

In [1]:

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""""
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& Senior Python Dev.

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Responsabile Cost&Qualità
Ex Ingegnere Hardware Baseband
Web: Curriculum pubblico
contributi scientifici: Ricerca
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-> Collab: Andrea Casula, Paolo Pintus
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Professor Paolo Pintus (Assistant Professor dipartimento di Fisica)
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Relazione con il professore: Relatore di tesi del 2022
Ruolo - Ricercatore a tempo determinato
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Relazione con il professore: Relatore di tesi del 2022
Ruolo - Professore associato
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Settore scientifico disciplinare - ING-INF/02 CAMPI ELETTROMAGNETICI

DATE: February 2025 NAME OF THE PROJECT: REAL TIME BLE TRACKER SYSTEM

""""
```

Out[1]:

```
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25 NAME OF THE PROJECT: REAL TIME BLE TRACKER SYSTEM\n\n"
```

In [2]:

```
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
from SALib.analyze import rbd_fast
```

```
# this is where you define your own model, procedure, experiment...
from SALib.test_functions import Ishigami
import numpy as np
```

In [3]:

```
# this is where you define your own model, procedure, experiment...
from SALib.test_functions import Ishigami
import numpy as np

def run_model(x1, x2):
    """
    COPY HERE YOUR OWN CODE

    the function takes as input 1 sample
    returns 1 or more output
    """
    lmbd_0 = 1.55 * (np.math.pow(10,-6)) # central wavelength is 1550 nanometers

    n1 = 2.848 # effective index of the grating teeth
    n2 = 2.534 # effective index of the grating slots
    # Delete from HERE =====

    # As an example, we'll look at the famous Ishigami function
    # (A and B are specific parameters for the Ishigami function)
    A = 7
    B = 0.1
    y = np.arcsin(((n1*x1+n2-x1*n2)*x2- lmbd_0)/x2 )
    y_deg = 180/np.pi * y
    # ===== TO HERE and replace with your own piece of code

    return y_deg

def conv_study(n, Y, X):
    # take n samples among the num_samples, without replacement
    subset = np.random.choice(num_samples, size=n, replace=False)
    return rbd_fast.analyze(problem=problem,
                            Y=Y[subset],
                            X=X[subset])['S1']

def bootstrap(problem, Y, X):
    """
    Calculate confidence intervals of rbd-fast indices
    1000 draws
    returns 95% confidence intervals of the 1000 indices

    problem : dictionnary as SALib uses it
    X : SA input(s)
    Y : SA output(s)
    """
    all_indices = []

    for i in range(1000):
        X_new = np.zeros(X.shape)
        Y_new = np.zeros(Y.shape).flatten()
```

```

# draw with replacement
tirage_indices = np.random.randint(0, high=Y.shape[0], size = Y.shape[0])

for j, index in enumerate(tirage_indices):
    X_new[j,:] = X[index, :]
    Y_new[j] = Y[index]

all_indices.append(rbd_fast.analyze(problem=problem, Y=Y_new, X=X_new)['S1'])

means = np.array([i.mean() for i in np.array(all_indices).T])
stds = np.array([i.std() for i in np.array(all_indices).T])
return np.array([means - 2 * stds, means + 2 * stds])

```

```

In [4]: # STATE THE PROBLEM DICTIONNARY
# what will be varying (=inputs) ? in what bounds ?
FF_min = 0.2
FF_max = 0.7
g_period_power = (np.math.pow(10,-9)) # structure grating period is 660 nanometers
g_period_min = 620 * g_period_power # 620 nm
g_period_max = 690 * g_period_power # 690 nm
problem = {
    'num_vars': 2,
    'names': ['FillFactor', 'GratingPeriod'],
    'bounds': [[FF_min,FF_max ],
               [g_period_min, g_period_max]]
}
# say we want to draw 150 samples

num_samples = 150
# we draw a Latin Hypercube Sampling (LHS) that is fitted for an RBD FAST analysis
# (other sampling methods available in the library though)

from SALib.sample.latin import sample

all_samples = sample(problem, num_samples)

# run your model, procedure, experiment for each sample of your sampling

# unpack all_samples into 3 vectors x1, x2, x3
x1, x2 = all_samples.T

```

```

In [5]: # run the model, all samples at the same time
ishigami_results = run_model(x1, x2)

# Let us look at the analyze method

rbd_fast.analyze(problem=problem, Y=ishigami_results, X=all_samples)
# storing the first order indices of the analyze method
si1 = rbd_fast.analyze(problem=problem, Y=ishigami_results, X=all_samples)['S1']

# make nice plots with the indices (looks good on your presentations)
# do not use the plotting tools of SALib, they are made for the method of Morris ...

```

```

In [6]: fig, ax = plt.subplots()
fig.set_size_inches(18, 5)

ax.tick_params(labelsize=18)

```

```

# ===== X-AXIS =====
ax.set_xticks(np.arange(problem['num_vars']))
ax.set_xticklabels(problem['names'])
ax.set_xlim(xmin=-0.5, xmax=problem['num_vars'] - 0.5)

# ===== Y-AXIS =====
ax.set_ylabel('RBD-FAST\nSensitivity indices for Grating Coupler : Fill Factor and P

# ===== BARS REPRESENTING THE SENSITIVITY INDICES =====
ax.bar(np.arange(problem['num_vars']), si1,
       color='DarkSeaGreen');

#in striped grey : not significant indices
ax.fill_between(x=[-0.5, 5.5], y1=-0.1, y2=0.1, color='grey', alpha=0.2, hatch='//',

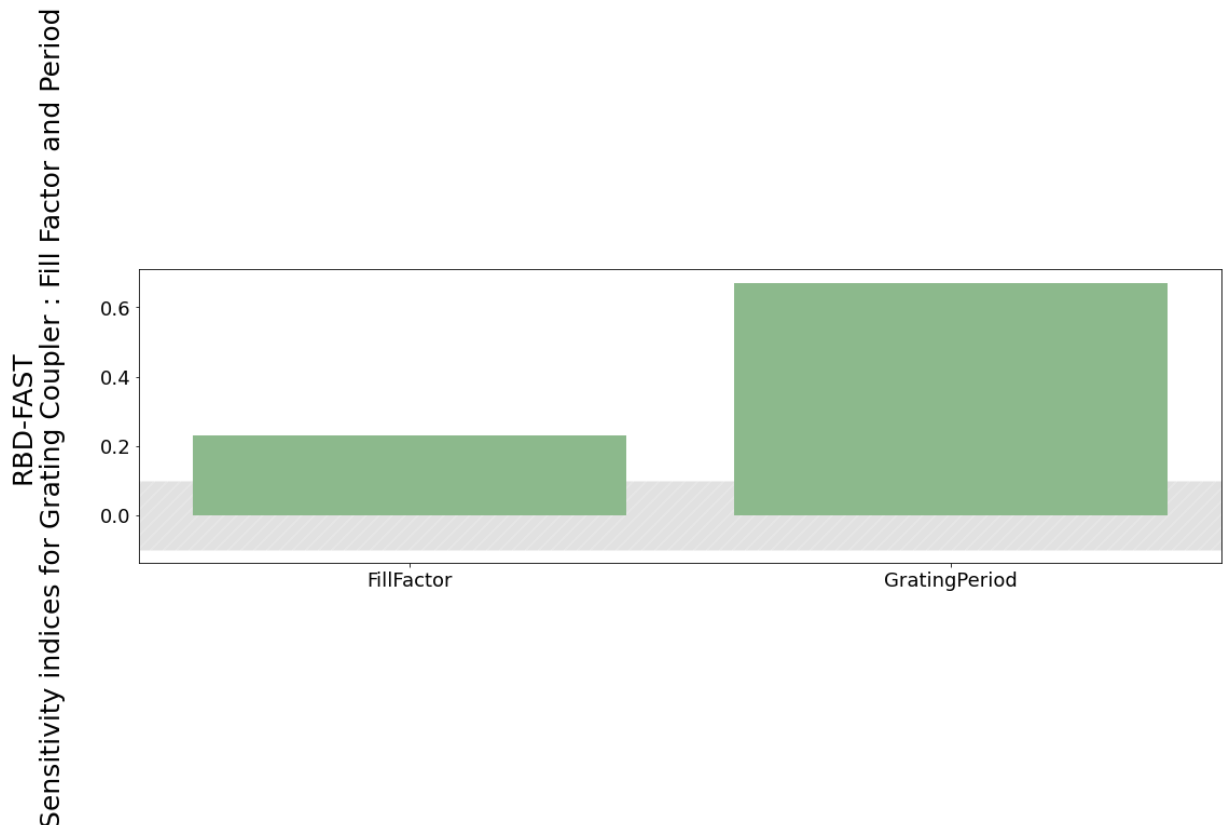
# take a closer look to understand interactions (looks even better on your presentati
# this part can be done without analyzing with SALib, just with the ouput from the s

fig, ax = plt.subplots()
fig.set_size_inches(8, 7)

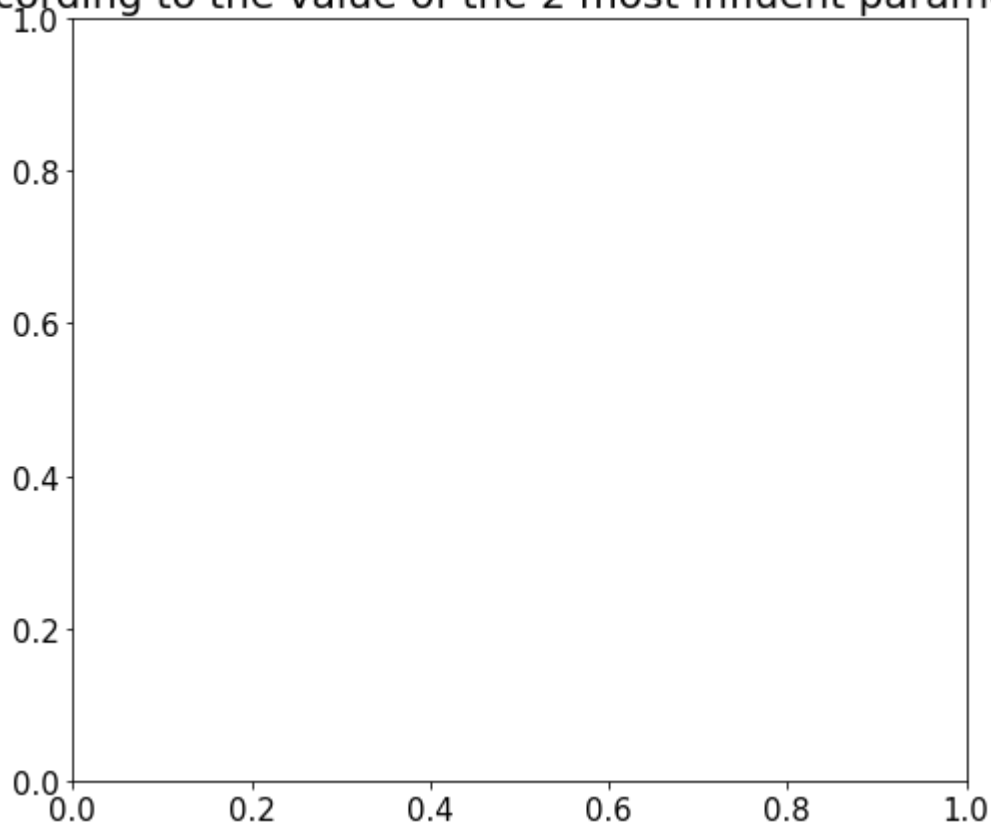
ax.tick_params(labelsize=15)
ax.set_title('Output of the model studied (Bragg condition)\n'
            'according to the value of the 2 most influent parameters',
            fontsize=20)

```

Out[6]: Text(0.5, 1.0, 'Output of the model studied (Bragg condition)\naccording to the value of the 2 most influent parameters')



Output of the model studied (Bragg condition)
according to the value of the 2 most influent parameters



In [7]:

```
fig, ax = plt.subplots()
fig.set_size_inches(18, 5)

ax.tick_params(labelsize=18)

# ===== X-AXIS =====
ax.set_xticks(np.arange(problem['num_vars']))
ax.set_xticklabels(problem['names'])
ax.set_xlim(xmin=-0.5, xmax=problem['num_vars'] - 0.5)

# ===== Y-AXIS =====
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ax.set_title('Output of the model studied (Bragg condition)\n'
            'according to the value of the 2 most influent parameters',
            fontsize=20)
```

```

# ===== SCATTER =====
size = np.ones(num_samples) * 75

sc = ax.scatter(x1, x2,
                c=ishigami_results,
                s=size,
                vmin=ishigami_results.min(),
                vmax=ishigami_results.max(),
                cmap='seismic',
                edgecolor=None)

# ===== X-AXIS =====
ax.set_xlim(xmin=problem['bounds'][0][0], xmax=problem['bounds'][0][1])
ax.set_xlabel('Parameter 1 -> Fill Factor [nanometers / nanometers]', fontsize=20)

# ===== Y-AXIS =====
ax.set_ylim(ymin=problem['bounds'][1][0], ymax=problem['bounds'][1][1])
ax.set_ylabel('Parameter 2 -> Grating Period [1e-07 meters]', fontsize=20)

# ===== COLORBAR =====
ticks = np.arange(ishigami_results.min(), ishigami_results.max(), 5)
cb = plt.colorbar(sc, ticks=ticks);
cb.ax.set_yticklabels([str(round(i,1)) for i in ticks])

fig.tight_layout()

all_indices = np.array([conv_study(n=n, Y=ishigami_results, X=all_samples)
                        for n in np.arange(50, num_samples + 1, 5)])

# convergence check
fig, ax = plt.subplots()
fig.set_size_inches(15,8)

ax.set_title('Convergence of the sensitivity indexes', fontsize=20)

ax.tick_params(labelsize=16)

ax.set_xlim(xmin=0, xmax=(num_samples - 50)//10)
ax.set_xticks(np.arange(0,(num_samples - 50)//10 +1,5))
ax.set_xticklabels([str(i) for i in range(50,num_samples+1,50)])

ax.set_ylim(ymin=-0.15, ymax=1)

for p,param in enumerate(problem['names']):
    ax.plot(all_indices[:,p], linewidth=3, label=param)

ax.fill_between(x=[0,(num_samples - 50)//10], y1=-0.15, y2=0.1, color='grey', alpha=
               label='REMINDER : not significant')

ax.legend(fontsize=20, ncol=4)

# Get bootstrap confidence intervals for each index

bootstrap_conf = bootstrap(problem=problem, Y=ishigami_results, X=all_samples)

```

```

# make nice plots with the indices (looks good on your presentations)
# do not use the plotting tools of SALib, they are made for the method of Morris ...

fig, ax = plt.subplots()
fig.set_size_inches(8,4)

ax.tick_params(labelsize=18)

# ===== X-AXIS =====
ax.set_xticks(np.arange(problem['num_vars']))
ax.set_xticklabels(problem['names'])
ax.set_xlim(xmin=-0.5, xmax=problem['num_vars'] - 0.5)

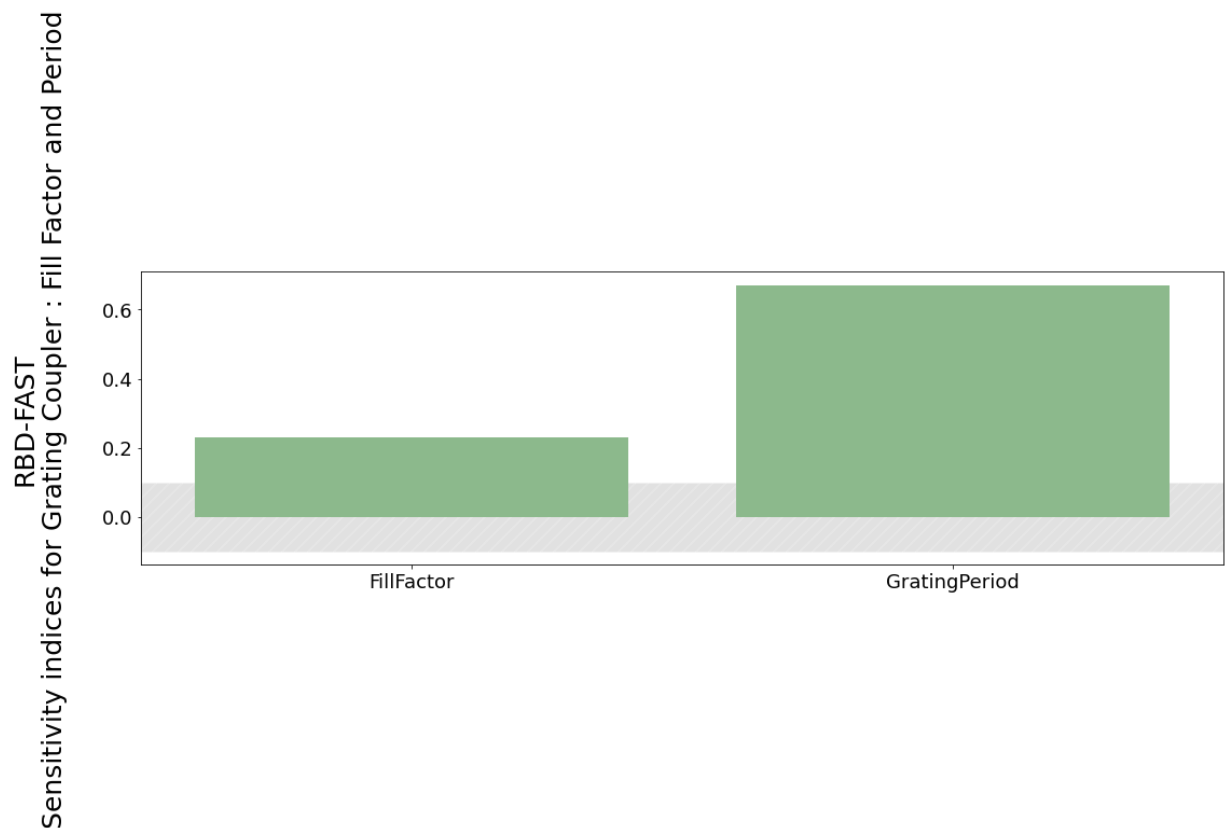
# ===== Y-AXIS =====
ax.set_ylabel('Sensitivity indices\n', fontsize=25)

# ===== BARS REPRESENTING THE SENSITIVITY INDICES =====
ax.bar(np.arange(problem['num_vars']), si1,
       color='DarkSeaGreen');

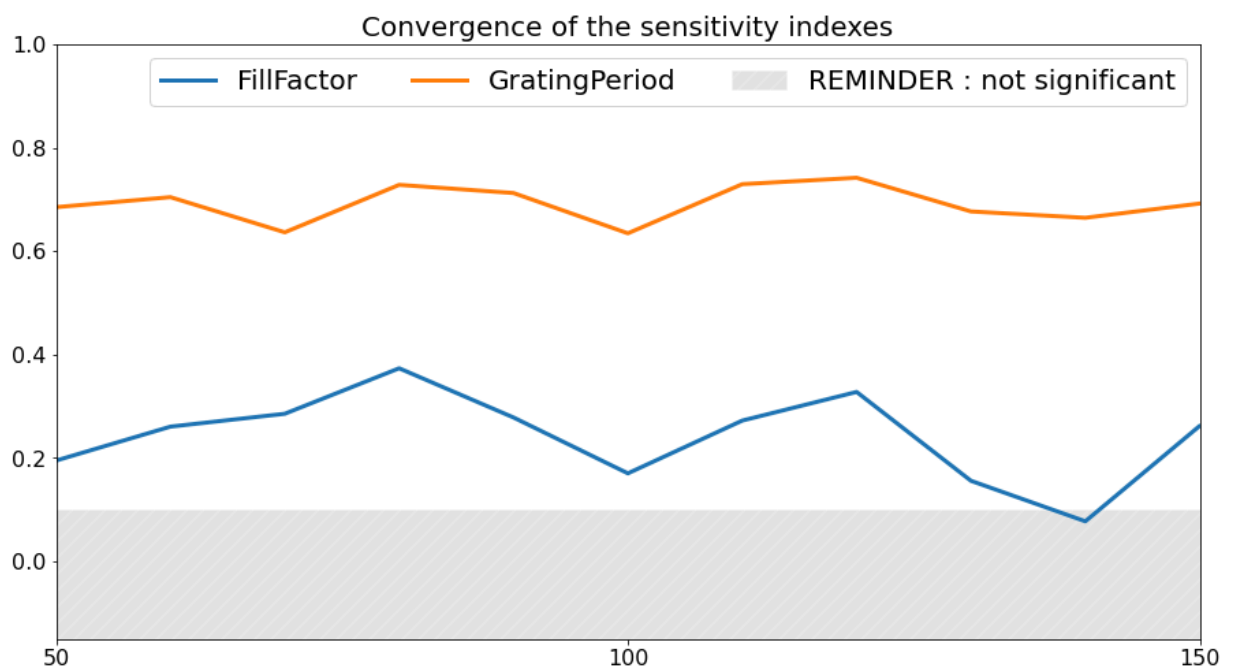
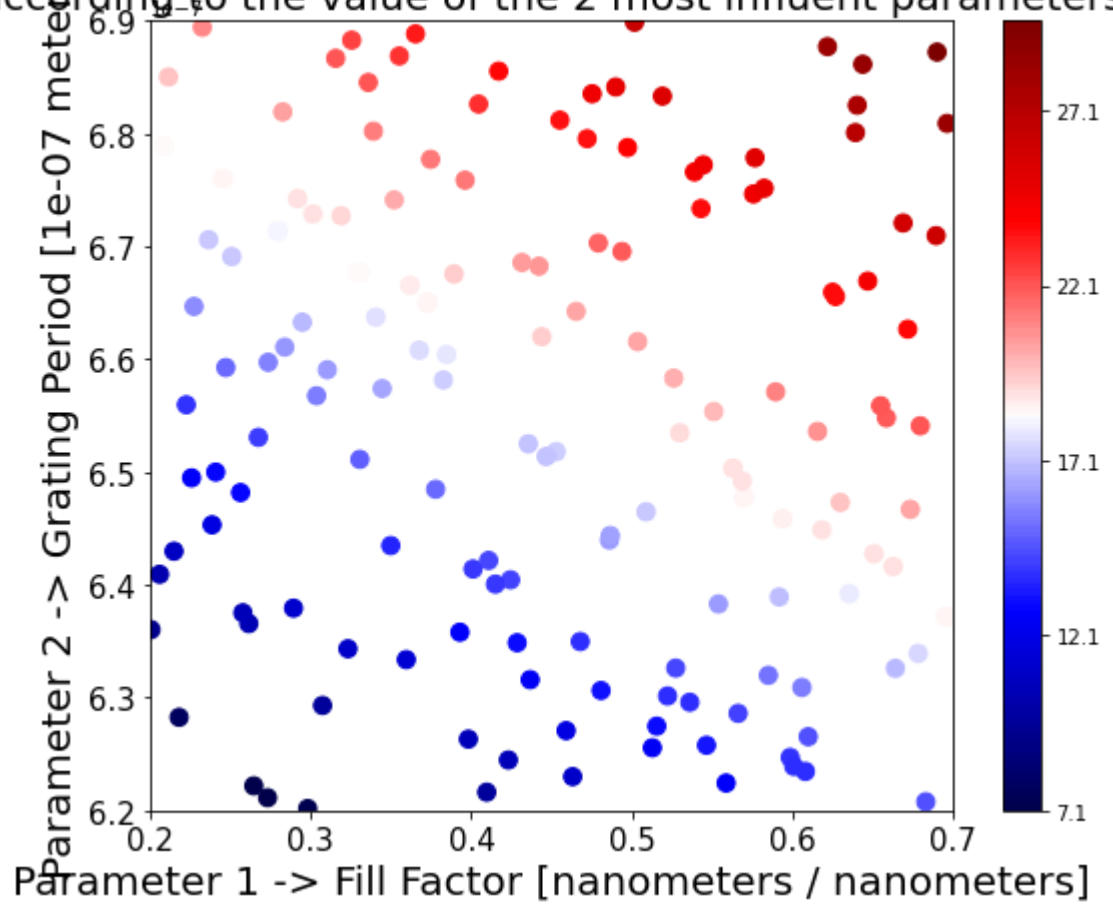
# ===== LINES REPRESENTING THE BOOTSTRAP "CONFIDENCE INTERVALS" =====
for j in range(problem['num_vars']):
    ax.plot([j, j], [bootstrap_conf[0, j], bootstrap_conf[1, j]],
            'k')

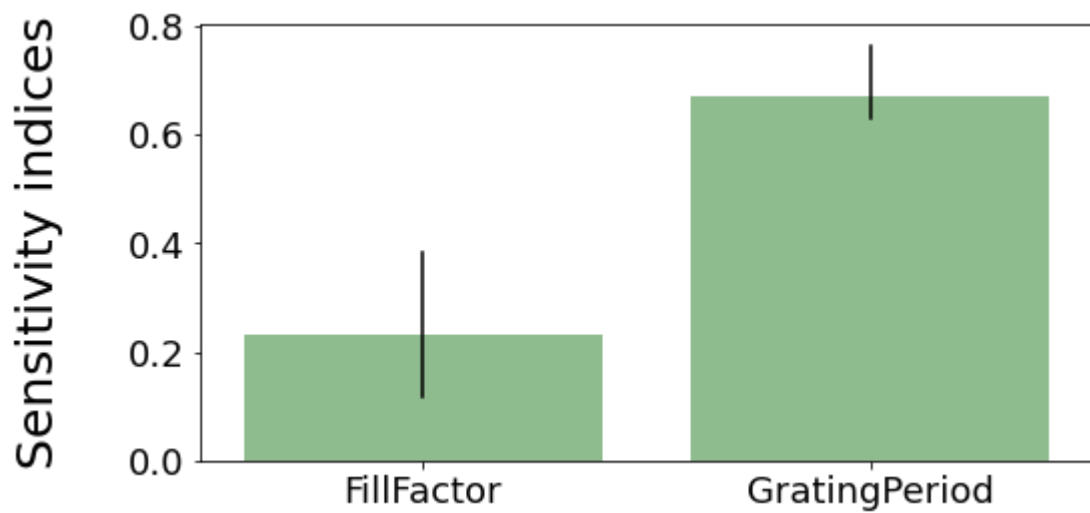
#ax.fill_between(x=[-0.5, 5.5], y1=-0.1, y2=0.1, color='grey', alpha=0.2, hatch='//')

```



Output of the model studied (Bragg condition)
according to the value of the 2 most influent parameters





In [8]: #####

In [9]:

```
# this is where you define your own model, procedure, experiment...
from SALib.test_functions import Ishigami
import numpy as np

def run_model(x1, x2):
    """
    COPY HERE YOUR OWN CODE

    the function takes as input 1 sample
    returns 1 or more output
    """
    #lmbd_0 = 1.55 * (np.math.pow(10,-6)) # central wavelength is 1550 nanometers
    # lmbd_0 is the new x1
    FF=0.6
    n1 = 2.848 # effective index of the grating teeth
    n2 = 2.534 # effective index of the grating slots
    # Delete from HERE =====

    # As an example, we'll look at the famous Ishigami function
    # (A and B are specific parameters for the Ishigami function)
    A = 7
    B = 0.1
    y = np.arcsin(((n1*FF+n2-FF*n2)*x2- x1)/x2 )
    y_deg = 180/np.pi * y
    # ===== TO HERE and replace with your own piece of code

    return y_deg

def conv_study(n, Y, X):
    # take n samples among the num_samples, without replacement
    subset = np.random.choice(num_samples, size=n, replace=False)
    return rbd_fast.analyze(problem=problem,
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    Calculate confidence intervals of rbd-fast indices
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    """
    all_indices = []

    for i in range(1000):
        X_new = np.zeros(X.shape)
        Y_new = np.zeros(Y.shape).flatten()
        # draw with replacement
        tirage_indices = np.random.randint(0, high=Y.shape[0], size = Y.shape[0])

        for j, index in enumerate(tirage_indices):
            X_new[j,:] = X[index, :]
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        all_indices.append(rbd_fast.analyze(problem=problem, Y=Y_new, X=X_new)['S1'])

    means = np.array([i.mean() for i in np.array(all_indices).T])
    stds = np.array([i.std() for i in np.array(all_indices).T])
    return np.array([means - 2 * stds, means + 2 * stds])

```

In [10]:

```

# STATE THE PROBLEM DICTIONNARY
# what will be varying (=inputs) ? in what bounds ?

g_period_power = (np.math.pow(10,-9)) # structure grating period is 660 nanometers
g_period_min = 620 * g_period_power # 620 nm
g_period_max = 690 * g_period_power # 690 nm
lmbd_BAND_C_min = 1530*g_period_power
lmbd_BAND_C_max = 1565*g_period_power

problem = {
    'num_vars': 2,
    'names': ['Wavelength', 'GratingPeriod'],
    'bounds': [[lmbd_BAND_C_min,lmbd_BAND_C_max ],
               [g_period_min, g_period_max]]
}
# say we want to draw 150 samples

num_samples = 150
# we draw a Latin Hypercube Sampling (LHS) that is fitted for an RBD FAST analysis
# (other sampling methods available in the library though)

from SALib.sample.latin import sample

all_samples = sample(problem, num_samples)

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# unpack all_samples into 3 vectors x1, x2, x3
x1, x2 = all_samples.T

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# run the model, all samples at the same time
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# Let us look at the analyze method

rbd_fast.analyze(problem=problem, Y=ishigami_results, X=all_samples)
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si1 = rbd_fast.analyze(problem=problem, Y=ishigami_results, X=all_samples)['S1']

# make nice plots with the indices (looks good on your presentations)
# do not use the plotting tools of SALib, they are made for the method of Morris ...

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In [12]:

```

import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
from SALib.analyze import rbd_fast

# this is where you define your own model, procedure, experiment...
from SALib.test_functions import Ishigami
import numpy as np
fig, ax = plt.subplots()
fig.set_size_inches(18, 5)

ax.tick_params(labelsize=18)

# ===== X-AXIS =====
ax.set_xticks(np.arange(problem['num_vars']))
ax.set_xticklabels(problem['names'])
ax.set_xlim(xmin=-0.5, xmax=problem['num_vars'] - 0.5)

# ===== Y-AXIS =====
ax.set_ylabel('Sensitivity indices for Grating Coupler : Wavelength and Period\n', f

# ===== BARS REPRESENTING THE SENSITIVITY INDICES =====
ax.bar(np.arange(problem['num_vars']), si1,
       color='DarkSeaGreen');

#in striped grey : not significant indices
ax.fill_between(x=[-0.5, 5.5], y1=-0.1, y2=0.1, color='grey', alpha=0.2, hatch='///',

# take a closer look to understand interactions (looks even better on your presentati
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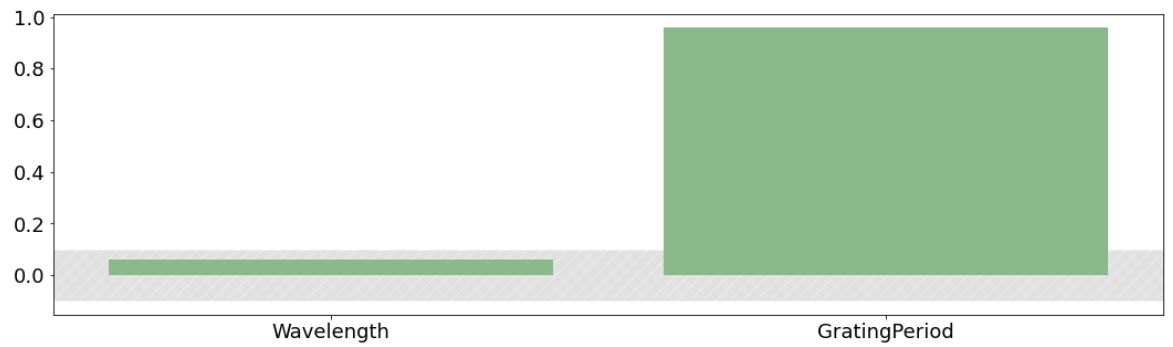
fig, ax = plt.subplots()
fig.set_size_inches(8, 7)

ax.tick_params(labelsize=15)
ax.set_title('Maximum Radiation Angle\n'
            'according to the value of the 2 most influent parameters',
            fontsize=20)

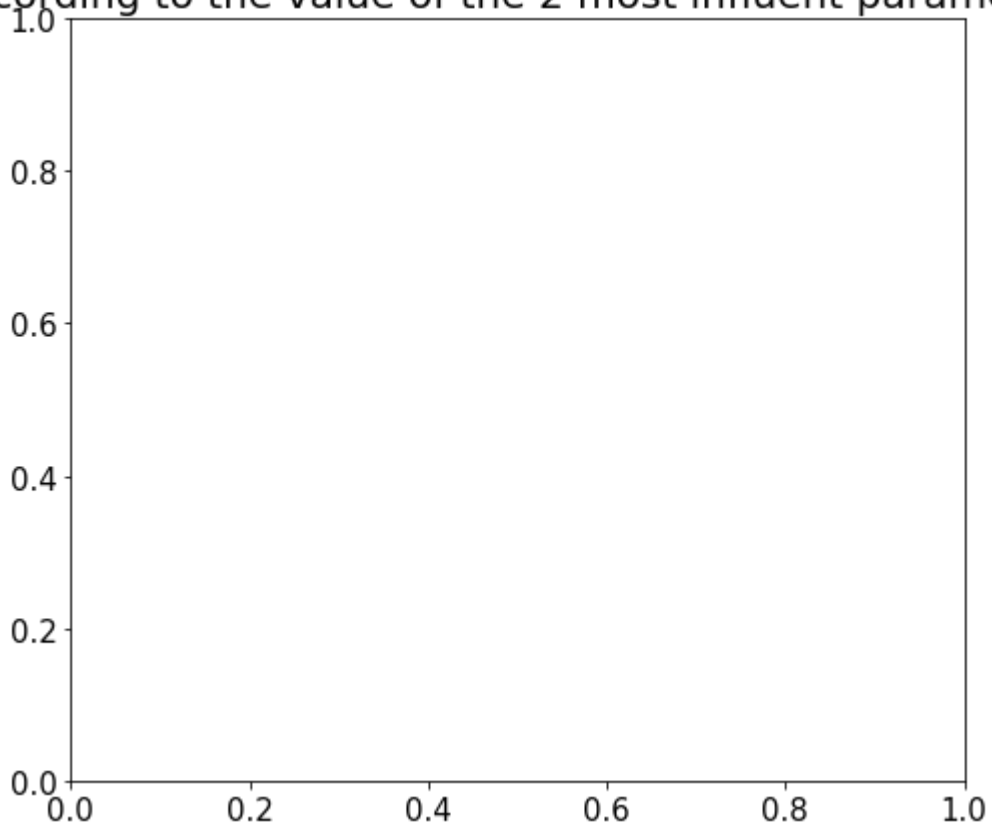
```

Out[12]: Text(0.5, 1.0, 'Maximum Radiation Angle\naccording to the value of the 2 most influent parameters')

Sensitivity indices for Grating Coupler : Wavelength and Period



Maximum Radiation Angle
according to the value of the 2 most influent parameters



In [13]: #####

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