

# General\_Results

August 16, 2017

```
In [1]: from IPython.display import HTML
HTML('''<script>
code_show=true;
function code_toggle() {
  if (code_show){
    $('div.input').hide();
  } else {
    $('div.input').show();
  }
  code_show = !code_show
}
$( document ).ready(code_toggle);
</script>''')
```

```
Out[1]: <IPython.core.display.HTML object>
```

```
In [208]: HTML('''
<style>
    .yourDiv {position: fixed;top: 100px; right: 0px;
              background: white;
              height: 100%;
              width: 175px;
              padding: 10px;
              z-index: 10000}

</style>
<script>
function showthis(url) {
    window.open(url, "pres",
        "toolbar=yes,scrollbars=yes,resizable=yes,top=10,left=400,width=500,height=400");
    return(false);
}
</script>

<div class=yourDiv>
    <h4>MENU</h4><br>
    <a href=#Data>1. Data</a><br>
    <a href=#FoodFeedOtherAcross>2. Across Farm Sizes</a><br>
    <a href=#FoodFeedOtherWithin>3. Within Farm Sizes</a><br><br>
''')
```

```

    <a href=#Top>Top</a><br>
    <a href="javascript:code_toggle()">Toggle Code On/Off</a><br>
    <a href=#LeftOff>Left Off Here</a><br>
    <a href='https://vinnyricciardi.github.io/farmsize_site/'>Site Index</a><br>
</div>
'''
)

```

Out[208]: <IPython.core.display.HTML object>

## General Results

This page overviews our general results:

- The main findings are that in our sample of 58 countries, farms < 2 ha produce 1% of the total food supply, while farms < 200 ha produce 72% of the total food supply. The largest differences (in terms of effect size) in the amount of food produced were between the very large farm and the smallest farm categories in terms of how much food each produced.
- This study also measures what percentage of each farm size class's crop production goes towards food, feed, processing, seed, waste, or other. The farm size category with the largest percentage of food produced compared to other categories within thier farm size group, are from 100 to 1000 ha, where 70% of thier crop production goes towards food.
- 55% of crop production on farms < 2 ha goes towards food, while 16% goes towards feed, and the remainder either goes towards other, waste, processing, or seed (respectively).
- The farm size contributing to the largest amount of food waste (post-harvest loss, not consumer based food waste) are farms between 50-100 ha.

```

In [183]: import warnings
          warnings.filterwarnings('ignore')
          import seaborn as sns
          import pandas as pd
          import numpy as np
          import matplotlib.pyplot as plt
          from matplotlib import cm
          import copy
          import matplotlib.gridspec as gridspec
          from collections import OrderedDict
          from pivottablejs import pivot_ui # python setup.py install --user
          plt.style.use('seaborn-muted')
          %matplotlib inline

```

```

In [119]: def read_data(path):

          data = pd.read_csv(path, low_memory=False)
          data = data.loc[data['data_unit'] == 't']
          data['Farm_Sizes'] = pd.cut(data['fs_class_max'],
                                     bins=[0, 1, 2, 5, 10, 20, 50,
                                            100, 200, 500, 1000, 100000])

```

```

global variables
variables = OrderedDict([('Farm_Sizes', 'Farm_Sizes'),
                          ('production_Food_kcal', 'Food'),
                          ('production_Feed_kcal', 'Feed'),
                          ('production_Seed_kcal', 'Seed'),
                          ('production_Waste_kcal', 'Waste'),
                          ('production_Processing_kcal', 'Processing'),
                          ('production_Other_kcal', 'Other')])

data = data.loc[:, variables.keys()]
data.columns = variables.values()

data = data.replace(0.0, np.nan) # there were many zero values

return data

```

```

In [120]: def data_stats(path):

    path = PATH
    data = pd.read_csv(path, low_memory=False)
    data = data.loc[data['data_unit'] == 't']
    number_records = len(data)
    number_countries = len(data.NAME_0.unique())

    return number_records, number_countries

```

```

In [121]: def piv(data, func=np.nansum):

    pivot = pd.pivot_table(data,
                            index=['Farm_Sizes'],
                            values=variables.values()[1:],
                            aggfunc=func)

    return pivot

```

```

In [122]: def perc(data, how='within'):

    if how is 'within':

        pivot = piv(data)
        pivot = pivot.transpose()

        for variable in pivot.columns:
            pivot[variable] = pivot[variable] / pivot[variable].sum()

    return pivot.transpose()

```

```

elif how is 'cumsum':

    pivot = piv(data)

    for variable in pivot.columns:
        pivot[variable] = pivot[variable] / pivot[variable].sum()

    pivot = pivot.cumsum(axis=0)

    return pivot

elif how is 'across':

    pivot = piv(data)

    for variable in pivot.columns:
        pivot[variable] = pivot[variable] / pivot[variable].sum()

    return pivot.transpose()

else:

    print 'Require how argument'

```

In [123]: `def plot_stacked_bar(data, how='within', fig_=True, ax=None):`

```

txt1 = ['Feed', 'Food', 'Other', 'Processing', 'Seed', 'Waste']
txt2 = ['< 1', '1 to 2', '2 to 5', '5 to 10', '10 to 20',
        '20 to 50', '50 to 100', '100 to 200', '200 to 500',
        '500 to 1000', '> 1000']
txt3 = ['< 1', '2 to 5', '10 to 20', '50 to 100', '200 to 500', '> 1000']

if how is 'within':

    legend_txts = copy.copy(txt1)
    labels_txts = copy.copy(txt2)
    cmap = cm.get_cmap('Set3')
    kind = 'bar'

elif how is 'across':

    legend_txts = copy.copy(txt2)
    labels_txts = copy.copy(txt1)
    cmap = cm.get_cmap('YlGnBu')
    kind = 'bar'

```

```

elif how is 'cumsum':

    legend_txts = copy.copy(txt1)
    labels_txts = copy.copy(txt3)
    cmap = cm.get_cmap('Set3')
    kind = 'area'

else:
    pass

if fig_ is True:

    fig = plt.figure(figsize=[10, 5], facecolor='white')
    ax = fig.add_subplot(111)

else:
    pass

data.plot(kind=kind,
          stacked=True,
          cmap=cmap,
          alpha=0.9,
          linewidth=0,
          grid=False,
          ax=ax)

# Axis main
ax.set_axis_bgcolor("#d6d7e5")
ax.set_clip_on(False)
box = ax.get_position()
ax.set_position([box.x0, box.y0, box.width * 0.8, box.height])

# Legend
legend_txts_r = copy.deepcopy(legend_txts)
legend_txts_r.reverse()
handles, labels = ax.get_legend_handles_labels()
legend = ax.legend(handles[::-1], labels[::-1],
                  loc='center left',
                  frameon=1,
                  bbox_to_anchor=(1, 0.5))

for i in xrange(len(legend_txts_r)):
    legend.get_texts()[i].set_text(legend_txts_r[i])

frame = legend.get_frame()
frame.set_color('white')

# Axis particulars

```

```

ax.set_xticklabels(labels_txts)
ax.set_xticklabels(ax.xaxis.get_majorticklabels(), rotation=45)

if how is 'within':
    ax.set_xlabel('Farm Sizes (ha)')
    ax.set_ylabel('Percentage\n')
    ax.set_ylim([0, 1])
    ax.set_title('Type of production per farm size\n', fontsize=12)

elif how is 'across':
    ax.set_xlabel('Category')
    ax.set_ylabel('Percentage\n')
    ax.set_ylim([0, 1])
    ax.set_title('Type of production across farm size\n', fontsize=12)

elif how is 'cumsum':
    ax.set_xlabel('Farm Sizes (ha)')
    ax.set_ylabel('Percentage\n')
    ax.set_title('Type of production per farm size: Cumulative\n', fontsize=12)

if fig_ is True:
    return plt.show()

else:
    return ax

```

```

In [124]: PATH = '/Users/Vinny_Ricciardi/Documents/Data_Library_Big/Survey/Global/Farm_Size/Data
df = read_data(PATH)

```

```

In [125]: df_within = perc(df, how='within')
df_across = perc(df, how='across')
df_cumsum = perc(df, how='cumsum')
df_raw = piv(df)

```

```

In [126]: tmp1 = df_within.copy()
tmp1['Type'] = 'Within'

tmp2 = df_across.copy()
tmp2 = tmp2.transpose()
tmp2['Type'] = 'Across'

tmp3 = df_cumsum.copy()
tmp3['Type'] = 'Cumsum'

tmp4 = df_raw.copy()
tmp4['Type'] = 'Raw'

tmp = pd.concat([tmp1, tmp2, tmp3, tmp4])

```

```
tmp = tmp.reset_index()
tmp['Farm_Sizes'] = tmp['Farm_Sizes'].str.replace('(', '')
tmp['Farm_Sizes'] = tmp['Farm_Sizes'].str.replace(']', '')
```

## Data

For an overview of the data, see the [data coverage notebook](#). Here is a [link](#) to an interactive pivot table so you can explore the data. Use 'type' to change whether you are looking at the percentage of crop produced for each category 'within' a farm size group, 'across' groups, via a 'cumulative' percentage across groups, or raw kcal produced. Use the dropdown labeled 'Food' to recalculate based on another production category.

```
In [127]: pivot_ui(tmp,
                  rows=['Farm_Sizes'],
                  cols=['Type'],
                  aggregatorName='Sum',
                  vals=['Food'],
                  rendererName='Col Heatmap',
                  )

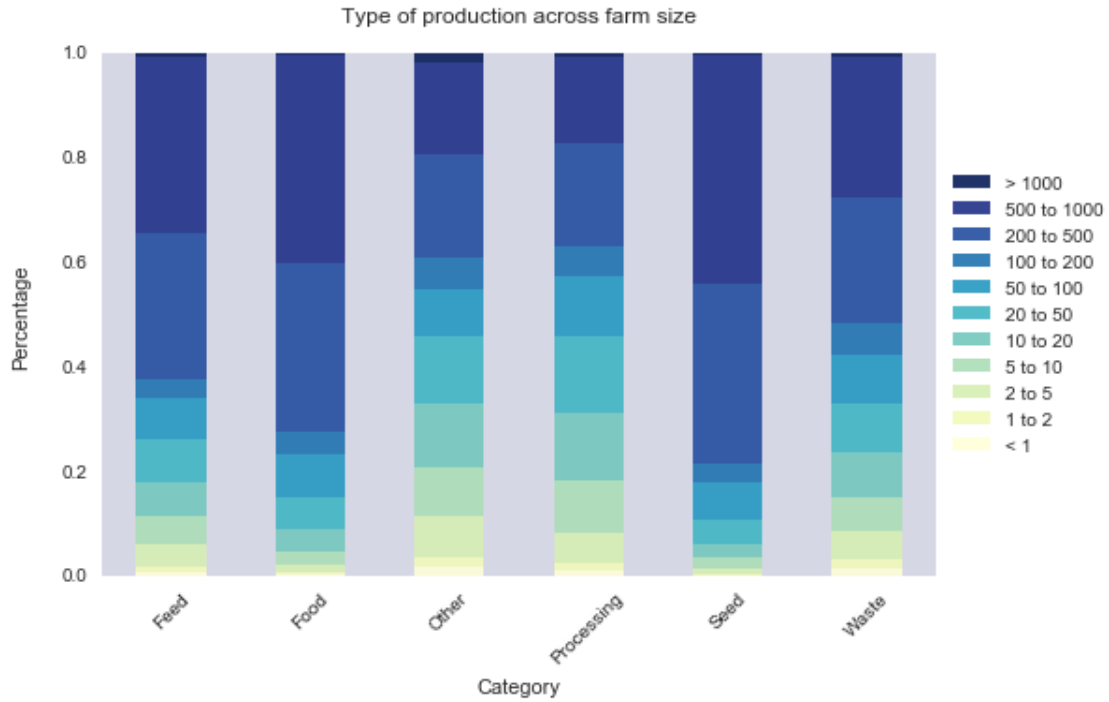
# Enable for pivot table interactivity within this Jupyter notebook
# Here's a nice example of how to change the html output file for prettier output:
# https://pivottable.js.org/examples/montreal_2014.html
```

```
Out[127]: <IPython.lib.display.IFrame at 0x1120601d0>
```

## Food Feed Other by Farm Size

This plot shows the percentage that each farm size contributes to each category (e.g., food, feed, other, etc.) For example, farms between 500 to 1000 ha produce 40.0% of the total food supply in our sample.

```
In [128]: plot_stacked_bar(df_across, how='across', fig_=True)
```

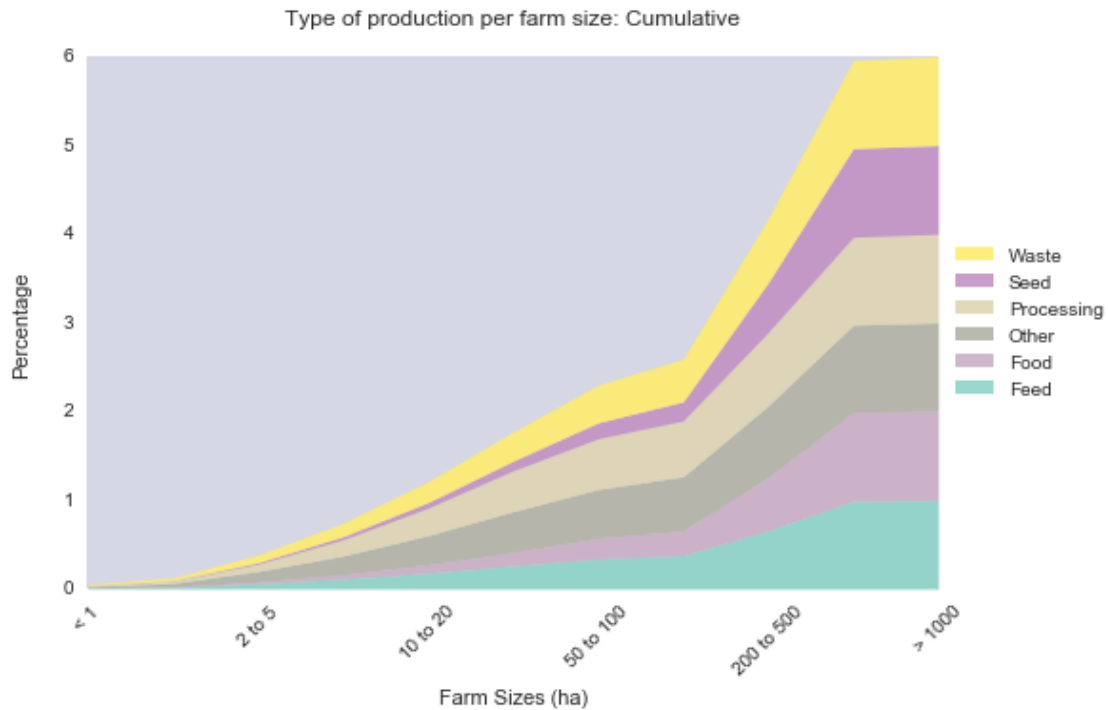


```
In [129]: under_two = round(100 * df_cumsum.iloc[2,1], 1)
          under_fifty = round(100 * df_cumsum.iloc[5,1], 1)
```

This plot also shows the percentage that each farm size contributes to each category but in cumulative percentages. For example, farms under 2 ha produce 2.1% of the total food supply in our sample and farms 50 ha and under produce 15.2% of the total food supply in our sample. Again, here is a [link](#) to an interactive pivot table so you can explore the data in more detail.

```
In [130]: plot_stacked_bar(df_cumsum, how='cumsum', fig=True)
```





```
In [131]: def factor_plot(data, id_var='Farm_Sizes'):

    data = pd.melt(data, id_vars='Farm_Sizes')
    data = data.loc[data['value'] > 0.5]
    data['log'] = np.log(data['value'])

    fs_order = ['(0, 1]', '(1, 2]', '(2, 5]', '(5, 10]',
                '(10, 20]', '(20, 50]', '(50, 100]', '(100, 200]', '(200, 500]',
                '(500, 1000]', '(1000, 1000000)']

    if id_var is 'Farm_Sizes':

        col = 'variable'
        x = 'Farm_Sizes'
        y = 'log'
        order = fs_order

    else:

        data = data.sort('Farm_Sizes')
        data['Farm_Sizes'] = pd.Categorical(data['Farm_Sizes'], fs_order)
        data = data.sort('Farm_Sizes')
        col = 'Farm_Sizes'
        x = 'variable'
```

```

y = 'log'
order = None

g = sns.factorplot(x=x, y=y,
                  col=col,
                  data=data,
                  kind='box',
                  col_wrap=2,
                  color='#55a868',
                  fliersize=1,
                  aspect = 1.5,
                  order=order)

g.fig.subplots_adjust(wspace=0.2, hspace=0.3)

titles = data.variable.unique()

fs_txt = ['< 1', '1 to 2', '2 to 5', '5 to 10', '10 to 20',
          '20 to 50', '50 to 100', '100 to 200', '200 to 500',
          '500 to 1000', '> 1000']

if id_var is 'Farm_Sizes':

    for ax, title in zip(g.axes.flat, titles):

        ax.set_title(title)
        ax.set_ylabel('Production \n ln (kcal)')
        ax.set_xticklabels(fs_txt)
        ax.set_xticklabels(ax.xaxis.get_majorticklabels(), rotation=45)
        ax.set_xlabel('\nFarm Sizes (ha)')

else:

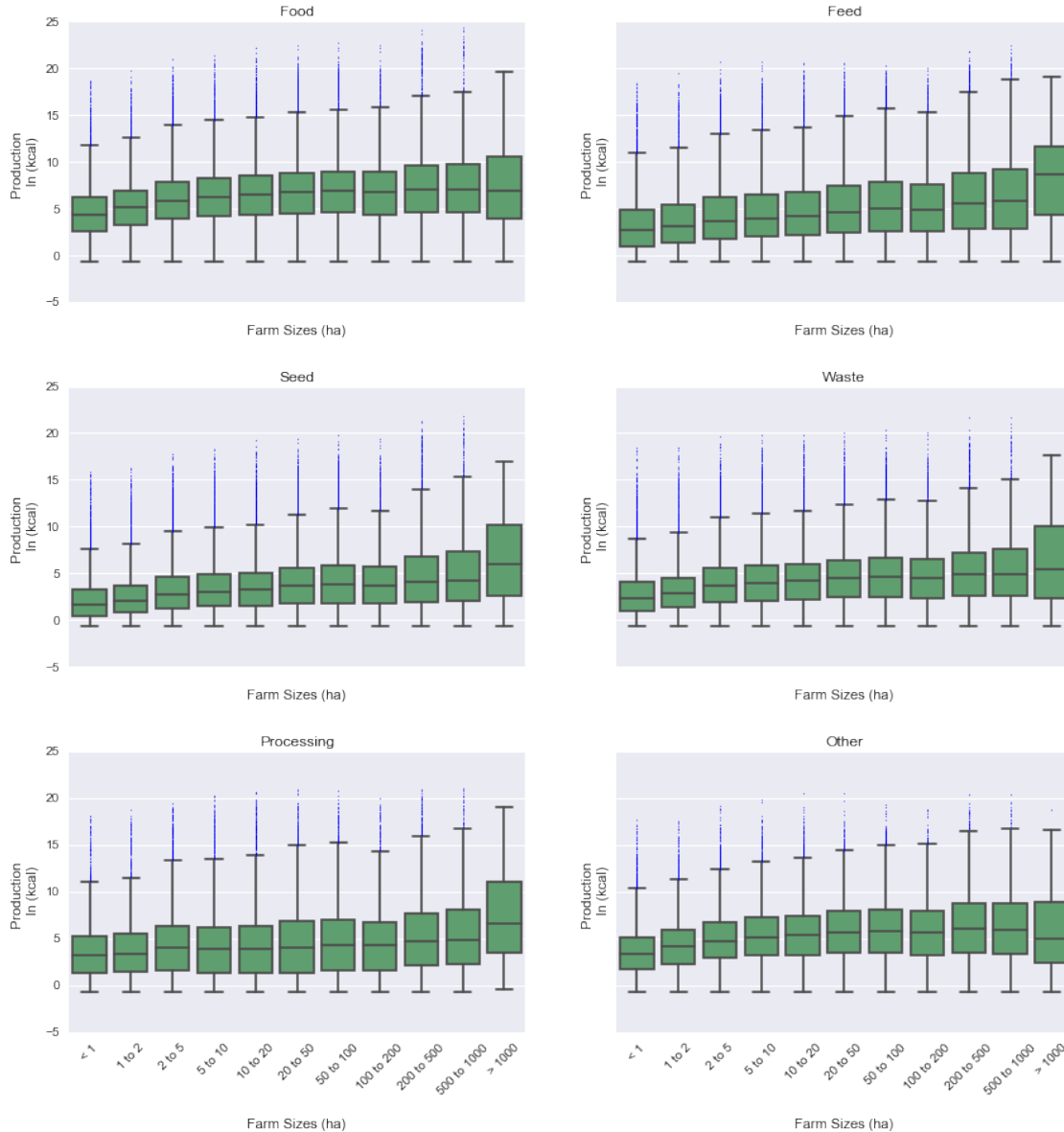
    for ax, title in zip(g.axes.flat, fs_txt):

        ax.set_title(title)
        ax.set_ylabel('Production \n ln (kcal)')
        ax.set_xticklabels(ax.xaxis.get_majorticklabels(), rotation=45)
        ax.set_xlabel('\nPoduction Category')

```

In order to better understand whether there are differences between the farm size groups, these boxplots show the total amount of crops produced (in logged kcal for standardization). Across all production categories (e.g., food, feed, other, etc.) there are no visual differences in the means and confidence intervals. But, there are many of outliers in each plot that indicate high variance.

```
In [132]: factor_plot(df, id_var='Farm_Sizes')
```



Since many of the plots have large variance and skewness, here are density plots comparing farm size classes per each category. We can see that the largest farm size group has different distributions from the remaining groups.

To do: - Fix color map to diverging

```
In [392]: tmp = df.copy()
          tmp = pd.melt(tmp, id_vars='Farm_Sizes')
          tmp = tmp.dropna()
          cols = np.sort(tmp['variable'].unique())
          fs = ['(0, 1]', '(1, 2]', '(2, 5]', '(5, 10]',
                '(10, 20]', '(20, 50]', '(50, 100]',
                '(100, 200]', '(200, 500]', '(500, 1000]']
```

```

'(1000, 100000]']

fig = plt.figure(figsize=[20,10])

for j in xrange(len(cols)):

    ax = fig.add_subplot(2, 3, j+1)
    tmp1 = tmp.loc[tmp['variable'] == cols[j]]

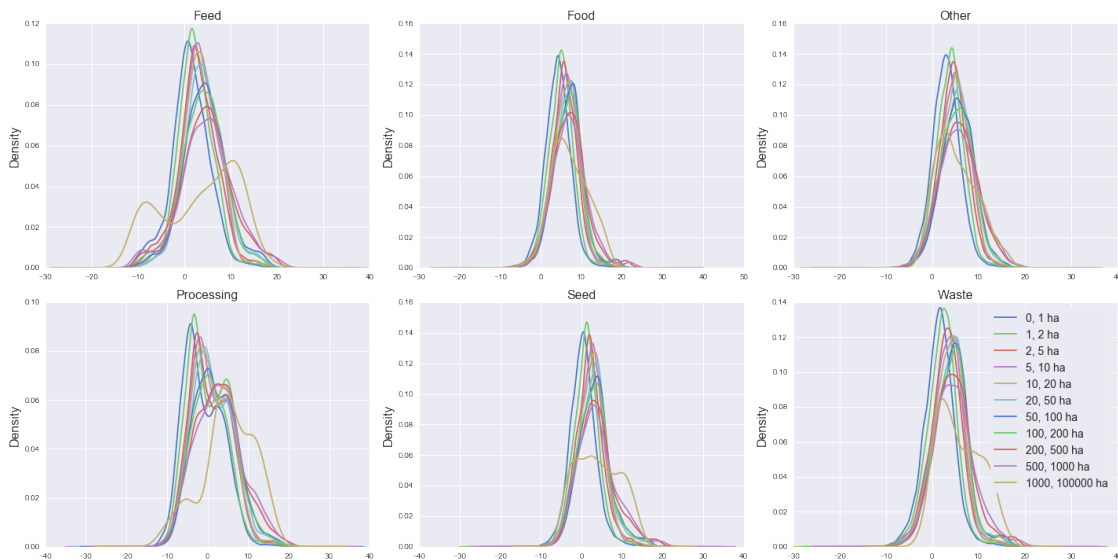
    for i in xrange(len(fs)):

        tmp2 = tmp1.loc[tmp1['Farm_Sizes'] == fs[i]]
        np.log(tmp2['value']).plot(kind='kde',
                                   label=fs[i][1:-1] + ' ha',
                                   ax=ax)

    ax.set_title(cols[j], fontsize=16)
    ax.set_ylabel('Density', fontsize=16)

    legend = ax.legend(frameon=True, fontsize=14)
    plt.tight_layout()
    plt.show()

```



In [133]: `number_records, number_countries = data_stats(PATH)`

Since our sample is very large (12 countries which comprise of 130760 total records due to subnational units and crop types) we cannot use p-values to determine significance between the amount of production by varying farm sizes. To circumvent the large dataset issue, the effect size was used to determine differences between farm sizes rather than statistical significance. Cohen's

d was calculated via taking the (mean of farmsize 1 - mean of farmsize 2) / (standard deviation of farmsize 1) then taking the absolute value. The relative significance was calculated by thresholds according to [Sullivan and Feinn 2012](#).

Where the Cohen d's effect size values correspond to percent non-overlapping observations, as in:

Relative Size	Effect Size	Percentile	% of Non-overlap
	0	50	0
Small	0.2	58	15
Medium	0.5	69	33
Large	0.8	79	47
	1.0	84	55
	1.5	93	71
	2.0	97	81

```
In [134]: def cohens_d(data1, data2, how='within'):

    cols = data1.columns
    check = []

    for i in xrange(0, len(cols)):

        for j in xrange(1, len(cols)):

            col_name = str(cols[i]) + '_' + str(cols[j])
            col_name_r = str(cols[j]) + '_' + str(cols[i])

            if col_name in check:

                pass

            elif cols[i] is cols[j]:

                pass

            else:

                data1[col_name] = ((data1[cols[i]] - data1[cols[j]]) / data2[cols[i]])
                check.append(col_name)
                check.append(col_name_r)

    data = data1.iloc[:, len(cols):]

    data = data.reset_index()

    if how is 'within':
```

```

        data = pd.melt(data, id_vars='Farm_Sizes', value_name='cohens_d')

    else:

        data = pd.melt(data, id_vars='index', value_name='cohens_d')

    data['cohens_d_level'] = np.where(data['cohens_d'] <= 0.2, 'small',
                                     np.where(data['cohens_d'] >= 0.8, 'large',
                                              'medium'))

    return data

In [464]: def cohens_interp(func=np.mean, index='index'):

    tmp = pd.pivot_table(out,
                        index=index,
                        columns='cohens_d_level',
                        values='cohens_d',
                        aggfunc=func)

    tmp = tmp.reset_index()

    try:
        tmp = tmp.sort('large', ascending=False)
        tmp.columns = ['Category', 'large d', 'medium d', 'small d']

    except:
        tmp = tmp.sort('medium', ascending=False)
        tmp.columns = ['Farm Sizes', 'medium d', 'small d']

    return tmp

```

Here are the biggest differences between farm size groups. As expected, there are large coefficients (effect size) between small and large farm classes.

```

In [437]: data = df.copy()
        data['Farm_Sizes'] = data['Farm_Sizes'].astype(str)
        means = pd.pivot_table(data, columns='Farm_Sizes', aggfunc=np.nanmean)
        sds = pd.pivot_table(data, columns='Farm_Sizes', aggfunc=np.nanstd)
        out = cohens_d(means, sds, how='across')

        tmp = out.copy()
        tmp['Farm_Sizes'] = tmp['Farm_Sizes'].str.replace(']', '')
        tmp['Farm_Sizes'] = tmp['Farm_Sizes'].str.replace('_', ' and ')
        tmp.columns = ['Category', 'Farm sizes compared', 'Cohens d', 'Relative effect size']
        tmp = tmp.sort('Cohens d', ascending=False)
        tmp.head(10)

```

```

Out[437]:      Category      Farm sizes compared      Cohens d      Relative effect size
58      Seed  (0, 1) and (500, 1000)  48.488213      large

```

55	Food	(0, 1) and (500, 1000)	31.978579	large
112	Seed	(1, 2) and (500, 1000)	31.014690	large
40	Seed	(0, 1) and (200, 500)	20.790542	large
109	Food	(1, 2) and (500, 1000)	17.098681	large
37	Food	(0, 1) and (200, 500)	14.102206	large
94	Seed	(1, 2) and (200, 500)	13.287843	large
268	Seed	(2, 5) and (500, 1000)	9.832608	large
54	Feed	(0, 1) and (500, 1000)	8.674037	large
91	Food	(1, 2) and (200, 500)	7.530880	large

On average, the largest effect size were in the seed and food categories.

```
In [439]: cohens_interp(func=np.mean)
```

```
Out[439]:
```

	Category	large d	medium d	small d
4	Seed	7.540147	0.474176	0.067408
1	Food	6.551549	0.483279	0.052100
0	Feed	2.248732	0.418498	0.060634
3	Processing	1.957038	0.392405	0.056883
5	Waste	1.542108	0.382661	0.059074
2	Other	1.182823	0.277370	0.057448

The seed and food categories also had the highest frequency of large effect sizes between farm size groups.

```
In [430]: cohens_interp(func=len)
```

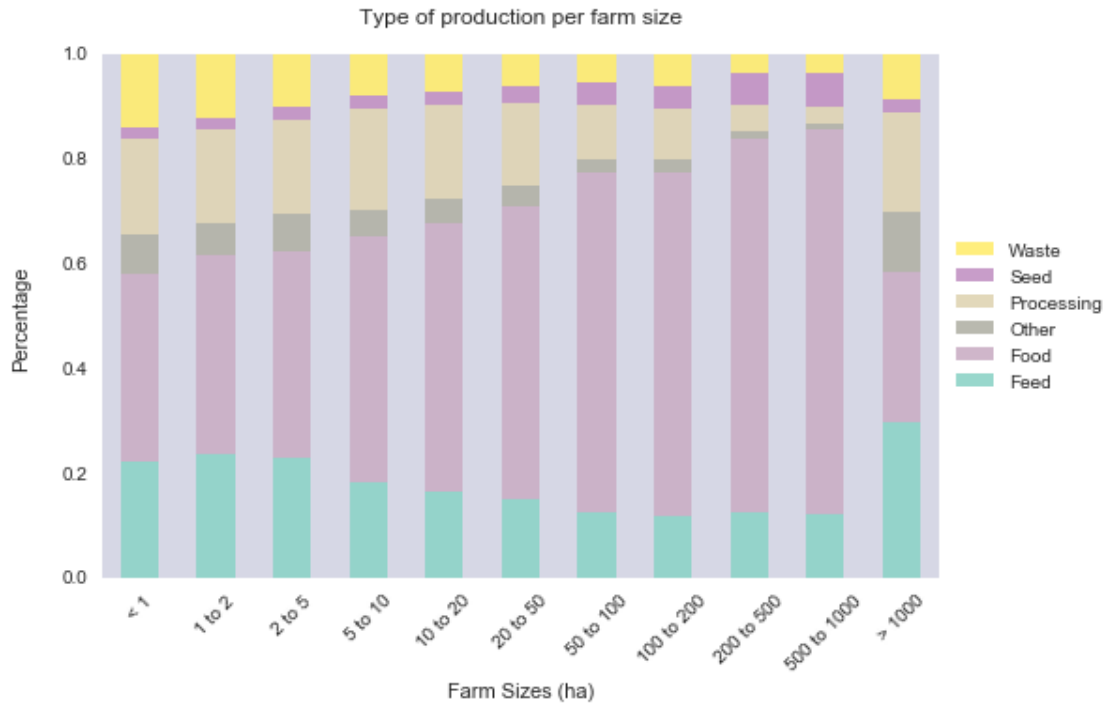
```
Out[430]:
```

	Category	large d	medium d	small d
4	Seed	21.0	8.0	26.0
1	Food	15.0	15.0	25.0
0	Feed	12.0	12.0	31.0
5	Waste	6.0	13.0	36.0
2	Other	5.0	12.0	38.0
3	Processing	4.0	19.0	32.0

#### Food Feed Other within Farm Size Groups

This plot shows the percentage of Food, Feed, Seed, Waste, Processing, and Other for each farm size category. For example, {{one\_food}} % of crop production for farms under 1 ha is food, while 6% of their crop production is waste.

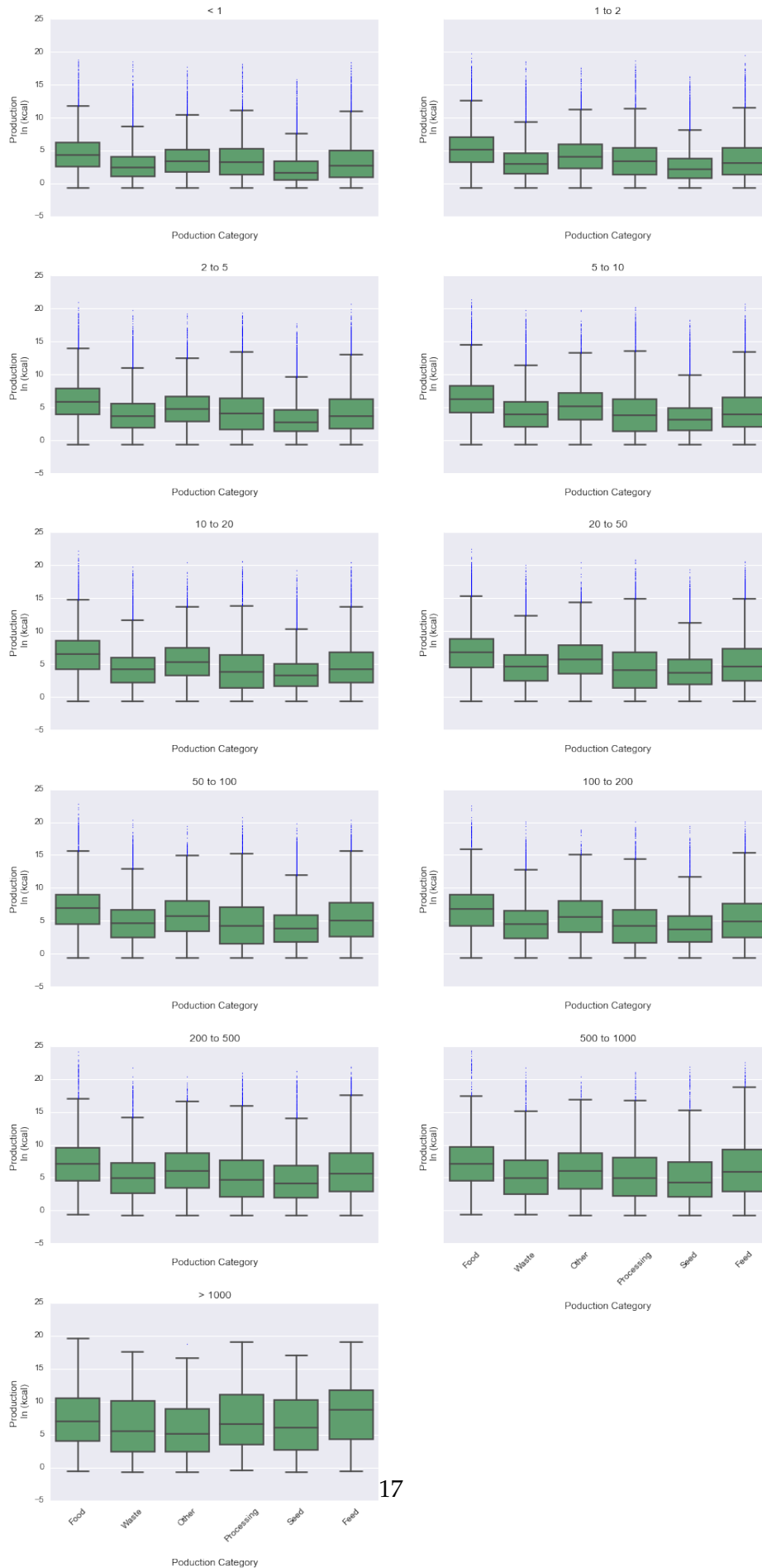
```
In [385]: plot_stacked_bar(df_within, how='within', fig_=True)
```



In order to better understand whether there are differences within how a given farm size group allocates their crop production, these boxplots show the total amount of crops produced (in logged kcal for standardization). As in the previous analysis, across all farm size categories there are no visual differences in the means and confidence intervals. But, there are many of outliers in each plot that indicate high variance.

```
In [386]: factor_plot(df, id_var='variable')
```





Since many of the plots have large variance and skewness, here are density plots comparing farm size classes per each category. We can see that the largest farm size group has different distributions from the remaining groups.

To do: - Fix color map to diverging

```
In [384]: tmp = df.copy()
fs = ['(0, 1]', '(1, 2]', '(2, 5]', '(5, 10]',
      '(10, 20]', '(20, 50]', '(50, 100]',
      '(100, 200]', '(200, 500]', '(500, 1000]',
      '(1000, 1000000]']
tmp = tmp.dropna()
cols = tmp.columns

fig = plt.figure(figsize=[20,10])

for j in xrange(len(fs)):

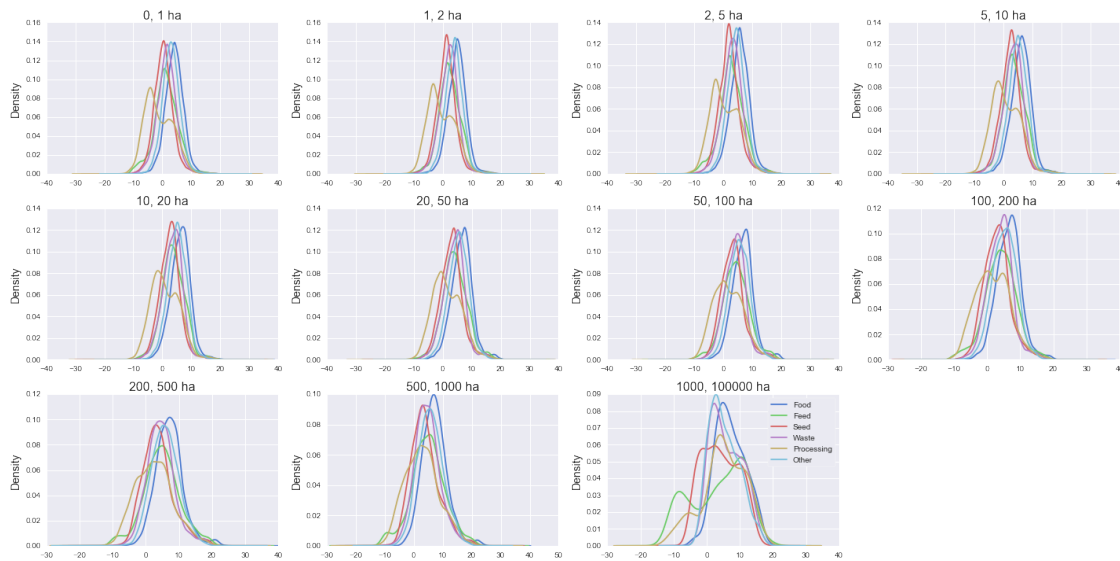
    tmp1 = tmp.loc[tmp['Farm_Sizes'] == fs[j]]

    ax = fig.add_subplot(3, 4, j+1)

    for i in xrange(1, len(cols)):
        np.log(tmp1[cols[i]]).plot(kind='kde',
                                   label=cols[i],
                                   ax=ax)

    ax.set_title(fs[j][1:-1] + ' ha', fontsize=16)
    ax.set_ylabel('Density', fontsize=14)

legend = ax.legend(frameon=True)
plt.tight_layout()
plt.show()
```



We calculated the effect size for the differences within each farm size group, but across production categories (e.g., food, feed, other). There were only medium differences in effect size when looking at how each farm size group allocated their production.

```
In [445]: means = pd.pivot_table(df, index='Farm_Sizes', aggfunc=np.nanmean)
          sds = pd.pivot_table(df, index='Farm_Sizes', aggfunc=np.nanstd)
          out = cohens_d(means, sds, how='within')

          tmp = out.copy()
          tmp = tmp.loc[out['cohens_d_level'] != 'small']
          tmp['Farm_Sizes'] = tmp['Farm_Sizes'].str.replace(']', '')
          tmp['variable'] = tmp['variable'].str.replace('_', ' and ')
          tmp.columns = ['Farm sizes', 'Categories compared', 'Cohens d', 'Relative effect size']
          tmp = tmp.sort('Cohens d', ascending=False)
          out = out.reset_index()
          tmp.head(10)
```

```
Out[445]:
```

	Farm sizes	Categories compared	Cohens d	Relative effect size
9	(500, 1000)	Feed and Food	0.399468	medium
8	(200, 500)	Feed and Food	0.395566	medium
119	(500, 1000)	Other and Seed	0.349541	medium
6	(50, 100)	Feed and Food	0.314126	medium
7	(100, 200)	Feed and Food	0.300623	medium
164	(1000, 100000)	Seed and Waste	0.237525	medium
154	(0, 1)	Seed and Waste	0.217207	medium
155	(1, 2)	Seed and Waste	0.204633	medium

On average, the largest effect size were in the larger farm size groups.

```
In [470]: cohens_interp(func=np.mean, index='Farm_Sizes').head()
```

```
Out[470]:
```

	Farm Sizes	medium d	small d
7	(200, 500]	0.395566	0.068049
10	(500, 1000]	0.374505	0.070297
9	(50, 100]	0.314126	0.047100
3	(100, 200]	0.300623	0.039264
4	(1000, 100000]	0.237525	0.062113

The most frequent differences between categories within a farm size group was for the 500 to 1000 ha group.

```
In [472]: cohens_interp(func=len, index='Farm_Sizes').head()
```

```
Out[472]:
```

	Farm Sizes	medium d	small d
10	(500, 1000]	2.0	13.0
0	(0, 1]	1.0	14.0
1	(1, 2]	1.0	14.0
3	(100, 200]	1.0	14.0
4	(1000, 100000]	1.0	14.0

Global Estimates

```
In [ ]: data = df.copy()
```

```
In [ ]: data.head()
```

Left Off

To Do: - Need to calculate bootstraps and jackknife estimates

```
In [ ]:
```