

Untitled

September 13, 2021

```
[14]: import pandas as pd
      from pandasql import sqldf
```

1 Methods

This was an investigation to determine the growth (weight gain) of coconut rhinoceros beetle (CRB) larva fed four different diets. Initial set up used CRB eggs placed in small vials, allowed to hatch and monitored the growth of the newly emerged larvae. The results from the study showed that the above two diets were the most ideal for larvae growth (Figs.1, 2). The other two “diets” (1) sphagnum peat moss (SM) and (2) “pure” (PF) frass pellets (obtained by manually separating out some of the frass pellets found in the FCT field sample) and then pulverized to form a powder were both poor diets for CRB larvae. Many of the larvae fed PF only survived from 2-3 weeks with little or no gain in weight, while the SM diet was even less supportive of larva growth dying within a week. Moreover, though both CM and FCT out performed PF and SM, CM showed higher larva mortality near the end of the study than FCT. Larva fed FCT also appeared to gain weight faster than CM but after 9 weeks larva fed CM appeared to have a higher overall increase in mean weight.

2 Data Preparation

2.1 Read Initial Spreadsheet

```
[15]: df = pd.read_excel('Diet study Means.xlsx')
      df
```

```
[15]:
```

	Treatment	Larvae ID	Hatch Date	Date	Weight (mg)	\
0	FCT 5/11/2021	1	5/24/2021 L	2021-05-24 00:00:00	203	
1	FCT 5/11/2021	2	5/24/2021 L	2021-05-24 00:00:00	114	
2	FCT 5/11/2021	3	5/24/2021 L	2021-05-24 00:00:00	256	
3	FCT 5/11/2021	4	5/24/2021 L	2021-05-24 00:00:00	145	
4	FCT 5/11/2021	5	5/24/2021 L	2021-05-24 00:00:00	236	
..	
168	CM 5/17/2021	4	5/25/2021 L	2021-05-25 00:00:00	85	
169	CM 5/17/2021	5	EMPTY	EMPTY	EMPTY	
170	CM 5/17/2021	6	5/25/2021 L	2021-05-25 00:00:00	152	
171	CM 5/17/2021	7	5/25/2021 L	2021-05-25 00:00:00	192	
172	CM 5/17/2021	8	5/25/2021 L	2021-05-25 00:00:00	138	

	Hatch Date.1	Date.1	Weight (mg).1	Hatch Date.2	\
0	6/2/2021 L	2021-06-02 00:00:00	420	6/9/2021 L	
1	6/2/2021 L	2021-06-02 00:00:00	381	6/9/2021 L	
2	6/2/2021 L	2021-06-02 00:00:00	385	6/9/2021 L	
3	6/2/2021 L	2021-06-02 00:00:00	424	6/9/2021 L	
4	6/2/2021 L	2021-06-02 00:00:00	425	6/9/2021 L	
..	
168	6/1/2021 D	2021-06-01 00:00:00	94	DEAD	
169	EMPTY	EMPTY	EMPTY	EMPTY	
170	6/1/2021 L	2021-06-01 00:00:00	284	6/8/2021 D	
171	6/1/2021 L	2021-06-01 00:00:00	210	6/8/2021 L	
172	6/1/2021 D	2021-06-01 00:00:00	211	DEAD	

	Date.2	...	Weight (mg).12	Hatch Date.13	Date.13	\
0	2021-06-09 00:00:00	...	NaN	NaN	NaN	
1	2021-06-09 00:00:00	...	NaN	NaN	NaN	
2	2021-06-09 00:00:00	...	NaN	NaN	NaN	
3	2021-06-09 00:00:00	...	NaN	NaN	NaN	
4	2021-06-09 00:00:00	...	NaN	NaN	NaN	
..	
168	DEAD	...	NaN	NaN	NaN	
169	EMPTY	...	NaN	NaN	NaN	
170	2021-06-08 00:00:00	...	NaN	NaN	NaN	
171	2021-06-08 00:00:00	...	NaN	NaN	NaN	
172	DEAD	...	NaN	NaN	NaN	

	Weight (mg).13	Hatch Date.14	Date.14	Weight (mg).14	Hatch Date.15	Date.15	\
0	NaN	NaN	NaN	NaN	NaN	NaN	
1	NaN	NaN	NaN	NaN	NaN	NaN	
2	NaN	NaN	NaN	NaN	NaN	NaN	
3	NaN	NaN	NaN	NaN	NaN	NaN	
4	NaN	NaN	NaN	NaN	NaN	NaN	
..	
168	NaN	NaN	NaN	NaN	NaN	NaN	
169	NaN	NaN	NaN	NaN	NaN	NaN	
170	NaN	NaN	NaN	NaN	NaN	NaN	
171	NaN	NaN	NaN	NaN	NaN	NaN	
172	NaN	NaN	NaN	NaN	NaN	NaN	

	Weight (mg).15
0	NaN
1	NaN
2	NaN
3	NaN
4	NaN
..	...
168	NaN

```

169         NaN
170         NaN
171         NaN
172         NaN

```

```
[173 rows x 50 columns]
```

2.2 Delete Unnecessary Columns

```
[16]: # Remove columns with "Date" heading because the data are redundant
```

```

cols = df.columns
for col in cols:
    if col.startswith('Date'):
        df.drop(labels=col, axis='columns', inplace=True)

# Drop all columns which do not contain data

df.dropna(how='all', axis='columns', inplace=True)

df

```

```

[16]:      Treatment  Larvae ID  Hatch Date  Weight (mg)  Hatch Date.1 \
0      FCT 5/11/2021         1  5/24/2021 L         203  6/2/2021 L
1      FCT 5/11/2021         2  5/24/2021 L         114  6/2/2021 L
2      FCT 5/11/2021         3  5/24/2021 L         256  6/2/2021 L
3      FCT 5/11/2021         4  5/24/2021 L         145  6/2/2021 L
4      FCT 5/11/2021         5  5/24/2021 L         236  6/2/2021 L
..      ...
168     CM 5/17/2021         4  5/25/2021 L          85  6/1/2021 D
169     CM 5/17/2021         5      EMPTY      EMPTY      EMPTY
170     CM 5/17/2021         6  5/25/2021 L         152  6/1/2021 L
171     CM 5/17/2021         7  5/25/2021 L         192  6/1/2021 L
172     CM 5/17/2021         8  5/25/2021 L         138  6/1/2021 D

      Weight (mg).1  Hatch Date.2  Weight (mg).2  Hatch Date.3  Weight (mg).3  ... \
0          420  6/9/2021 L          686  6/15/2021 L          1494  ...
1          381  6/9/2021 L          390  6/15/2021 L          1102  ...
2          385  6/9/2021 L          768  6/15/2021 L          1460  ...
3          424  6/9/2021 L          548  6/15/2021 L          1210  ...
4          425  6/9/2021 L          778  6/15/2021 L          1442  ...
..      ...
168          94      DEAD      DEAD      DEAD      DEAD  ...
169      EMPTY      EMPTY      EMPTY      EMPTY      EMPTY  ...
170          284  6/8/2021 D          334      DEAD      DEAD  ...
171          210  6/8/2021 L          188  6/15/2021 D          34  ...
172          211      DEAD      DEAD      DEAD      DEAD  ...

```

	Hatch Date.7	Weight (mg).7	Hatch Date.8	Weight (mg).8	Hatch Date.9	\
0	7/14/2021	L 2834	7/20/2021	L 3084	7/28/2021	L
1	7/14/2021	L 1969	7/20/2021	L 1924	7/28/2021	L
2	7/14/2021	L 3006	7/20/2021	L 3458	7/28/2021	L
3	7/14/2021	L 3163	7/20/2021	L 3467	7/28/2021	L
4	7/14/2021	L 2401	7/20/2021	L 2702	7/28/2021	L
..
168	DEAD	DEAD	DEAD	DEAD	DEAD	DEAD
169	EMPTY	EMPTY	EMPTY	EMPTY	EMPTY	EMPTY
170	DEAD	DEAD	DEAD	DEAD	DEAD	DEAD
171	DEAD	DEAD	DEAD	DEAD	DEAD	DEAD
172	DEAD	DEAD	DEAD	DEAD	DEAD	DEAD

	Weight (mg).9	Hatch Date.10	Weight (mg).10	Hatch Date.11	Weight (mg).11
0	3454	8/4/2021	L 5260	8/10/2021	L 5613
1	1715	8/4/2021	L 2594	8/10/2021	L 3463
2	3820	8/4/2021	L 5077	8/10/2021	L 5943
3	3347	8/4/2021	L 5320	8/10/2021	L 6045
4	2688	8/4/2021	L 4542	8/10/2021	L 5572
..
168	DEAD	DEAD	DEAD	DEAD	DEAD
169	EMPTY	EMPTY	EMPTY	EMPTY	EMPTY
170	DEAD	DEAD	DEAD	DEAD	DEAD
171	DEAD	DEAD	DEAD	DEAD	DEAD
172	DEAD	DEAD	DEAD	DEAD	DEAD

[173 rows x 26 columns]

2.3 Normalize Data

```
[17]: df.columns
```

```
[17]: Index(['Treatment', 'Larvae ID', 'Hatch Date', 'Weight (mg)', 'Hatch Date.1',
        'Weight (mg).1', 'Hatch Date.2', 'Weight (mg).2', 'Hatch Date.3',
        'Weight (mg).3', 'Hatch Date.4', 'Weight (mg).4', 'Hatch Date.5',
        'Weight (mg).5', 'Hatch Date.6', 'Weight (mg).6', 'Hatch Date.7',
        'Weight (mg).7', 'Hatch Date.8', 'Weight (mg).8', 'Hatch Date.9',
        'Weight (mg).9', 'Hatch Date.10', 'Weight (mg).10', 'Hatch Date.11',
        'Weight (mg).11'],
        dtype='object')
```

```
[18]: # Create a new dataframe "data" for first "Hatch Date", "Weight (mg)" columns

mylist = ['Treatment', 'Larvae ID', 'Hatch Date', 'Weight (mg)']
data = df[mylist]
data.columns = ['treatment', 'larva_id', 'hatch_date', 'mass']
```

```
# Append rows containing values from subsequent "Hatch Date.*", "Weight (mg).*"
↳ columns

for i in range(1,12):
    mylist = ['Treatment', 'Larvae ID', f'Hatch Date.{i}', f'Weight (mg).{i}']
    df_temp = df[mylist]
    df_temp.columns = ['treatment', 'larva_id', 'hatch_date', 'mass']
    data = data.append(df_temp, ignore_index=True)

data
```

```
[18]:
```

	treatment	larva_id	hatch_date	mass
0	FCT 5/11/2021	1	5/24/2021 L	203
1	FCT 5/11/2021	2	5/24/2021 L	114
2	FCT 5/11/2021	3	5/24/2021 L	256
3	FCT 5/11/2021	4	5/24/2021 L	145
4	FCT 5/11/2021	5	5/24/2021 L	236
...
2071	CM 5/17/2021	4	DEAD	DEAD
2072	CM 5/17/2021	5	EMPTY	EMPTY
2073	CM 5/17/2021	6	DEAD	DEAD
2074	CM 5/17/2021	7	DEAD	DEAD
2075	CM 5/17/2021	8	DEAD	DEAD

[2076 rows x 4 columns]

```
[19]: # Split hatch_date column

data['observation_date'] = data.apply(lambda x : x['hatch_date'].split(" ")[0],
↳axis='columns')
data['status'] = data.apply(lambda x : x['hatch_date'][-1], axis='columns')
data.drop('hatch_date', axis='columns', inplace=True)
data['status'].replace({'L':'larva', 'D':'dead', 'Y':'empty', 'G':'egg'},
↳inplace=True)

# Recode 'status'

data['unique_larva_id'] = data.apply(lambda x: f"{x['treatment']}_
↳{x['larva_id']}", axis='columns')
#data['unique_larva_id'] = data.apply(lambda x: x['treatment'].replace(' ', '_
↳'), axis='columns')
data.drop('larva_id', axis='columns', inplace=True)

# Recode 'treatment'
```

```

data['treatment'] = data.apply(lambda x: x['treatment'].split(' ')[0],
    ↪axis='columns')
data['treatment'].replace({'FCT':'frass', 'CM':'manure', 'SM':'peat moss', 'PF':
    ↪'fecal pellets'}, inplace=True)

# Add a new record for setup date of each insect

new_rows = []
for unique_larva_id in data.unique_larva_id.unique():
    new_rows.append({'unique_larva_id':unique_larva_id,
        'observation_date': unique_larva_id.split(' ')[1],
        'treatment': unique_larva_id.split(' ')[0],
        'status':'egg', 'mass':10})
data = data.append(new_rows, ignore_index=True, sort=False)
data['treatment'].replace({'FCT':'frass', 'CM':'manure', 'SM':'peat moss', 'PF':
    ↪'fecal pellets'}, inplace=True)

# Delete rows where mass equals EMPTY or DEAD

data = data[data['mass'] != 'EMPTY']
data = data[data['mass'] != 'DEAD']

# sort

data = data.sort_values(by=['unique_larva_id', 'observation_date'])
data = data[['unique_larva_id','treatment','observation_date','status','mass']]

# write CSV filr

data.to_csv('data.csv', index=False)

data

```

```

[19]:
   unique_larva_id  treatment  observation_date  status  mass
2203    CM 5/11/2021 1    manure      5/11/2021    egg    10
127     CM 5/11/2021 1    manure      5/25/2021  larva   186
300     CM 5/11/2021 1    manure      6/1/2021   larva   381
646     CM 5/11/2021 1    manure      6/15/2021  larva  1307
819     CM 5/11/2021 1    manure      6/22/2021  larva  1548
...
84      SM 5/17/2021 8    peat moss    5/25/2021  larva    32
257     SM 5/17/2021 8    peat moss      6/2/2021  dead    37
2161    SM 5/17/2021 9    peat moss    5/17/2021    egg    10
85      SM 5/17/2021 9    peat moss    5/25/2021  larva    45
258     SM 5/17/2021 9    peat moss      6/2/2021  dead    21

```

[1200 rows x 5 columns]

3 Analysis

```
[20]: # get the last observation record for each larva
```

```
df_endpoint = data.groupby('unique_larva_id').tail(1)
df_endpoint.sort_values(['status', 'unique_larva_id'])
```

```
[20]:
```

	unique_larva_id	treatment	observation_date	status	mass
1166	CM 5/11/2021 2	manure	7/6/2021	dead	43
480	CM 5/11/2021 8	manure	6/8/2021	dead	165
481	CM 5/12/2021 1	manure	6/8/2021	dead	341
317	CM 5/12/2021 10	manure	6/1/2021	dead	172
311	CM 5/12/2021 4	manure	6/1/2021	dead	191
...
1766	FCT 5/17/2021 5	frass	8/4/2021	larva	4443
1767	FCT 5/17/2021 6	frass	8/4/2021	larva	4800
1769	FCT 5/17/2021 8	frass	8/4/2021	larva	5142
469	PF 5/17/2021 4	fecal pellets	6/9/2021	larva	141
470	PF 5/17/2021 5	fecal pellets	6/9/2021	larva	201

[173 rows x 5 columns]

```
[21]: # Summarize the status variable values for each treatment
```

```
# egg: egg did not hatch
# dead: larva died before end of experiment
# larva: larva was alive at end of experiment

q = """
SELECT treatment, status, COUNT(*) AS N
FROM df_endpoint
GROUP BY treatment, status
ORDER BY treatment, status;
"""
sqldf(q, globals())
```

```
[21]:
```

	treatment	status	N
0	fecal pellets	dead	30
1	fecal pellets	egg	9
2	fecal pellets	larva	2
3	frass	egg	10
4	frass	larva	31
5	manure	dead	10
6	manure	egg	11

```

7      manure  larva  25
8      peat moss  dead  23
9      peat moss  egg  22

```

[22]: *# Calculate mean weight for surviving larvae*

```

q = """
SELECT treatment, COUNT(*) AS N, AVG(mass) AS mean, MIN(mass) as min, MAX(mass) AS max
FROM df_endpoint
WHERE status == 'larva'
GROUP BY treatment;
"""
sqldf(q, globals())

```

```

[22]:      treatment    N      mean    min    max
0 fecal pellets    2  171.000000  141    201
1          frass   31  4832.612903 1965   7592
2          manure  25  2549.040000  108   7335

```

3.1 Treatment effect on egg hatch

treatment	hatched	unhatched	proportion hatched
frass	31	10	0.76
manure	35	11	0.76
fecal pellets	32	9	0.78
peat moss	23	22	0.51

3.2 Treatment effect on larval survival

treatment	alive	dead	proportion surviving
frass	31	0	1.00
manure	25	10	0.71
fecal pellets	2	30	0.06
peat moss	0	23	0.00

3.3 Treatment effect on larval mass at end of experiment

treatment	N	mean {mg}	range (mg)
frass	31	4833	1965-7592
manure	25	2549	108-7335
fecal pellets	2	171	141-201
peat moss	0	-	-