```
*********
   * Penguin Analysis
2
   * Demonstration Do File
   *********
4
5
   * So many projects have the same, or similar, workflow.
6
7
   * have a question ->
8
  * get data ->
  * process and clean data ->
10
   * analyze data ->
11
   * visualize data ->
12
   * make conclusions
13
14
   /* do files are useful to preserve
15
   a record of your work. They help
   to keep an audit trail of the
   decisions that you have made. */
19
   /* do files thus serve as a way of creating an
   automated, replicable and documented workflow
21
   as well as finding and minimizing errors */
22
23
   * A `*` character at the beginning of a line makes that
24
   line a comment
25
   /* You can also use asterisk slash to denote multiple
26
   lines of comment */
27
   *********
28
   * get data
29
   *********
30
   * a good workflow habit is to
32
   * always--or at least frequently--
   * work from your raw data.
34
35
   * i.e. run your script so you are always--
36
   * or at least often--opening your raw data,
37
   * cleaning the data, creating new variables,
38
   * and then running analyses.
39
40
```

```
clear all // clear the workspace
41
42
   * get data from web
43
44
   use "penguins.dta", clear
45
46
   **********
47
   * take a look at the data
48
   *********
49
50
   * NB if you have a lot of variables, the commands below
51
   will produce a lot of (too much) output
52
   * you may need to `describe` or `codebook` specific
53
   variables
54
   describe // describe the variables
56
   codebook // full descriptions of all the variables;
57
   produces a lot of output
58
   *********
59
   * descriptive statistics
60
   *********
61
62
   summarize // descriptive statistics for all variables
63
64
   summarize body mass g // descriptive statistics for this
65
   variable
66
   tabulate species // tabulate this categorical variable
67
68
   * dtable is a useful new command
69
   * for producing tables of descriptive statistics
   * be sure to denote indicator variables with an `i.`
71
72
   dtable culmen length mm body mass g i.species
73
74
   ********
75
   * data wrangling
76
   *********
77
78
```

```
* find variables of interest
79
80
   lookfor mass // look for a variable w a particular keyword
81
82
   * sometimes it is useful to `keep` only the variables in
83
   which you have an interest
   * to reduce the size of the data set
84
85
   * recode variables
86
87
   generate big penguin = body mass g > 4000 // create a
88
   big penguin variable
89
   tabulate big penguin
90
91
   *********
92
   * ANOVA
93
   *********
94
95
   oneway body_mass_g species, tabulate
96
97
   **********
98
   * regression
99
   ********
100
101
   regress culmen length mm body mass q
102
103
   est store M1 // store regression estimates
104
105
   regress culmen_length_mm body_mass_g i.species
106
107
   est store M2 // store regression estimates
108
109
   * /// indicates that a command spans multiple lines
110
111
   etable, estimates(M1 M2) /// nicely formatted table of
112
   regression estimates
   cstat( r b) /// beta's only
113
   showstars showstarsnote // show stars and note
114
115
116
   *********
   * graph
117
```

```
**************

************

graph bar body_mass_g, over(species) // bar graph

twoway scatter culmen_length_mm body_mass_g // scatterplot

twoway scatter culmen_length_mm body_mass_g // scatterplot

123
124
125
126
127
128
```