

Optimize the power conversion efficiency of organic photovoltaic solar cells

Introduction

Organic photovoltaic (OPV) are 'plastic' solar cells that can be made cheaply and easily as you can use techniques like roll to roll printing, inject printing and spray coating. Current generation solar cells take several years of use before they payback the energy required in their manufacture, OPVs are so efficient that their energy payback is only 24hours. Power conversion efficiencies (PCEs) of OPVs are now around 14%. To commercialise them, we need to figure out how best to manufacture them.

Organic photovoltaic devices have a sandwich architecture. The bottom layers Al/Mg and LiF are the bottom electrode. The important part is the bulk heterojunction, shown in red in the figure below, which comprises of a low band gap polymer which is the electron donor and fullerene which is the electron acceptor. Addition of an additive helps with forming and bridging separate nanodomains of donor and acceptor. Solar cells work by using light to form an exciton which then separates into an electron-hole pair and you want these to be separated from each other, which is why you want separate nanodomain of donor and acceptor. The top of the solar cell is PEDOT:PSS (a conducting polymer) and ITO (indium tin oxide), a see-through electrode, which together act as the top electrode.

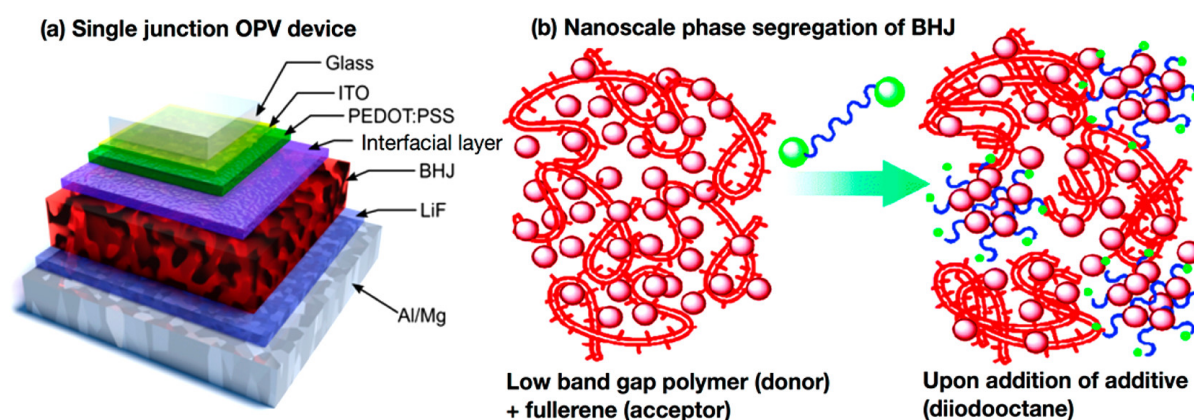


Figure 1. (a) Schematic of single junction organic photovoltaic (OPV) devices, showing the bulk heterojunction (BHJ; in red), and the multiple interfacial layers in the device. (b) Schematic of BHJ morphology: in this case, a low band gap polymer donor and a fullerene acceptor undergoing nanoscale phase segregation into discrete nanoscale domains of donor and acceptor. The use of an additive is often purported to assist in nanodomain formation, as shown here. Taken from [ACS Nano 2018, 12, 7434–7444]

The task

The task is to optimise the construction of this type of solar cell. Donor weight percentage is a measure of the ratio of donor to acceptor in the heterojunction. Total solution concentration is the concentration of the spin-coating solution. Bulk heterojunction spin-coating speed is a measure of how fast you spin the device when coating it with the bulk heterojunction mixture. Processing

additive is the amount of additive (diiodooctane) added to the mixture. The thickness of a spun film is determined by the spin speed, solvent vapour pressure and solution viscosity, as both the donor weight percentage and total solution concentration can affect viscosity, the first three factors can all affect the thickness of the final BHJ layer. The additive (diiodooctane) increases the drying time for the film, helping to separate the heterojunction out into nanodomains of donor and acceptor rich areas.

Factors selected:

Name	Factors	Factor range	No. of levels
Donor	Donor weight percentage	10-55 (wt %)	4
Conc.	Total solution concentration	10-25 (mg/mL)	4
Spin	Bulk heterojunction spin-case speed	600 - 3000 (rpm)	4
Add.	Processing additive	0-12 (vol %)	4

We shall use the shortened names from the table above.

Files:

1. `solar_cells_1.csv` results from the first experiment, fractional factorial, 4 factors and 4 levels, here we have 16 experiments, one failed to solidify.
2. `'solar_cells_2.csv'` has results from the second experiment, a fractional factorial, 3 factors and 3 levels. This covers a smaller range.

The data is taken from: "How To Optimize Materials and Devices via Design of Experiments and Machine Learning: Demonstration Using Organic Photovoltaics", Bing Cao, Lawrence A. Adutwum, Anton O. Oliynyk, Erik J. Luber, Brian C. Olsen, Arthur Mar, and Jillian M. Buriak, ACS Nano 2018, 12, 7434–7444

First we import our packages

```
In [1]: # for dataframes
import pandas as pd

# for pictures
import matplotlib.pyplot as plt
# for maths
import numpy as np

## Some code in doenuit needs updating, so use this to ignore the warnings
import warnings
warnings.filterwarnings('ignore')

# make sure these are in the same directory as this file
import doenuit
import designer
```

Read in the first experiment's data

```
In [2]: df=pd.read_csv('solar_cells_1.csv')
df
```

Out[2]:

	experiment #	donor percentage	total concentration	spin speed	additive	PCE	std of PCE (%)	number of devices
0	NaN	% (wt)	mg/mL	rpm	vol %	%	NaN	NaN
1	1-1	10	20	3000	2	0.05	5.0	14.0
2	1-2	10	25	1000	8	3.24	11.0	10.0
3	1-3	10	10	600	0	0.016	16.0	14.0
4	1-4	10	15	2000	12	0.0004	4.0	10.0
5	1-5	25	20	600	12	7.14	13.0	8.0
6	1-6	25	15	1000	2	3.22	32.0	8.0
7	1-7	25	10	3000	8	0.00033	7.0	14.0
8	1-8	25	25	2000	0	7.21	17.0	11.0
9	1-9	40	10	1000	12	1.85	5.0	3.0
10	1-10	40	20	2000	8	6.16	28.0	12.0
11	1-11	40	25	600	2	3.9	8.0	11.0
12	1-12	40	15	3000	0	2.27	35.0	9.0
13	1-13	55	10	2000	2	1.16	4.0	3.0
14	1-14	55	15	600	8	3.18	12.0	10.0
15	1-15	55	20	1000	0	3.89	10.0	13.0
16	1-16	55	25	3000	12	NaN	NaN	NaN

Set up input and response dataframes

We must drop the last experiment, as these devices didn't set.

```
In [3]: inputs = pd.DataFrame({
    'Donor %': [float(x) for x in df.iloc[1:-1,1]],
    'Conc.': [float(x) for x in df.iloc[1:-1,2]],
    'Spin': [float(x) for x in df.iloc[1:-1,3]],
    'Add.': [float(x) for x in df.iloc[1:-1,4]]})
inputs
```

```
Out[3]:
```

	Donor %	Conc.	Spin	Add.
0	10.0	20.0	3000.0	2.0
1	10.0	25.0	1000.0	8.0
2	10.0	10.0	600.0	0.0
3	10.0	15.0	2000.0	12.0
4	25.0	20.0	600.0	12.0
5	25.0	15.0	1000.0	2.0
6	25.0	10.0	3000.0	8.0
7	25.0	25.0	2000.0	0.0
8	40.0	10.0	1000.0	12.0

	Donor %	Conc.	Spin	Add.
9	40.0	20.0	2000.0	8.0
10	40.0	25.0	600.0	2.0
11	40.0	15.0	3000.0	0.0
12	55.0	10.0	2000.0	2.0
13	55.0	15.0	600.0	8.0
14	55.0	20.0	1000.0	0.0

```
In [4]: responses = pd.DataFrame({'PCE': [float(x) for x in df['PCE'][1:-1]]})
responses
```

```
Out[4]:
```

	PCE
0	0.05000
1	3.24000
2	0.01600
3	0.00040
4	7.14000
5	3.22000
6	0.00033
7	7.21000
8	1.85000
9	6.16000
10	3.90000
11	2.27000
12	1.16000
13	3.18000
14	3.89000

Task 1. Create a linear (main factors only) model

Create a linear model, i.e. a model that has just the main effects (also known as a first order model or main effects model) Fit your linear model to the first experiment's data and calculate R2 and Q2 for your fitted model. Then answer the questions.

```
In [5]: # this selects which columns in inputs to use
input_selector = range(len(inputs.columns))

this_model, R2, temp_tuple, _ = doenut.calculate_R2_and_Q2_for_models(
    inputs,
    # input dataframe
    responses,
```

```

# responses dataframe
input_selector=input_selector,
# which columns in input to fit
response_selector=[0],
# This selects the zeroth column of responses
use_scaled_inputs=True,
# we scale the inputs so the coefficients
# are comparable
do_scaling_here=True
# scale inputs inside this function
)

```

```
new_model, predictions, ground_truth, coeffs, R2s, R2, Q2= temp_tuple
```

Input terms are ['Donor %', 'Conc.', 'Spin', 'Add.']

Input Responses are ['PCE']

Selected Response is PCE

Selected input terms: ['Donor %', 'Conc.', 'Spin', 'Add.']

Averaging replicates

Input data is 15 points long

We are using 15 data points

Left out data point 0:	R2 = 0.627	Ave. Error = -2.86
Left out data point 1:	R2 = 0.643	Ave. Error = -2.3
Left out data point 2:	R2 = 0.568	Ave. Error = 1.59
Left out data point 3:	R2 = 0.604	Ave. Error = -2.3
Left out data point 4:	R2 = 0.638	Ave. Error = 3.78
Left out data point 5:	R2 = 0.627	Ave. Error = 1.6
Left out data point 6:	R2 = 0.558	Ave. Error = -0.0576
Left out data point 7:	R2 = 0.643	Ave. Error = 3.87
Left out data point 8:	R2 = 0.599	Ave. Error = -0.189
Left out data point 9:	R2 = 0.599	Ave. Error = 2.3
Left out data point 10:	R2 = 0.672	Ave. Error = -3.01
Left out data point 11:	R2 = 0.607	Ave. Error = 0.816
Left out data point 12:	R2 = 0.591	Ave. Error = -0.533
Left out data point 13:	R2 = 0.619	Ave. Error = -1.38
Left out data point 14:	R2 = 0.615	Ave. Error = -1.43

R2 overall is 0.604

Mean of test set: 2.885782

Mean being used: 2.885782

Sum of squares of the residuals (explained variance) is 72.3948507524448

Sum of squares total (total variance) is 87.23748999604001

Q2 is 0.17

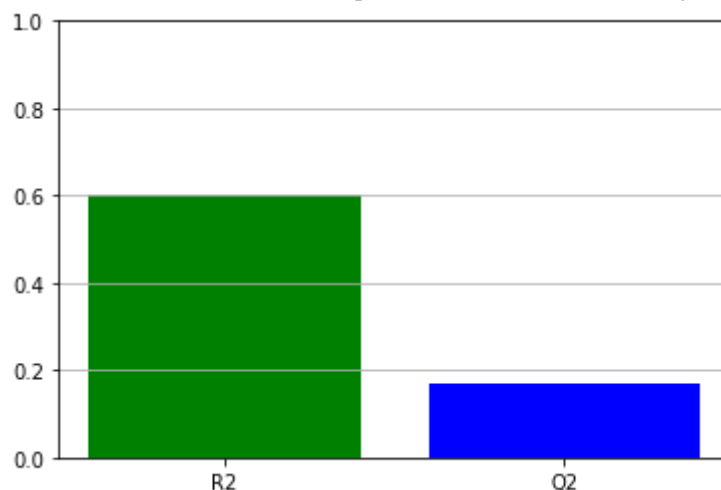
Response PCE R2 is 0.604

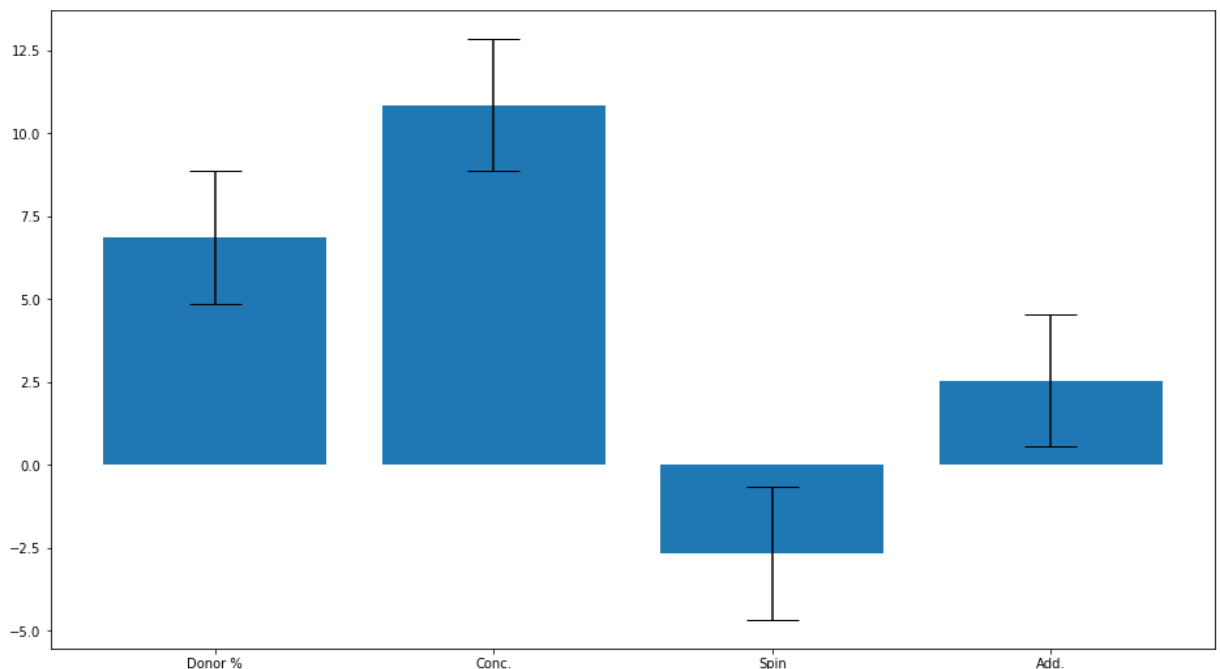
Input selector was range(0, 4)

Input_selector was: [0, 1, 2, 3]

Average coefficients are: [6.85475012 10.85602224 -2.66214206 2.5458804]

Coefficient labels are: ['Donor %', 'Conc.', 'Spin', 'Add.']





Task 2. Create a quadratic, parsimonious and hierarchical model

Task 2. Create a **quadratic**, **parsimonious** and **hierarchical** model. Starting with a quadratic model, and making sure that all models are hierarchical, optimise the model by removing **only** the **statistically insignificant** terms. Keep a note of the terms removed and the R^2 and Q^2 values.

First we must expand the input dataframe to include the higher order terms.

```
In [6]: sat_source_list = []
source_list = []
sat_inputs_orig, sat_source_list = doenum.add_higher_order_terms(
    inputs,
    add_squares=True,
    add_interactions=True,
    column_list=[])

```

Input array has columns ['Donor %', 'Conc.', 'Spin', 'Add.']

Adding square terms:

Donor %**2

Conc.**2

Spin**2

Add.**2

Adding interaction terms:

Donor %*Conc.

Donor %*Spin

Donor %*Add.

Conc.*Spin

Conc.*Add.

Spin*Add.

Full saturated quadratic model:

This contains all the main terms and all the square terms.

```
In [7]: input_selector = [0, 1, 2, 3,
                          4, 5, 6, 7]

```

```
scaled_model, R2, temp_tuple, _ =doenut.tune_model(
    sat_inputs_orig,
    responses,
    input_selector=input_selector,
    response_selector=[0]
)
```

```
new_model, predictions, ground_truth, coeffs, R2s, R2, Q2= temp_tuple
```

Input terms are ['Donor %', 'Conc.', 'Spin', 'Add.', 'Donor %**2', 'Conc.**2', 'Spin**2', 'Add.**2', 'Donor %*Conc.', 'Donor %*Spin', 'Donor %*Add.', 'Conc.*Spin', 'Conc.*Add.', 'Spin*Add.']

Input Responses are ['PCE']

Selected Response is PCE

Selected input terms: ['Donor %', 'Conc.', 'Spin', 'Add.', 'Donor %**2', 'Conc.**2', 'Spin**2', 'Add.**2']

Have found no replicates

Input data is 15 points long

We are using 15 data points

Left out data point 0:	R2 = 0.799	Ave. Error = -0.849
Left out data point 1:	R2 = 0.816	Ave. Error = 0.19
Left out data point 2:	R2 = 0.831	Ave. Error = 3.42
Left out data point 3:	R2 = 0.869	Ave. Error = -4.02
Left out data point 4:	R2 = 0.872	Ave. Error = 4.05
Left out data point 5:	R2 = 0.846	Ave. Error = -3.35
Left out data point 6:	R2 = 0.796	Ave. Error = 0.587
Left out data point 7:	R2 = 0.838	Ave. Error = 3.59
Left out data point 8:	R2 = 0.842	Ave. Error = -2.39
Left out data point 9:	R2 = 0.8	Ave. Error = 1.37
Left out data point 10:	R2 = 0.92	Ave. Error = -4.38
Left out data point 11:	R2 = 0.816	Ave. Error = -0.473
Left out data point 12:	R2 = 0.819	Ave. Error = 1.49
Left out data point 13:	R2 = 0.83	Ave. Error = 1.78
Left out data point 14:	R2 = 0.817	Ave. Error = -0.841

R2 overall is 0.815

Mean of test set: 2.885782

Mean being used: 2.885782

Sum of squares of the residuals (explained variance) is 102.55309297319681

Sum of squares total (total variance) is 87.23748999604001

Q2 is -0.176

Response PCE R2 is 0.816

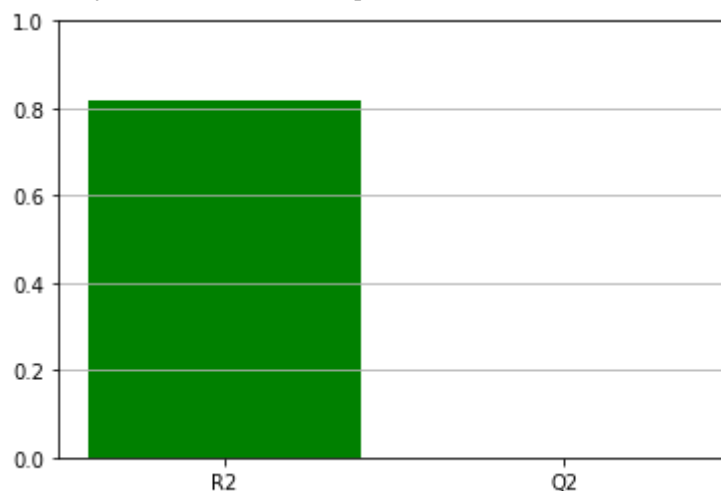
Input selector was [0, 1, 2, 3, 4, 5, 6, 7]

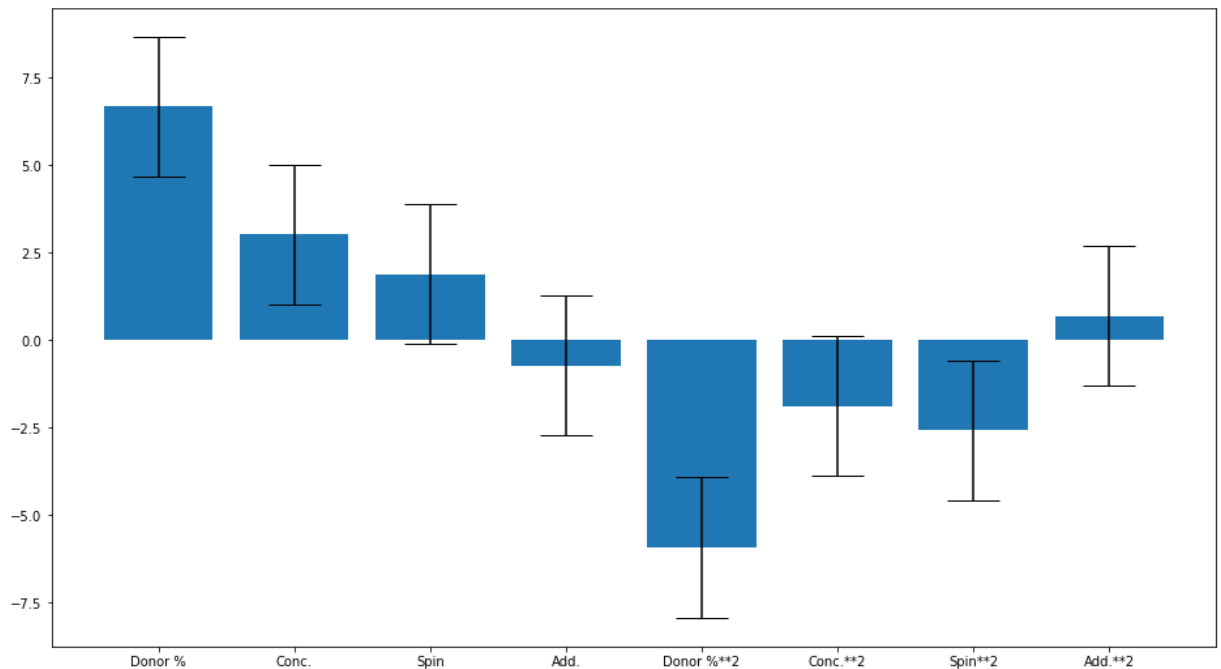
Input_selector was: [0, 1, 2, 3, 4, 5, 6, 7]

Average coefficients are: [6.68270726 3.02883412 1.87771844 -0.72988262 -5.93271432 -1.89506711

-2.58602063 0.69844306]

Coefficient labels are: ['Donor %', 'Conc.', 'Spin', 'Add.', 'Donor %**2', 'Conc.**2', 'Spin**2', 'Add.**2']





```
In [8]: input_selector = [0, 1, 2,
                          4, 5, 6]
scaled_model, R2, temp_tuple, _ = doanut.tune_model(
    sat_inputs_orig,
    responses,
    input_selector=input_selector,
    response_selector=[0]
)

new_model, predictions, ground_truth, coeffs, R2s, R2, Q2 = temp_tuple
```

Input terms are ['Donor %', 'Conc.', 'Spin', 'Add.', 'Donor %**2', 'Conc.**2', 'Spin**2', 'Add.**2', 'Donor %*Conc.', 'Donor %*Spin', 'Donor %*Add.', 'Conc.*Spin', 'Conc.*Add.', 'Spin*Add.']
 Input Responses are ['PCE']

Selected Response is PCE

Selected input terms: ['Donor %', 'Conc.', 'Spin', 'Donor %**2', 'Conc.**2', 'Spin**2']

Have found no replicates

Input data is 15 points long

We are using 15 data points

Left out data point 0: R2 = 0.797	Ave. Error = -0.906
Left out data point 1: R2 = 0.812	Ave. Error = -0.04
Left out data point 2: R2 = 0.812	Ave. Error = 1.92
Left out data point 3: R2 = 0.833	Ave. Error = -2.57
Left out data point 4: R2 = 0.855	Ave. Error = 3.23
Left out data point 5: R2 = 0.833	Ave. Error = -1.71
Left out data point 6: R2 = 0.792	Ave. Error = 0.229
Left out data point 7: R2 = 0.824	Ave. Error = 2.92
Left out data point 8: R2 = 0.824	Ave. Error = -1.39
Left out data point 9: R2 = 0.789	Ave. Error = 0.757
Left out data point 10: R2 = 0.919	Ave. Error = -4.4
Left out data point 11: R2 = 0.812	Ave. Error = -0.16
Left out data point 12: R2 = 0.814	Ave. Error = 1.33
Left out data point 13: R2 = 0.821	Ave. Error = 1.17
Left out data point 14: R2 = 0.812	Ave. Error = -0.467

R2 overall is 0.813

Mean of test set: 2.885782

Mean being used: 2.885782

Sum of squares of the residuals (explained variance) is 58.245326009154304

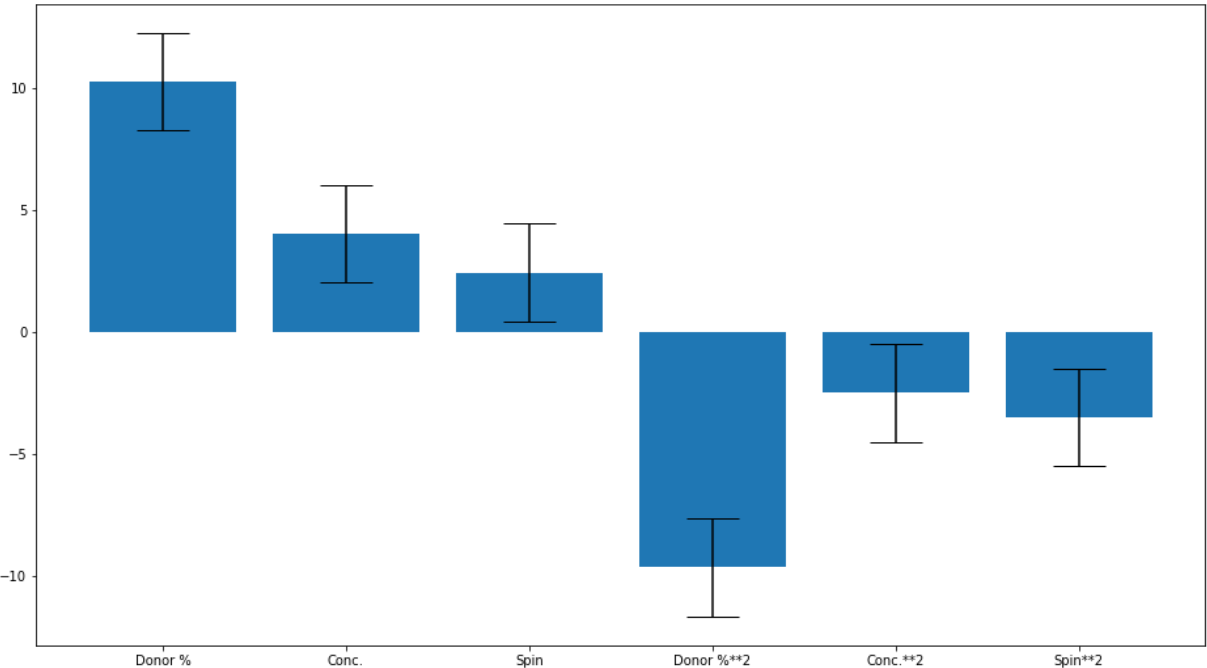
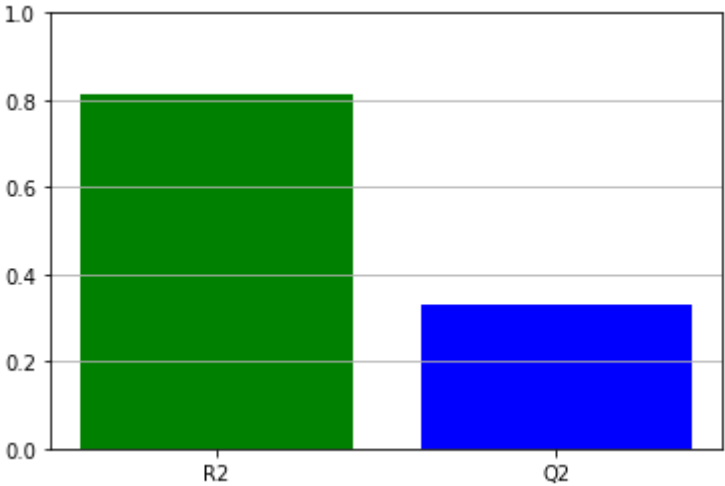
Sum of squares total (total variance) is 87.23748999604001

Q2 is 0.332

Response PCE R2 is 0.813

Input selector was [0, 1, 2, 4, 5, 6]


```
Input_selector was: [0, 1, 2, 3, 4, 5]
Average coefficients are: [10.24376537  4.00601457  2.41963174 -9.66908521 -2.524063
76 -3.51264286]
Coefficient labels are: ['Donor %', 'Conc.', 'Spin', 'Donor %**2', 'Conc.**2', 'Spin
**2']
```



15 datapoints so 14 DoF.

Starting from quadratic model

No. of terms	DoF	term removed	factor removed	R^2	Q^2
8	6			0.815	-0.176
7	7	8	'Add.**2	0.813	0.0863
6	8	3	Add.	0.813	0.332

this is the model with no statistically insignificant terms. It's heirarchical.

The Q2 is better than the main effects only model

Task 3. Create a parsimonious interaction model

Create hierarchical parsimonious interaction model. Starting with a interaction model, and making sure that all models are hierarchical, optimise the model by removing only the statistically insignificant terms. Keep a note of the terms removed and the Q^2 and R^2 values.

```
In [ ]: input_selector = [0, 1, 2, 3,
                        9, 10]

scaled_model, R2, temp_tuple, _ = doenut.tune_model(
    sat_inputs_orig,
    responses,
    input_selector=input_selector,
    response_selector=[0]
)

new_model, predictions, ground_truth, coeffs, R2s, R2, Q2 = temp_tuple
```

15 terms so 14 DoF

Starting from square model

No. of terms	DoF	term removed	factor removed	R^2	Q^2
10	4			0.811	-1.79
9	5	12	Conc.*Add.	0.813	0.0863
8	6	13	'Spin*Add.'	0.798	-0.555
7	7	11	Conc*Spin.	0.777	0.0133
6	8	8	Donor*Conc.	0.761	0.315

this is the model with no statistically insignificant terms. It's heirarchical.

The Q^2 is better than the main effects only model, but not as good as the square terms.

Task 4: Combine data from both experiments and train a parsimonious model

```
In [9]: df=pd.read_csv('solar_cells_2.csv')
df
```

```
Out[9]:
```

	experiment #	donor %	total concentration	spin speed	PCE	thickness	number of devices
0	NaN	wt %	mg/mL	rpm	%	nm	NaN
1	2-1	20	20	1500	6.32	73	5.0
2	2-2	27	20	1500	7.21	77	11.0
3	2-3	20	25	1500	6.83	126	6.0
4	2-4	27	25	1500	6.96	131	6.0
5	2-5	20	23	1000	7.77	109	4.0
6	2-6	27	23	1000	6.87	136	4.0
7	2-7	20	23	2000	6.43	76	8.0
8	2-8	27	23	2000	7.65	88	7.0

	experiment #	donor %	total concentration	spin speed	PCE	thickness	number of devices
9	2-9	25	20	1000	7.43	115	4.0
10	2-10	25	25	1000	6.88	135	8.0
11	2-11	25	20	2000	7.32	104	7.0
12	2-12	25	25	2000	7.21	126	8.0
13	2-13	25	23	1500	7.4	129	7.0

```
In [10]: inputs_2 = pd.DataFrame({
    'Donor %': [float(x) for x in df.iloc[1:-1,1]],
    'Conc.': [float(x) for x in df.iloc[1:-1,2]],
    'Spin': [float(x) for x in df.iloc[1:-1,3]]})
inputs_2
```

```
Out[10]:
```

	Donor %	Conc.	Spin
0	20.0	20.0	1500.0
1	27.0	20.0	1500.0
2	20.0	25.0	1500.0
3	27.0	25.0	1500.0
4	20.0	23.0	1000.0
5	27.0	23.0	1000.0
6	20.0	23.0	2000.0
7	27.0	23.0	2000.0
8	25.0	20.0	1000.0
9	25.0	25.0	1000.0
10	25.0	20.0	2000.0
11	25.0	25.0	2000.0

```
In [11]: responses_2 = pd.DataFrame({'PCE': [float(x) for x in df['PCE'][1:-1]]})
responses_2
```

```
Out[11]:
```

	PCE
0	6.32
1	7.21
2	6.83
3	6.96
4	7.77
5	6.87
6	6.43
7	7.65
8	7.43
9	6.88

PCE**10** 7.32**11** 7.21In [12]: `inputs[['Donor %', 'Conc.', 'Spin']]`

Out[12]:

	Donor %	Conc.	Spin
0	10.0	20.0	3000.0
1	10.0	25.0	1000.0
2	10.0	10.0	600.0
3	10.0	15.0	2000.0
4	25.0	20.0	600.0
5	25.0	15.0	1000.0
6	25.0	10.0	3000.0
7	25.0	25.0	2000.0
8	40.0	10.0	1000.0
9	40.0	20.0	2000.0
10	40.0	25.0	600.0
11	40.0	15.0	3000.0
12	55.0	10.0	2000.0
13	55.0	15.0	600.0
14	55.0	20.0	1000.0

In [13]: `new_inputs = pd.concat([inputs[['Donor %', 'Conc.', 'Spin']], inputs_2], axis=0)`
`new_responses = pd.concat([responses, responses_2], axis=0)`

In [14]: `new_inputs`

Out[14]:

	Donor %	Conc.	Spin
0	10.0	20.0	3000.0
1	10.0	25.0	1000.0
2	10.0	10.0	600.0
3	10.0	15.0	2000.0
4	25.0	20.0	600.0
5	25.0	15.0	1000.0
6	25.0	10.0	3000.0
7	25.0	25.0	2000.0
8	40.0	10.0	1000.0
9	40.0	20.0	2000.0
10	40.0	25.0	600.0

	Donor %	Conc.	Spin
11	40.0	15.0	3000.0
12	55.0	10.0	2000.0
13	55.0	15.0	600.0
14	55.0	20.0	1000.0
0	20.0	20.0	1500.0
1	27.0	20.0	1500.0
2	20.0	25.0	1500.0
3	27.0	25.0	1500.0
4	20.0	23.0	1000.0
5	27.0	23.0	1000.0
6	20.0	23.0	2000.0
7	27.0	23.0	2000.0
8	25.0	20.0	1000.0
9	25.0	25.0	1000.0
10	25.0	20.0	2000.0
11	25.0	25.0	2000.0

```
In [15]: sat_source_list = []
source_list = []
sat_inputs_2, sat_source_list = doenum.add_higher_order_terms(
    new_inputs,
    add_squares=True,
    add_interactions=True,
    column_list=[])

```

Input array has columns ['Donor %', 'Conc.', 'Spin']
Adding square terms:
Donor %**2
Conc.**2
Spin**2
Adding interaction terms:
Donor %*Conc.
Donor %*Spin
Conc.*Spin

Saturated model: 9 terms

```
In [16]: input_selector = [0, 1, 2,
                          3,4,5,
                          6,7,8]

scaled_model, R2, temp_tuple, _ =doenum.tune_model(
    sat_inputs_2,
    new_responses,
    input_selector=input_selector,
    response_selector=[0]
)

new_model, predictions, ground_truth, coeffs, R2s, R2, Q2= temp_tuple

```

Input terms are ['Donor %', 'Conc.', 'Spin', 'Donor %**2', 'Conc.**2', 'Spin**2', 'Donor %*Conc.', 'Donor %*Spin', 'Conc.*Spin']
 Input Responses are ['PCE']

Selected Response is PCE

Selected input terms: ['Donor %', 'Conc.', 'Spin', 'Donor %**2', 'Conc.**2', 'Spin**2', 'Donor %*Conc.', 'Donor %*Spin', 'Conc.*Spin']

Have found no replicates

Input data is 27 points long

We are using 27 data points

Left out data point 0:	R2 = 0.851	Ave. Error = -1.01
Left out data point 1:	R2 = 0.909	Ave. Error = -2.8
Left out data point 2:	R2 = 0.759	Ave. Error = 13.7
Left out data point 3:	R2 = 0.896	Ave. Error = -2.5
Left out data point 4:	R2 = 0.91	Ave. Error = 1.77
Left out data point 5:	R2 = 0.919	Ave. Error = -2.62
Left out data point 6:	R2 = 0.926	Ave. Error = 5.98
Left out data point 7:	R2 = 0.891	Ave. Error = 0.493
Left out data point 8:	R2 = 0.911	Ave. Error = -3.07
Left out data point 9:	R2 = 0.896	Ave. Error = -0.901
Left out data point 10:	R2 = 0.926	Ave. Error = -3.49
Left out data point 11:	R2 = 0.901	Ave. Error = -2.85
Left out data point 12:	R2 = 0.89	Ave. Error = 1.02
Left out data point 13:	R2 = 0.906	Ave. Error = 2.29
Left out data point 14:	R2 = 0.889	Ave. Error = 0.127
Left out data point 0:	R2 = 0.851	Ave. Error = 0.722
Left out data point 1:	R2 = 0.909	Ave. Error = 0.469
Left out data point 2:	R2 = 0.759	Ave. Error = 1.13
Left out data point 3:	R2 = 0.896	Ave. Error = -0.258
Left out data point 4:	R2 = 0.91	Ave. Error = 1.92
Left out data point 5:	R2 = 0.919	Ave. Error = -0.342
Left out data point 6:	R2 = 0.926	Ave. Error = -0.0549
Left out data point 7:	R2 = 0.891	Ave. Error = 0.741
Left out data point 8:	R2 = 0.911	Ave. Error = 0.793
Left out data point 9:	R2 = 0.896	Ave. Error = 0.156
Left out data point 10:	R2 = 0.926	Ave. Error = 1.35
Left out data point 11:	R2 = 0.901	Ave. Error = 0.614

R2 overall is 0.89

Mean of test set: 4.746915925925926

Mean being used: 4.746915925925926

Sum of squares of the residuals (explained variance) is 294.06908219945797

Sum of squares total (total variance) is 206.3168644580518

Q2 is -0.425

Response PCE R2 is 0.896

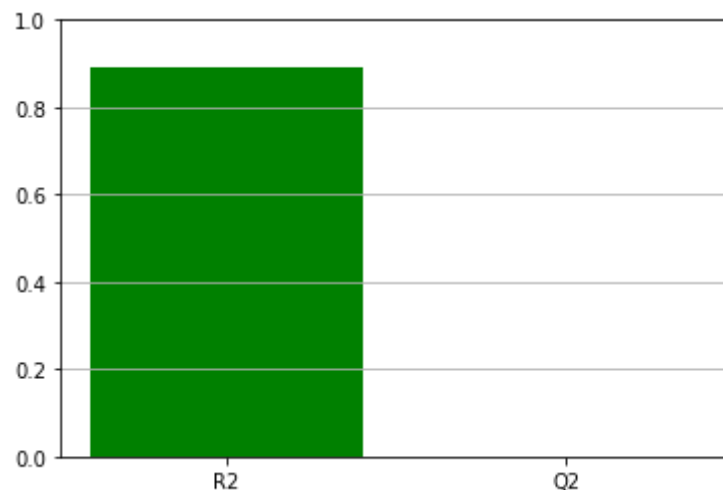
Input selector was [0, 1, 2, 3, 4, 5, 6, 7, 8]

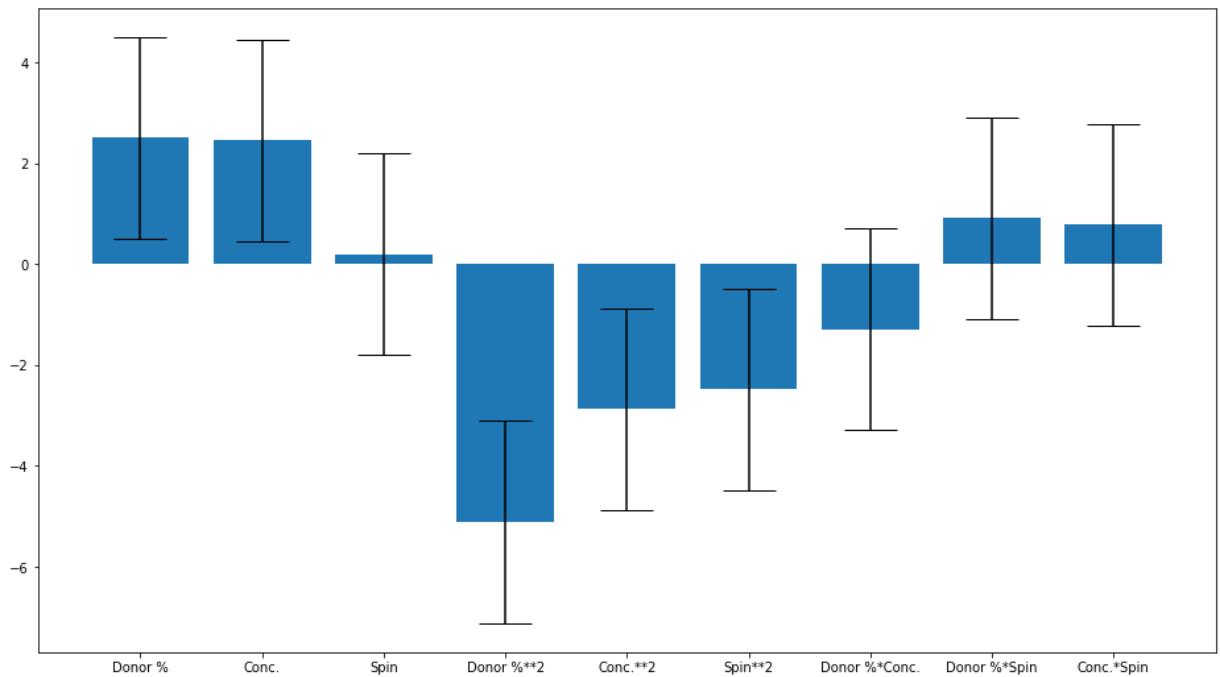
Input_selector was: [0, 1, 2, 3, 4, 5, 6, 7, 8]

Average coefficients are: [2.50008245 2.4484955 0.19604689 -5.11218527 -2.8778947 -2.48480065

-1.28965495 0.90939463 0.78220542]

Coefficient labels are: ['Donor %', 'Conc.', 'Spin', 'Donor %**2', 'Conc.**2', 'Spin**2', 'Donor %*Conc.', 'Donor %*Spin', 'Conc.*Spin']





Optimised parsimonious model

```
In [17]: input_selector = [0, 1, 2,
                          3, 4, 5]
```

```
scaled_model, R2, temp_tuple, _ = doanut.tune_model(
    sat_inputs_2,
    new_responses,
    input_selector=input_selector,
    response_selector=[0]
)
```

```
new_model, predictions, ground_truth, coeffs, R2s, R2, Q2 = temp_tuple
```

Input terms are ['Donor %', 'Conc.', 'Spin', 'Donor %**2', 'Conc.**2', 'Spin**2', 'Donor %*Conc.', 'Donor %*Spin', 'Conc.*Spin']
 Input Responses are ['PCE']

Selected Response is PCE

Selected input terms: ['Donor %', 'Conc.', 'Spin', 'Donor %**2', 'Conc.**2', 'Spin**2']

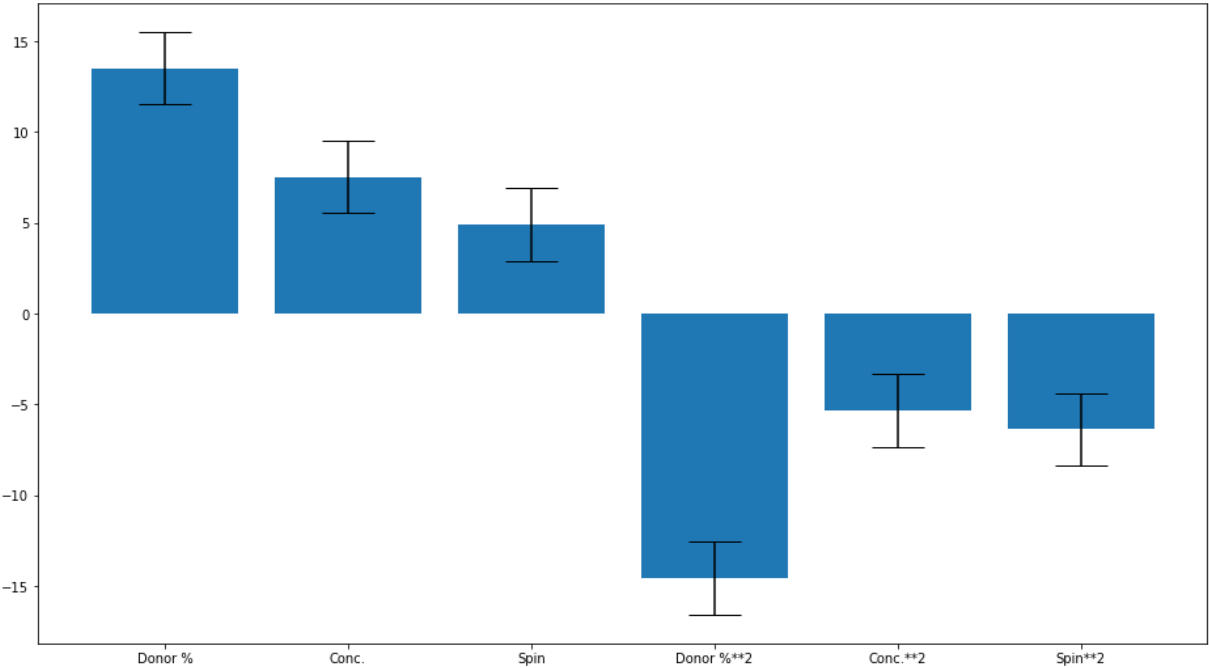
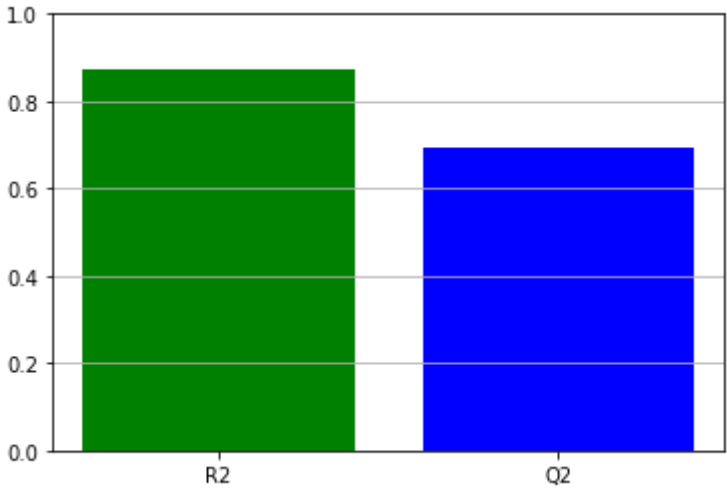
Have found no replicates

Input data is 27 points long

We are using 27 data points

Left out data point 0: R2 = 0.856	Ave. Error = -1.01
Left out data point 1: R2 = 0.869	Ave. Error = -0.921
Left out data point 2: R2 = 0.867	Ave. Error = 2.28
Left out data point 3: R2 = 0.889	Ave. Error = -3.22
Left out data point 4: R2 = 0.891	Ave. Error = 2.07
Left out data point 5: R2 = 0.892	Ave. Error = -2.53
Left out data point 6: R2 = 0.855	Ave. Error = 0.63
Left out data point 7: R2 = 0.865	Ave. Error = 0.597
Left out data point 8: R2 = 0.874	Ave. Error = -1.67
Left out data point 9: R2 = 0.869	Ave. Error = -0.689
Left out data point 10: R2 = 0.921	Ave. Error = -3.98
Left out data point 11: R2 = 0.864	Ave. Error = -0.207
Left out data point 12: R2 = 0.87	Ave. Error = 1.83
Left out data point 13: R2 = 0.879	Ave. Error = 1.75
Left out data point 14: R2 = 0.872	Ave. Error = -0.627
Left out data point 0: R2 = 0.856	Ave. Error = 0.137
Left out data point 1: R2 = 0.869	Ave. Error = 0.216
Left out data point 2: R2 = 0.867	Ave. Error = 0.46

Left out data point 3: $R^2 = 0.889$ Ave. Error = -0.326
Left out data point 4: $R^2 = 0.891$ Ave. Error = 1.96
Left out data point 5: $R^2 = 0.892$ Ave. Error = -0.415
Left out data point 6: $R^2 = 0.855$ Ave. Error = 0.37
Left out data point 7: $R^2 = 0.865$ Ave. Error = 0.799
Left out data point 8: $R^2 = 0.874$ Ave. Error = 0.909
Left out data point 9: $R^2 = 0.869$ Ave. Error = 0.0224
Left out data point 10: $R^2 = 0.921$ Ave. Error = 1.37
Left out data point 11: $R^2 = 0.864$ Ave. Error = 0.499
 R^2 overall is 0.871
Mean of test set: 4.746915925925926
Mean being used: 4.746915925925926
Sum of squares of the residuals (explained variance) is 62.93998023442563
Sum of squares total (total variance) is 206.3168644580518
 Q^2 is 0.695
Response PCE R^2 is 0.871
Input_selector was [0, 1, 2, 3, 4, 5]
Input_selector was: [0, 1, 2, 3, 4, 5]
Average coefficients are: [13.51406635 7.52612545 4.90682899 -14.54539532 -5.32766778
 -6.36236656]
Coefficient labels are: ['Donor %', 'Conc.', 'Spin', 'Donor %**2', 'Conc.**2', 'Spin**2']

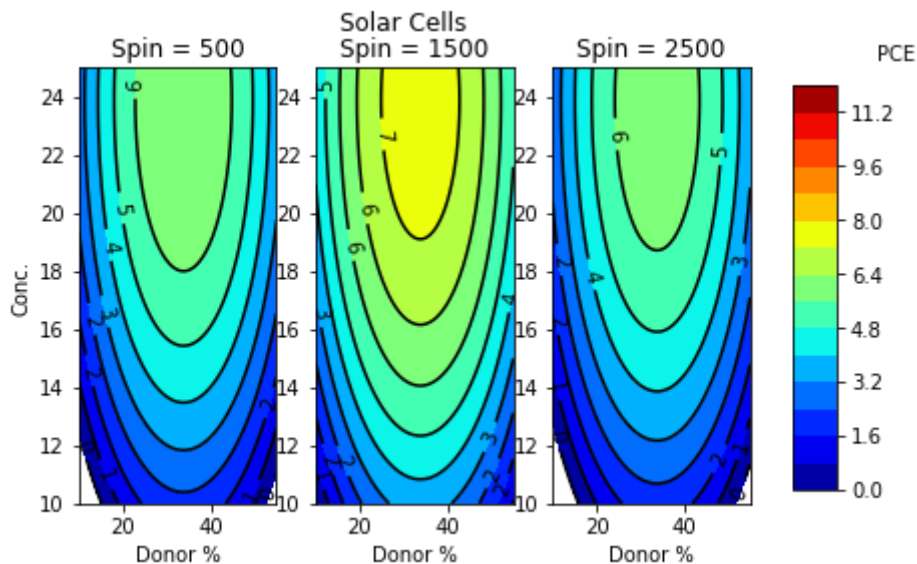


```
In [18]: 27-1-9
Out[18]: 17
```


In []:

Task 5. Method 1: Plot a 4-D contour plot and read the values off:

<Figure size 1440x864 with 0 Axes>



Task 5. Method 2. Run the model on the input values:

```
In [20]: question_5p2=pd.DataFrame({'A':{'Donor %': 20, 'Conc.': 12, 'Spin': 500},
    'B':{'Donor %': 40, 'Conc.': 16, 'Spin': 1500},
    'C':{'Donor %': 35, 'Conc.': 22, 'Spin': 1500},
    'D':{'Donor %': 45, 'Conc.': 18, 'Spin': 2500},
    'E':{'Donor %': 20, 'Conc.': 17, 'Spin': 2500}}).T
```

```
In [21]: question_5p2
```

```
Out[21]:
```

	Donor %	Conc.	Spin
A	20	12	500
B	40	16	1500
C	35	22	1500
D	45	18	2500
E	20	17	2500

```
In [22]: question_5p2.index
```

```
Out[22]: Index(['A', 'B', 'C', 'D', 'E'], dtype='object')
```

```
In [23]: sat_source_list = []
source_list = []
sat_inputs_q5, sat_source_list = doenut.add_higher_order_terms(
    question_5p2,
    add_squares=True,
    add_interactions=True,
    column_list=[],
    verbose=False)

results, _ =doenut.predict_from_model(
    scaled_model,
    sat_inputs_q5,
    input_selector)
letters = [x for x in question_5p2.index]
[print(f"{letters[i]}:\t{results[i]}") for i in range(len(letters))];
```

```
A: 2.105294443842542
B: 6.111167642320282
```

C: 7.610495288984239
D: 4.653001005905132
E: 3.969517548689133

Answer C is above 7.

In []:

In []:

In []: