Week12_Assignment

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```
## -- Attaching core tidyverse packages ------------------------ tidyverse 2.0.0 --
## v dplyr     1.1.4     v readr     2.1.5
## v forcats   1.0.0     v stringr   1.5.1
## v ggplot2   3.5.1     v tibble    3.2.1
## v lubridate 1.9.4     v tidyr     1.3.1
## v purrr     1.0.4
## -- Conflicts ------------------------------------------ tidyverse_conflicts() --
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()    masks stats::lag()
## i Use the conflicted package (<http://conflicted.r-lib.org/>) to force all conflicts to become errors
```

1. For Loop Basics (30 pts)

1a.

```
## [1] 3
## [1] 6
## [1] 9
## [1] 12
## [1] 15
```

1b.

```
## [1] 4.84
## [1] 7.7
## [1] 21.12
## [1] 2.64
```

1c.

```
## [1] "robin"
## [1] "woodpecker"
## [1] "blue jay"
## [1] "sparrow"
```

1d.

```
## [1]  5.309292 13.854424 38.484510
```

1e.

```
## [1] 3.85 5.28 4.48
```

2. Size Estimates by Name (30 pts)

```
## Rows: 500 Columns: 2
## -- Column specification --------------------------------------------------------
## Delimiter: ","
## chr (1): species
## dbl (1): lengths
##
## i Use `spec()` to retrieve the full column specification for this data.
## i Specify the column types or set `show_col_types = FALSE` to quiet this message.
```

2a.

```
## [1] 24341.68 27017.90 67453.38 22114.19 53884.76 52026.34
```

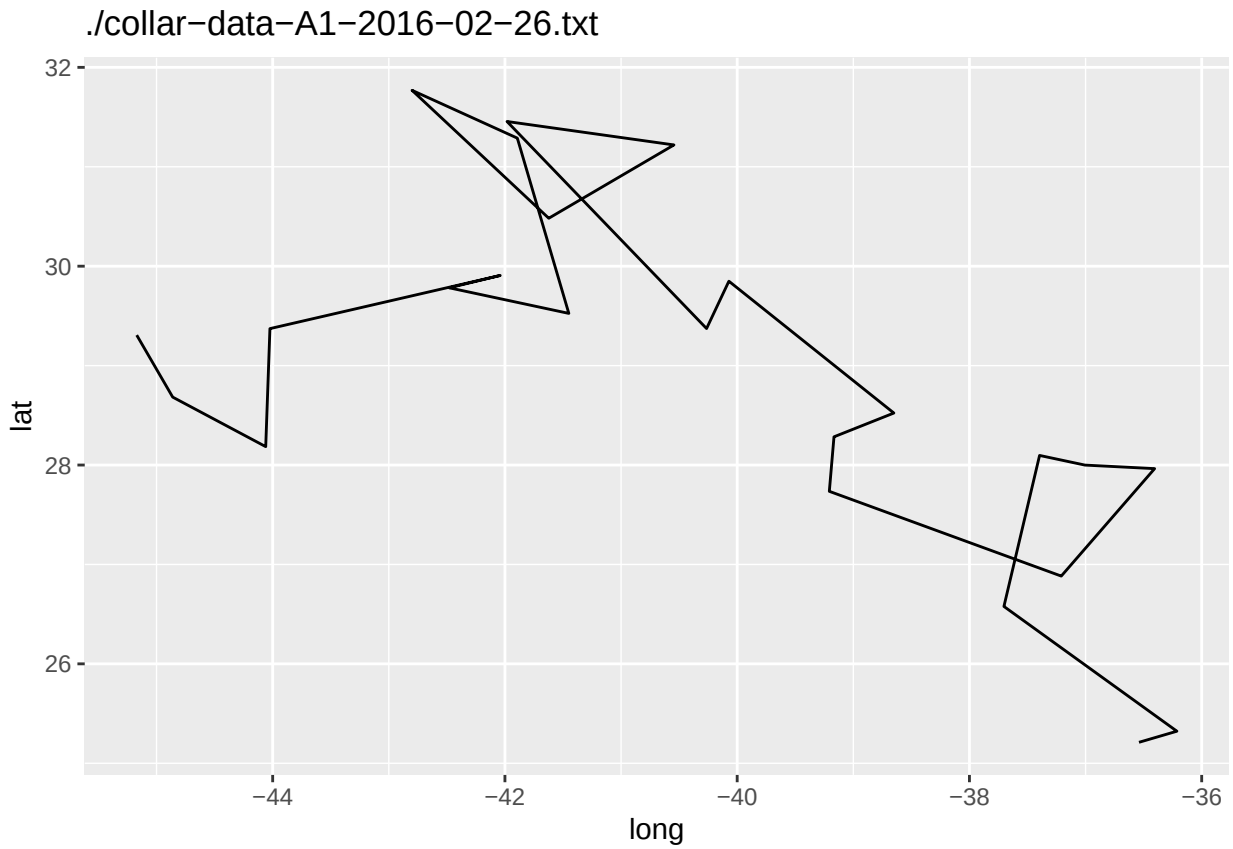
2b.

```
## # A tibble: 6 x 3
##   species      lengths masses
##   <chr>          <dbl>  <dbl>
## 1 Stegosauria     18.5 24342.
## 2 Ankylosauria    16.4 27018.
## 3 Ankylosauria    23.7 67453.
## 4 Sauropoda       23.9 22114.
## 5 Ankylosauria    21.7 53885.
## 6 Ankylosauria    21.4 52026.
```

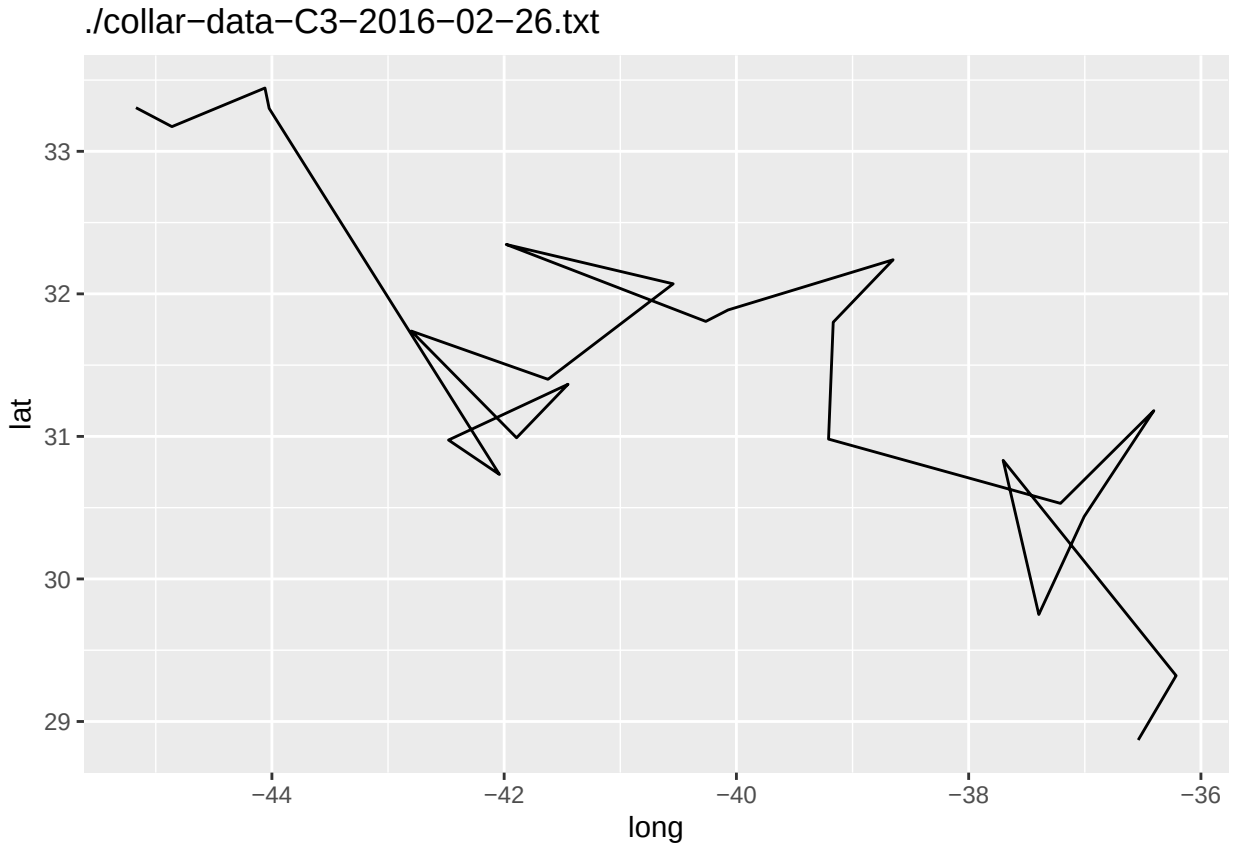
2c.

```
## # A tibble: 4 x 2
##   species      avg_mass
##   <chr>           <dbl>
## 1 Ankylosauria   46819.
## 2 Sauropoda      16104.
## 3 Stegosauria    31924.
## 4 Theropoda      45572.
```

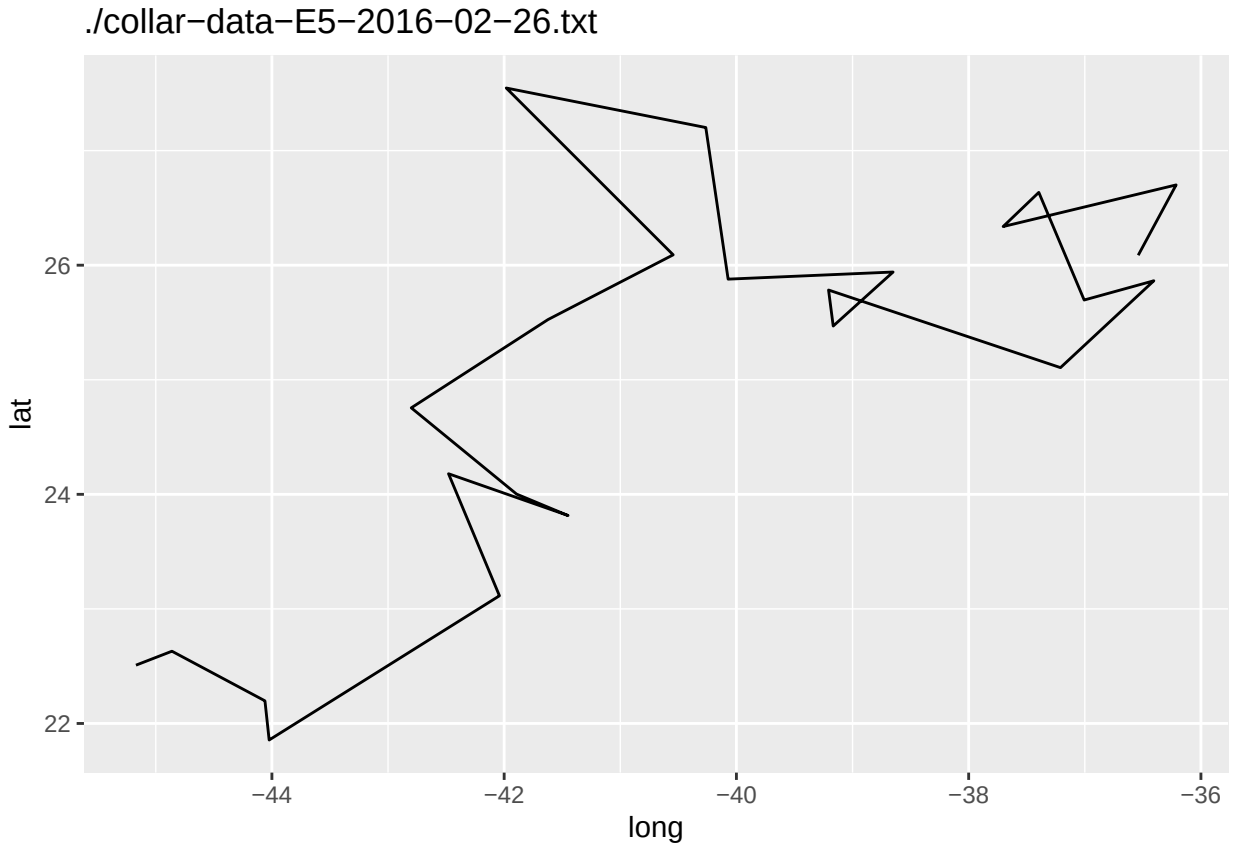
3. Multi-file Analysis (40 pts)

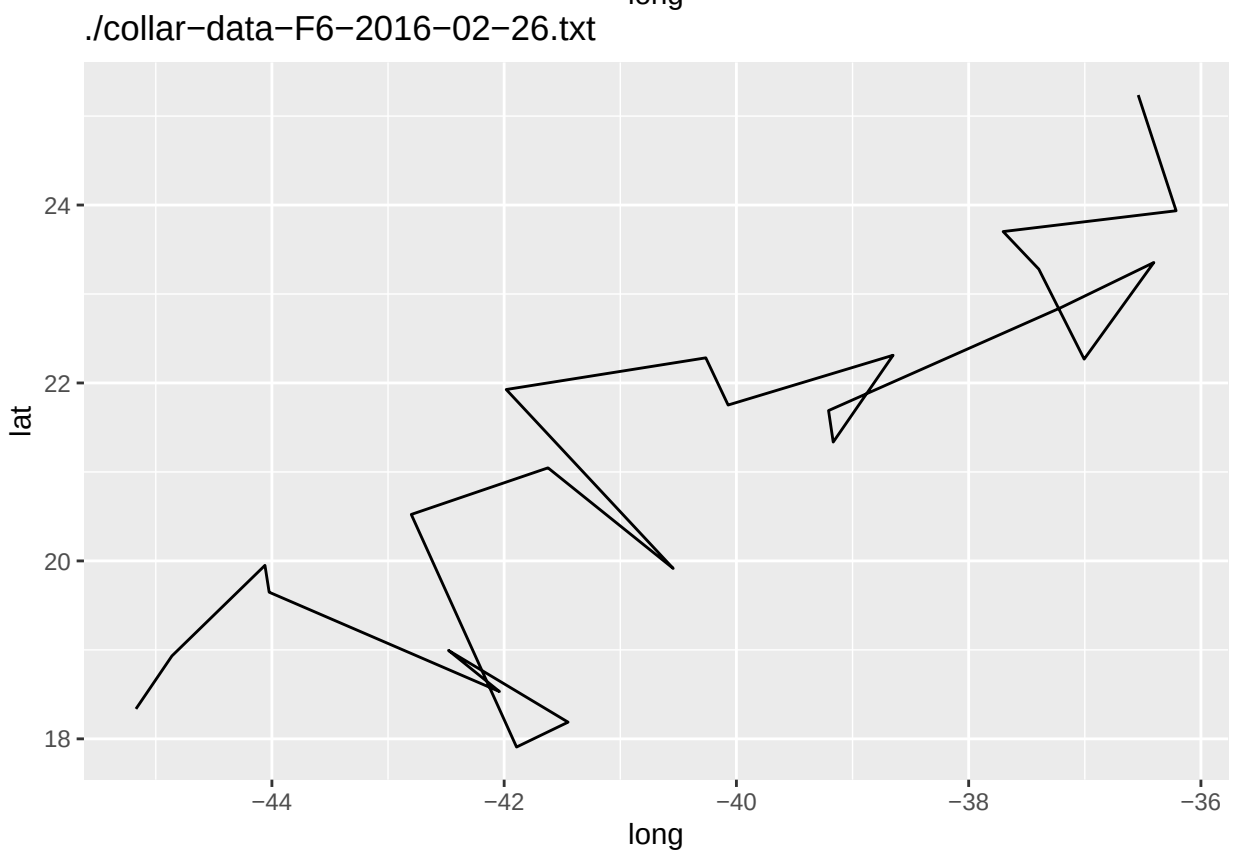


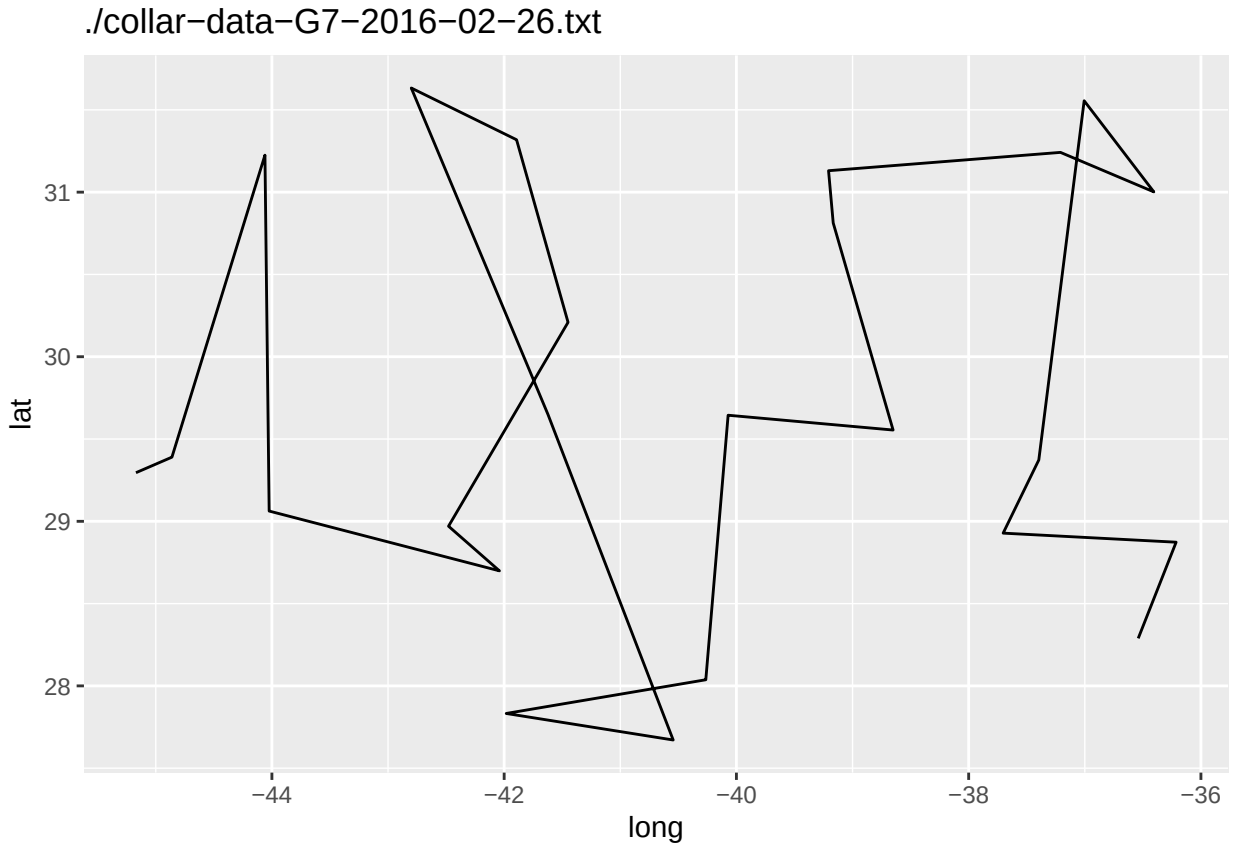
3a.

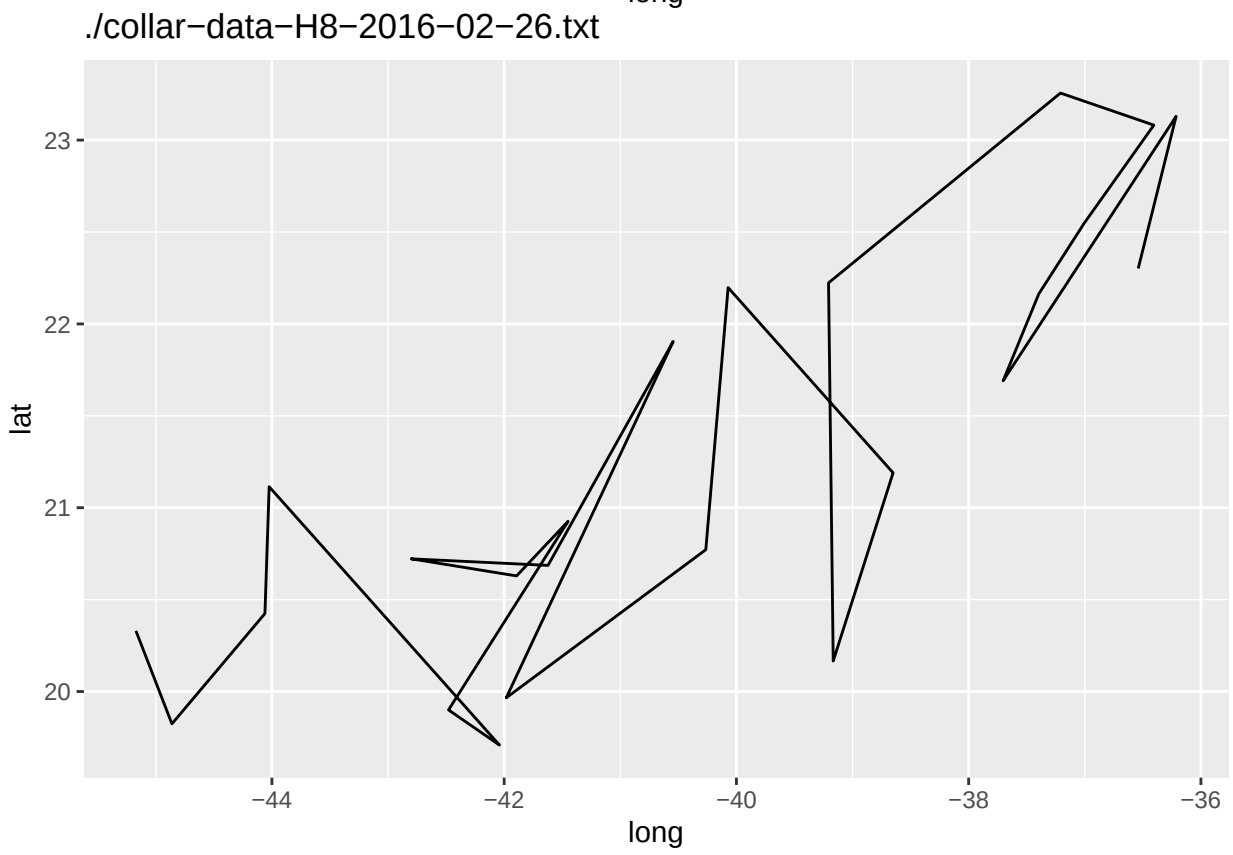

./collar−data−C3−2016−02−26.txt
lat
long
33
32
31
30
29
−44
−42
−40
−38
−36

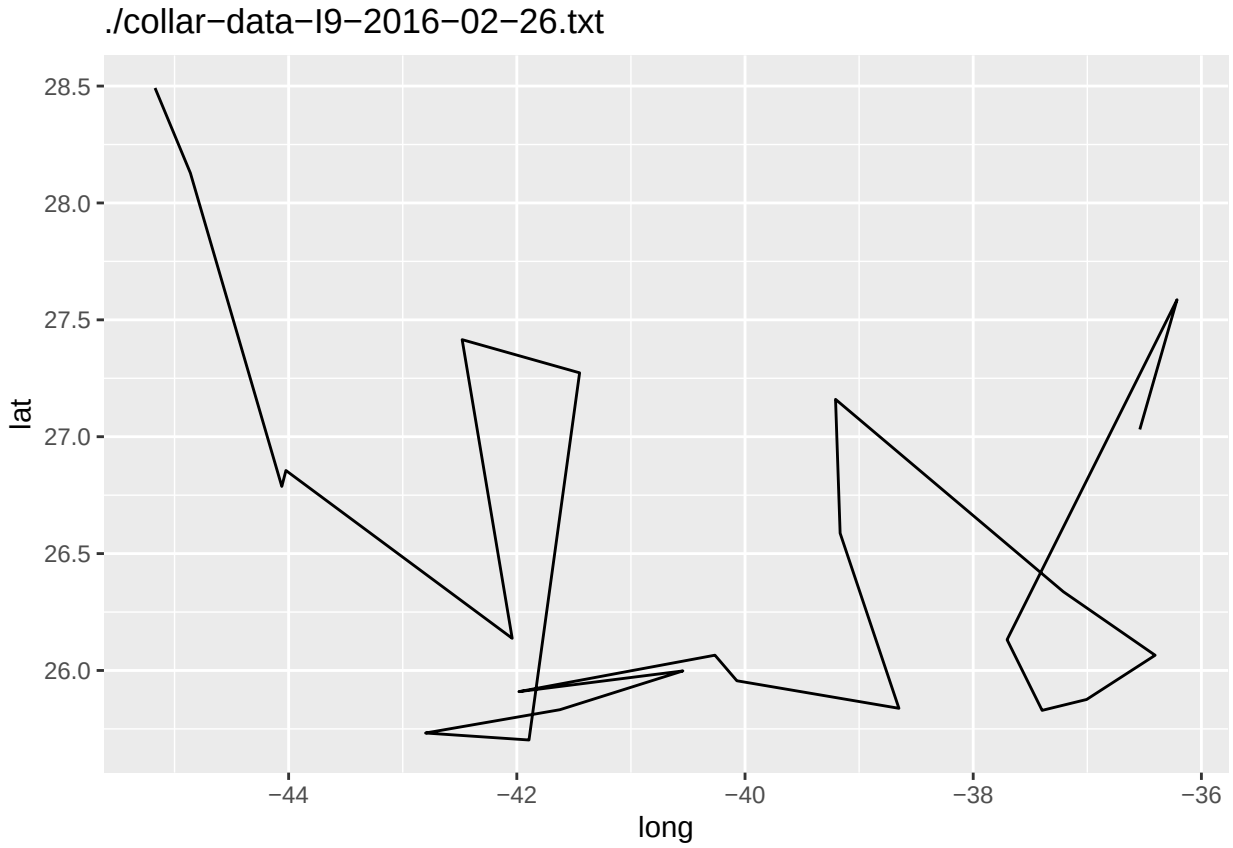

./collar−data−D4−2016−02−26.txt
lat
long
24
23
22
−44
−42
−40
−38
−36


./collar−data−E5−2016−02−26.txt
26
24
22
lat
−44
−42
−40
−38
−36
long


./collar−data−F6−2016−02−26.txt
24
22
20
18
lat
−44
−42
−40
−38
−36
long


./collar−data−G7−2016−02−26.txt
31
30
29
28
lat
−44
−42
−40
−38
−36
long

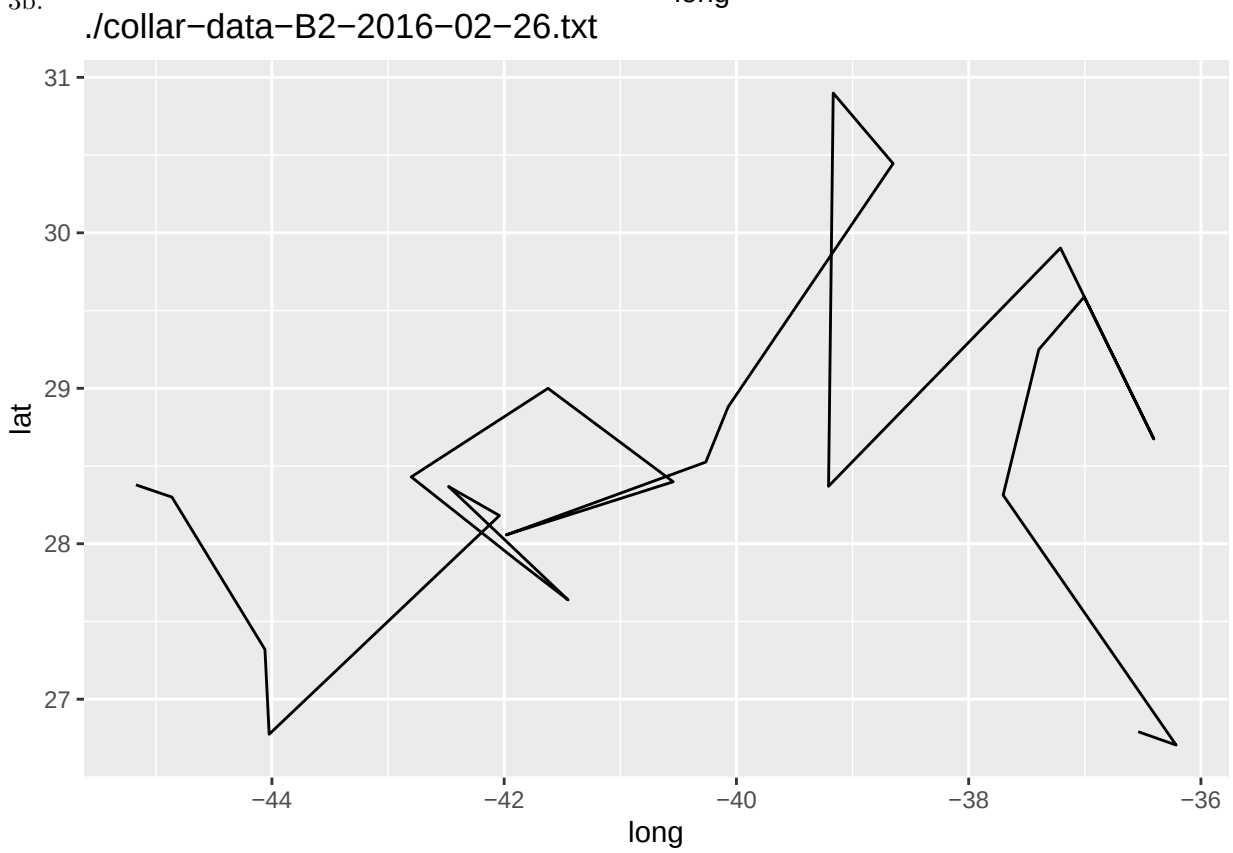

./collar−data−H8−2016−02−26.txt
23
22
21
20
lat
−44
−42
−40
−38
−36
long

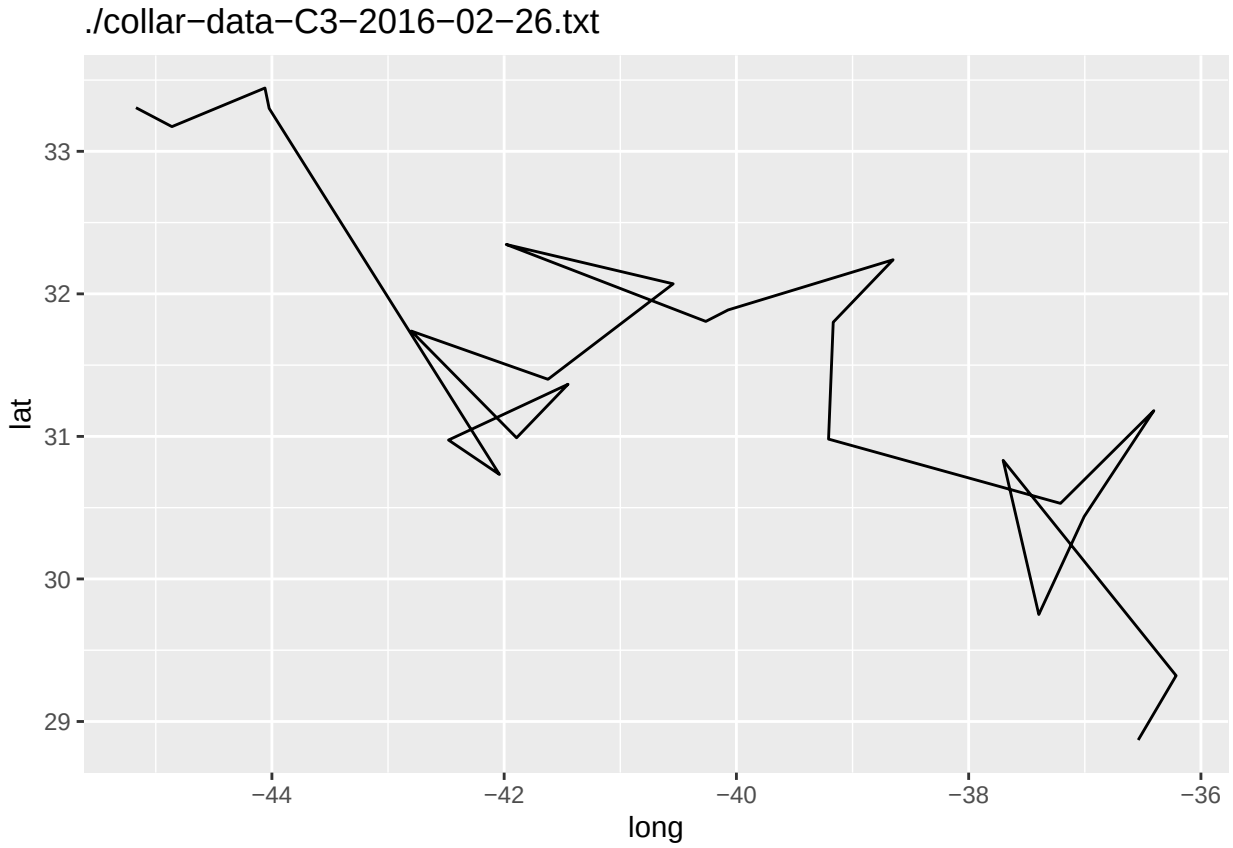

./collar−data−I9−2016−02−26.txt
28.5
28.0
27.5
27.0
26.5
26.0
lat
−44
−42
−40
−38
−36
long

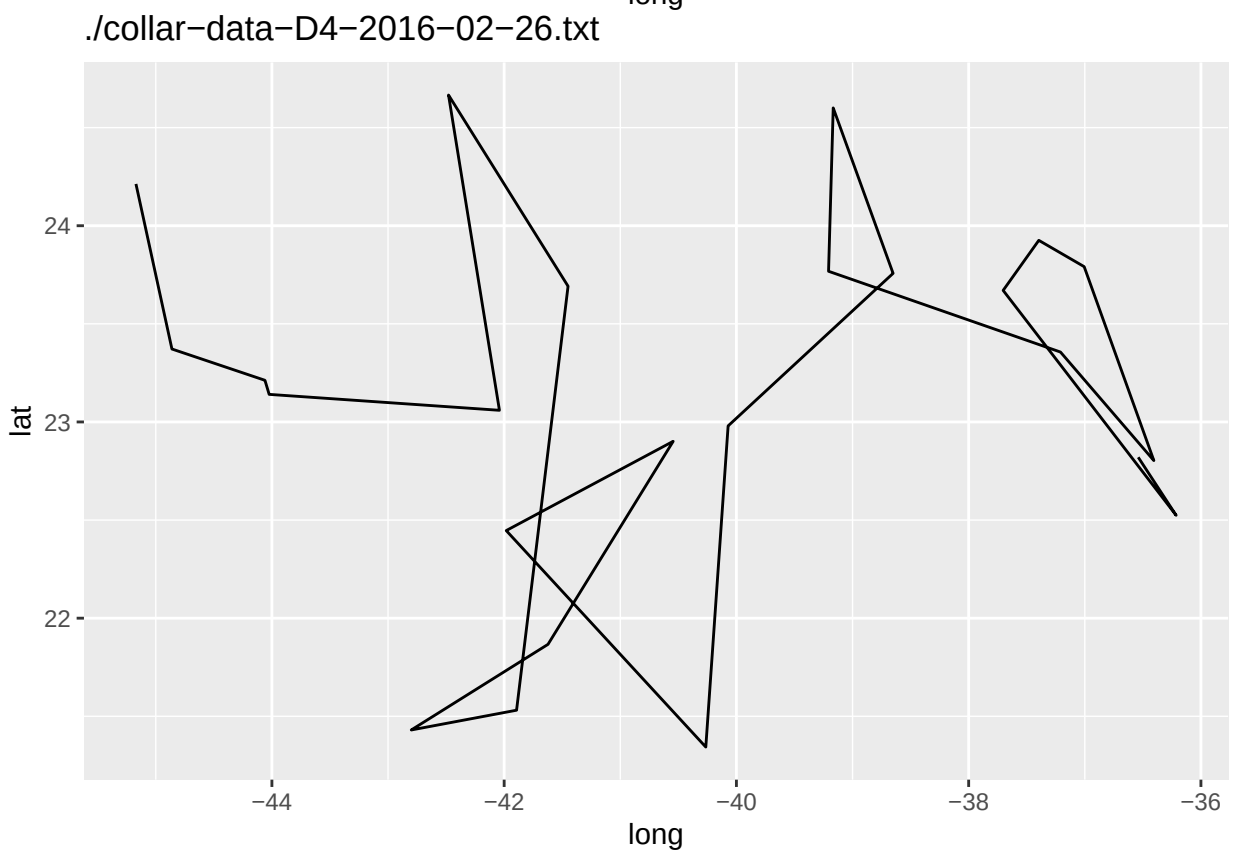

./collar−data−J10−2016−02−26.txt
27
26
25
lat
−44
−42
−40
−38
−36
long

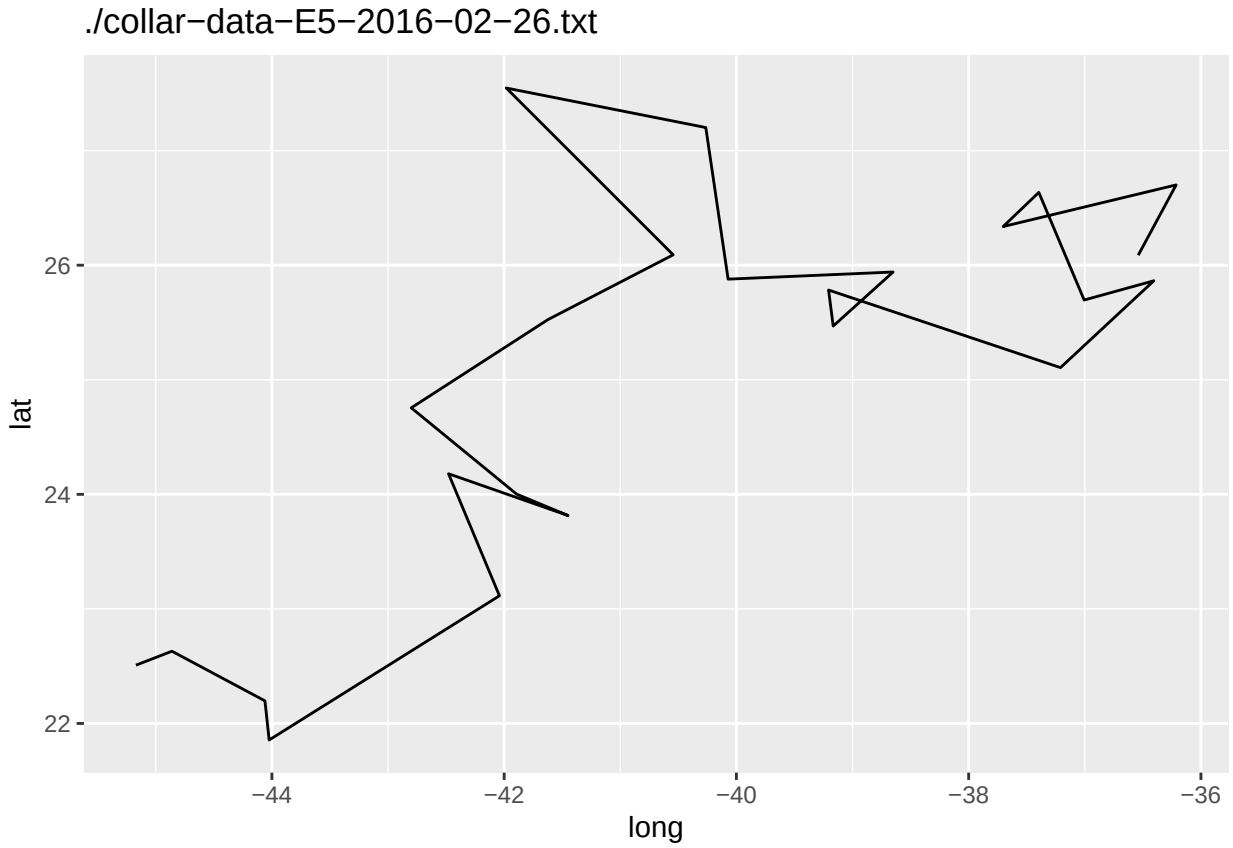

./collar−data−A1−2016−02−26.txt
32
30
28
26
lat
−44
−42
−40
−38
−36
long

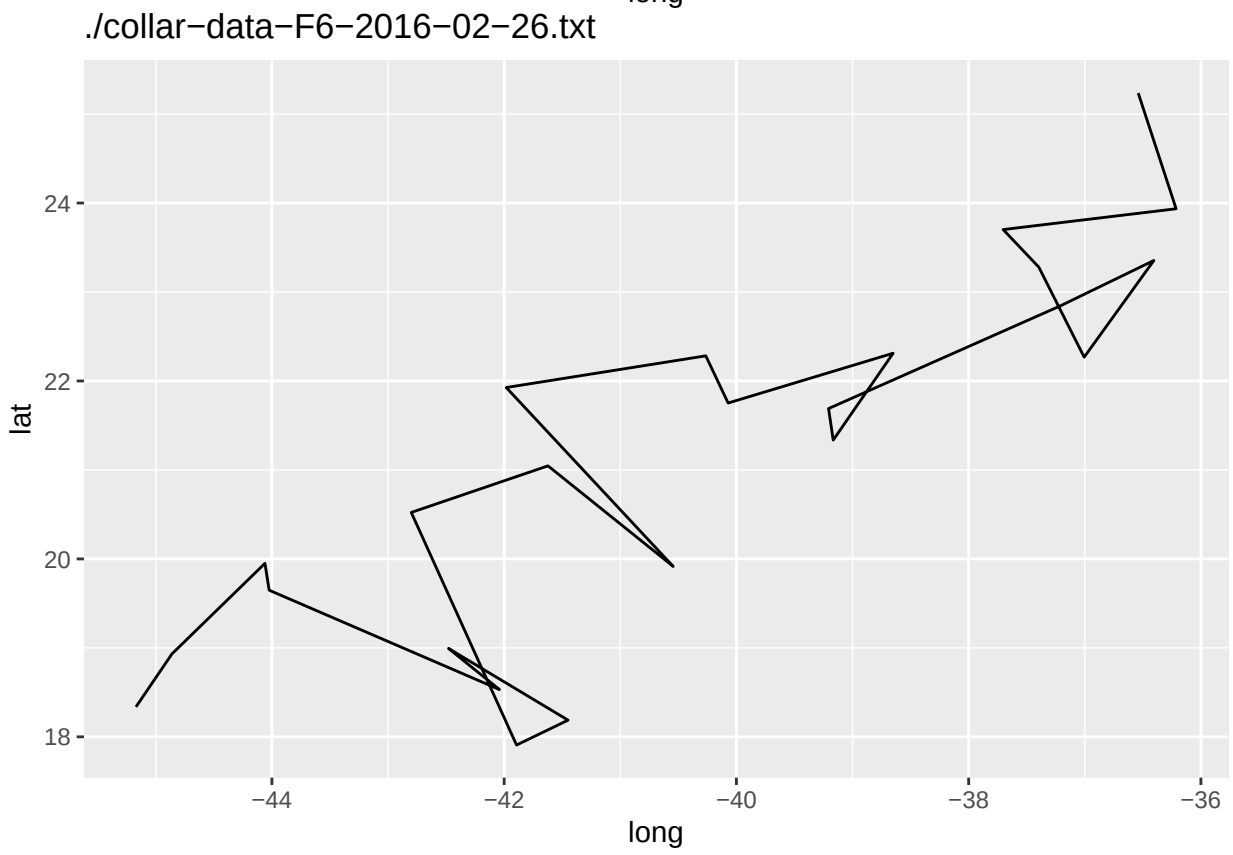
3b.

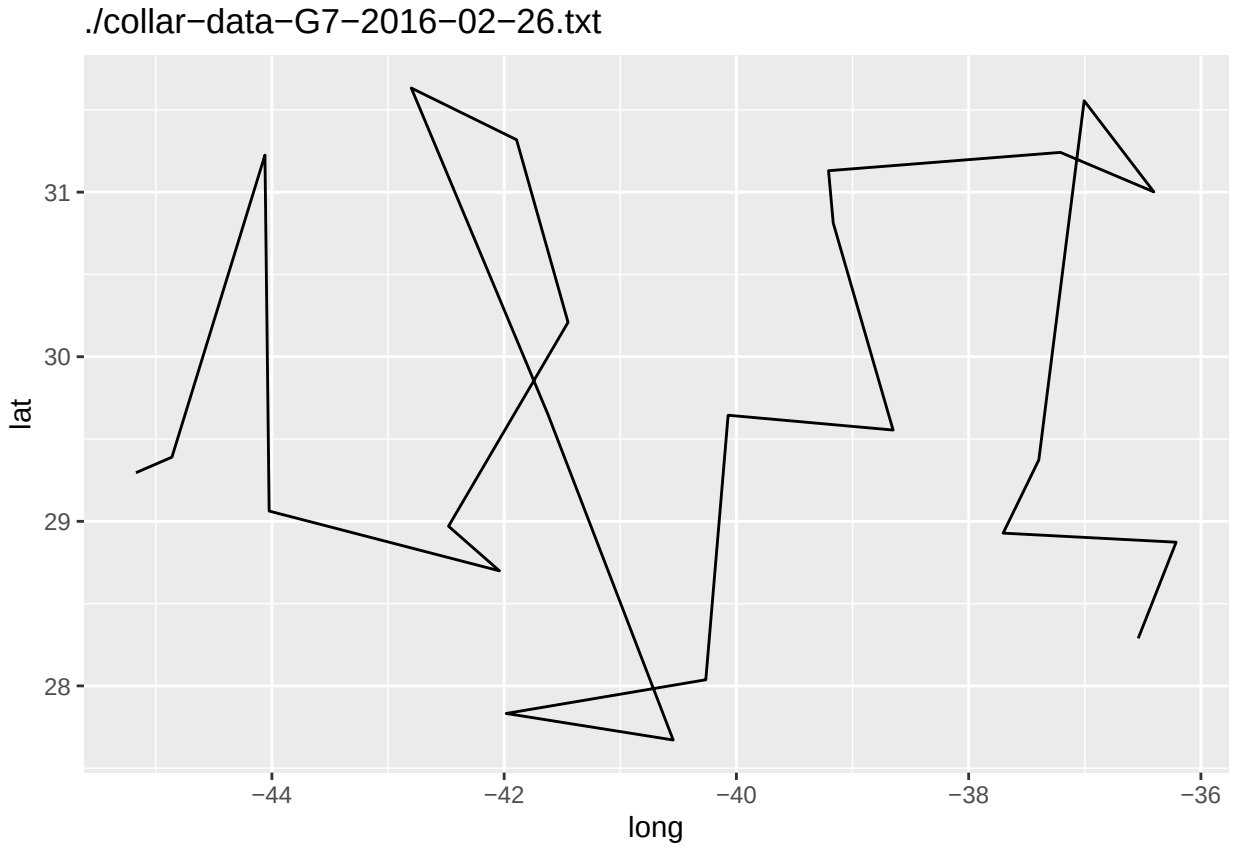

./collar−data−B2−2016−02−26.txt
31
30
29
28
27
lat
−44
−42
−40
−38
−36
long

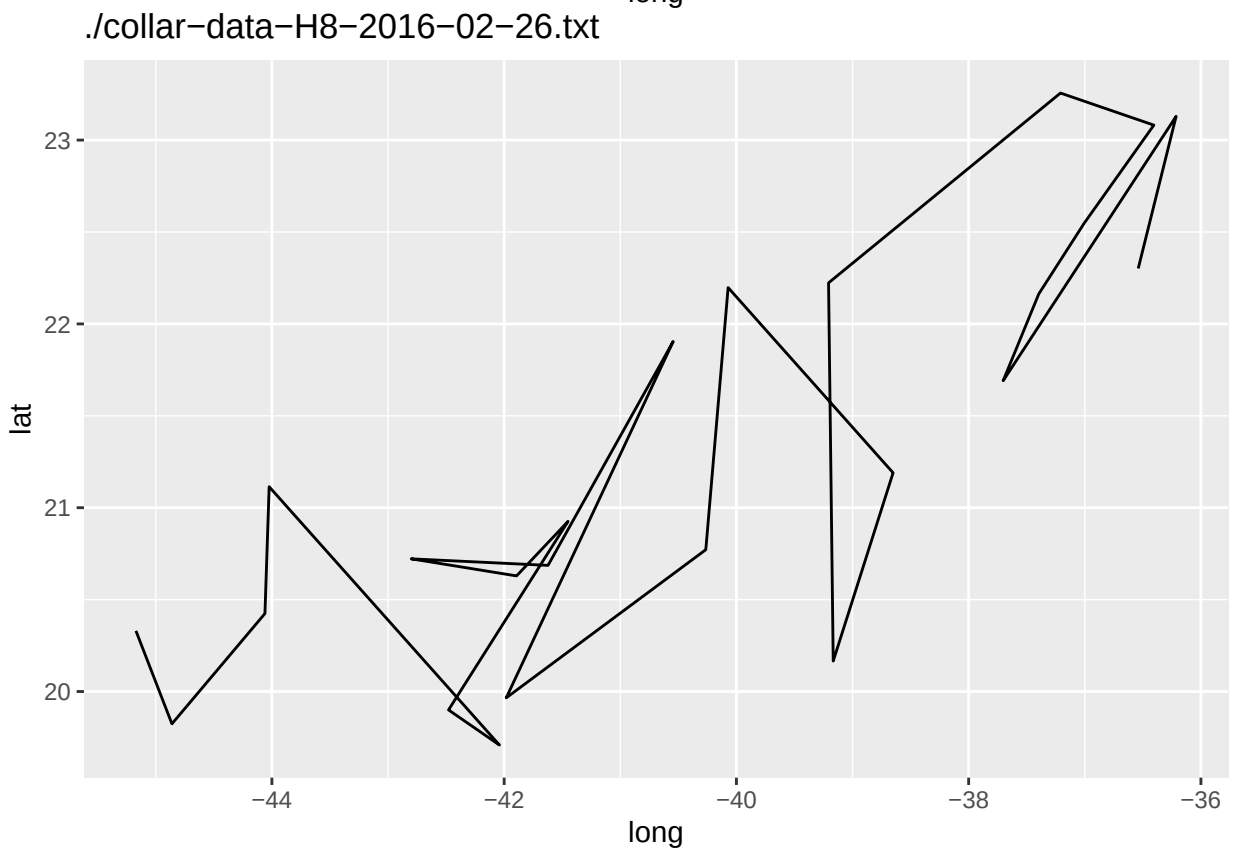

./collar−data−C3−2016−02−26.txt
lat
long
33
32
31
30
29
−44
−42
−40
−38
−36

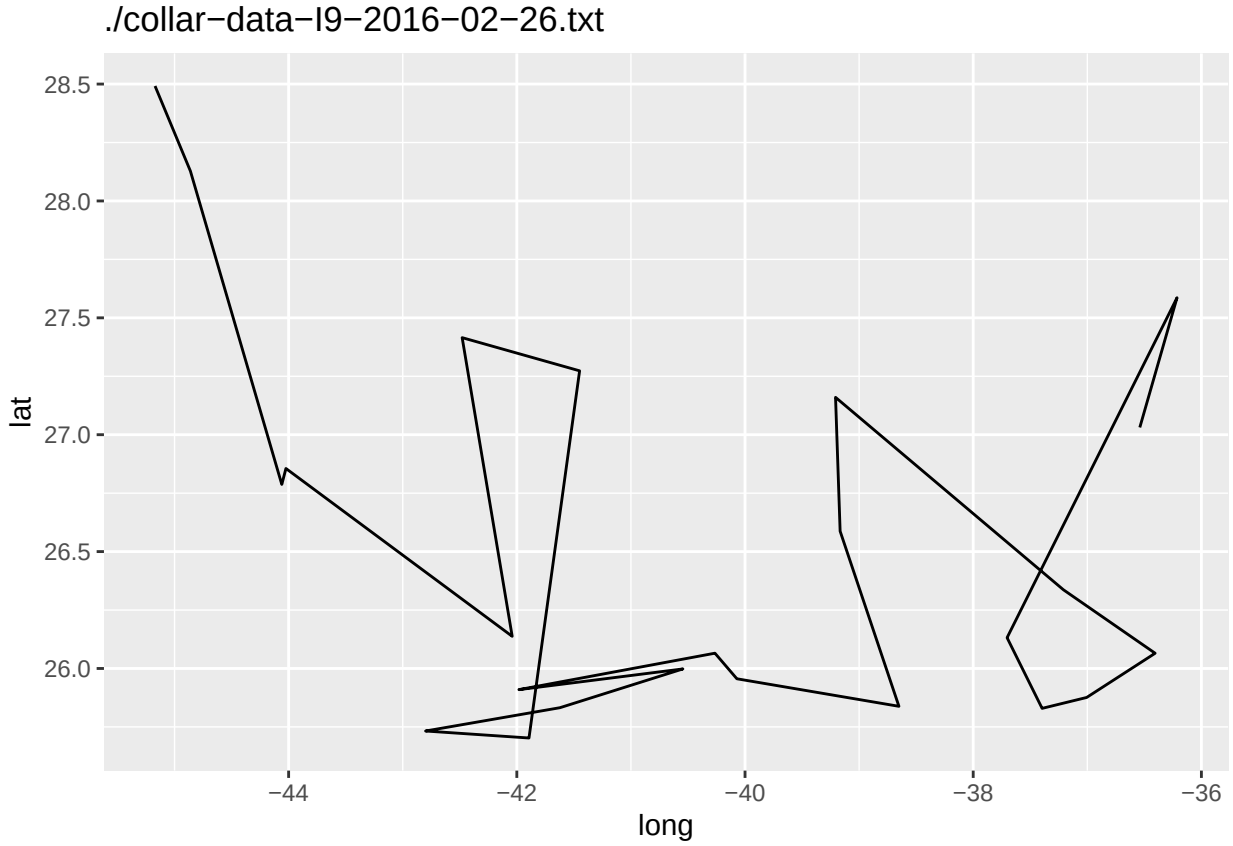

./collar−data−D4−2016−02−26.txt
lat
long
24
23
22
−44
−42
−40
−38
−36

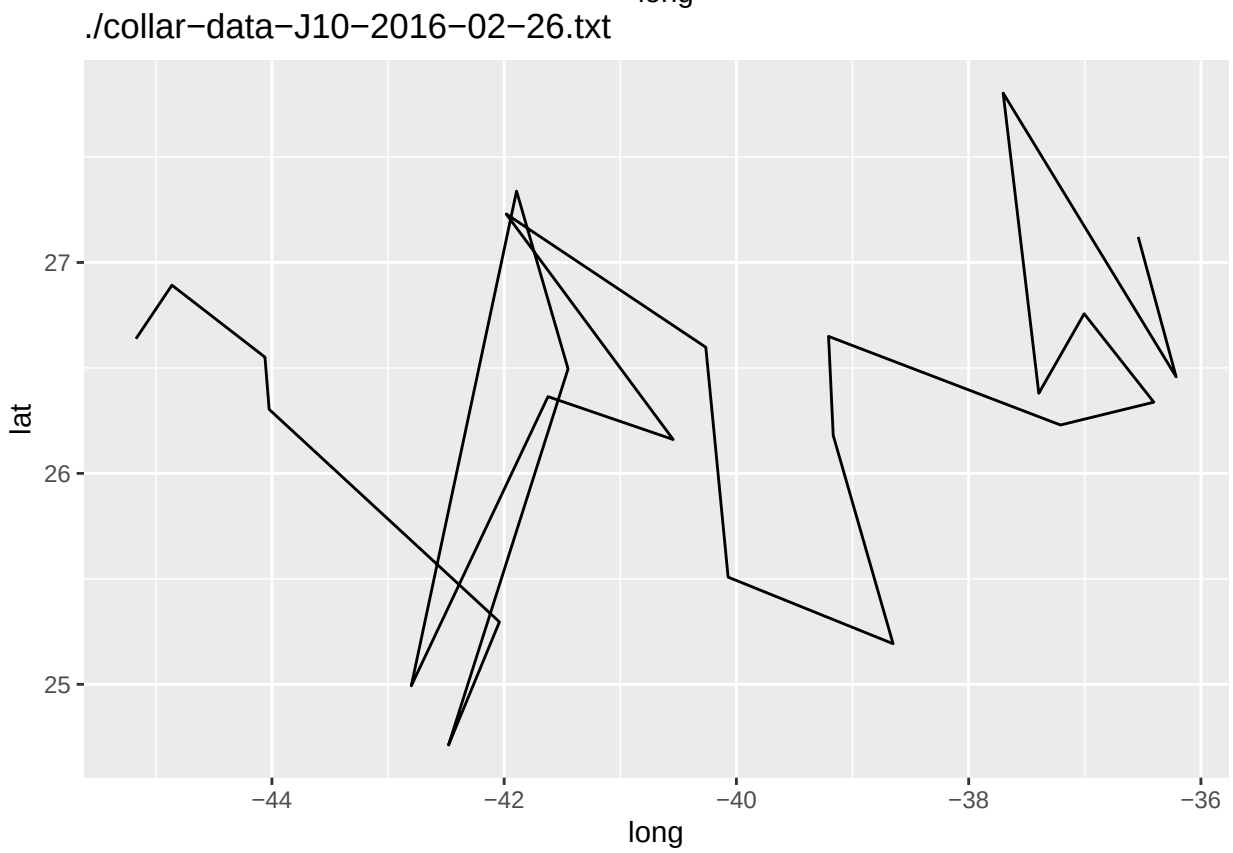

./collar−data−E5−2016−02−26.txt
lat
26
24
22
−44
−42
−40
−38
−36
long


./collar−data−F6−2016−02−26.txt
lat
24
22
20
18
−44
−42
−40
−38
−36
long


./collar−data−G7−2016−02−26.txt
31
30
29
28
lat
−44
−42
−40
−38
−36
long


./collar−data−H8−2016−02−26.txt
23
22
21
20
lat
−44
−42
−40
−38
−36
long


./collar−data−I9−2016−02−26.txt
28.5
28.0
27.5
27.0
26.5
26.0
lat
−44
−42
−40
−38
−36
long


./collar−data−J10−2016−02−26.txt
27
26
25
lat
−44
−42
−40
−38
−36
long

```
##                           file_name  max_lat  min_lat observations
## 1   ./collar-data-A1-2016-02-26.txt 31.76912 25.21080           24
## 2   ./collar-data-B2-2016-02-26.txt 30.89907 26.70509           24
## 3   ./collar-data-C3-2016-02-26.txt 33.44421 28.86998           24
## 4   ./collar-data-D4-2016-02-26.txt 24.66598 21.34315           24
## 5   ./collar-data-E5-2016-02-26.txt 27.54663 21.85565           24
## 6   ./collar-data-F6-2016-02-26.txt 25.23623 17.90788           24
## 7   ./collar-data-G7-2016-02-26.txt 31.63272 27.67120           24
## 8   ./collar-data-H8-2016-02-26.txt 23.25601 19.70875           24
## 9   ./collar-data-I9-2016-02-26.txt 28.49172 25.70252           24
## 10 ./collar-data-J10-2016-02-26.txt 27.80325 24.71200           24
```

4. DNA or RNA (20 points)

4b.

```
## [1] "DNA"
```

```
## [1] "RNA"
```

```
## [1] "UNKNOWN"
```

4c.

```
## [1] "DNA"
## [1] "RNA"
## [1] "UNKNOWN"
## [1] "RNA"
## [1] "RNA"
```

4d.

```
## Warning: ‘as.tibble()‘ was deprecated in tibble 2.0.0.
## i Please use ‘as_tibble()‘ instead.
## i The signature and semantics have changed, see ‘?as_tibble‘.
## This warning is displayed once every 8 hours.
## Call ‘lifecycle::last_lifecycle_warnings()‘ to see where this warning was
## generated.
```

```
## # A tibble: 5 x 2
##   type    sequence
##   <chr>   <chr>
## 1 DNA     ttgaatgccttacaactgatcattacacaggcggcatgaagcaaaaatatactgtgaaccaatgcaggcg
## 2 RNA     gauuauuccccacaaagggaguggguauuaggagcugcaucauuuacaagagcagaauguuucaaaugcau
## 3 UNKNOWN gaaagcaagaaaaggcaggcgaggaagggaagaaggggggggaaacc
## 4 RNA     guuuccuacaguauuugaugagaaugagaguuuacuccuggaagauaaauauuagaauguuuacaacugc~
## 5 RNA     gauaaggaagaugaagacuuucaggaaucuaauaaaaugcacuccaugaauggauucauguaugggaau~
```

4e. OPTIONAL

```
##                                 ttgaatgccttacaactgatcattacacaggcggcatgaagcaaaaatatactgtgaaccaatgcaggcg
##                                                                                                  "DNA"
##                                 gauuauuccccacaaagggaguggguauuaggagcugcaucauuuacaagagcagaauguuucaaaugcau
##                                                                                                  "RNA"
##                                                                  gaaagcaagaaaaggcaggcgaggaagggaagaaggggggggaaacc
##                                                                                               "UNKNOWN"
## guuuccuacaguauuugaugagaaugagaguuuacuccuggaagauaauauuuagaauguuuacaacugcaccugaucagguggauaaggaagaugaagacu
##                                                                                                  "RNA"
##                       gauaaggaagaugaagacuuucaggaaucuaauaaaaaugcacuccaugaauggauucauguauggggaaucagccgggu
##                                                                                                  "RNA"
```

4f. OPTIONAL

```
## # A tibble: 5 x 2
## # Rowwise:
##   type    sequences
##   <chr>   <chr>
## 1 DNA     ttgaatgccttacaactgatcattacacaggcggcatgaagcaaaaatatactgtgaaccaatgcaggcg
## 2 RNA     gauuauuccccacaaagggaguggguauuaggagcugcaucauuuacaagagcagaauguuucaaaugcau
## 3 UNKNOWN gaaagcaagaaaaggcaggcgaggaagggaagaaggggggggaaacc
## 4 RNA     guuuccuacaguauuugaugagaaugagaguuuacuccuggaagauaauauuuagaauguuuacaacugc~
## 5 RNA     gauaaggaagaugaagacuuucaggaaucuaauaaaaaugcacuccaugaauggauucauguauggggaau~
```