

Factorial designs: principles and applications

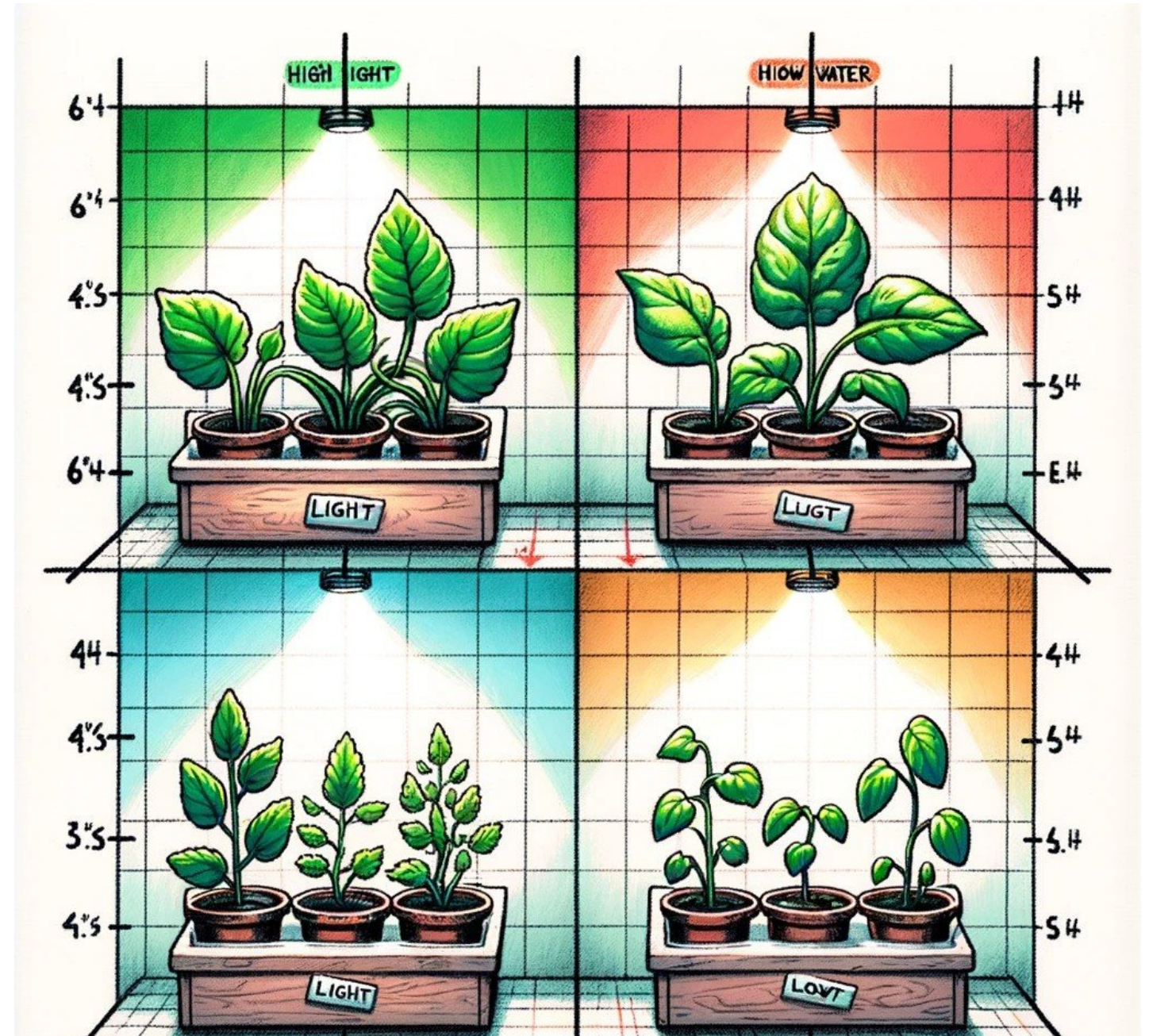
EXPERIMENTAL DESIGN IN PYTHON



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Understanding factorial design

- Study *multiple independent variables/factors* in one experiment
- Test *every combination* of factor levels
- Discover direct effects and *interactions* between factors



¹ Image Generated with DALL·E 3

Factorial design data example

- Factor 1 (`Light_Condition`) - two levels: `Full Sunlight` and `Partial Shade`
- Factor 2 (`Fertilizer_Type`) - two levels: `Synthetic` and `Organic`
- Numeric response/dependent/outcome variable: `Growth_cm`

```
plant_growth_data.head()
```

	<code>Plant_ID</code>	<code>Light_Condition</code>	<code>Fertilizer_Type</code>	<code>Growth_cm</code>
0	1	<code>Full Sunlight</code>	<code>Synthetic</code>	16.489735
1	2	<code>Partial Shade</code>	<code>Organic</code>	18.361689
2	3	<code>Full Sunlight</code>	<code>Synthetic</code>	18.039459
3	4	<code>Full Sunlight</code>	<code>Organic</code>	12.682425
4	5	<code>Full Sunlight</code>	<code>Organic</code>	21.480601

Organizing data to visualize interactions

```
plant_growth = pd.pivot_table(plant_growth_data,  
                               values='Growth_cm',  
                               index='Light_Condition',  
                               columns='Fertilizer_Type',  
                               aggfunc='mean')
```

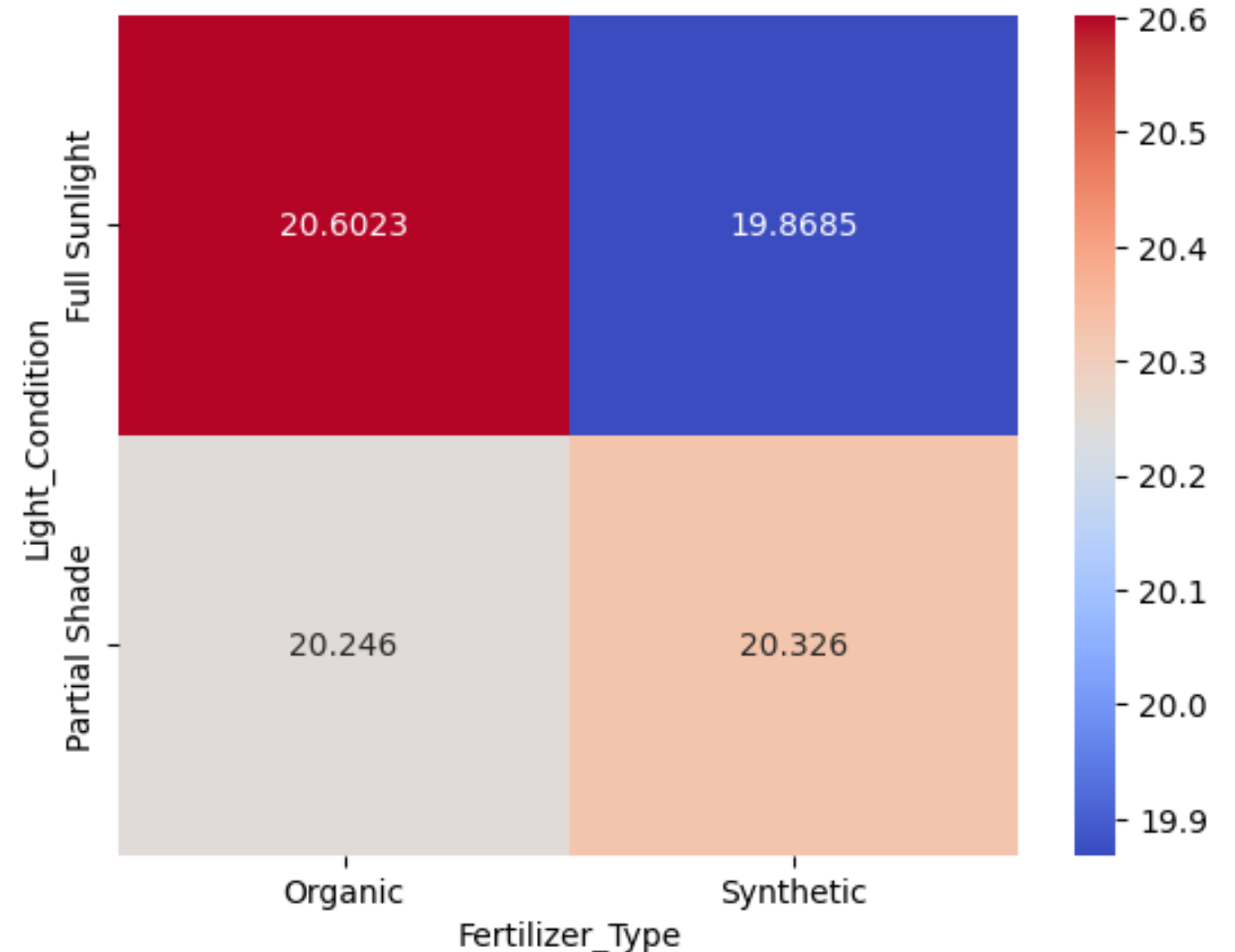
plant_growth

Light_Condition	Organic	Synthetic
Full Sunlight	20.602	19.869
Partial Shade	20.246	20.326

Visualize interactions with heatmap

```
import seaborn as sns
import matplotlib.pyplot as plt
sns.heatmap(plant_growth,
            annot=True,
            cmap='coolwarm',
            fmt='g')

plt.show()
```



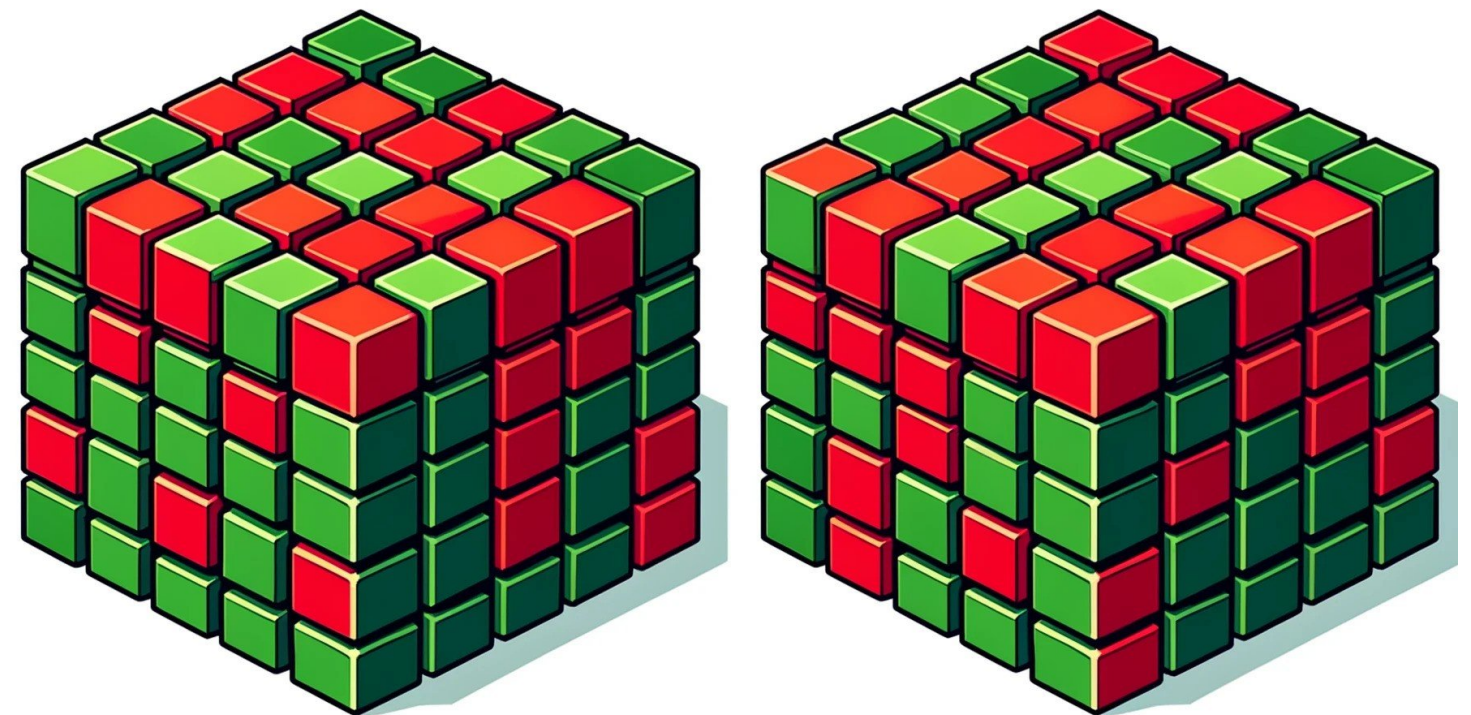
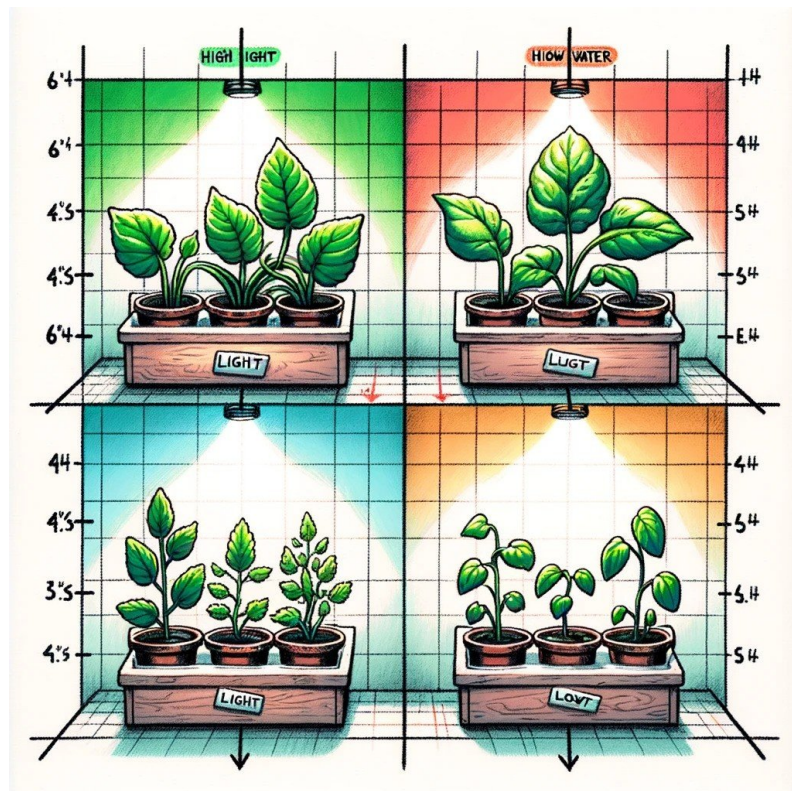
Interpreting interactions

Light_Condition	Organic	Synthetic
Full Sunlight	20.602	19.869
Partial Shade	20.246	20.326

- **Interactions:** how the effect of one factor varies with the level of another factor
- Significant interaction → *factors do not work independently*

Factorial designs vs. randomized block designs

- Multiple treatments and interactions
- Dissect complex multi-variable effects and interactions
- Can require more subjects
- Group similar subjects in randomized designs
- Control within-block variance
- Each treatment is tested within every block



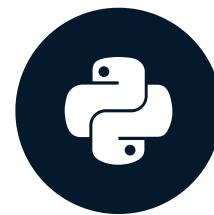
¹ Images Generated with DALL·E 3

Let's practice!

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Randomized block design: controlling variance

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Understanding blocking

- *Reduce variance* by grouping similar units
- Each block receives all treatments
- Focus on treatment effects, *controlling for block effects*

Block design data example

```
athletes.head()
```

	Athlete_ID	Initial_Fitness_Level	Muscle_Gain_kg
0	113	Beginner	3.225102
1	30	Advanced	3.976548
2	183	Intermediate	5.165449
3	200	Beginner	2.188297
4	194	Beginner	4.724162

Implementing randomized block design

- Use `.groupby()` to shuffle within blocks

```
blocks = athletes.groupby('Initial_Fitness_Level').apply(
    lambda x: x.sample(frac=1)
)
blocks = blocks.reset_index(drop=True)
blocks
```

	Athlete_ID	Initial_Fitness_Level	Muscle_Gain_kg
0	198	Advanced	5.742
1	146	Advanced	6.248
2	157	Advanced	6.049
..
198	164	Intermediate	6.134
199	178	Intermediate	6.591

Implemented randomized blocks

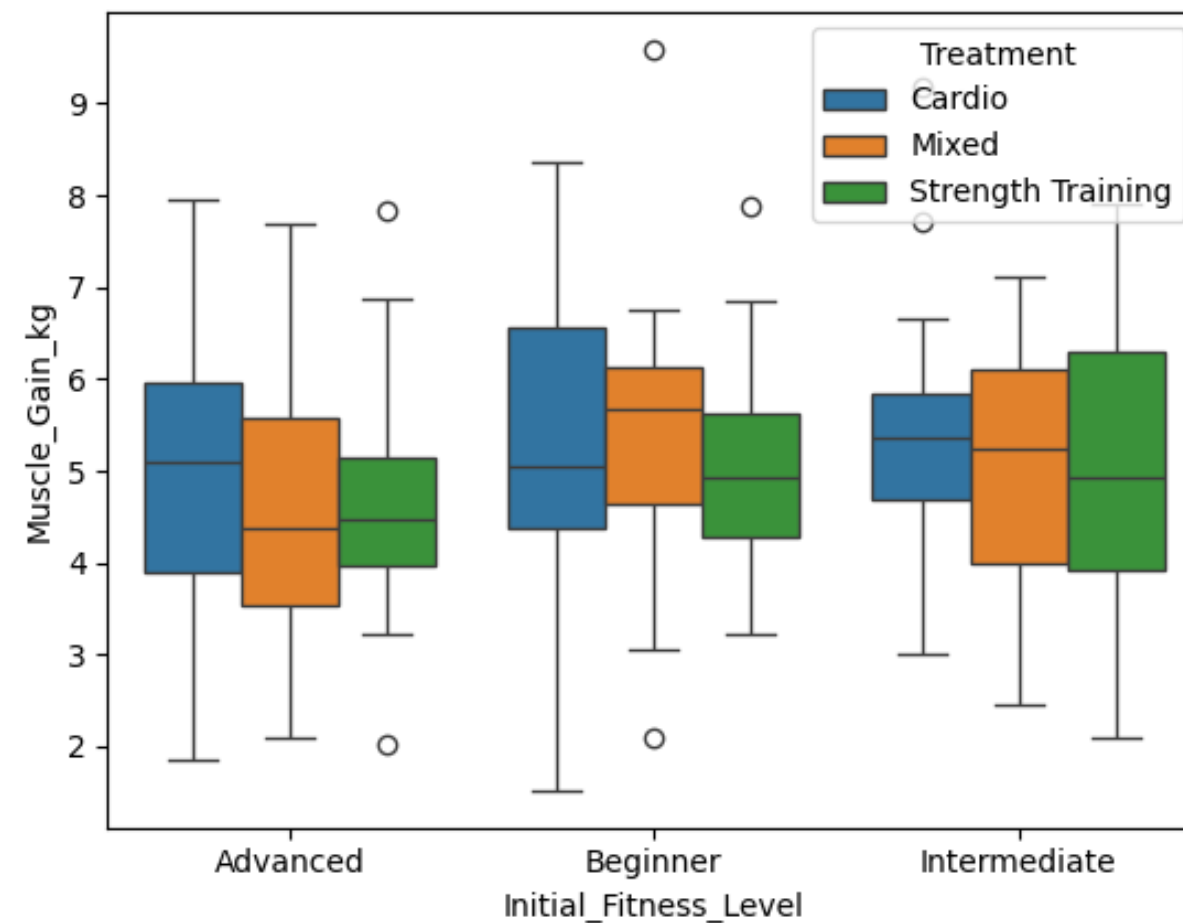
- `numpy.random.choice()` for random treatment assignment within blocks

```
blocks['Treatment'] = np.random.choice(
    ['Cardio', 'Strength Training', 'Mixed'],
    size=len(blocks))
blocks.sample(n=5)
```

	Athlete_ID	Initial_Fitness_Level	Muscle_Gain_kg	Treatment
87	194	Beginner	4.724	Cardio
54	3	Advanced	3.731	Strength Training
177	80	Intermediate	6.758	Mixed
146	183	Intermediate	5.165	Strength Training
60	190	Advanced	3.763	Cardio

Visualizing treatment effects within blocks

```
import seaborn as sns
sns.boxplot(x='Initial_Fitness_Level', y='Muscle_Gain_kg', hue='Treatment', data=blocks)
plt.show()
```



ANOVA within blocks

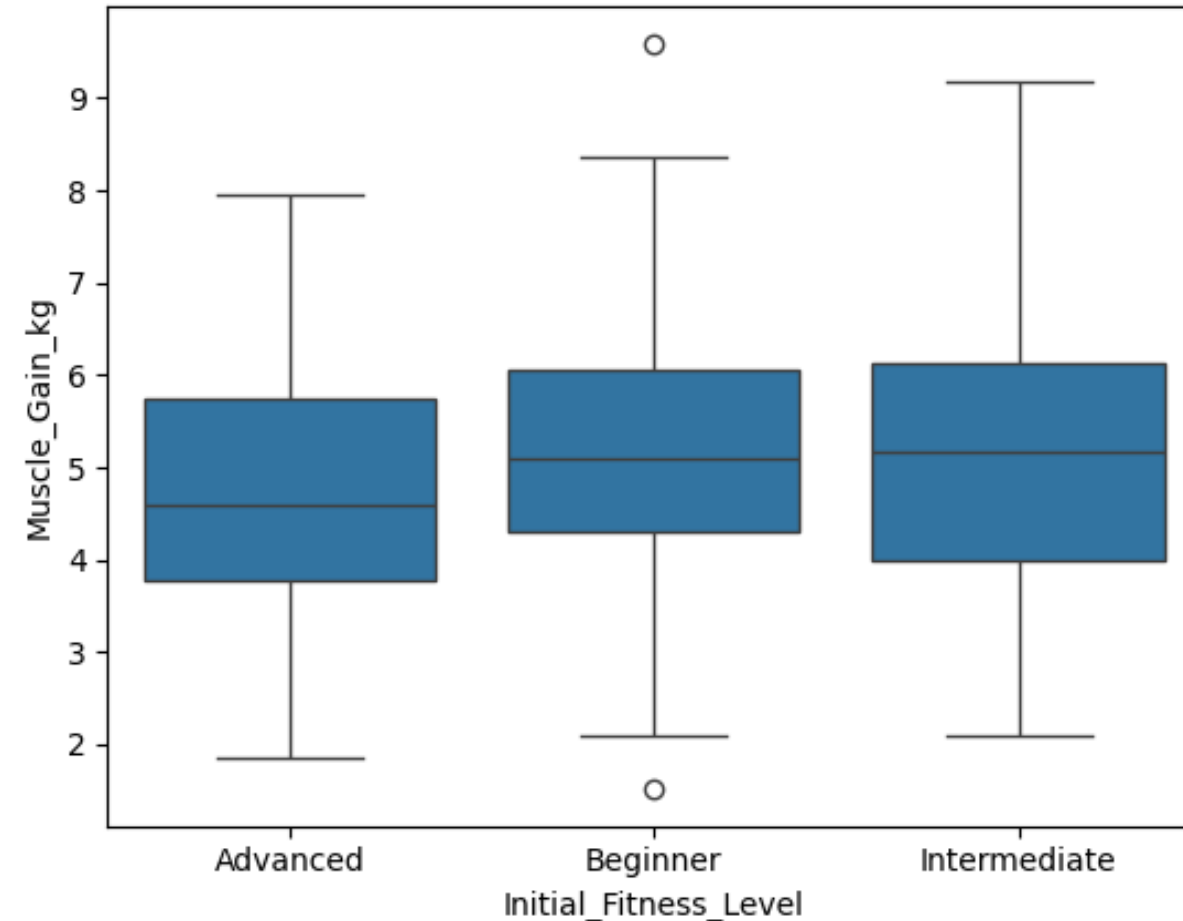
- Assume a significance level α of 0.05

```
from scipy.stats import f_oneway
blocks.groupby('Initial_Fitness_Level').apply(
    lambda x: f_oneway(x[x['Treatment'] == 'Cardio']['Muscle_Gain_kg'],
                        x[x['Treatment'] == 'Mixed']['Muscle_Gain_kg'],
                        x[x['Treatment'] == 'Strength Training']['Muscle_Gain_kg'])
)
```

```
Block
Initial_Fitness_Level
Advanced      (0.7951054385317405, 0.4555687666120679)
Beginner      (0.1085790370950905, 0.8972754969684291)
Intermediate  (0.5678877824942661, 0.5698403547950377)
dtype: object
```

Visualizing effects across blocks

```
import seaborn as sns
sns.boxplot(x='Initial_Fitness_Level', y='Muscle_Gain_kg', data=blocks)
plt.show()
```



ANOVA between blocks

```
f_oneway(  
    blocks[blocks['Initial_Fitness_Level'] == "Advanced"]['Muscle_Gain_kg'],  
    blocks[blocks['Initial_Fitness_Level'] == "Beginner"]['Muscle_Gain_kg'],  
    blocks[blocks['Initial_Fitness_Level'] == "Intermediate"]['Muscle_Gain_kg']  
)
```

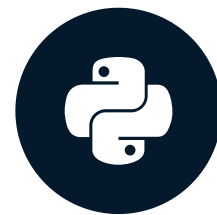
```
F_onewayResult(statistic=2.325058605244051, pvalue=0.10045536062209368)
```

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Covariate adjustment in experimental design

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Introduction to covariates

- **Covariates:** potentially affect experiment results but aren't primary focus
- Importance in reducing confounding
- Impact on precision and validity of results
- **Example:** Impact of teaching method on test scores

Does the **teaching method** impact **scores**?

Control Group



Prior Knowledge

Treatment Group



No Prior Knowledge

Prior Knowledge = Covariate

Experimental data example

```
exp_plant_data = plant_growth_data[['Plant_ID', 'Fertilizer_Type', 'Growth_cm']]
```

	Plant_ID	Light_Condition	Fertilizer_Type	Growth_cm
0	1	Full Sunlight	Synthetic	16.489735
1	2	Partial Shade	Organic	18.361689
2	3	Full Sunlight	Synthetic	18.039459
3	4	Full Sunlight	Organic	12.682425
4	5	Full Sunlight	Organic	21.480601

Covariate data example

```
covariate_data
```

	Plant_ID	Watering_Days_Per_Week
0	1	6
1	2	6
2	3	4
3	4	3
4	5	7

Combining experimental data with covariates

```
merged_plant_data = pd.merge(exp_plant_data, covariate_data, on='Plant_ID')
```

	Plant_ID	Fertilizer_Type	Growth_cm	Watering_Days_Per_Week
0	1	Synthetic	16.489735	6
1	2	Organic	18.361689	6
2	3	Synthetic	18.039459	4
3	4	Organic	12.682425	3
4	5	Organic	21.480601	7

Adjusting for covariates

```
from statsmodels.formula.api import ols
model = ols('Growth_cm ~ Fertilizer_Type + Watering_Days_Per_Week',
            data=merged_plant_data).fit()
model.summary()
```

OLS Regression Results

```
=====
Dep. Variable:          Growth_cm    R-squared:                0.011
Model:                  OLS          Adj. R-squared:           -0.006
Method:                 Least Squares    F-statistic:             0.6370
No. Observations:       120          Prob (F-statistic):       0.531 <---
Df Residuals:           117          Log-Likelihood:         -360.45
Df Model:                2           AIC:                      726.9
Covariance Type:        nonrobust      BIC:                      735.3
=====
```

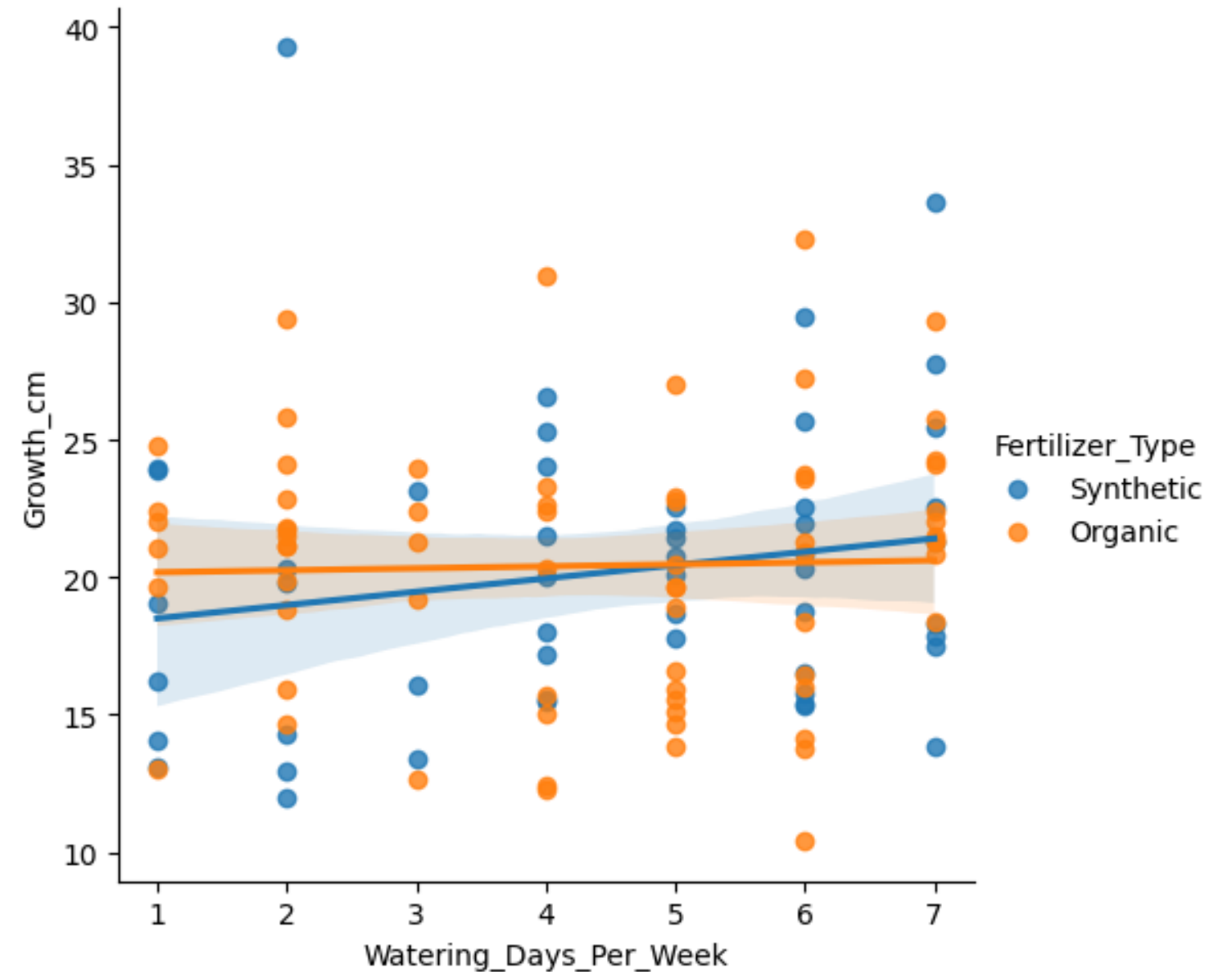

Further exploring ANCOVA results

	coef	std err	t	P> t	[0.025	0.975]
<hr/>						
Intercept	19.3373	1.150	16.820	0.000	17.060	21.614
Fertilizer_Type[T.Synthetic]	-0.2796	0.913	-0.306	0.760 <--	-2.088	1.528
Watering_Days_Per_Week	0.2507	0.229	1.097	0.275 <--	-0.202	0.703
=====						
Omnibus:	14.446	Durbin-Watson:	1.992			
Prob(Omnibus):	0.001	Jarque-Bera (JB):	18.267			
Skew:	0.675	Prob(JB):	0.000108			
Kurtosis:	4.352	Cond. No.	13.3			
=====						

Visualizing treatment effects with covariate adjustment

```
import seaborn as sns
import matplotlib.pyplot as plt
sns.lmplot(x='Watering_Days_Per_Week',
           y='Growth_cm',
           hue='Fertilizer_Type',
           data=merged_plant_data)

plt.show()
```



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