# **US Baby Names 1880-2010**

The United States Social Security Administration (SSA) has made available data on the frequency of baby names from 1880 through the present. Hadley Wickham, an author of several popular R packages, has often made use of this data set in illustrating data manipulation in R.

```
In [4]: names.head(10)
Out[4]:
      name sex births year
      Mary F 7065 1880
0
      Anna F
               2604 1880
1
      Emma F 2003 1880
2
3 Elizabeth F 1939 1880
    Minnie F 1746 1880
5 Margaret F 1578 1880
6
      Ida F 1472 1880
     Alice F 1414 1880
7
    Bertha F 1320 1880
8
9
     Sarah F
               1288 1880
```

There are many things you might want to do with the data set:

- Visualize the proportion of babies given a particular name (your own, or another name) over time.
- Determine the relative rank of a name.
- Determine the most popular names in each year or the names with largest increases or decreases.
- Analyze trends in names: vowels, consonants, length, overall diversity, changes in spelling, first and last letters
- Analyze external sources of trends: biblical names, celebrities, demographic changes

Using the tools we've looked at so far, most of these kinds of analyses are very straightforward, so I will walk you through many of them. I encourage you to download and explore the data yourself. If you find an interesting pattern in the data, I would love to hear about it.

As of this writing, the US Social Security Administration makes available data files, one per year, containing the total number of births for each sex/name combination. The raw archive of these files can be obtained here:

```
http://www.ssa.gov/oact/babynames/limits.html
```

In the event that this page has been moved by the time you're reading this, it can most likely be located again by Internet search. After downloading the "National data" file names.zip and unzipping it, you will have a directory containing a series of files like yob1880.txt. I use the UNIX head command to look at the first 10 lines of one of the files (on Windows, you can use the more command or open it in a text editor):

```
In [367]: !head -n 10 names/yob1880.txt
Mary, F, 7065
Anna, F, 2604
Emma, F, 2003
Elizabeth, F, 1939
Minnie, F, 1746
Margaret, F, 1578
Ida, F, 1472
Alice,F,1414
Bertha, F, 1320
Sarah, F, 1288
```

As this is a nicely comma-separated form, it can be loaded into a DataFrame with pandas.read csv:

```
In [368]: import pandas as pd
In [369]: names1880 = pd.read csv('names/yob1880.txt', names=['name', 'sex', 'births'])
In [370]: names1880
Out[370]:
<class 'pandas.core.frame.DataFrame'>
Int64Index: 2000 entries, 0 to 1999
Data columns:
         2000 non-null values
name
         2000 non-null values
sex
births
         2000 non-null values
dtypes: int64(1), object(2)
```

These files only contain names with at least 5 occurrences in each year, so for simplicity's sake we can use the sum of the births column by sex as the total number of births in that year:

```
In [371]: names1880.groupby('sex').births.sum()
Out[371]:
sex
F
        90993
       110493
Name: births
```

Since the data set is split into files by year, one of the first things to do is to assemble all of the data into a single DataFrame and further to add a year field. This is easy to do using pandas.concat:

```
# 2010 is the last available year right now
years = range(1880, 2011)
pieces = []
columns = ['name', 'sex', 'births']
for year in years:
    path = 'names/yob%d.txt' % year
    frame = pd.read csv(path, names=columns)
    frame['year'] = year
    pieces.append(frame)
```

```
# Concatenate everything into a single DataFrame
names = pd.concat(pieces, ignore index=True)
```

There are a couple things to note here. First, remember that concat glues the DataFrame objects together row-wise by default. Secondly, you have to pass ignore index=True because we're not interested in preserving the original row numbers returned from read csv. So we now have a very large DataFrame containing all of the names data:

Now the names DataFrame looks like:

```
In [373]: names
Out[373]:
<class 'pandas.core.frame.DataFrame'>
Int64Index: 1690784 entries, 0 to 1690783
Data columns:
name
       1690784 non-null values
       1690784 non-null values
sex
births 1690784 non-null values
year 1690784 non-null values
dtypes: int64(2), object(2)
```

With this data in hand, we can already start aggregating the data at the year and sex level using groupby or pivot table, see Figure 2-4:

```
In [374]: total births = names.pivot table('births', rows='year',
                                          cols='sex', aggfunc=sum)
In [375]: total births.tail()
Out[375]:
sex
year
2006 1896468 2050234
2007 1916888 2069242
2008 1883645 2032310
2009 1827643 1973359
2010 1759010 1898382
In [376]: total births.plot(title='Total births by sex and year')
```

Next, let's insert a column prop with the fraction of babies given each name relative to the total number of births. A prop value of 0.02 would indicate that 2 out of every 100 babies was given a particular name. Thus, we group the data by year and sex, then add the new column to each group:

```
def add prop(group):
    # Integer division floors
    births = group.births.astype(float)
    group['prop'] = births / births.sum()
    return group
names = names.groupby(['year', 'sex']).apply(add prop)
```

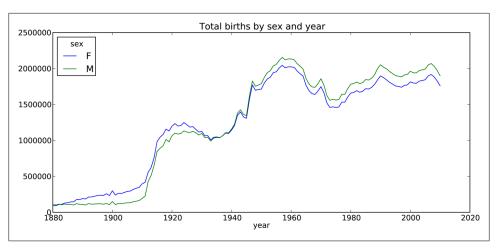


Figure 2-4. Total births by sex and year



Remember that because births is of integer type, we have to cast either the numerator or denominator to floating point to compute a fraction (unless you are using Python 3!).

The resulting complete data set now has the following columns:

```
In [378]: names
Out[378]:
<class 'pandas.core.frame.DataFrame'>
Int64Index: 1690784 entries, 0 to 1690783
Data columns:
name
         1690784 non-null values
sex
         1690784 non-null values
         1690784 non-null values
births
         1690784 non-null values
year
         1690784 non-null values
prop
dtypes: float64(1), int64(2), object(2)
```

When performing a group operation like this, it's often valuable to do a sanity check, like verifying that the prop column sums to 1 within all the groups. Since this is floating point data, use np.allclose to check that the group sums are sufficiently close to (but perhaps not exactly equal to) 1:

```
In [379]: np.allclose(names.groupby(['year', 'sex']).prop.sum(), 1)
Out[379]: True
```

Now that this is done, I'm going to extract a subset of the data to facilitate further analysis: the top 1000 names for each sex/year combination. This is yet another group operation:

```
def get top1000(group):
    return group.sort index(by='births', ascending=False)[:1000]
```

```
grouped = names.groupby(['year', 'sex'])
    top1000 = grouped.apply(get top1000)
If you prefer a do-it-yourself approach, you could also do:
    pieces = []
    for year, group in names.groupby(['year', 'sex']):
        pieces.append(group.sort index(by='births', ascending=False)[:1000])
    top1000 = pd.concat(pieces, ignore index=True)
The resulting data set is now quite a bit smaller:
    In [382]: top1000
    Out[382]:
    <class 'pandas.core.frame.DataFrame'>
    Int64Index: 261877 entries, 0 to 261876
    Data columns:
              261877 non-null values
    name
    sex
              261877 non-null values
    births
             261877 non-null values
    year 261877 non-null values
             261877 non-null values
    dtypes: float64(1), int64(2), object(2)
```

We'll use this Top 1,000 data set in the following investigations into the data.

## **Analyzing Naming Trends**

With the full data set and Top 1,000 data set in hand, we can start analyzing various naming trends of interest. Splitting the Top 1,000 names into the boy and girl portions is easy to do first:

```
In [383]: boys = top1000[top1000.sex == 'M']
In [384]: girls = top1000[top1000.sex == 'F']
```

Simple time series, like the number of Johns or Marys for each year can be plotted but require a bit of munging to be a bit more useful. Let's form a pivot table of the total number of births by year and name:

```
In [385]: total births = top1000.pivot table('births', rows='year', cols='name',
                                             aggfunc=sum)
```

Now, this can be plotted for a handful of names using DataFrame's plot method:

```
In [386]: total births
Out[386]:
<class 'pandas.core.frame.DataFrame'>
Int64Index: 131 entries, 1880 to 2010
Columns: 6865 entries, Aaden to Zuri
dtypes: float64(6865)
In [387]: subset = total births[['John', 'Harry', 'Mary', 'Marilyn']]
In [388]: subset.plot(subplots=True, figsize=(12, 10), grid=False,
                      title="Number of births per year")
   . . . . . :
```

See Figure 2-5 for the result. On looking at this, you might conclude that these names have grown out of favor with the American population. But the story is actually more complicated than that, as will be explored in the next section.

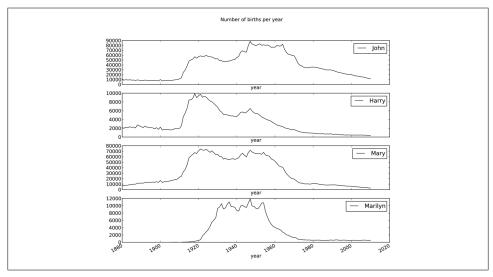


Figure 2-5. A few boy and girl names over time

### Measuring the increase in naming diversity

One explanation for the decrease in plots above is that fewer parents are choosing common names for their children. This hypothesis can be explored and confirmed in the data. One measure is the proportion of births represented by the top 1000 most popular names, which I aggregate and plot by year and sex:

```
In [390]: table = top1000.pivot table('prop', rows='year',
                                      cols='sex', aggfunc=sum)
In [391]: table.plot(title='Sum of table1000.prop by year and sex',
                     yticks=np.linspace(0, 1.2, 13), xticks=range(1880, 2020, 10))
```

See Figure 2-6 for this plot. So you can see that, indeed, there appears to be increasing name diversity (decreasing total proportion in the top 1,000). Another interesting metric is the number of distinct names, taken in order of popularity from highest to lowest, in the top 50% of births. This number is a bit more tricky to compute. Let's consider just the boy names from 2010:

```
In [392]: df = boys[boys.year == 2010]
In [393]: df
Out[393]:
<class 'pandas.core.frame.DataFrame'>
Int64Index: 1000 entries, 260877 to 261876
Data columns:
```

```
name
         1000 non-null values
sex
         1000 non-null values
births
         1000 non-null values
         1000 non-null values
year
prop
         1000 non-null values
dtypes: float64(1), int64(2), object(2)
```

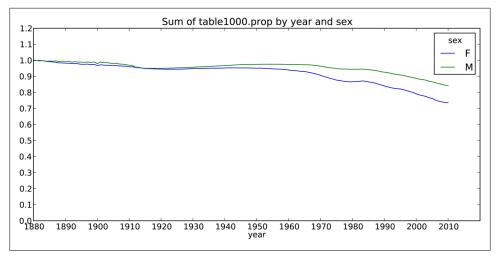


Figure 2-6. Proportion of births represented in top 1000 names by sex

After sorting prop in descending order, we want to know how many of the most popular names it takes to reach 50%. You could write a for loop to do this, but a vectorized NumPy way is a bit more clever. Taking the cumulative sum, cumsum, of prop then calling the method searchsorted returns the position in the cumulative sum at which 0.5 would need to be inserted to keep it in sorted order:

```
In [394]: prop cumsum = df.sort_index(by='prop', ascending=False).prop.cumsum()
In [395]: prop cumsum[:10]
Out[395]:
260877
         0.011523
260878
         0.020934
260879
          0.029959
260880
         0.038930
260881
         0.047817
260882
         0.056579
260883
          0.065155
260884
         0.073414
260885
         0.081528
260886
         0.089621
In [396]: prop cumsum.searchsorted(0.5)
Out[396]: 116
```

Since arrays are zero-indexed, adding 1 to this result gives you a result of 117. By contrast, in 1900 this number was much smaller:

```
In [397]: df = boys[boys.year == 1900]
In [398]: in1900 = df.sort index(by='prop', ascending=False).prop.cumsum()
In [399]: in1900.searchsorted(0.5) + 1
Out[399]: 25
```

It should now be fairly straightforward to apply this operation to each year/sex combination; groupby those fields and apply a function returning the count for each group:

```
def get quantile count(group, q=0.5):
    group = group.sort index(by='prop', ascending=False)
    return group.prop.cumsum().searchsorted(q) + 1
diversity = top1000.groupby(['year', 'sex']).apply(get quantile count)
diversity = diversity.unstack('sex')
```

This resulting DataFrame diversity now has two time series, one for each sex, indexed by year. This can be inspected in IPython and plotted as before (see Figure 2-7):

```
In [401]: diversity.head()
Out[401]:
sex
year
1880 38 14
1881 38
         14
1882 38 15
1883 39 15
1884 39 16
```

In [402]: diversity.plot(title="Number of popular names in top 50%")

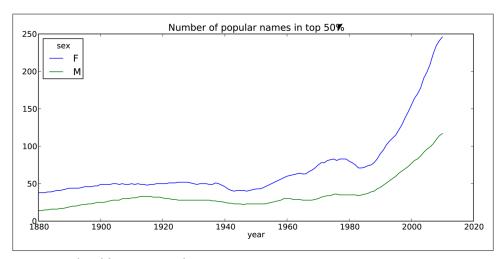


Figure 2-7. Plot of diversity metric by year

As you can see, girl names have always been more diverse than boy names, and they have only become more so over time. Further analysis of what exactly is driving the diversity, like the increase of alternate spellings, is left to the reader.

#### The "Last letter" Revolution

In 2007, a baby name researcher Laura Wattenberg pointed out on her website (http: //www.babynamewizard.com) that the distribution of boy names by final letter has changed significantly over the last 100 years. To see this, I first aggregate all of the births in the full data set by year, sex, and final letter:

```
# extract last letter from name column
get last letter = lambda x: x[-1]
last letters = names.name.map(get last letter)
last letters.name = 'last letter'
table = names.pivot table('births', rows=last letters,
                          cols=['sex', 'year'], aggfunc=sum)
```

Then, I select out three representative years spanning the history and print the first few rows:

```
In [404]: subtable = table.reindex(columns=[1910, 1960, 2010], level='year')
In [405]: subtable.head()
Out[405]:
Sex
            1910 1960 2010 1910
year
                                     1960
                                            2010
last letter
         108376 691247 670605 977
                                     5204 28438
b
          NaN 694 450 411 3912 38859
C
             5
                  49
                          946 482 15476 23125
d
            6750 3729 2607 22111 262112
                                           44398
          133569 435013 313833 28655 178823 129012
```

Next, normalize the table by total births to compute a new table containing proportion of total births for each sex ending in each letter:

```
In [406]: subtable.sum()
Out[406]:
sex year
    1910
         396416
    1960
           2022062
         1759010
    2010
    1910
         194198
    1960
           2132588
    2010 1898382
```

In [407]: letter prop = subtable / subtable.sum().astype(float)

With the letter proportions now in hand, I can make bar plots for each sex broken down by year. See Figure 2-8:

```
import matplotlib.pyplot as plt
```

```
fig, axes = plt.subplots(2, 1, figsize=(10, 8))
letter_prop['M'].plot(kind='bar', rot=0, ax=axes[0], title='Male')
letter prop['F'].plot(kind='bar', rot=0, ax=axes[1], title='Female',
                      legend=False)
```

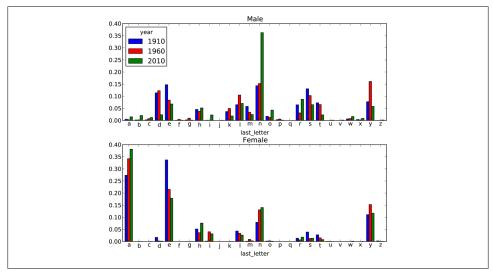


Figure 2-8. Proportion of boy and girl names ending in each letter

As you can see, boy names ending in "n" have experienced significant growth since the 1960s. Going back to the full table created above, I again normalize by year and sex and select a subset of letters for the boy names, finally transposing to make each column a time series:

```
In [410]: letter prop = table / table.sum().astype(float)
In [411]: dny ts = letter prop.ix[['d', 'n', 'y'], 'M'].T
In [412]: dny ts.head()
Out[412]:
year
1880 0.083055 0.153213 0.075760
1881 0.083247 0.153214 0.077451
1882 0.085340 0.149560 0.077537
1883 0.084066 0.151646 0.079144
1884 0.086120 0.149915 0.080405
```

With this DataFrame of time series in hand, I can make a plot of the trends over time again with its plot method (see Figure 2-9):

```
In [414]: dny ts.plot()
```

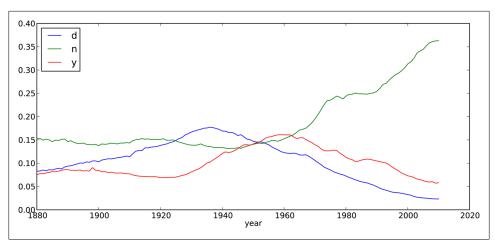


Figure 2-9. Proportion of boys born with names ending in d/n/y over time

### Boy names that became girl names (and vice versa)

Another fun trend is looking at boy names that were more popular with one sex earlier in the sample but have "changed sexes" in the present. One example is the name Lesley or Leslie. Going back to the top1000 dataset, I compute a list of names occurring in the dataset starting with 'lesl':

```
In [415]: all names = top1000.name.unique()
In [416]: mask = np.array(['lesl' in x.lower() for x in all names])
In [417]: lesley like = all names[mask]
In [418]: lesley like
Out[418]: array([Leslie, Lesley, Leslee, Lesli, Lesly], dtype=object)
```

From there, we can filter down to just those names and sum births grouped by name to see the relative frequencies:

```
In [419]: filtered = top1000[top1000.name.isin(lesley like)]
In [420]: filtered.groupby('name').births.sum()
Out[420]:
name
Leslee
            1082
Lesley
           35022
Lesli
             929
Leslie
          370429
Lesly
           10067
Name: births
```

Next, let's aggregate by sex and year and normalize within year:

```
In [421]: table = filtered.pivot table('births', rows='year',
                                       cols='sex', aggfunc='sum')
In [422]: table = table.div(table.sum(1), axis=0)
In [423]: table.tail()
Out[423]:
sex
year
2006 1 NaN
     1 NaN
2007
2008 1 NaN
2009 1 NaN
2010 1 NaN
```

Lastly, it's now easy to make a plot of the breakdown by sex over time (Figure 2-10): In [425]: table.plot(style={'M': 'k-', 'F': 'k--'})

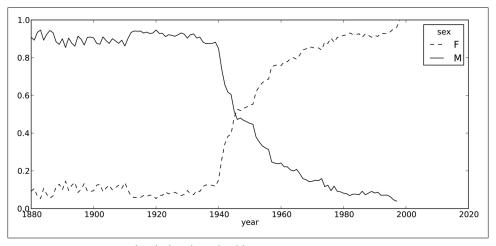


Figure 2-10. Proportion of male/female Lesley-like names over time

## Conclusions and The Path Ahead

The examples in this chapter are rather simple, but they're here to give you a bit of a flavor of what sorts of things you can expect in the upcoming chapters. The focus of this book is on *tools* as opposed to presenting more sophisticated analytical methods. Mastering the techniques in this book will enable you to implement your own analyses (assuming you know what you want to do!) in short order.

