Non-directed interactions



Directed interactions



Section 3. Covariance and Correlation

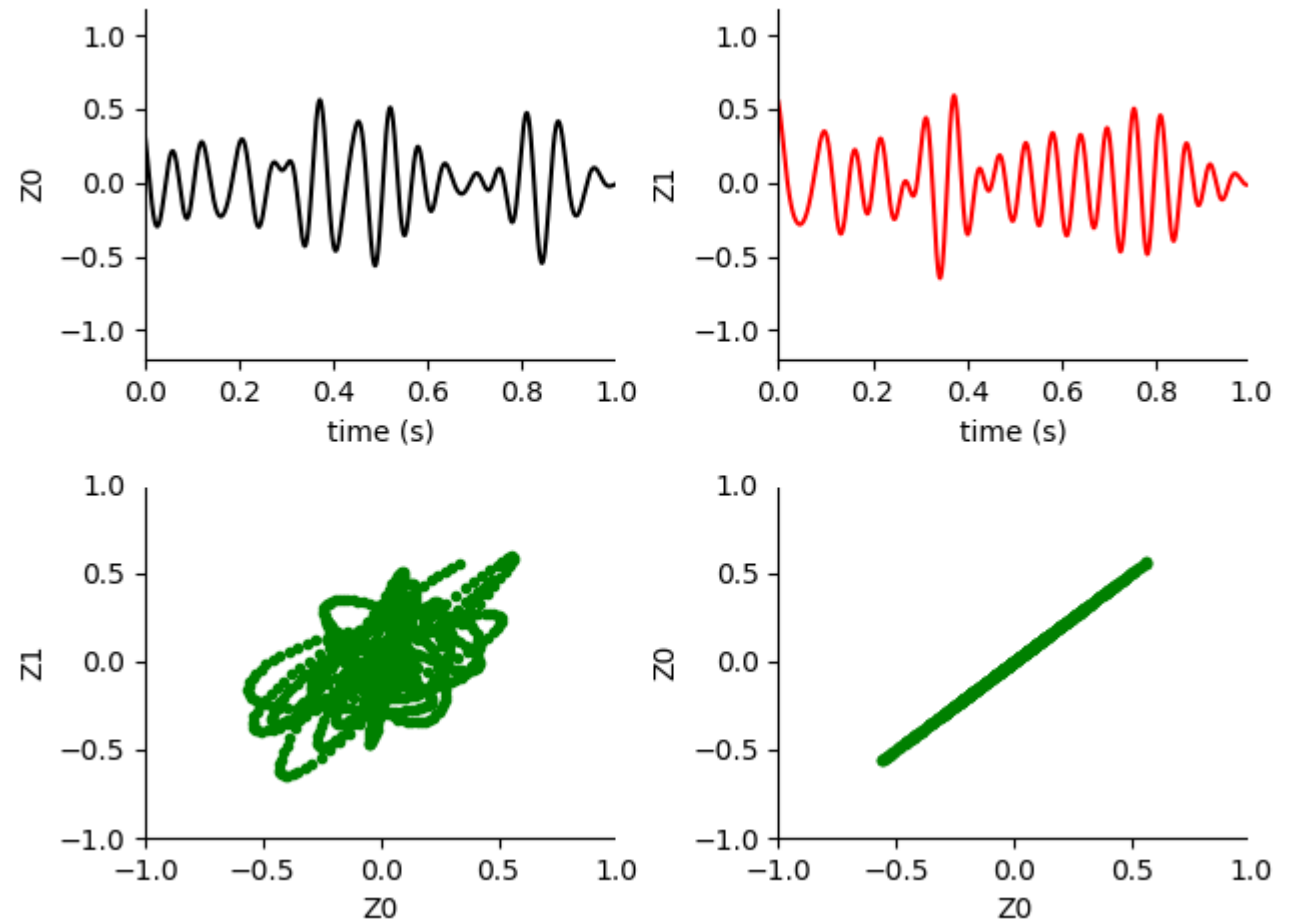
Covariance

What is covariance? It is a number that shows similarity between signals.

```
# mixing matrix
A = np.array([[1.0, 0.7], \
              [0.3, 1.0]])

# mixing
Z = np.dot(A, S)
Z0 = Z[0, :]
Z1 = Z[1, :]

# covariance
r = np.sum(((Z0 - np.mean(Z0)) *
            (Z1 - np.mean(Z1)))) / N
```



See, “L08_covariance_and_correlation.py”

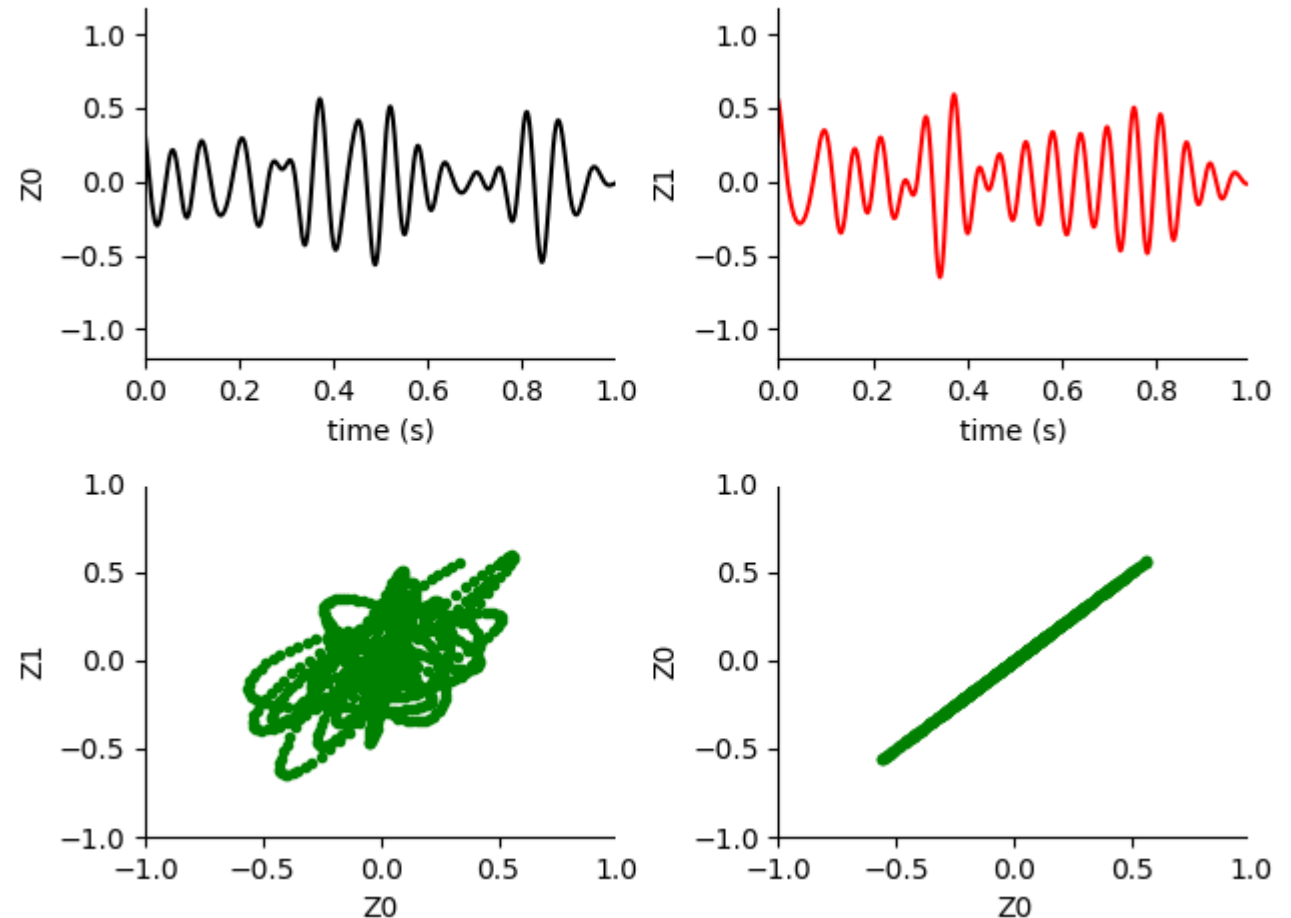
Correlation

What is correlation? Correlation is the normalized covariance.

```
# mixing matrix
A = np.array([[1.0, 0.7], \
              [0.3, 1.0]])

# mixing
Z = np.dot(A, S)
Z0 = Z[0, :]
Z1 = Z[1, :]

# correlation, range [-1, 1]
r = np.sum(((Z0 - np.mean(Z0)) *
            (Z1 - np.mean(Z1))) /
            (np.std(Z0) * np.std(Z1))) / N
```



See, “L08_covariance_and_correlation.py”

Section 4. Regression and Partial correlation

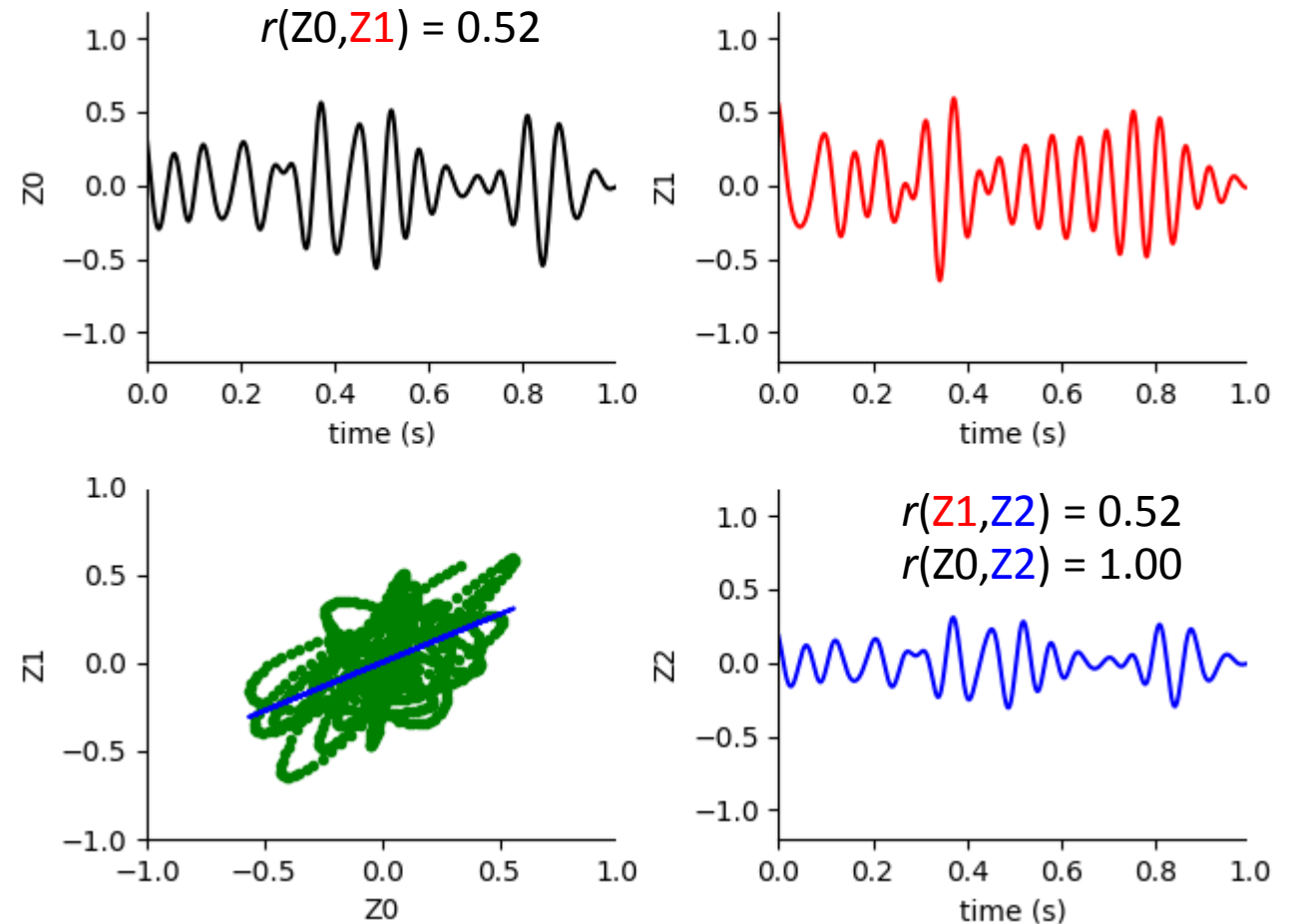
Regression

What is regression? Regression tries to represent y as a weighted version of x .

```
# mixing matrix
A = np.array([[1.0, 0.7], \
              [0.3, 1.0]])
```

```
# mixing
Z = np.dot(A, S)
Z0 = Z[0, :]
Z1 = Z[1, :]
```

```
# regression
p = np.polyfit(Z0, Z1, 1)
Z2 = p[0] * Z0 + p[1]
```



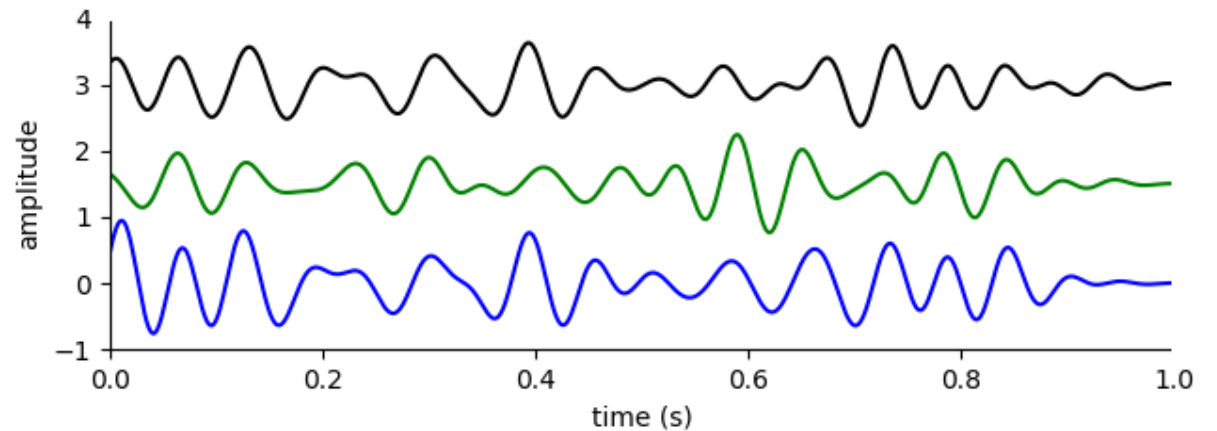
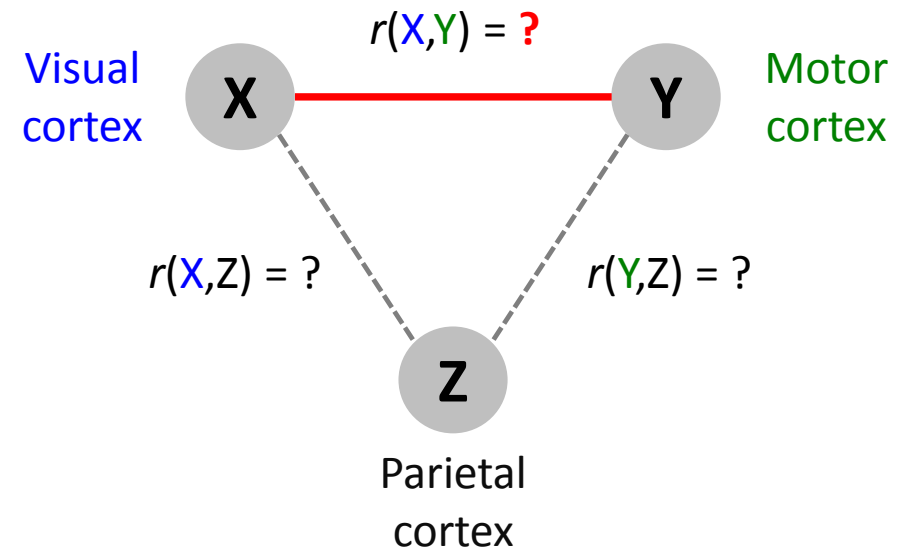
See, “L08_regression.py”

Partial correlation (1/3)

What is partial correlation? It is correlation between X and Y with regressed out Z.

```
# mixing matrix
A = np.array([[1.0, 0.7, 0.3], \
              [0.3, 1.0, 0.1], \
              [0.5, 0.5, 1.0]])

# mixing
U = np.dot(A, S)
X = U[0, :] # visual cortex
Y = U[1, :] # motor cortex
Z = U[2, :] # parietal cortex
```



See, “L08_partial_correlation.py”

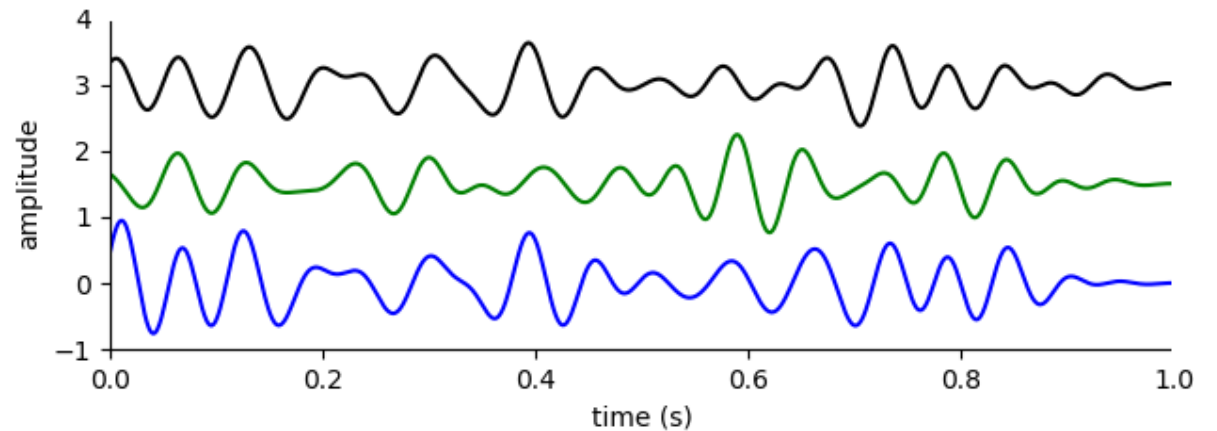
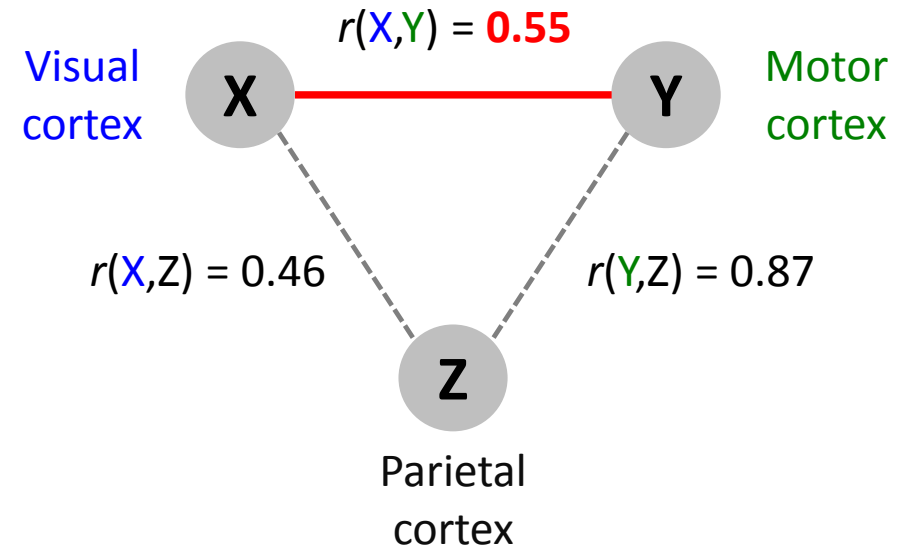
Partial correlation (2/3)

We can compute correlation coefficients between all pairs.

```
# mixing matrix
A = np.array([[1.0, 0.7, 0.3], \
              [0.3, 1.0, 0.1], \
              [0.5, 0.5, 1.0]])

# mixing
U = np.dot(A, S)
X = U[0, :] # visual cortex
Y = U[1, :] # motor cortex
Z = U[2, :] # parietal cortex

# correlation
rXY = np.corrcoef(X, Y)[0, 1]
rXZ = np.corrcoef(X, Z)[0, 1]
rYZ = np.corrcoef(Y, Z)[0, 1]
```



See, “L08_partial_correlation.py”

Partial correlation (3/3)

Can we assess correlation between X and Y without influence of Z?

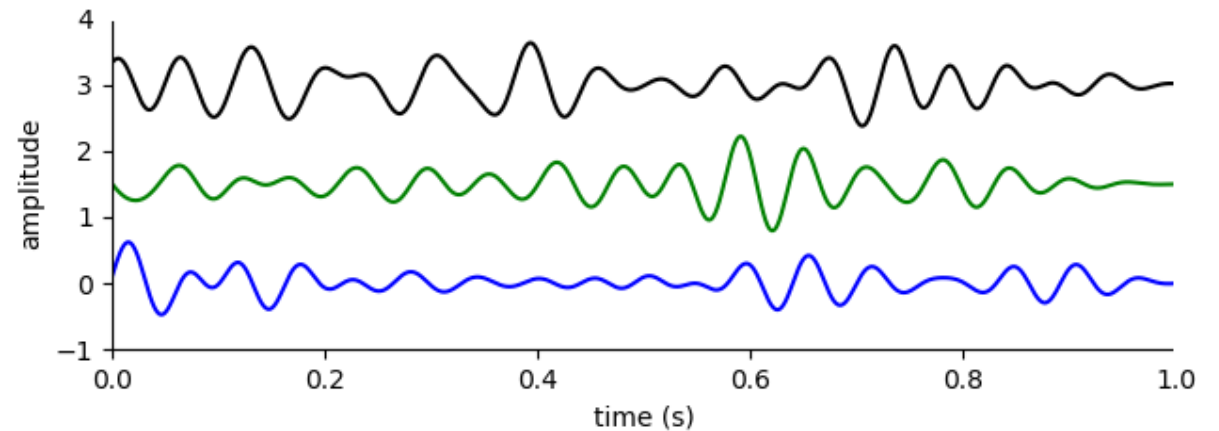
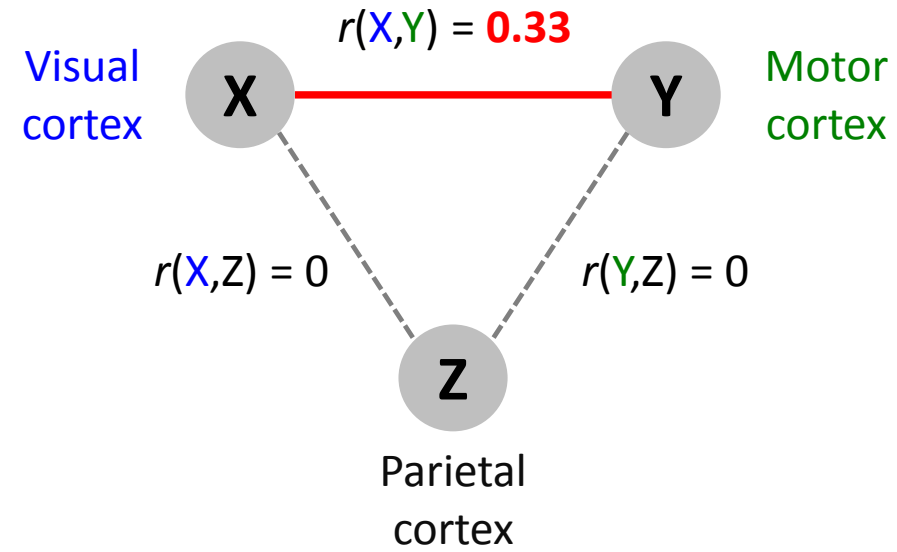
```
# fitting and regressing out
```

```
p = np.polyfit(Z, X, 1)
fX = X - (p[0] * Z + p[1])
```

```
p = np.polyfit(Z, Y, 1)
fY = Y - (p[0] * Z + p[1])
```

```
# correlation
```

```
rXY = np.corrcoef(fX, fY)[0, 1]
rXZ = np.corrcoef(fX, Z)[0, 1]
rYZ = np.corrcoef(fY, Z)[0, 1]
```

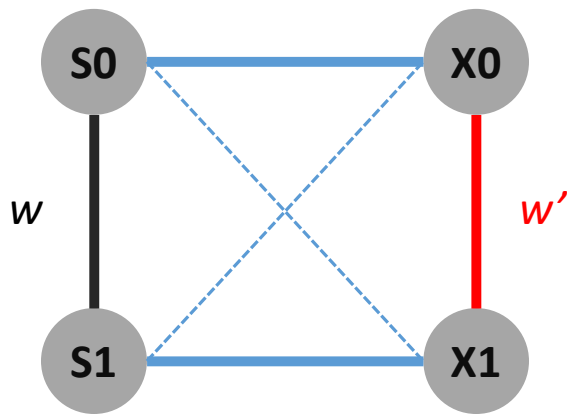


See, “L08_partial_correlation.py”

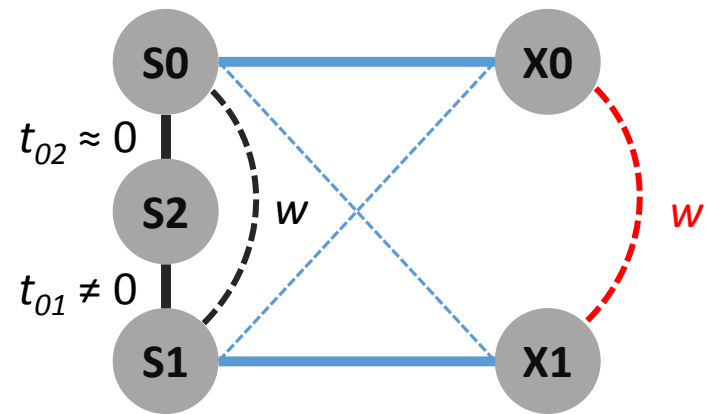
Section 5. Zero-lag interactions

Interactions

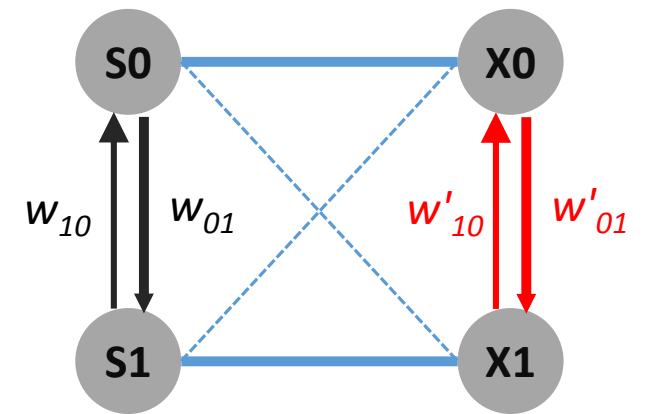
Zero-lag interactions



Non-zero-lag interactions



Causal interactions



Signal parameters

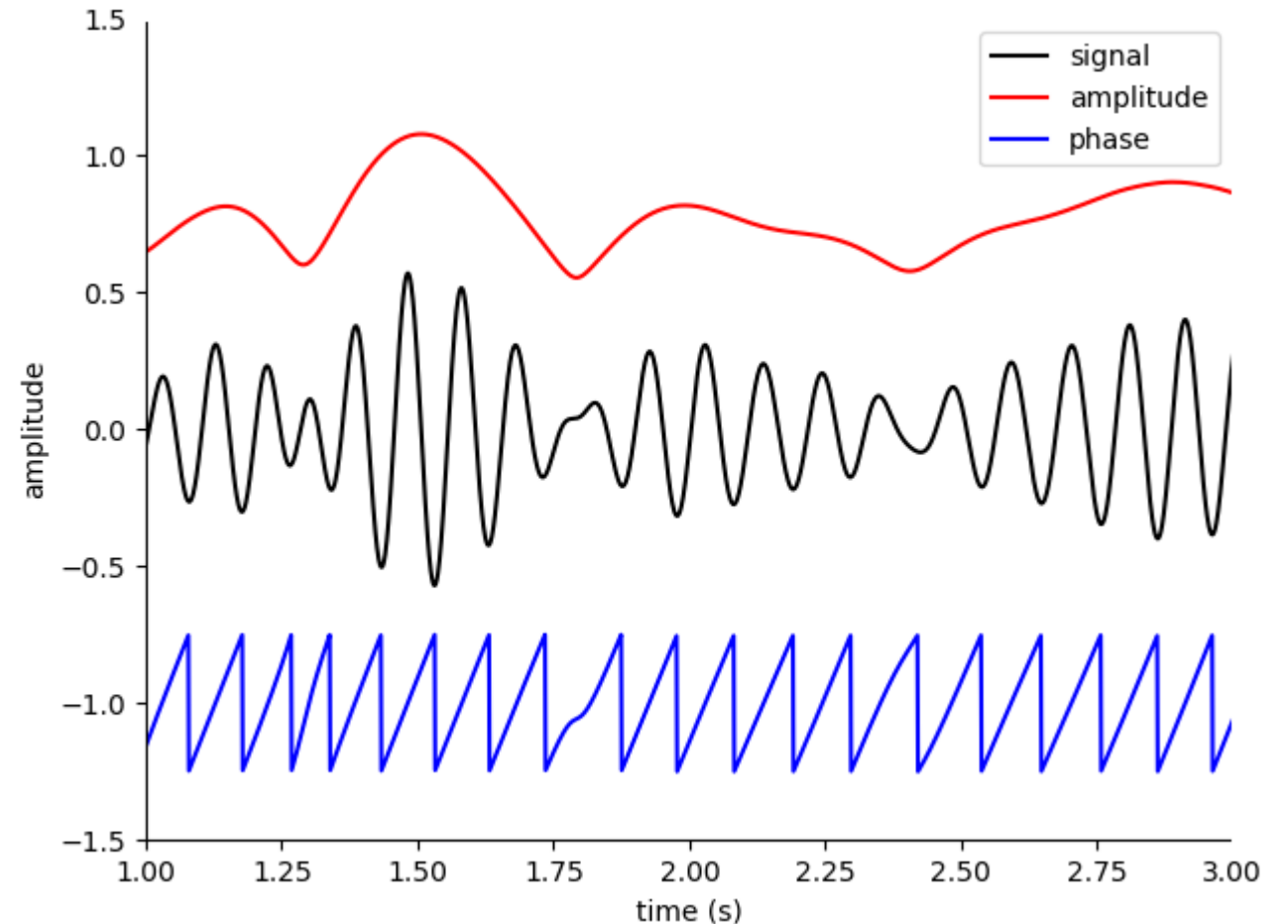
Signal can be represented by its amplitude and phase.

```
# signal
S = np.random.randn(1, N)

# filtering
[b, a] = signal.butter(4,
                      [8.0 / (fs/2), 12.0 / (fs/2)],
                      'bandpass')
x = signal.filtfilt(b, a, S)

# amplitude
AX = np.abs(signal.hilbert(x))

# phase
PX = np.angle(signal.hilbert(x))
```



See, “L08_zero_lag_interactions_signal.py”

I. Amplitude-amplitude correlations

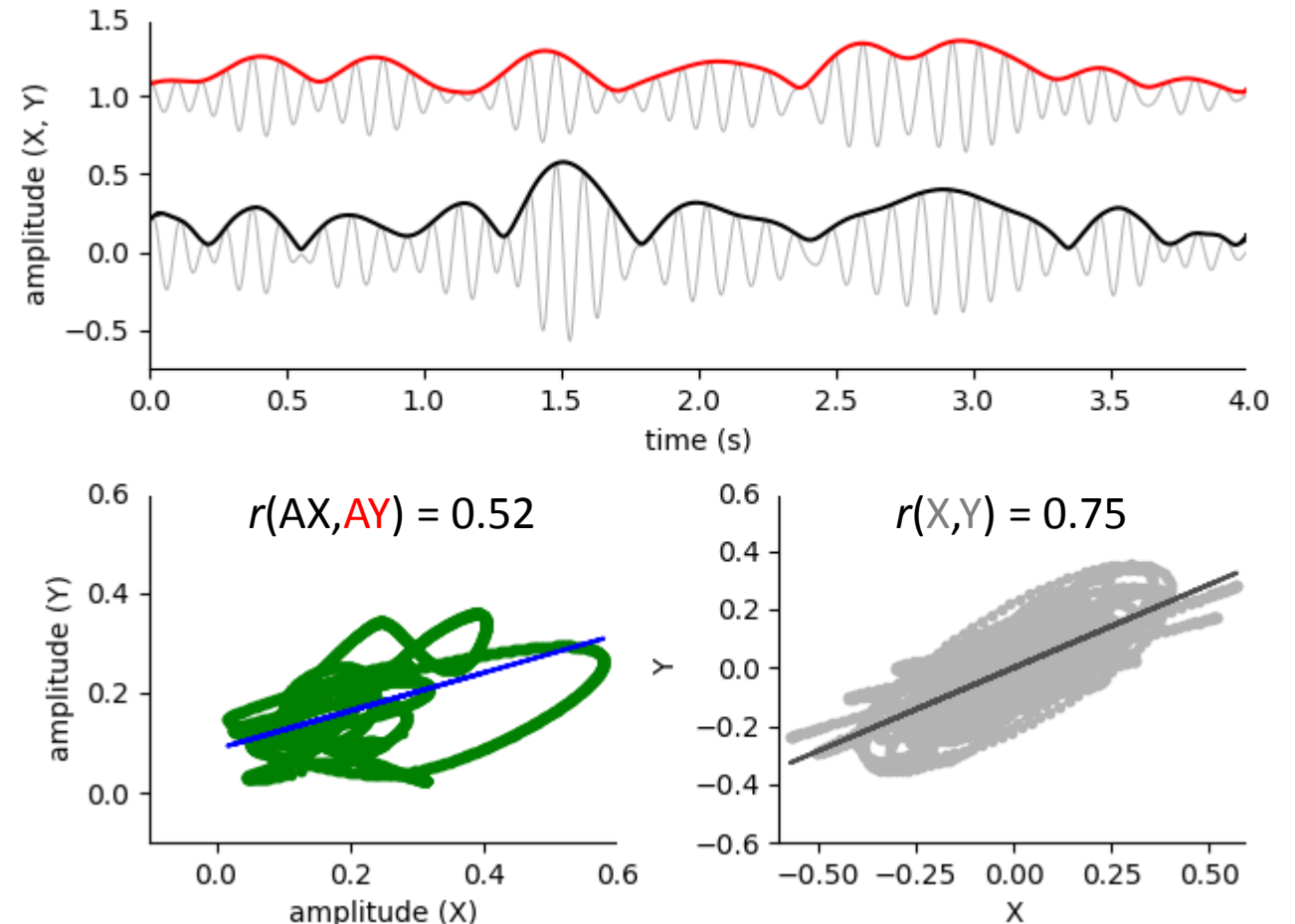
Correlation between amplitudes of two signals.

```
# amplitude
AX = np.abs(signal.hilbert(X))
AY = np.abs(signal.hilbert(Y))

# linear fit (amplitudes)
p = np.polyfit(AX, AY, 1)
AU = p[0] * AX + p[1]

# linear fit (signal)
p = np.polyfit(X, Y, 1)
U = p[0] * X + p[1]

# correlation
rAA = np.corrcoef(AX, AY)[0, 1]
rXY = np.corrcoef(X, Y)[0, 1]
```



See, “L08_zero_lag_interactions_amplitude.py”

II. Phase-phase coupling (1/2)

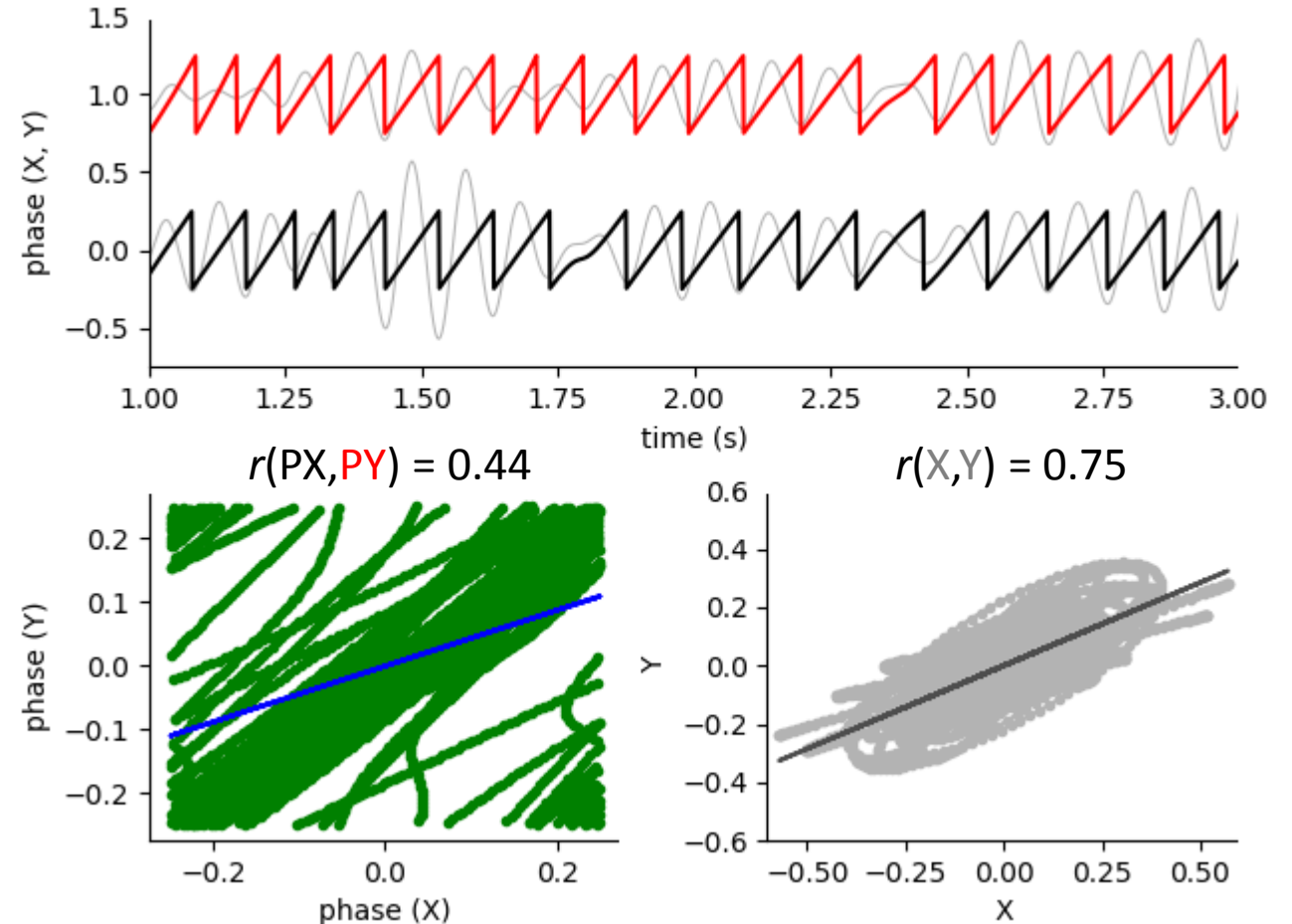
Coupling between phases of two signals.

```
# phase
PX = np.angle(signal.hilbert(X))
PY = np.angle(signal.hilbert(Y))

# linear fit (phases)
p = np.polyfit(PX, PY, 1)
PU = p[0] * PX + p[1]

# linear fit (signal)
p = np.polyfit(X, Y, 1)
U = p[0] * X + p[1]

# correlation
rPP = np.corrcoef(PX, PY)[0, 1]
rXY = np.corrcoef(X, Y)[0, 1]
```



See, “L08_zero_lag_interactions_phase.py”

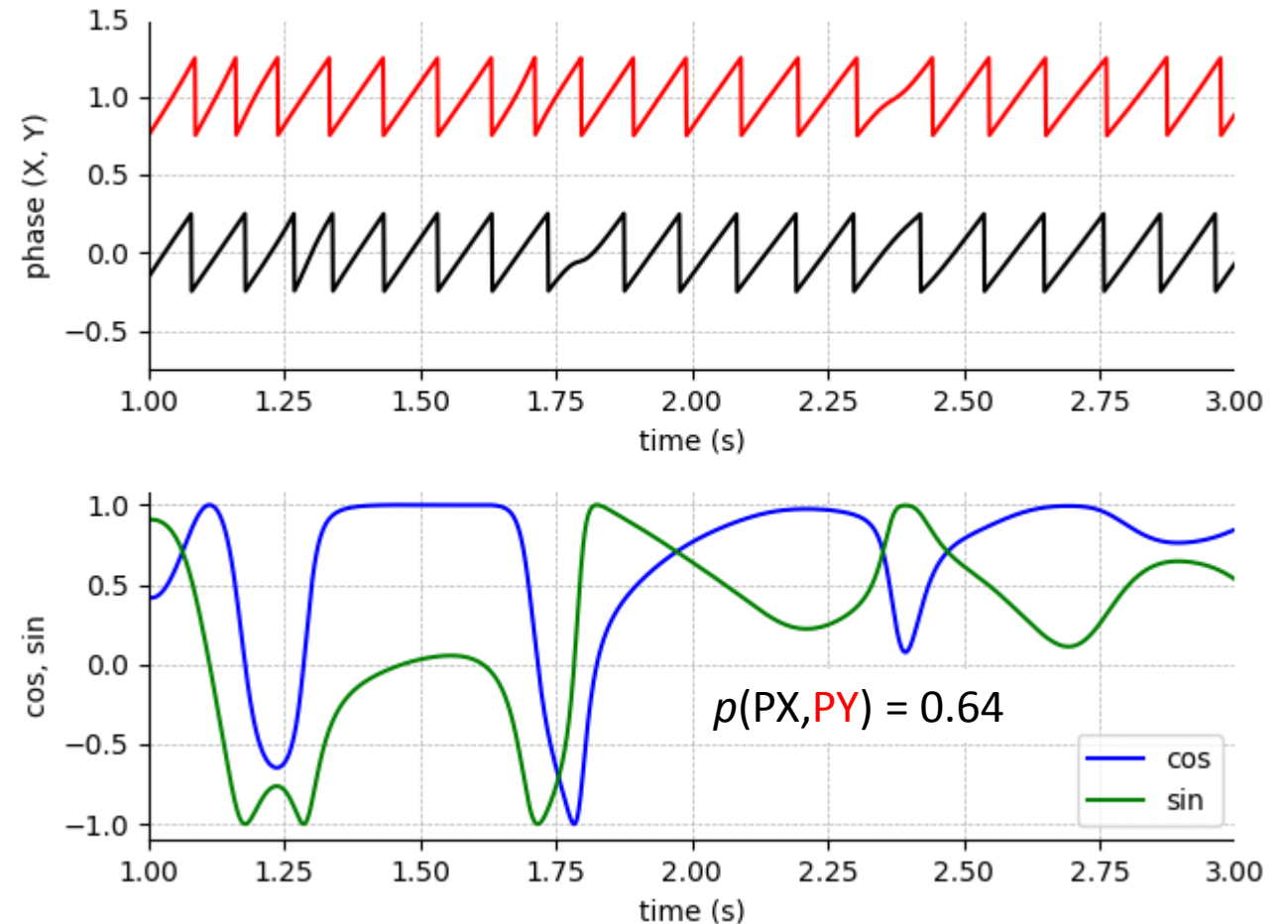
II. Phase-phase coupling (2/2)

Coupling between phases of two signals can be assessed via phase-locking value.

```
# phase
PX = np.angle(signal.hilbert(X))
PY = np.angle(signal.hilbert(Y))

# phase-locking value
p = np.abs(np.sum(np.exp(1j * (PX - PY))) / N)

# phase-locking value via sine
p = np.abs(np.sum(np.cos(PX - PY) +
                  1j * np.sin(PX - PY)) / N)
```



See, “L08_zero_lag_interactions_phase.py”

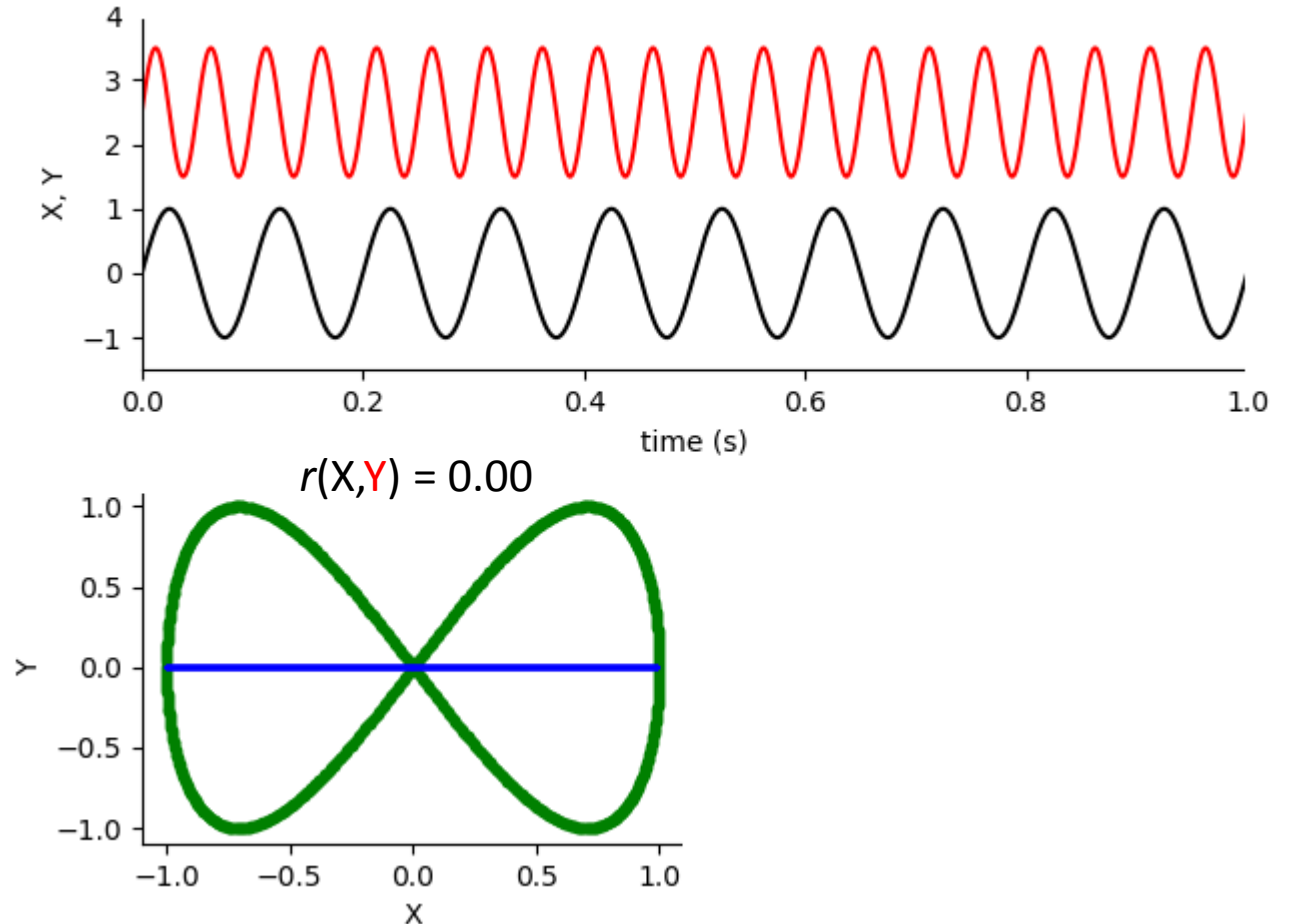
III. Cross-frequency phase-phase coupling (1/3)

How could we quantify interactions between two signals at different frequencies?

```
# signal
ratio = 2
f0 = 10
X = np.sin(2 * np.pi * f0 * t)
Y = np.sin(2 * np.pi * (f0 * ratio) * t)

# linear relationship
p = np.polyfit(X, Y, 1)
U = p[0] * X + p[1]

# correlation
r = np.corrcoef(X, Y)[0, 1]
```



See, “L08_zero_lag_interactions_cf.py”

III. Cross-frequency phase-phase coupling (2/3)

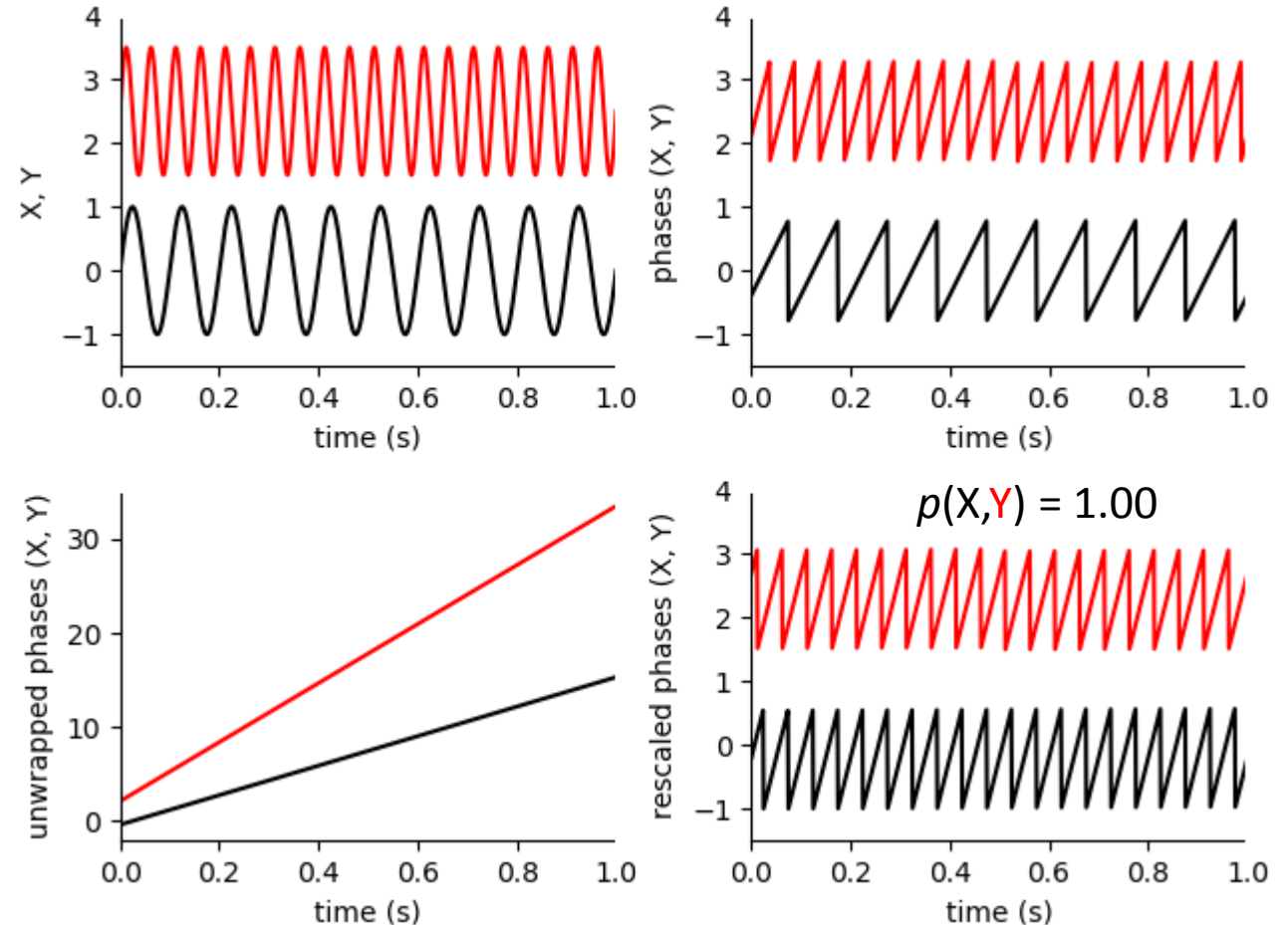
Rescaling the phase time series could solve this problem.

```
# get phase
LF_ph = np.angle(signal.hilbert(X))
HF_ph = np.angle(signal.hilbert(Y))

# unwrap phases
LF_unwrap_ph = np.unwrap(LF_ph)
HF_unwrap_ph = np.unwrap(HF_ph)

# rescale phase
LF_res_ph = (LF_unwrap_ph % (2 * np.pi /
    ratio)) * ratio
HF_res_ph = (HF_unwrap_ph % (2 * np.pi))

# compute PLV
p = np.abs(np.sum(np.exp(1j *
    (LF_res_phase - HF_res_phase)))) / N
```



See, “L08_zero_lag_interactions_cf.py”

III. Cross-frequency phase-phase coupling (3/3)

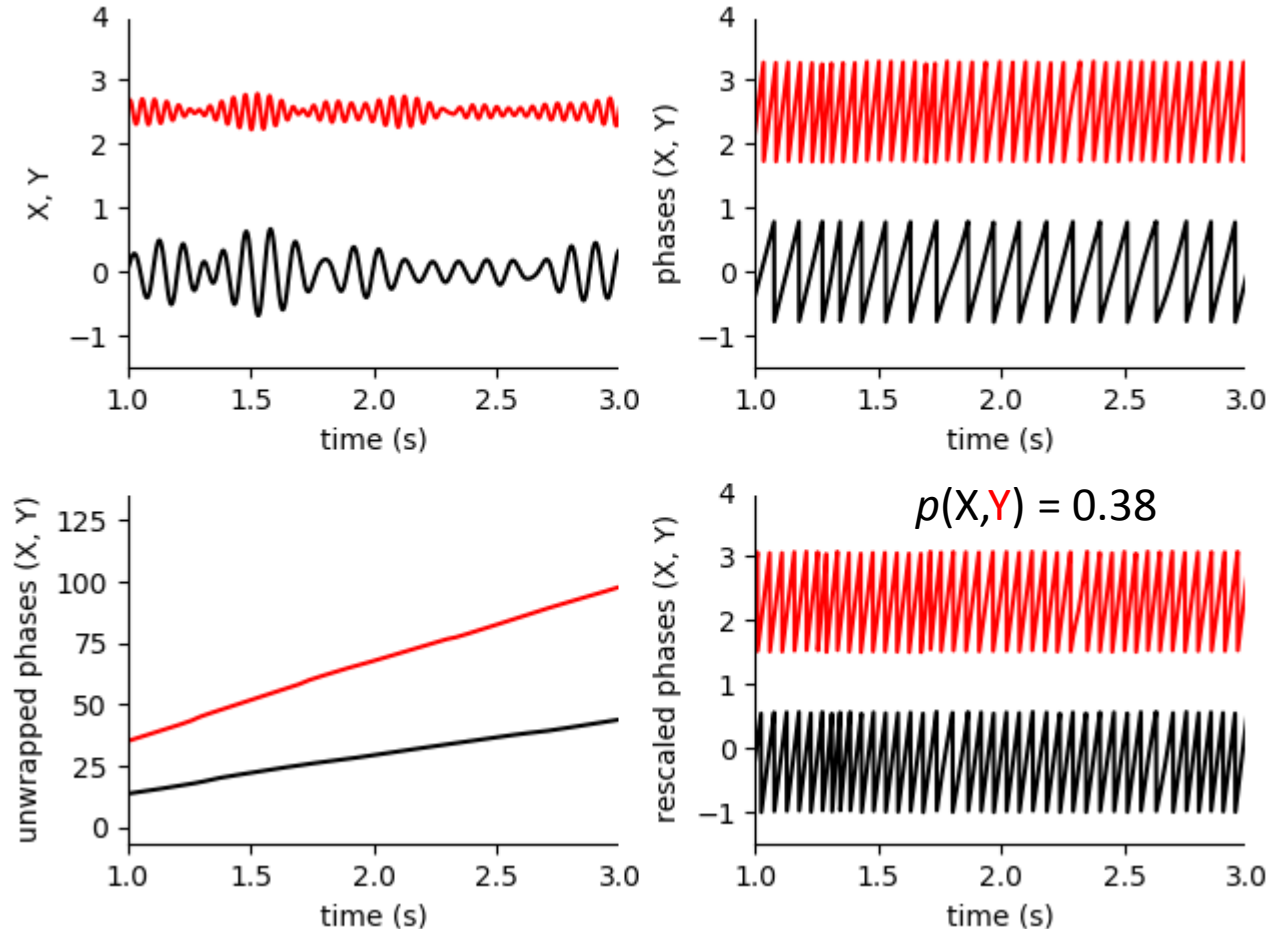
How do it work for realistic data?

```
# get phase
LF_ph = np.angle(signal.hilbert(X))
HF_ph = np.angle(signal.hilbert(Y))

# unwrap phases
LF_unwrap_ph = np.unwrap(LF_ph)
HF_unwrap_ph = np.unwrap(HF_ph)

# rescale phase
LF_res_ph = (LF_unwrap_ph % (2 * np.pi /
    ratio)) * ratio
HF_res_ph = (HF_unwrap_ph % (2 * np.pi))

# compute phase-locking value
p = np.abs(np.sum(np.exp(1j *
    (LF_res_phase - HF_res_phase)))) / N
```



See, “L08_zero_lag_interactions_cf.py”

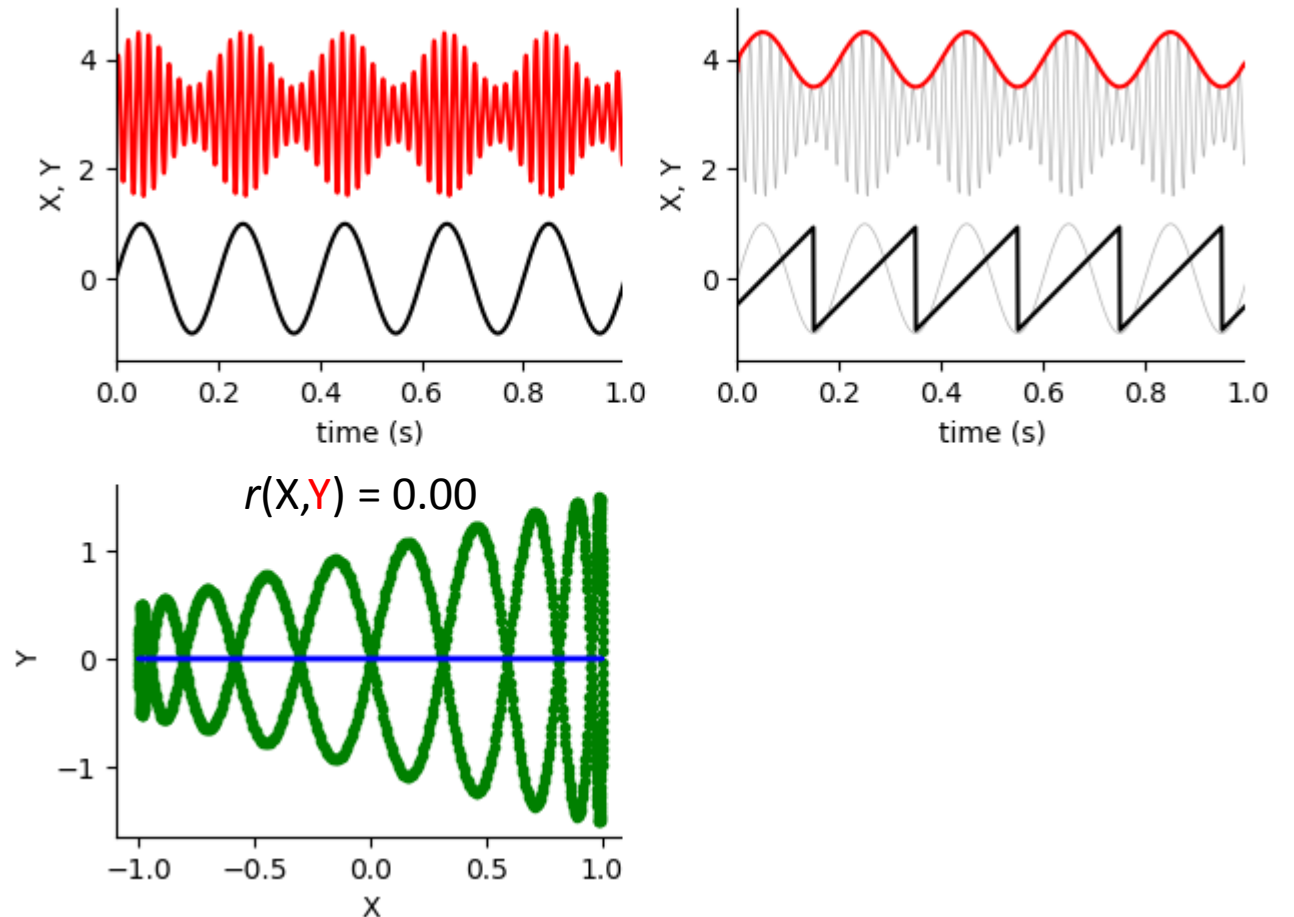
IV. Phase-amplitude modulation (1/2)

There are cases when phase of slow signal modulates amplitude of fast signal.

```
# signal
X = np.sin(2 * np.pi * 5 * t)
Y = np.sin(2 * np.pi * 50 * t) *
(1 + 0.5 * np.sin(2 * np.pi * 5 * t)) # AM

# linear relationship
p = np.polyfit(X, Y, 1)
U = p[0] * X + p[1]

# correlation
r = np.corrcoef(X, Y)[0, 1]
```



See, “L08_zero_lag_interactions_pac.py”

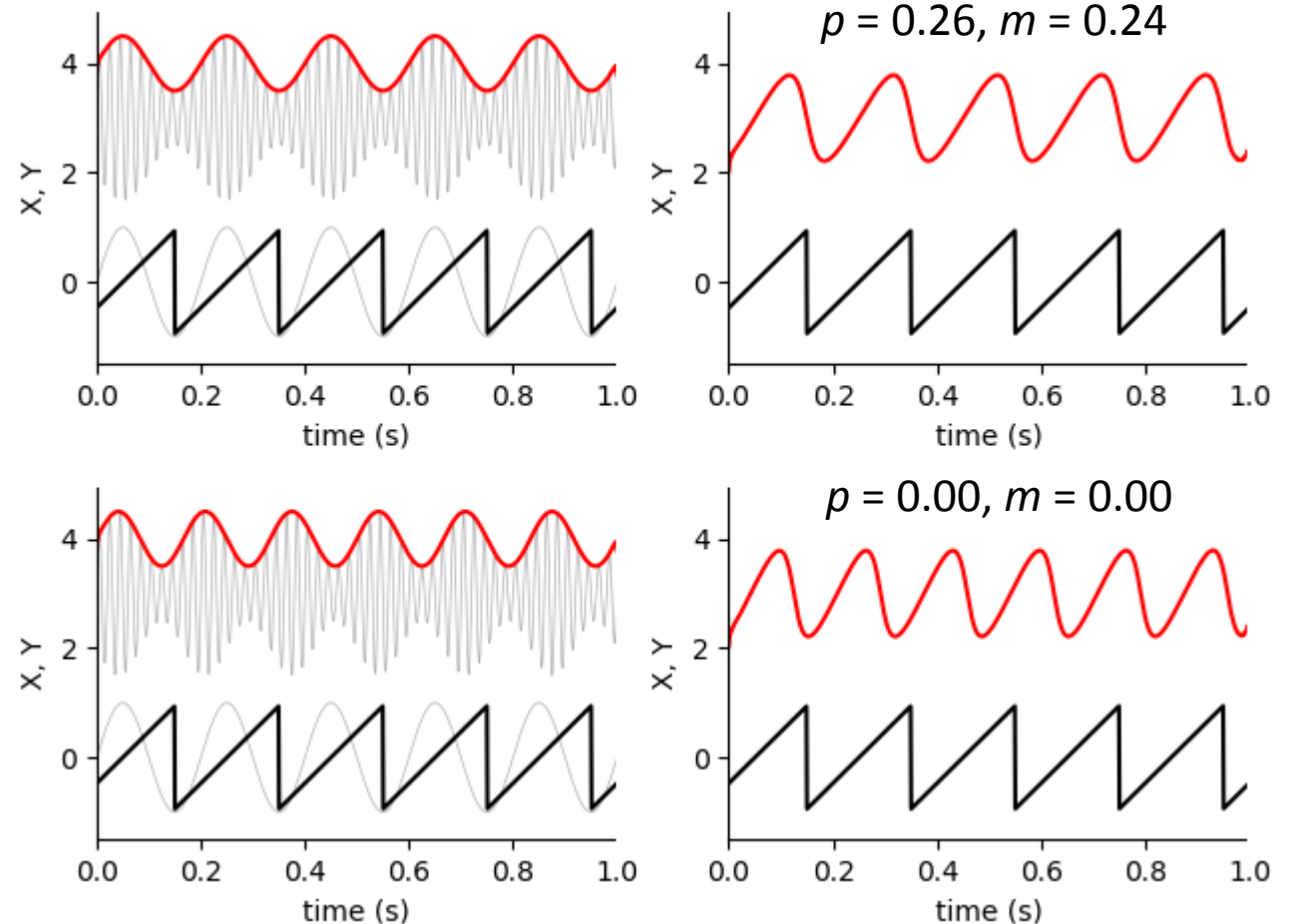
IV. Phase-amplitude modulation (2/2)

How could we detect phase-amplitude modulation?

```
# phase of slow and amplitude of fast
PX = np.angle(signal.hilbert(X))
AY = np.abs(signal.hilbert(Y))

# * approach 1: intuitive
PA = np.angle(signal.hilbert(AY))
p = np.abs(np.sum(np.exp(1j * (PX - PA)))) / N

# * approach 2: modulation index
Z = AY * np.exp(1j * PX)
m = np.abs(np.mean(Z)) / np.sqrt(np.mean(AY**2))
```

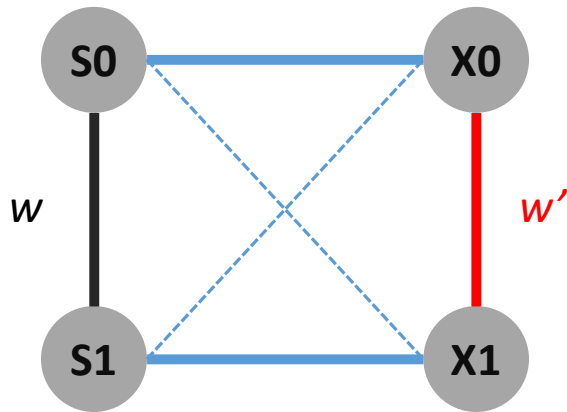


See, “L08_zero_lag_interactions_pac.py”

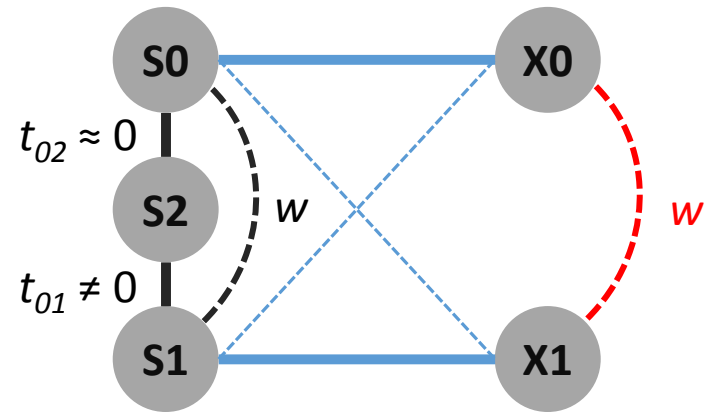
Section 6. Non-zero-lag interactions

Interactions

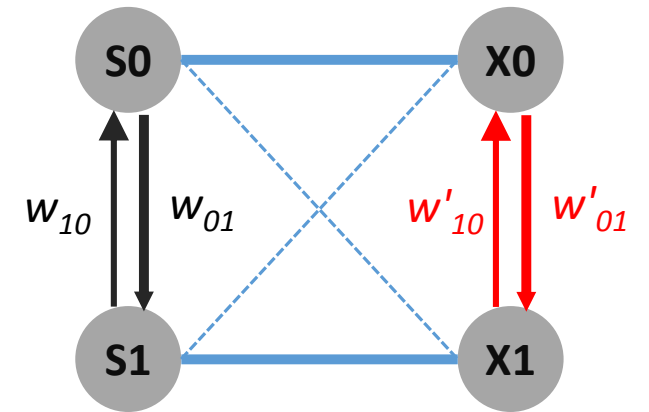
Zero-lag interactions



Non-zero-lag interactions



Causal interactions



I. Amplitude-amplitude correlations (1/2)

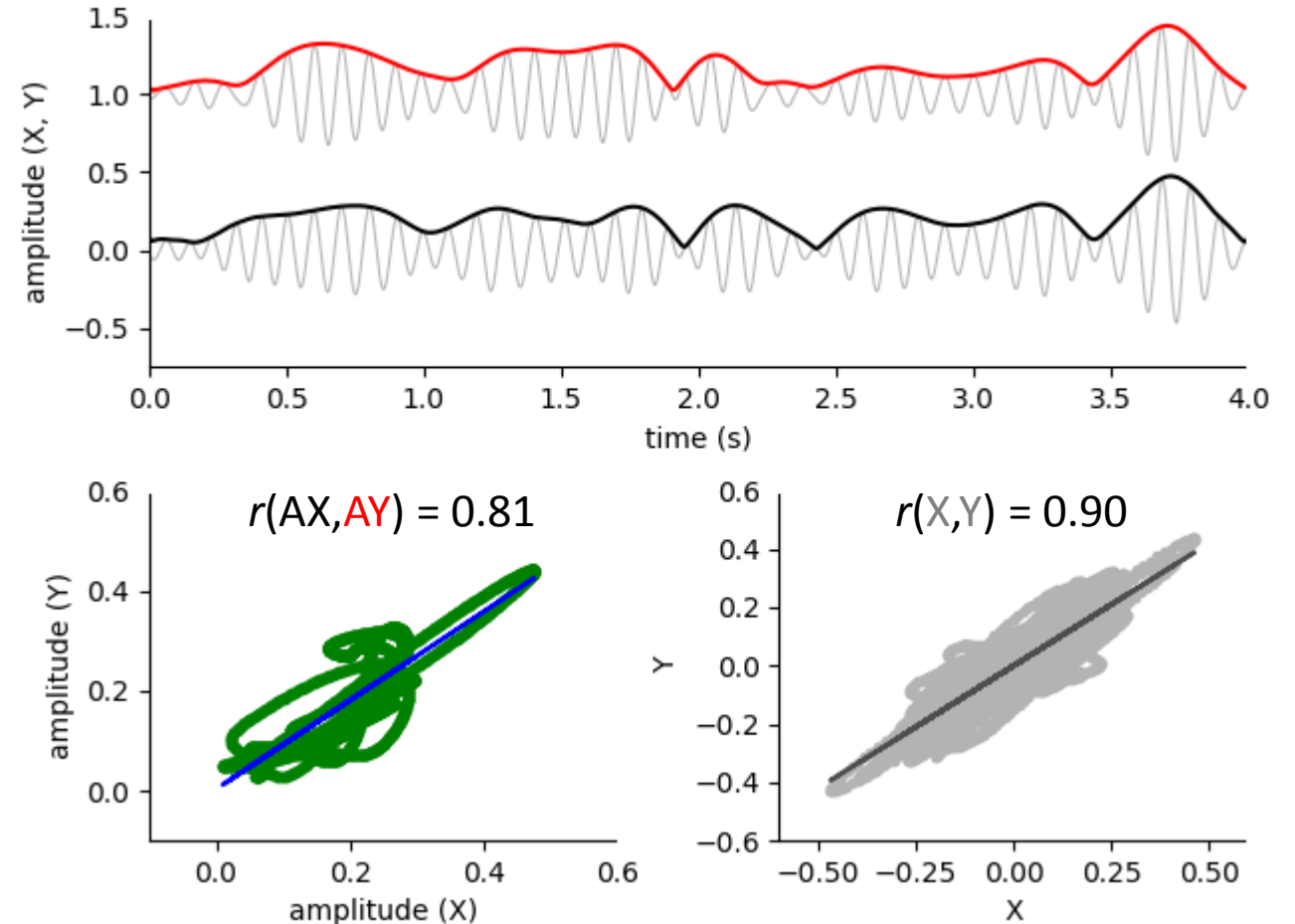
Correlation between amplitudes of two signals via third source.

```
# mixing matrix
A = np.array([[1.0, 0.0, 2.0], \
              [0.0, 1.0, 2.0], \
              [0.0, 0.0, 1.0]])

# amplitude
AX = np.abs(signal.hilbert(X))
AY = np.abs(signal.hilbert(Y))

# linear fit
p = np.polyfit(AX, AY, 1)
fAY = p[0] * AX + p[1]
p = np.polyfit(X, Y, 1)
fY = p[0] * X + p[1]

# correlation
rAA = np.corrcoef(AX, AY)[0, 1]
rXY = np.corrcoef(X, Y)[0, 1]
```



See, “L08_non_zero_lag_interactions_amplitude.py”

I. Amplitude-amplitude correlations (2/2)

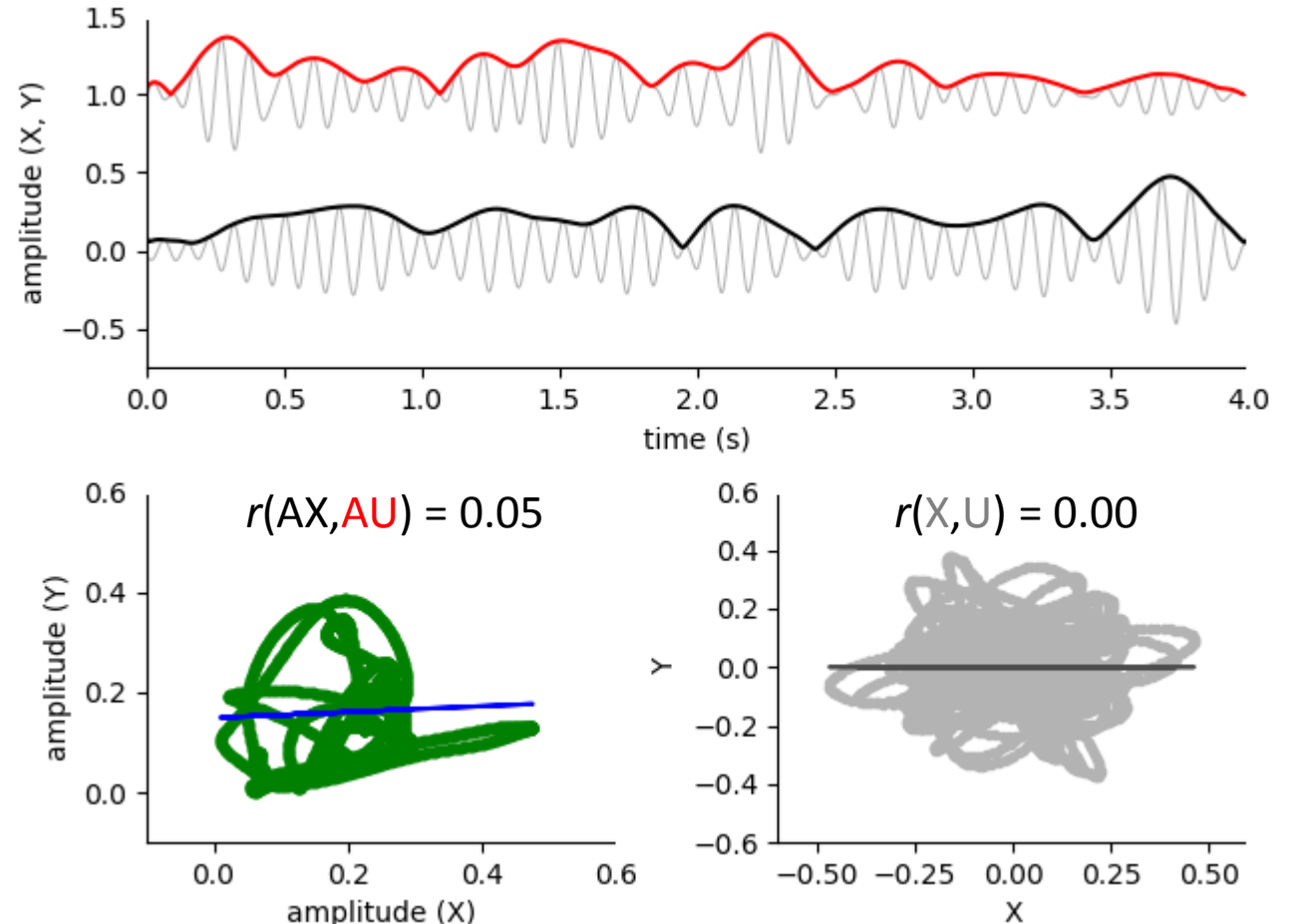
Correlation between amplitudes of two signals with regressed out a common source.

```
# regression
b = np.sum((X / np.sum(X ** 2)) * Y)
U = Y - X * b
```

```
# amplitude
AX = np.abs(signal.hilbert(X))
AU = np.abs(signal.hilbert(U))
```

```
# linear fit
p = np.polyfit(AX, AU, 1)
AU = p[0] * AX + p[1]
p = np.polyfit(X, U, 1)
fU = p[0] * X + p[1]
```

```
# correlation
rAA = np.corrcoef(AX, AU)[0, 1]
rXU = np.corrcoef(X, U)[0, 1]
```

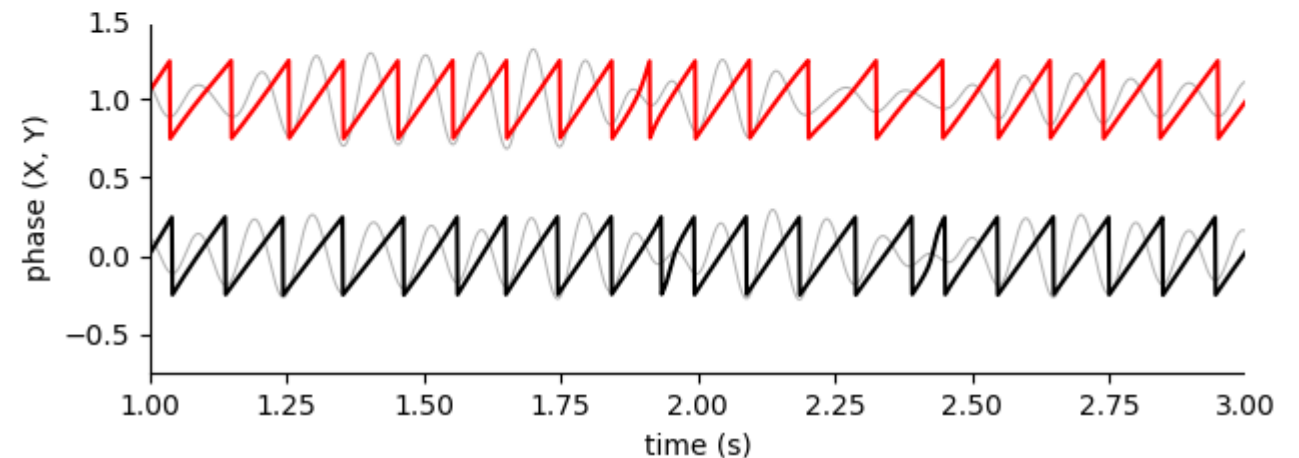
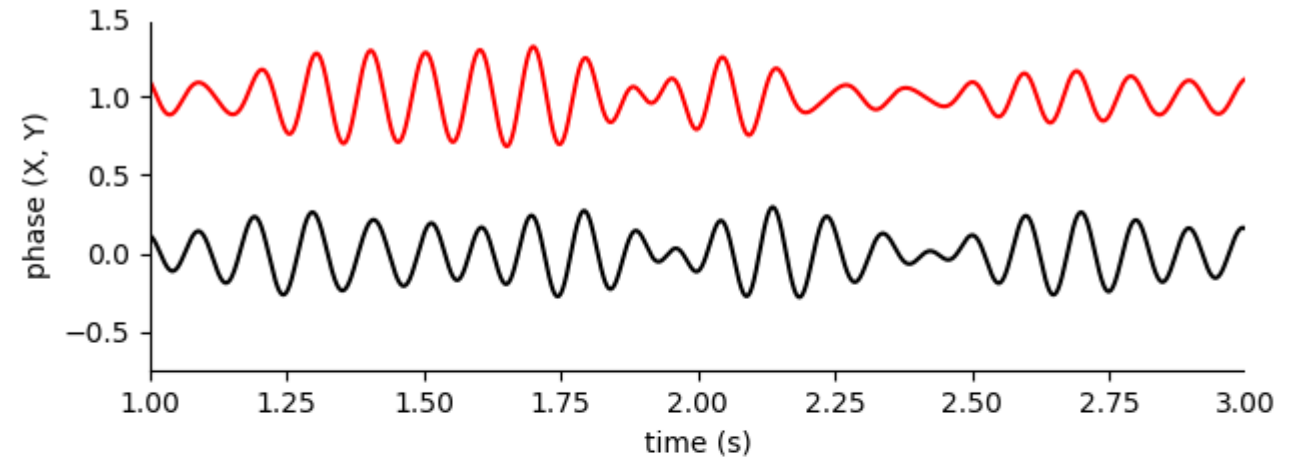


See, “L08_non_zero_lag_interactions_amplitude.py”

II. Phase-phase coupling (1/2)

Coupling between phases of two signals can be assessed via phase-locking value.

```
# phase  
PX = np.angle(signal.hilbert(X))  
PY = np.angle(signal.hilbert(Y))
```



See, “L08_non_zero_lag_interactions_phase.py”

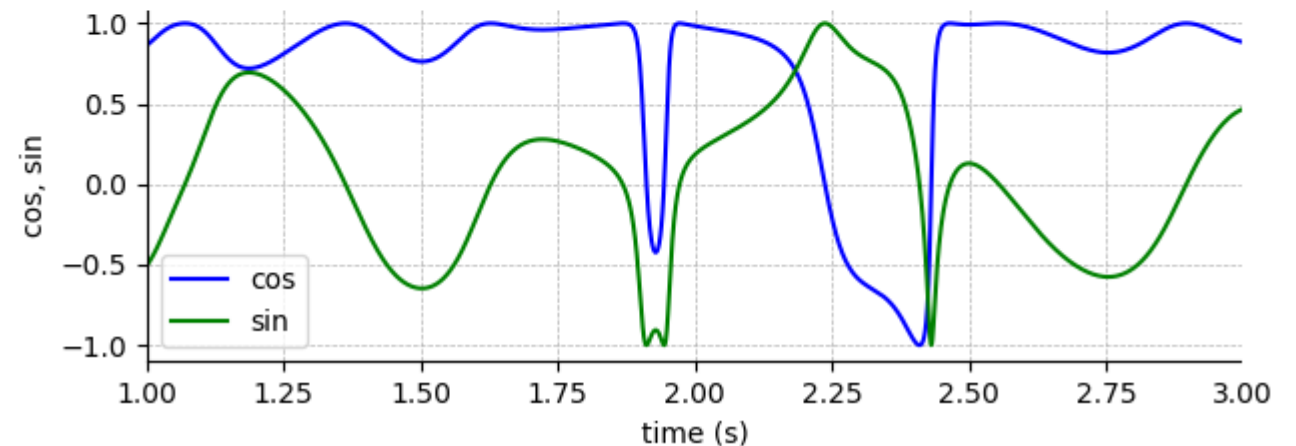
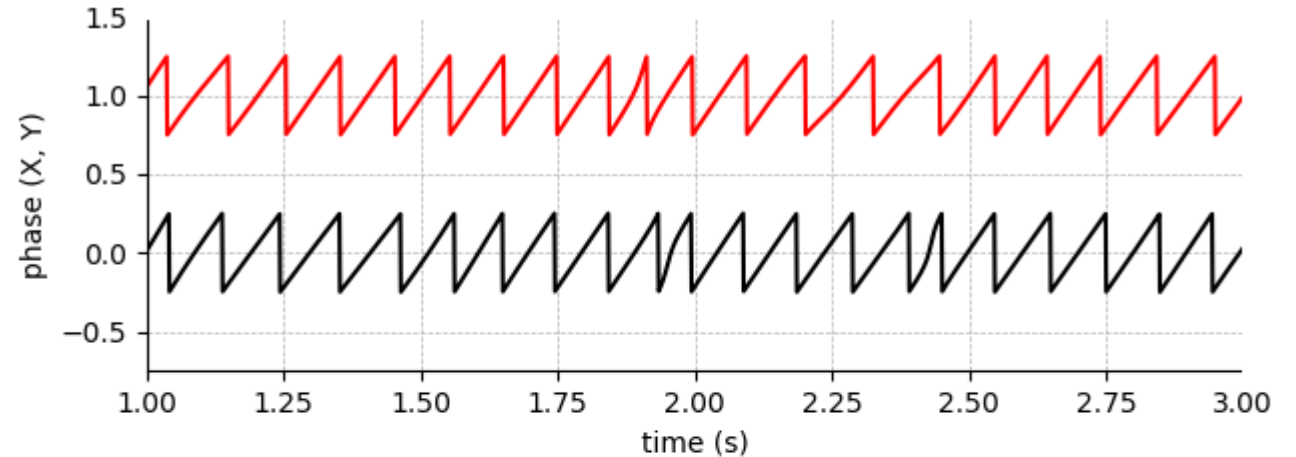
II. Phase-phase coupling (2/2)

Coupling between phases of two signals can be assessed via phase-locking value.

```
# phase
PX = np.angle(signal.hilbert(X))
PY = np.angle(signal.hilbert(Y))

# phase-locking value
p = np.abs(np.sum(np.exp(1j * (PX - PY))) / N)

# phase-locking value
pr = np.real(np.sum(np.exp(1j * (PX - PY))) / N)
pi = np.imag(np.sum(np.exp(1j * (PX - PY))) / N)
```



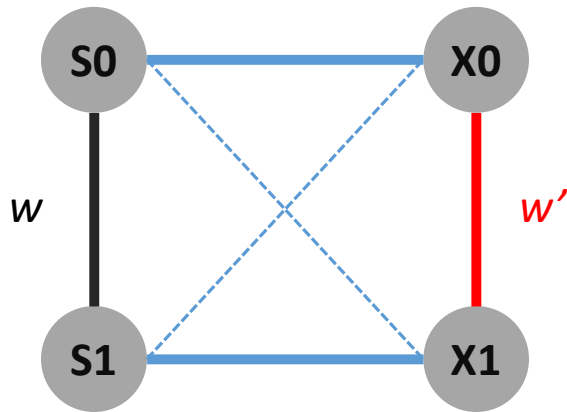
$$p(PX, PY) = 0.75 \quad p_{re}(PX, PY) = 0.75 \quad p_{im}(PX, PY) = 0.00$$

See, “L08_non_zero_lag_interactions_phase.py”

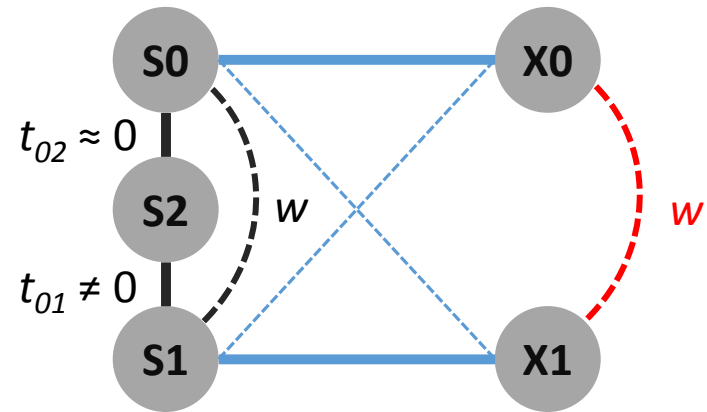
Section 7. Causal interactions

Interactions

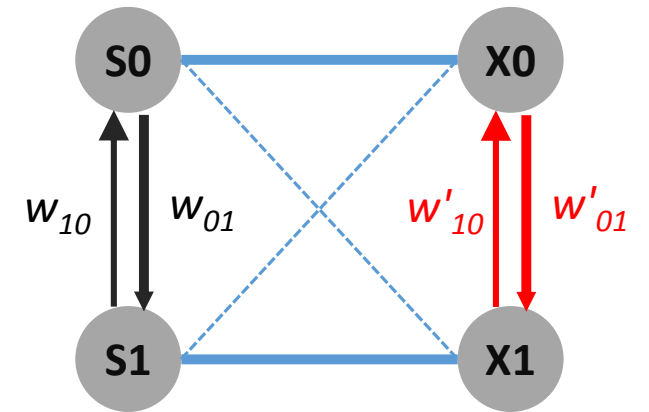
Zero-lag interactions



Non-zero-lag interactions



Causal interactions



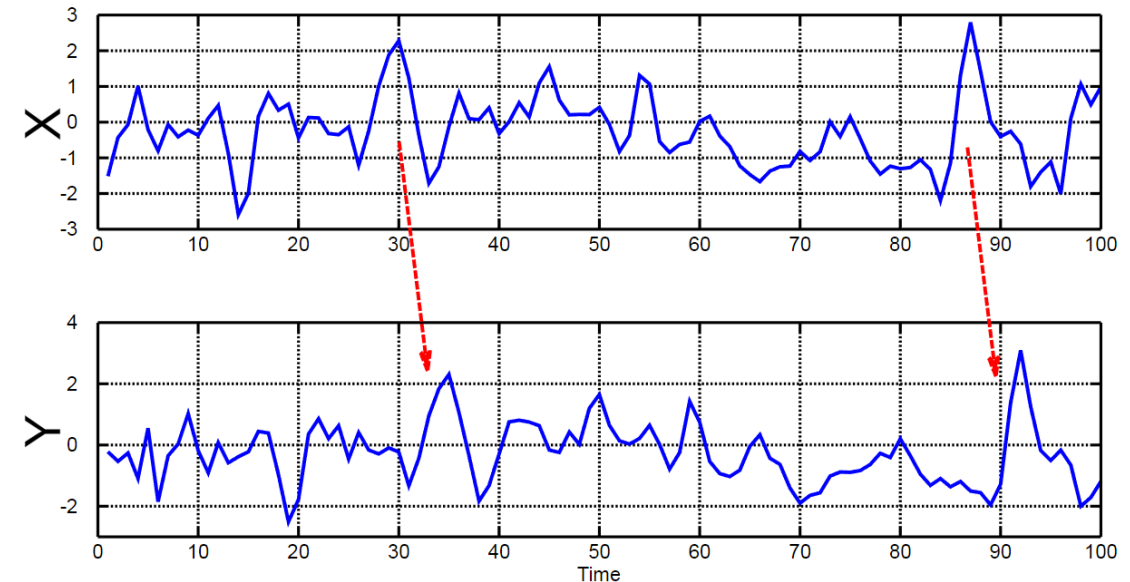
Granger causality introduction

The Granger causality test is a statistical hypothesis test for determining whether one time series is useful in **forecasting** another.

Intuition

We say that a variable X that evolves over time Granger-causes another evolving variable Y

if predictions of the value of Y based on its own past values and on the past values of X **are better** than predictions of Y based only on its own past values.



https://en.m.wikipedia.org/wiki/Granger_causality
http://www.scholarpedia.org/article/Granger_causality

Granger causality (1/2)

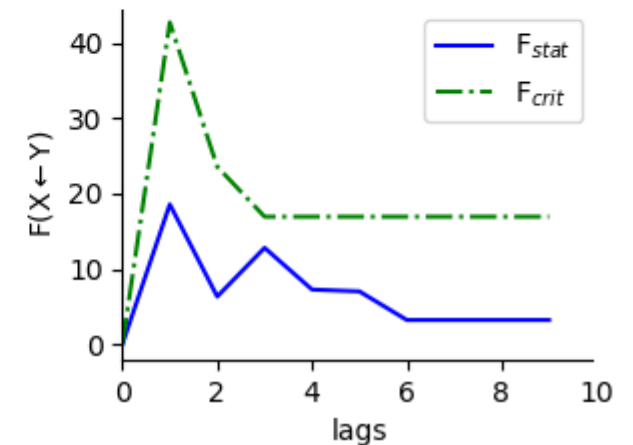
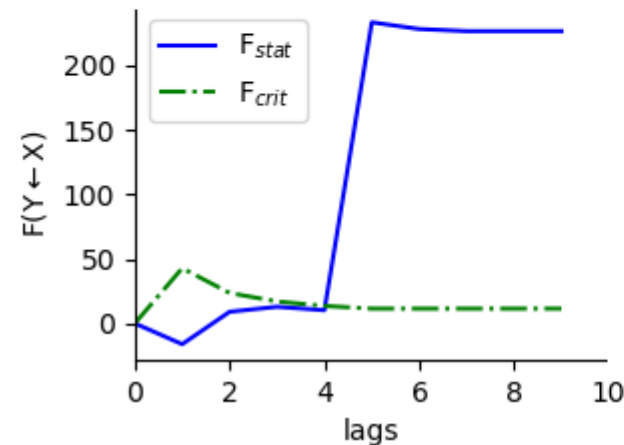
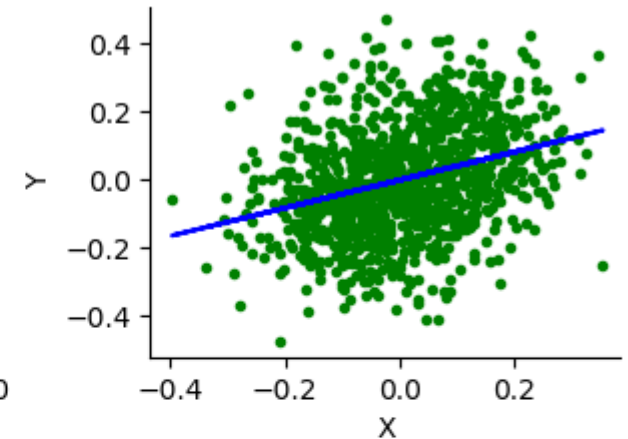
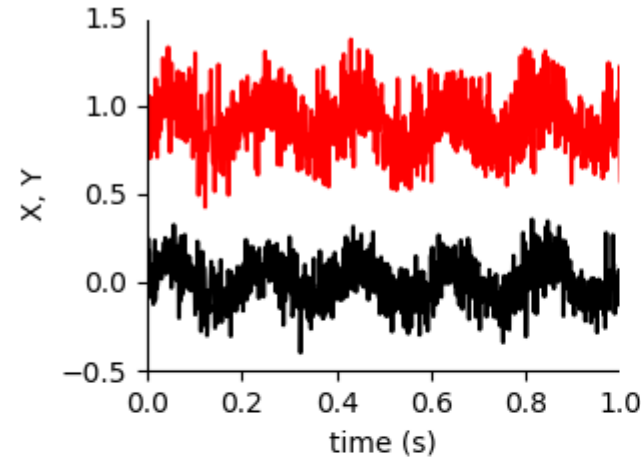
Does Y granger cause X?

```
# signal
lag = 5
X = 0.1 * np.sin(2 * np.pi * 5 * t) +
    0.1 * np.random.randn(N)
Y = np.concatenate((X[lag:], X[:lag])) +
    0.1 * np.random.randn(N)
```

```
# Granger causality
max_lags = 10
```

```
for max_lag in range(1, max_lags):
    # Y <- X
    F_stat, F_crit = granger_causality(X, Y,
                                       1e-10, max_lag)

    # X <- Y
    F_stat, F_crit = granger_causality(Y, X,
                                       1e-10, max_lag)
```



See, "L08_causal_interactions.py"

Granger causality (2/2)

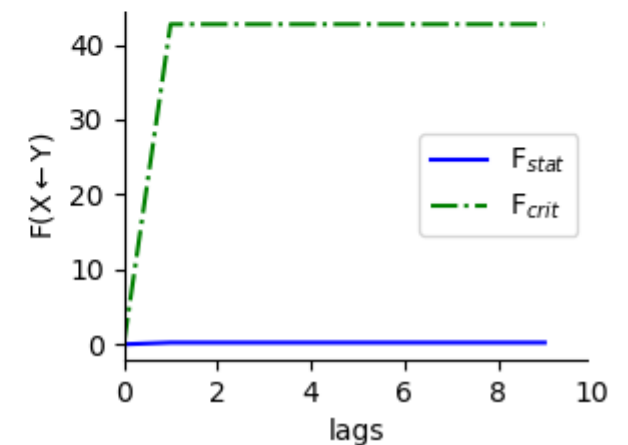
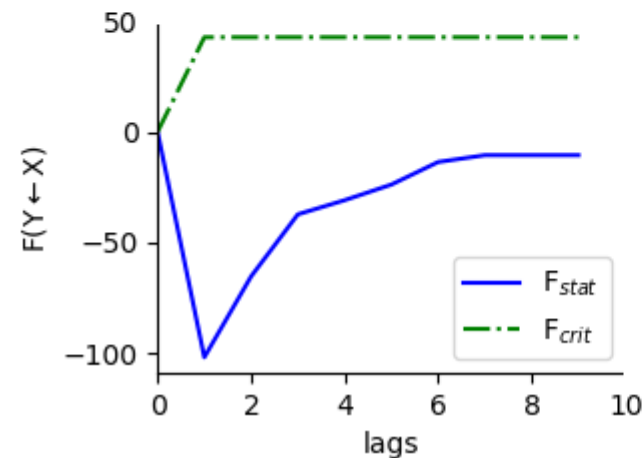
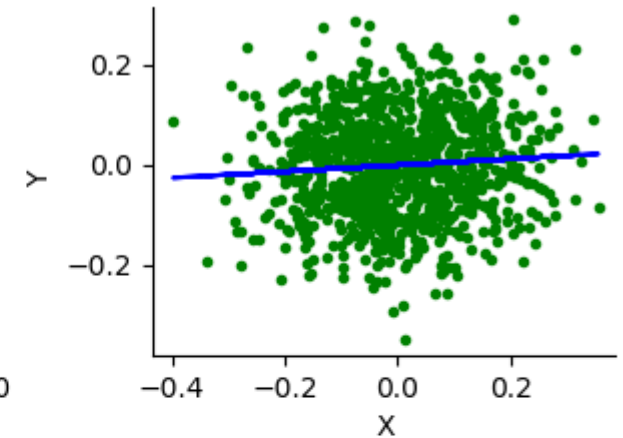
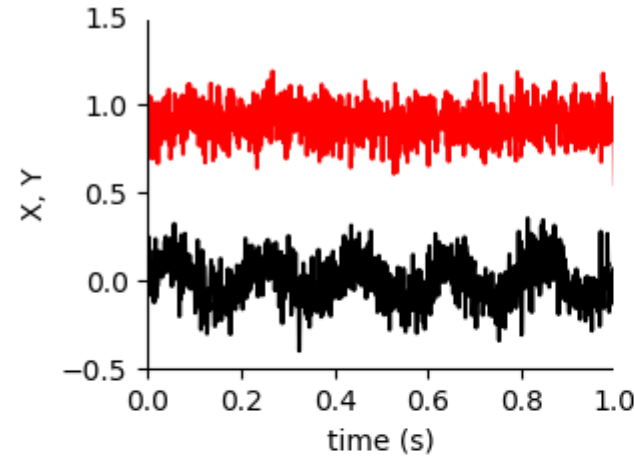
Does Y granger cause X?

```
# signal
X = 0.1 * np.sin(2 * np.pi * 5 * t) +
    0.1 * np.random.randn(N)
Y = 0.1 * np.random.randn(N)

# Granger causality
max_lags = 10

for max_lag in range(1, max_lags):
    # Y <- X
    F_stat, F_crit = granger_causality(X, Y,
                                       1e-10, max_lag)

    # X <- Y
    F_stat, F_crit = granger_causality(Y, X,
                                       1e-10, max_lag)
```



See, "L08_causal_interactions.py"

Granger algorithm (1/3)

How does it work?

```
def granger_causality(x, y, alpha, max_lag):  
  
    # fit restricted RSS model  
    x_lag, RSS_R = fit_restricted_rss_model(x, max_lag)  
  
    # fit full RSS model  
    y_lag, RSS_F = fit_full_rss_model(x, y, x_lag, max_lag)  
  
    # compare models  
    F_stat, F_crit = compare_rss_models(x_lag, y_lag, RSS_R, RSS_F, alpha)
```

See, “L08_causal_interactions.py”

Granger algorithm (2/3)

How does it work?

```
def fit_restricted_rss_model(x, max_lag):  
  
    # over lags  
    for i in range(1, (max_lag+1)):  
        Y = x[i:T]  
        X = [(np.ones(T-i, 1), np.zeros(T-i, i))] # T = len(x)  
        for j in range(1, (i+1)):  
            X[:, j] = x[(i-j):(T-j)] # lags x  
  
        # compute residuals  
        b = np.linalg.lstsq(X, Y) # regression coefficients  
        r = Y - np.dot(X, b[0]) # Y - X*b  
  
        # compute the bayesian information criterion and init model  
        BIC[i-1] = T*np.log(np.cov(r)*((T-2)/T)) + (i+1)*np.log(T)  
        RSS_R[i-1] = np.cov(r)*(T-2) # error covariance  
  
    # get best model  
    x_lag = np.argmin(BIC)
```

See, “L08_causal_interactions.py”

Granger algorithm (3/3)

How does it work?

```
def fit_full_rss_model(x, y, x_lag, max_lag):

    # over lags
    for i in range(1, (max_lag+1)):
        Y = x[(i+x_lag):T]
        X = [(np.ones(T-(i+x_lag), 1), np.zeros(T-(i+x_lag), x_lag+i))]
        for j in range(1, (x_lag+1)):
            X[:, j] = x[(i+x_lag-j):(T-j)] # lags x
        for j in range(1, (i+1)):
            X[:, (x_lag+j)] = y[(i+x_lag-j):(T-j)] # lags y
        # compute residuals
        b = np.linalg.lstsq(X, Y) # regression coefficients
        r = Y - np.dot(X, b[0]) # Y - X*b

        # compute the bayesian information criterion and init model
        BIC[i-1] = T*np.log(np.cov(r)*((T-2)/T)) + (i+1)*np.log(T)
        RSS_F[i-1] = np.cov(r)*(T-2) # error covariance

    # get best model
    y_lag = np.argmin(BIC)
```

See, “L08_causal_interactions.py”

VAR models

VAR(1) in two variables can be written in matrix form as,

$$\begin{bmatrix} y_{1,t} \\ y_{2,t} \end{bmatrix} = \begin{bmatrix} c_1 \\ c_2 \end{bmatrix} + \begin{bmatrix} A_{1,1} & A_{1,2} \\ A_{2,1} & A_{2,2} \end{bmatrix} \begin{bmatrix} y_{1,t-1} \\ y_{2,t-1} \end{bmatrix} + \begin{bmatrix} e_{1,t} \\ e_{2,t} \end{bmatrix}$$

or, equivalently, as the following system of two equations

$$\begin{aligned} y_{1,t} &= c_1 + A_{1,1}y_{1,t-1} + A_{1,2}y_{2,t-1} + e_{1,t} \\ y_{2,t} &= c_2 + A_{2,1}y_{1,t-1} + A_{2,2}y_{2,t-1} + e_{2,t} \end{aligned}$$

Properties of the **VAR model** are usually summarized using [Granger causality](#), [impulse responses](#), and [forecast error](#) variance decompositions.

https://en.m.wikipedia.org/wiki/Vector_autoregression

Literature

- **Python programming language**
 - <http://www.scipy-lectures.org/>, see “materials/L02_ScipyLectures.pdf”
- **Data analysis**
 - Cohen M., “Analyzing Neural Time Series Data: Theory and Practice”