

To replicate the analyses

1. Install R Version ($\geq 3.3.1$) from <https://cran.r-project.org>.
2. Install JAGS ($\geq 4.2.0$) from <http://sourceforge.net/projects/mcmc-jags/files/> to the default location.
3. If using Windows install Rtools ($\geq 3.3.1$) from <https://cran.r-project.org/bin/windows/Rtools/> choosing the default options.
4. Cut and paste the following code into the R console to quickly generate non-convergent results.

```
install.packages("devtools") # if not already installed  
devtools::install_github("poissonconsulting/klexr")
```

```
library(klexr)  
replicate_results()
```

5. Open the `results` folder in the working directory to view the plots which are saved as png files.
6. For more information type `?replicate_results`.