```
<head>
    <meta charset="utf-8">
    <title>Why R</title>
    <meta name="description" content="A epistemological answer to the obvious question. Why would anyon
    <meta name="author" content="Andy Choens, MSW">
    <meta name="apple-mobile-web-app-capable" content="yes" />
    <meta name="apple-mobile-web-app-status-bar-style" content="black-translucent" />
    <meta name="viewport" content="width=device-width, initial-scale=1.0, maximum-scale=1.0, user-scala</pre>
    <link rel="stylesheet" href="css/reveal.css">
    <link rel="stylesheet" href="css/theme/solarized.css" id="theme">
    <!-- Code syntax highlighting -->
    <link rel="stylesheet" href="lib/css/zenburn.css">
    <!-- Printing and PDF exports -->
    <script>
     var link = document.createElement( 'link' );
    link.rel = 'stylesheet';
     link.type = 'text/css';
     link.href = window.location.search.match( /print-pdf/gi ) ? 'css/print/pdf.css' : 'css/print/paper
    document.getElementsByTagName( 'head' )[0].appendChild( link );
    </script>
    <!--[if lt IE 9]>
    <script src="lib/js/html5shiv.js"></script>
    <![endif]-->
</head>
<body>
    <div class="reveal">
       <!-- Any section element inside this container is displayed as a slide -->
       <div class="slides">
           <section>
               <div style="display:inline-block; margin:0 auto; float:left; width:50%;">
                   <h1>Why</h1>
               <div style="display:inline-block; margin:0 auto; float:right; width:50%;">
                   <img src="svg/r.svg" height="75%" width="75%">
                       <small>Presented by: Andy Choens, MSW</small>
                   </div>
               <aside class="notes">
                   According
                       to <a href="https://en.wikipedia.org/wiki/R_%28programming_language%29">Wikiped
```

```
<code>
R is a programming language and software environment for
statistical computing and graphics. The R language is
widely used among statisticians and data miners for
developing statistical software and data
analysis. Polls, surveys of data miners, and studies of
scholarly literature databases show that R's popularity
has increased substantially in recent years.
         </code>
         >
             This is not a WHY NOT SAS presentation. This is a Why R
             presentation. By the end of the presentation, I hope to
             have demonstrated that R is a relevant language in the
             work we are doing today and will be even more relevant
             in years to come.
         </aside>
  </section>
  <!-- URLs to Remember ----- -->
  <section>
     <h2>URLs To Remember:</h2>
     <a href="http://choens.github.io/why-r/">http://choens.github.io/why-r/</a>
         <a href="https://github.com/choens/why-r/">https://github.com/choens/why-r/</a>
         <a href="http://cran.r-project.org/">http://cran.r-project.org/</a>
     <aside class="notes">
         <l
             <
                 The 1st link will let you see the presentation, as
                 it was presented here today.
             <1i>
                 The 2nd link will let you see all of the underlying
                 code used in today's presentation.
             <1i>>
                 Later in the presentation, I will show some demo
                 output files and demo code files. These resources can be
                 found in the examples folder on GitHub which you can
                 access by following the 2nd link.
             <
                 You are free to use any of these materials. This
                 presentation, slides, demo files, etc. are licensed
```

Any time you see text in this color of blue. It is a link. If you click it,

under the XXXX license.

you will go somewhere fun.

```
</aside>
</section>
<section>
  <h2>Bad Reasons To Learn / Use R </h2>
  <div style=" display:inline-block; margin:0 auto; float:left; width:60%;">
        Some
           guy with a beard likes it.
        Someone
           on the Internet said it was great.
        You
           read about it somewhere.
     </div>
  <div style="display:inline-block; margin:0 auto; float:right; width:40%;">
     <img src="svg/bad.svg" height="100%" width="100%">
  <aside class="notes">
     ul>
        Advance: 3x
        Let's get this out of the way. There are several bad reasons to use any too
        In the context of today's discussion, I came up with the following bad reas
     </aside>
</section>
<section>
  <h2>Good Reasons to Learn / Use R</h2>
  <div style="display:inline-block; margin:0 auto; float:left; width:40%;">
     <img src="svg/good.svg" height="100%" width="100%">
  </div>
  <div style="display:inline-block; margin:0 auto; float:right; width:60%;">
     <111>
        Ecosystem
        Literate Programming
        Open Source
        Future Opportunities
        Don't Take My Word For It
     </div>
  <aside class="notes">
     <l
        <
           Advance: The ecosystem of tools around R is nothing
           short of AMAZING and it is getting more amazing every
           single day, because ANYONE can contribute to the
           ecosystem.
        <
```

As of February 2, 2015: C the primary R repository, contains 6,391 packages. These are peer reviewed packages, developed by the community. <a href="http://www.bioconductor.org/packages/release/Bio an additional 936 packages. Just let that sink in. And while you do:

<

I'd like to tell you about where you can use R. You can use R from within SAS. You can use R from within SPSS. You can use R from within MATLAB, Oracle, Postgres, and Vertica. R skills are portable skills. Microsoft recently bought the R-based consulting firm Revolution Analytics, which could result in better integration of R in Microsoft produces such as SQL Server or even Office.

<1i>>

Advance: This presentation isn't long enough to take a deep dive into Literate Programming, but I do want to mention this idea briefly.

<

Literate programming is an idea from the 1980s, to interweave natural language and computer programming language. I'll give you concrete examples of this later, but the upshot is that it makes it possible to write a program that is both self-documenting AND is able to compile to a finished, publishable manuscript or report. No more cutting and pasting for an hour. No more messing up a table because you forgot a decimal point on the 9th row of the 3rd column. A self-documenting, transparent manuscript that can be published.

>

This is an idea from the 80s. The book, Literate Programming was published in 1984. This is an idea that is very influential in the R community and is not only considered to be current and cutting edge, it is something that people are working on today to make it easier and faster to produce.

>

Advance: One good reason to use R is because it is open source. Those 7,000+ packages? Only possible because R is open source and engages with the community.

>

IBM is unlikely to make integrate SAS into SPSS. They are competitors. Besides, owning SAS and SPSS on the same machine without a student discout is . . . prohibitively expensive. R is considered safe, common ground, because it is not a commercial entity.

```
<1i>>
               Advance: A perfectly reasonable, self-interested
               reason to use any tool professionally is because you
               think it will lead to future job opportunities.
           <
               Knowing how to use SAS effectively is a good way to
               make yourself eligible for a large number of job
               opportunities. Knowing how to use both R <emphasis>and</emphasis>
               SAS effectively makes you relevant to even more future
               job opportunities. Please note: I said AND. Not
               or. Not instead of. I said AND.
           <1i>
               Advance: But please. Don't take my word for it. See
               what some of these people have to say.
       </aside>
</section>
<!-- Section: Ecosystem ========= -->
   <div style="display:inline-block; margin:0 auto; float:left; width:50%;">
       <h1>Ecosystem</h1>
   </div>
   <div style="display:inline-block; margin:0 auto; float:right; width:50%;">
       <img src="svg/r.svg" height="75%" width="75%">
   </div>
   <aside class="notes">
       <l
           <
               The R ecosystem is vast. Because anyone can alter it,
               extend it or otherwise bend it to their will, they
               have.
           <
               R is for social science research, genetics,
               econometrics and betting on Football (both types).
           <1i>>
               I'm going to give you a peek into this vast
               ecosystem. But all I can't even begin to explain it
               all. There's too much out there and quite frankly, I
               don't understand half of it.
           >
               But don't worry. Neither will you.
           </aside>
</section>
```

```
<section>
   <h2>Packages</h2>
   The capabilities of R are extended through
       user-created packages, which allow specialized
       statistical techniques, graphical devices (<a href="http://cran.r-project.org/web/p">http://cran.r-project.org/web/p</a>
       import/export capabilities, reporting tools
       (<a hre="http://cran.r-project.org/web/packages/knitr/index.html">knitr</a>, <a hre
       etc. These packages are developed primarily in
       R, and sometimes in Java, C, C++ and Fortran.
   </section>
<section>
   <h2>Popular Package Places</h2>
   <l
       >
            <a href="http://cran.r-project.org/">Comprehensive R Archive Network (CRAN)</a>
       <1i>>
            <a href="http://www.bioconductor.org/">Bioconductor</a>
       <1i>>
           <a href="http://www.omegahat.org/">Omegahat</a>
       <1i>>
           <a href="http://r-forge.r-project.org/">R-Forge</a>
       >
            <a href="https://github.com/">GitHub</a>
       <aside class="notes">
       <l
            <1i>
               CRAN: The official R
               repositories. Wide ranging collection
               of packages. The growth of packages on
               this repo alone is staggering, and
               only increasing.
            <
               Bio: Provides packages for the
               analysis of genomic data. Bioconductor
               and R are the two repos I use the
               most. Packages here tend to be of very
               high quality, are updated, and well
               accepted by the broader
               community. There are over 7,000
               packages at these two sites alone.
           <1i>
               Omegahat: Older repo. Predates
               CRAN. Has more bleeding edge
               packages. For example, there is a
```

```
package to integrate R and Firefox,
               allowing you to run R within a
              webpage. Some duplication with CRAN.
           <
              R-Forge: Clone of the venerable Source
              Forge site. This page is mostly
              packages not yet approved for
              distribution via CRAN, or is the beta
              version of the next release. Use most
               of this at your own risk. Some very
               interesting packages.
           <
              GH: OMG. There are over 20,000 public
              repositories on GitHub, which appear
              to be R source code. Some of these are
              packages developed by the author for
              personal use. Some are analyses. Some
              are for various classes. The diversity
              here is incredible. I have no idea how
              to count only those repos on GH that
              are packages. I looked around, and
              nobody else does either.
           </aside>
</section>
<!-- To Infinity and Beyond! ----- -->
<section>
   <h2>To Infinity & Beyond!</h2>
   <img src ="img/fig_8_cran.png" width="640" style="position: relative; top:-150px; z-ind</pre>
   <a href="http://r4stats.com/articles/popularity/">r4stats.com/articles/popularity/<
   <aside class="notes">
       ul>
           <
               I want to plug the r4stats page, which is where I
               got a lot of the information I used in this
              presentation. The amount of information there is
               staggering and quite frankly somewhat obsessive.
           <
               R-Core is a powerful stats package. But I don't use
              R because of the brilliance of R-Core. I use R
              because of the packages.
           <1i>
              In 2009 John Fox wrote some code that calculates the
              number of R packages available on CRAN for each
              release of R. This image does not include
```

```
Bioconductor or any other R package
             repositories. Just the official CRAN packages.
          <1i>>
             For those keeping track at home, the super-imposed
             model is a quadratic parabola. More importantly is
             the continuing, rapid rise.
          <
             This single image is quite possibly one of the best
             reasons to use {\tt R} and one of the best reasons to run
             for the hills. With this many options available,
             there is duplication. Some packages are
             brilliant. Some are bizarre. Some are probably
             downright stupid. To leverage this incredible
             resource takes time and it takes expertise.
          </aside>
</section>
<!-- Two Examples ----- -->
<section>
   <h2>Packages Examples</h2>
   <h3><a href="http://cran.r-project.org/web/packages/dplyr/index.html">dplyr
          <h3><a href="http://cran.r-project.org/web/packages/ggplot2/index.html">ggp
          <h3><a href="http://cran.r-project.org/web/packages/sqlutils/index.html">sq
          <small>A fast, consistent tool for
             working with data frame like objects,
             both in memory and out of memory.</small>
          </t.d>
          <small>An implementation of the
             grammar of graphics in R. It combines
             the advantages of both base and
             lattice graphics: conditioning and
             shared axes are handled automatically,
             and you can still build up a plot step
             by step from multiple data sources. It
             also implements a sophisticated
             multidimensional conditioning system
             and a consistent interface to map data
             to aesthetic attributes.</small>
```

```
<t.d>
              <small>This package provides utilities
              for working with a library of SQL
              files.</small>
          <aside class="notes">
       <l
          <1i>>
              I'm not even going to pretend like I know what some
              of these packages do. Yes. There is actually a
              package on CRAN dedicated to fisheries science. And,
              I genuinely have NO idea what the ChIPpeakAnno
              package does. But with over 7,000 packages, even a
              social worker like me is bound to find something
              useful.
          </section>
<!-- Diverse Examples ----- -->
<section>
   <h2>Package ????</h2>
   <h3><a href="http://cran.r-project.org/web/packages/kobe/index.html">kobe-p
          <h3><a href="http://www.bioconductor.org/packages/release/bioc/html/ChIPpea
          <small>The tuna Regional Fisheries Management
                 Organisations (tRFMOs) use a common framework for
                 providing scintific advice, i.e. the Kobe II
                 Framework. This is based on maintaining fishing
                 mortal- ity below FMSY and stock biomass above
                 BMSY. This package provides methods for summarising
                 results from stock assessments and Management
                 Strategy Evaluations in the Kobe format.</small>
          <small>The package includes functions to retrieve
                 the sequences around the peak, obtain enriched Gene
                 Ontology (GO) terms, find the nearest gene, exon,
                 miRNA or custom features such as most conserved
                 elements and other transcription factor binding
                 sites supplied by users. Starting 2.0.5, new
                 functions have been added for finding the peaks with
                 bi-directional promoters with summary statistics
                 (peaksNearBDP), for summarizing the occurrence of
```

```
motifs in peaks (summarizePatternInPeaks) and for
                adding other IDs to annotated peaks or enrichedGO
                (addGeneIDs). This package leverages the biomaRt,
                IRanges, Biostrings, BSgenome, GO.db, multtest and
                stat packages.</small>
         <aside class="notes">
      <l
          <1i>
             I'm not even going to pretend like I know what some
             of these packages do. Yes. There is actually a
             package on CRAN dedicated to fisheries science. And,
             I genuinely have NO idea what the ChIPpeakAnno
             package does. But with over 7,000 packages, even a
             social worker like me is bound to find something
             useful.
         </section>
<section>
   <h2>Commercial Support</h2>
   <div style="display:inline-block; margin:0 auto; float:left; width:33%;">
      <h3>Programming Languages</h3>
      <l
         <small>
             C
             Java
             JMP
             Mathematica
             MATLAB
             Python
             SAS
             SPSS
             Statistica
             tableau
          </small>
      </div>
   <div style="display:inline-block; margin:0 auto; width:33%;">
      <h3>Database <br/> Vendors</h3>
      <l
         <small>
             Hadoop
             Oracle
             PostgreSQL
             Vertica
          </small>
      </div>
```

```
<h3>Business Intelligence</h3>
       ul>
           <small>
               Alteryx
              Jaspersoft
               Oracle Business Intelligence Enterprise Edition
               Pentaho
               SAP (and SAP HANA)
           </small>
       </div>
   <aside class="notes">
       <l
           <
              R bills itself as the lingua franca of
              statistics. This is done for several
              reasons.
           <1i>>
              It is fun to say lingua franca. Admit
              it. You're rolling your eyes, but you
              know I'm right.
           <
              More importantly, providers of
              commercial products nearly universally
               support it. You can use R from within
              SAS, SPSS, Oracle and Vertica. In
              fact, it is actually difficult to find
              a statistics package, database or
              business intelligence tool that
              doesn't support R.
           <
              That means R skills are portable. Even
              in an office that uses SPSS, it is
              possible to leverage R skills. If you
              later move to a job that uses SAS,
              your SPSS skills aren't very
              useful. Your R skills . . . are. Sorry
              for the pun. But I meant to do it.
           </aside>
</section>
<!-- Interface Choices ----- -->
<section>
   <h2>Interface Choices</h2>
   <img src="img/screenshots/RStudio.png" style="position: relative; top:-50px; left:-50px</pre>
   <img src="img/screenshots/ESS.png" style="position: relative; top:-512px; left:0; z-ind</pre>
   <img src="img/screenshots/Eclipse.png" style="position: relative; top:-974px; left:50px</pre>
```

<div style="display:inline-block; margin:0 auto; float:right; width:33%;">

```
<img src="img/screenshots/Rkward.png" style="position: relative; top:-1425px; left:100p</pre>
    <aside class="notes">
        <111>
            <1i>>
                Most commercial statistics packages
                come with a default interface of
                questionable quality. And if you don't
                like it . . . you can get over
                it. There probably aren't very many
                alternative choices.
            <
                R is different. In truth, the default
                R interface makes the SAS interface
                look positively utopic. But, because R
                is open source and is the lingua
                franca of the statistics community,
                there are more interfaces available
                for R than you have time to experiment
                with. And more are coming out every
                day.
            <1i>>
                I apologize for the color theme shown
                in all of the screenshots. I like
                it. It is called Solarized and is
                available in most pro-grade
                programming environments these days. I
                tend to use it anywhere / everywhere I
                can. Makes my eyes hurt less. All
                screenshots are taken from one of my
                laptops, and show a section of code I
                wrote in R and Emacs org-mode to
                assess a Kaggle data set about the
                Titanic disaster.
        </aside>
</section>
<section>
    <h2>Interface - RStudio</h2>
    <a href="img/screenshots/RStudio.png"><img src="img/screenshots/RStudio.png" width="192</pre>
    <aside class="notes">
        <l
            <
                RStudio is becoming THE interface for
                R. It presents the user with an
                interface roughly modeled after
                matlab. It is easy to use, available
                on Windows, Mac and Linux and has an
                active user base. This the interface I
                recommend to people if they don't
```

otherwise have an opinion.

```
</aside>
</section>
<section>
    <h2>Interface - ESS</h2>
    <a href="img/screenshots/ESS.png"><img src="img/screenshots/ESS.png" width="100%"></a>
    <aside class="notes">
        <111>
            <1i>
               ESS - Emacs Speaks Statistics. This is
               as bare bones as it gets. RStudio is
               an IDE. This is a text editor. In
               truth. It is THE text editor. All
               others are just playing catch-up. Fans
                of other text editors are welcome to
                email me to try to convince me
               otherwise.
            </aside>
</section>
<section>
    <h2>Interface - Eclipse/Statet</h2>
    <a href="img/screenshots/Eclipse.png"><img src="img/screenshots/Eclipse.png" width="100"</pre>
    <aside class="notes">
        <l
            <
                Eclipse is a popular open-source Java
               IDE that is used by Java developers
               here at DOH. The plugin for R is
               called Statet. Eclipse has plugins
               that allow it to connect to databases
                such as Oracle and Vertica, write Java
               / C applications, program in R, SAS,
               etc.
            <
               If all you want is an interface for R,
               you are probably better off with
               RStudio. If you want a high-quality
                interface to nearly everything. Use
               Emacs or Eclipse.
            </aside>
</section>
<section>
    <h2>Interface - JASP</h2>
    <a href="img/screenshots/JASP.png"><img src="img/screenshots/JASP.png" width="100%"></a
    <aside class="notes">
        ul>
```

```
<
          <
          </aside>
</section>
<!-- Literate Programming ----- -->
<section>
   <div style="display:inline-block; margin:0 auto; float:left; width:50%;">
       <h2>Literate Programming</h2>
   </div>
   <div style="display:inline-block; margin:0 auto; float:right; width:50%;">
       <img src="svg/r.svg" height="75%" width="75%">
   </div>
   <aside class="notes">
       <111>
          </aside>
</section>
<!-- MTCARS ----- -->
<section>
   <h2>MTCARS</h2>
   >
       The data was extracted from the 1974 Motor
       Trend US magazine, and comprises fuel
       consumption and 10 aspects of automobile
       design and performance for 32 automobiles
       (1973 - 1974 \text{ models}).
   </section>
<section>
 <h2>MTCARS - The Data</h2>
 <small>
   <!--begin.rcode
       data(mtcars)
       kable(head(mtcars))
       end.rcode-->
 </small>
   <aside class="notes">
       <l
        <1i>>
          So, this is the first 10 lines of the
          mtcars data set.
```

```
>
           We are going to focus on the mpg, cyl, and
           disp columns.
         </aside>
</section>
<section>
   <div style="display:inline-block; margin:0 auto; float:left; width:50%;">
       <h1>Open Source</h1>
       <h3>Free as in freedom, not beer.</h2>
   </div>
   <div style="display:inline-block; margin:0 auto; float:right; width:50%;">
       <img src="svg/r.svg" height="75%" width="75%">
   <aside class="notes">
       ul>
           <
              The term open source means different things to
              different people. In a nutshell, people interpret it
              to mean 'free'.
           <1i>>
              But it means much more than that. Open Source, or more
              formally, Free and Open Source Software aka FOSS, is an
              ideological method for the development, distribution
              and use of software.
           <1i>>
              Yeah. That's a little heavey. We're less than 10
              minutes into this and I'm pulling out the word
              ideology.
           <
              But it is this ideology that drives so many of the
              practical reasons people cite when choosing to use
              R.
           <
              The ecosystem, the integration with other tools, the
              veritable multitude of user interface options
              available for R. . . .
           <
              All of this is made possible by the ideological
              choices made by the R-Core team who lead the
              development of R.
           <1i>
              I don't think R is competing with SAS and SPSS because
              it is such a beautiful language. It isn't. It is
              competing with them because of the side affects of
```

```
the decision to develop R in an open, inclusive and
                                  transparent manner.
                          </aside>
</section>
<h2>Open Source Is Everywhere:</h2>
        <\td>
                          <img src="img/logos/firefox.png" height="128px"</pre>
                                             width="128px" style="vertical-align:middle;">
                          <img src="img/logos/adium.png" height="128px"</pre>
                                             width="128px" style="vertical-align:middle;">
                          <img src="img/logos/java.png" height="128px" width="128px"</pre>
                                             style="vertical-align:middle;">
                          <a href="https://filezilla-project.org/">Fil
                                  <a href="https://www.adium.im/">Adium</a></t
                                  <a href="https://www.java.com/en/download/">
                          <h3>Based On:</h3>
                                           <img src="img/logos/chrome.png" height="128px"</pre>
                                                      width="128px" style="vertical-align:middle;">
                                  <img src="img/logos/safari.jpg" height="128px"</pre>
                                                      width="128px" style="vertical-align:middle;">
                                  <img src="img/logos/android.png" height="128px"</pre>
                                                      width="128px" style="vertical-align:middle;">
                                  <
                                  <a href="http://www.google.com/chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/">Chrome/"
                                  <a href="https://www.apple.com/safari/">Safa
                                  <a href="https://www.android.com/">Android/
```

```
<aside class="notes"></aside>
</section>
<!-- DOH: Open Source ------ -->
<section>
  <h2>Even @ DOH:</h2>
  <t.r>
        <\td>
        <img src="img/logos/filezilla.png"</pre>
              height="128px" width="128px" style="vertical-align:middle;">
        <img src="img/logos/workbench.png" height="128px"</pre>
               width="128px" style="vertical-align:middle;">
        <img src="img/logos/7-zip.jpg" height="128px"</pre>
              width="128px" style="vertical-align:middle;">
        <t.r>
        <
        <a href="https://filezilla-project.org/">Filezil
        <a href="http://www.sql-workbench.net/">SQL Work
        <a href="http://www.7-zip.org/">7-Zip</a>
     <h3>Based On:</h3>
        <img src="img/logos/hp-vertica.jpeg" height="128px"</pre>
               width="128px" style="vertical-align:middle;">
        <img src="img/logos/SAS.jpg" height="128px"</pre>
              width="128px" style="vertical-align:middle;">
        <
        <a href="http://www.vertica.com/">Vertica</a></t</pre>
        <a href="http://www.sas.com/en_us/home.html">SAS
        <aside class="notes"></aside>
</section>
```

```
<section>
   <h1>Why?</h1>
   <aside class="notes"></aside>
</section>
<!-- The Four Freedoms ------ -->
   <h2> The Four Freedoms</h2>

    start ="0">

       <1i>>
          The freedom to run the program for any purpose.
       <1i>>
          The freedom to study how the program works, and change
          it to make it do what you wish.
       <1i>>
          The freedom to redistribute copies so you can help your
          neighbor.
       <1i>>
          The freedom to improve the program, and release your
          improvements (and modified versions in general) to the
          public, so that the whole community benefits.
       <br/>
   <img src="img/logos/fsf.org.png">
   <aside class="notes">
       <l
          <
              <a href="https://en.wikipedia.org/wiki/Free_software">Wikipedia</a>,
              lists the four freedoms associated with Free Software,
              as defined by the Free Sotware Foundation as:
          <
              Richard Stallman wrote these, thinking about
              software. But, at the time her was working at the AI
              lab at MIT, so science is not an unknown idea to him.
          <
              I believe these four freedoms fit in well with the
              scientific process.
          </aside>
</section>
<h2> The Four Freedoms: <small>Analytics Style</small></h2>

    start ="0">

       <1i>>
          The freedom to run the analysis for any purpose.
```

```
<1i>>
           The freedom to study how the analysis works, and change
           it to make it do what you wish.
       <1i>>
           The freedom to redistribute copies so you can help your
           neighbor.
       <1i>>
           The freedom to improve the analysis, and release your
           improvements (and modified versions in general) to the
           public, so that the whole community benefits.
       </section>
<div style="display:inline-block; margin:0 auto; float:left; width:50%;">
       <h2>Future Opportunities</h2>
   <div style="display:inline-block; margin:0 auto; float:right; width:50%;">
       <img src="svg/r.svg" height="75%" width="75%">
   </div>
   <aside class="notes">
       <111>
           </aside>
</section>
<!-- Accelerating Growth ----- -->
<section>
   <h2>Accelerating Growth</h2>
   <h3>Indeed.com</h3>
   \langle br/ \rangle
   <div style="width:50%; float:left;text-align: left;">
       <small><code>
     !"R D" !"A R" !"H R" !"R N" !toys !kids !" R Walgreen" !walmart !"HVAC R" !"R Bard" <br/>
     and ( "biostatistics" or "data analysis" or "data analyst"
           or "epidemiologist" <br/>
           or "healthcare analysis" or "healthcare analyst" <br/>
           or "statistical" <br/>
     )<br/>
     ,SAS
     !"system administrator" !"school age" !sata !firmware !scsi !raid !samsung !scandinav
     and ( "biostatistics" or "data analysis" or "data analyst"
           or "epidemiologist" <br/>
           or "healthcare analysis" or "healthcare analyst" <br/>
           or "statistical" <br/>>
     )
       </code></small>
```

```
</div>
             <div style="width:50%; float:right;">
                 <a href="http://www.indeed.com/jobtrends?q=R++%21%22R+D%22+%21%22A+R%22+%21%22H+R%2
                     <img width="540" height="300" src="http://www.indeed.com/trendgraph/jobgraph.pn</pre>
                 </a>
             </div>
             <aside class="notes">
                 <111>
                     </aside>
         </section>
         <div style="display:inline-block; margin:0 auto; float:left; width:50%;">
                 <h2>Others Say Similar Things</h2>
             </div>
             <div style="display:inline-block; margin:0 auto; float:right; width:50%;">
                 <img src="svg/r.svg" height="75%" width="75%">
             </div>
             <aside class="notes">
                 <111>
                     </aside>
         </section>
     </div>
 </div>
 <script src="lib/js/head.min.js"></script>
 <script src="js/reveal.js"></script>
  <script>
  // Full list of configuration options available at:
// https://github.com/hakimel/reveal.js#configuration
  Reveal.initialize({
      controls: true,
      progress: true,
      history: true,
      center: true,
      transition: 'slide', // none/fade/slide/convex/concave/zoom
      // Optional reveal.js plugins
      dependencies: [
          { src: 'lib/js/classList.js', condition: function() { return !document.body.classList; } }
          { src: 'plugin/markdown/marked.js', condition: function() { return !!document.querySelector
          { src: 'plugin/markdown/markdown.js', condition: function() { return !!document.querySelec
          { src: 'plugin/highlight/highlight.js', async: true, condition: function() { return !!docu
```