Process Book: Health Insurance Coverage in the United States

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Overview and Motivation

One thing that was discussed a lot during the last president election is Obamacare. The Patient Protection and Affordable Care Act (ACA), also commonly known as Obamacare, was a major and wide-sweeping healthcare reform initiative enacted by President Obama. The ACA was signed into law by Obama on March 23, 2010, and changes were gradually implemented over the next few years, though the majority of the provisions went into effect only on January 1, 2014. Trump criticized Obamacare a lot. So will the ACA survive into 2017? Does it depend on the results of the 2016 election? Whether ACA is as terrible as trump said?

To answer those questions, we are interested in finding out how health insurance coverage in the USA differs from state to state, its relationship to indicators of health performance, and whether it can be predicted by political factors. More specifically, we want to see the impact of Affordable care act on the insurance coverage rate.

Quantitatively, we want see the change over time and the difference between states. We want to know in which states did insurance coverage expand under Obamacare? Which states observed the greatest decline in their uninsured rate? How does insurance coverage differ from state to state? We will also control some variables like income, mortality rate and rates of major diseases.

And then, we want to see whether higher insurance rate really benefit people. And how does health insurance coverage relate to participation uptake in preventative measures and treatment options (such as vaccinations, mammograms)?

And finally, we want to know what people think about Obamacare. To realize this, we will do some text analysis and see how do the sentiments towards Obamacare differ from state to state, and how does it relate to voting patterns? If feasible, we plan to analyze sentiments from twitter data, whether Obamacare is mentioned alongside positive or negative words.

We will make use of ggplot2, spatial data techniques, and text mining techniques in our visualization project.

Data

We used these data sources for our project:

- 1. General state-level data (population, age distribution and income distribution). United States Census Bureau: https://www.census.gov/data.html The United States Census bureau provides dataset that include information like population, average income and age distribution.
- 2. Insurance coverage and health indicators datasets State data on the Affordable Care Act (ACA) (from US National Library of Medicine): https://aspe.hhs.gov/compilation-state-data-affordable-care-act This state-level dataset includes coverage rates of the ACA, growth and expansion status of the ACA, employer vs. individual market coverage, and Medicaid / Medicare numbers. (filename: states aca general)
- 3. State scorecard on various health indicators (from the Commonwealth Fund): http://datacenter.commonwealthfund.org/#ind=1/sc=1 This state-level dataset has collated information about the health performance of the states along various dimensions. It includes insurance coverage rates, participation rates in prevention and treatment activities, and some measures of mortality. It also provides some differentiating information by race and income, allowing us to explore issues of equity.
- 4. Health Insurance Marketplace Data (from US Department of Health and Human Services): https://www.kaggle.com/hhs/health-insurance-marketplace This dataset contains information on health and dental plans offered through the US Health Insurance Marketplace.
- 5. Twitter API data & New York Times API data For the text analysis, we use twitter API data and New York Times API data. For the twitter API, we will try to extract text data on individuals' thoughts and opinions on Obamacare, capturing positive and negative sentiments. And we downloaded the New York Times articles from LexisNexis and then use R to transform them into corpus that can be analysis with R.

Downloading Twitter API data

The more challenging part is the API data. At first we planned to download all tweets that mentioned ACA from 2008 to 2016. However, we realized that will be too much data and will be hard to clean up and analysis. So we decided to download the New York Times articles and do the long term text analysis based on the New York Times articles. For twitter API, we only focus on the most recent 1000 tweets and try to have a sense of people's opinion about Affordable care act after the election.

Code for downloading tweets

Get some tweets from Twitter to analyze and visualize Set up Twitter API: Selecting data including Obamacare, ACA, Affordable Care Act, and #ACA (n=1000)

```
library(httr)
# library(oauth)
library(ROAuth)
library(twitteR)
library(RCurl)
library(RJSONIO)
library(stringr)
# secretkey myapp <- oauth app('twitter', key = 'liLn6XJFenGjtvWFwi5LnDS1M',
# secret = 'dsCBm9Kyaeu9GMKlM9xwKl7eKmDn6qsjP31LQtwMGkF60QdLh6') #Get OAuth
# credentials twitter token <- oauth1.0 token(oauth endpoints('twitter'),
# myapp) Declare Twitter API Credentials
api_key <- "liLn6XJFenGjtvWFwi5LnDS1M" # From dev.twitter.com
api secret <- "dsCBm9Kyaeu9GMK1M9xwK17eKmDn6qsjP31LQtwMGkF60QdLh6" # From dev.twitter.com
token <- "772176811455381505-PYuNAEqhHFc02r83WS9Y5dnsZciIY5v" # From dev.twitter.com token secret <- '
# Create Twitter Connection
library("base64enc")
```

Data cleaning

Plots of states trends

We first cleaned our data and unified the name of some variables as the data sets have different sources. We had to perform substantial data cleaning, including transforming reshaping one of our datasets to a wide format from its following original state. This was pretty challenging maneuver to figure out in terms of the data wrangling.

Original states health indicator data

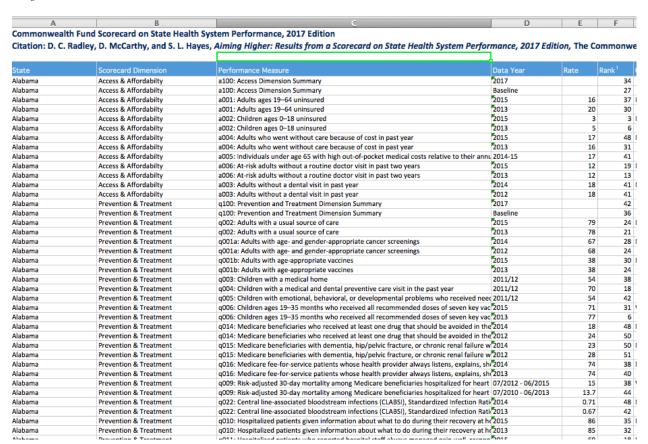


Figure 1:

We relabelled our variables extensively from the original excel sheets they came in, and came up with variable conversion lists such as the following. This helped us to get a bird-eye view of our available health insurance-related variables to work with. We were fortunate to find that we had much information in health and financial outcomes. Too much, in fact, so we tried to distill the variables down, highlighting the useful ones.

		NATION		State
Coverage		Uninsured Rate	2010	%
Gains		Uninsured Rate	2015	%
		Percentage Point Decrease in Uninsured Rate	2010-2015	%
		People Gaining Coverage	2010-2015	#
mployer		People with Employer Coverage	2015	#
Coverage		Individuals who Gained Coverage by Staying on their Parents' Plan Until Age 26	2013	#
	People with Lifetime	Total	2008-2010	
	Limits on health	Children	2008-2010	#
	benefits pre-	Adult Males	2008-2010	#
	ACA	Adult Females	2008-2010	#
	People with Private	Total	2013-2015	
	Coverage			#
	with No Cost	Children	2013-2015	#
	Sharing for Preventive	Adult Males	2013-2015	#
	Services	Adult Nales Adult Females	2013-2015	#
	Services	Average Annual Growth in Family Premiums for Employer Coverage	2000-2010	%
		Average Annual Growth in Family Premiums for	2010-2015	
		Employer Coverage		%
	-	Family Employer Premium Savings Compared to Continued Growth at pre-ACA Rate	2015	\$ \$
		·	2016	\$
		Total Group Market Consumers Benefiting from MLR Rebates Total Group Market Rebates	2012	# \$
		Total Group Market Consumers Benefiting from		
		MLR Rebates Total Group Market Rebates	2013	# \$
		Total Group Market Consumers Benefiting from MLR Rebates	2014	#
		Total Group Market Rebates Total Group Market Consumers Benefiting from		\$
		MLR Rebates	2015	#
		Total Group Market Rebates		\$
		Total Group Market Rebates	2012-2015	\$
∕ledicaid	People	Total Medicaid and CHIP Enrollment	July - September 2013	#
	Covered by	Total Medicaid and CHIP Enrollment	August 2016	#
	Medicaid or Total Medicaid/CHIP Child Enrollment CHIP Change in total Medicaid and CHIP Coverage		August 2016	#
	CHIP	Change in total Medicaid and CHIP Coverage Individuals Dually Eligible for full Medicaid and	2013-2016	#
		Medicare Benefits ("full duals")	December 2015	#
		Individuals Dually Eligible for partial Medicaid and Medicare Benefits ("partial duals")	December 2015	#
		Individuals Dually Eligible for Either Full or Partial Medicaid and Medicare Benefits	December 2015	#
		State Has Expanded	As of December 2016	Yes/No
		Increase in Number of People with Insurance Coverage due to Medicaid Expansion	2016, if all states expanded	#
	Projected Increase in	Cholesterol-Level Screening in Past 12 Months		
	People Receiving		Annual impact if all states expanded	#
	Preventive	Mammogram in Past 12 Months	Annual impact if all states expanded	#
	Care if State Expands	Papanicolaou Smear in Past 12 Months	Annual impact if all states	
	Medicaid Projected	Additional People with a Usual Source of Clinic	expanded Annual impact if all states	#
	Effects on Access to	Care Additional People Receiving All Needed Care in	expanded Annual impact if all states	#
	Care if State Expands	Past 12 Months	expanded	#
	Medicaid	Number of Additional Physician Visits Each Year	Annual impact if all states expanded	#
	Projected Effects on Health	Reduction in Number of People Experiencing Symptoms of Depression	Annual impact if all states expanded	#
	Outcomes if	Additional People Reporting Good, Very Good, or	Annual impact if all states	
	State	Excellent Health	expanded	#

NEW LABEL
state
unins_all_pct_10
unins_all_pct_15
unins_all_decr_pct
unins_all_decr
cov_emp
cov_parents_plan
list ii
lifetime_lim_preACA_tot
lifetime_lim_preACA_child
lifetime_lim_preACA_adultM
lifetime_lim_preACA_adultF
cov_private_tot
cov_private_child
cov_private_adultM
cov_private_adultF
premium_emp_avg_growth_pct_00to10
premium_emp_avg_growth_pct_10to15
premium_emp_savings_15
premium_emp_savings_16
MLR_rebate_beneficiaries_12
MLR_rebate_amt_12
MLR_rebate_beneficiaries_13
MLR rebate amt 13
MLR_rebate_beneficiaries_14
MLR_rebate_amt_14
inizit_resucc_dinit_1
MLR_rebate_beneficiaries_15
MLR_rebate_amt_15
MLR_rebate_amt_12to15
medicaid_enroll_13
medicaid enroll 16
medicaid_erroll_10 medicaid enroll child 13
medicaid_enroll_incr_13to16
medicald_effroil_incr_13t016
medicaid_full_duals
medicaid_partial_duals
medicaid_full_or_partial_duals
state_has_expanded
insurance_incr_medicaid
saraneeea.ea.ea
cholest_scr_incr_medicaid
mammogram_incr_medicaid
papsmear_incr_medicaid
clinic_care_incr_medicaid
all_care_incr_medicaid
 phycisian_visit_ann_incr_medicaid
priysisian_visic_arm_mor_medicald
depression_decr_medicaid
depression_decr_medicaid

	Expands	Reduction in Annual Number of Deaths	Annual impact if all states	
	Medicaid		expanded	#
		Reduction in Number of People with Catastrophic	Annual impact if all states	
		Out-of-Pocket Costs in a Typical Year	expanded	#
		Reduction in Number of People Borrowing to Pay	Annual impact if all states	
		Bills or Skipping Payments Due to Medical Bills	expanded	#
		Net Increase in Federal Spending	2016, if all states	
			expanded	\$M
		Reduction in Uncompensated Care	2016, if all states	
			expanded	\$M
		Share of People Potentially Eligible for Expansion		
		with Any Mental Illness or Substance Use Disorder	2010-2014	
				%
		Number of People Potentially Eligible for		
		Expansion with Any Mental Illness or Substance	2014	
		Use Disorder		#
Individual		Individuals with Pre-Existing Condition	2009	#
Market		Individuals Selecting a Marketplace Plan	2016	#
Coverage		Individuals with Marketplace Coverage	Q1 2016	#
		Individuals Receiving Tax Credits	Q1 2016	#
		Average Monthly Tax Credit Among those	01 2016	
		Receiving Tax Credits	Q1 2016	\$
		Individuals Receiving Cost Sharing Reductions	Q1 2016	#
		Average Number of Marketplace Plans Available	2017	#
		Marketplace Consumers who Could Select a Plan	2017	
		for less than \$75	2017	%
		Marketplace Consumers who Could Select a Plan	2017	
		for less than \$100	2017	%
		Individuals Purchasing Coverage Off-Marketplace	2016	
		who Could Quality for Tax Credits		#
		Rate Review Funds Awarded to the State	2010-2014	\$
		Health Insurance Enforcement and Consumer	2016	
		Protections Grant Awards		\$
Medicare		Medicare Enrollment	September 2016	#
		Medicare Beneficiaries benefitting from donut	2015	
		hole savings		#
		Total Medicare beneficiary savings from donut	2015	
		hole		\$
		Average per Medicare Beneficiary Donut Hole	2015	
		Savings		\$
		Part B Beneficiaries Using Free Preventive	2015	
		Services		#
		Share of Part B Beneficiaries Using Free	2015	
		Preventive Services		%
		% Change in Hospital Readmission Rates Among	2010-2015	
		Medicare Beneficiaries		%
		Avoided Readmissions	2015	#
		Number of Accountable Care Organizations	2016	#

deaths_ann_decr_medicaid
catastrophic_oop_ann_decr_medicaid
indebted_ppl_decr_medicaid
fed_spending_net_incr_inMil
uncompensated_care_decr_inMil
mental_substance_elig_medicaid_share
mental_substance_elig_medicaid
preexisting_condition_09
cov_mkt_plan_16
cov_mkt_16
receive_tax_credit
avg_tax_credit receive_cost_sharing
avg_num_mkt_plans_avail_17
cov_mkt_under75D_pct
cov_mkt_under100D_pct
cov_offmkt_elig_tax_credit
rate review funds to state
Tate_review_rands_to_state
HIECP_grant_award_to_state
medicare_enroll_16
medicare_benef_donuthole
medicare_benef_donuthole_savings_tot
medicare_benef_donuthole_savings_avg
medicare_partB_free_prev_services
medicare_partB_free_prev_services_share
medicare_incr_readmit_rate
medicare avoided readmit
accountable_care_org_num

a.unins_adult	
a.unins_child	
a.no_care_bc_cost_adult	
a high OOD valative und	oC.C
a.high_OOP_relative_und a.at risk no routine doc	
	tor_addit
a.no_dental_adult	
<pre>q.summary_prev_treat q.with_usual_care_adult</pre>	
q.with cancer screening	adult
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<pre>q.with_vaccines_adult a.with medical home ch</pre>	ild
q.witii_iiiedicai_iioiiie_tii	iiu
q.with_prev_medical_den	tal_child
q.with_mental_healthcare	e_child
q.with_vaccines_infant	
q.drug_should_avoid_med	dicare
a drug chould avaid 2	nditions medicare
q.drug_should_avoid_3co	nuitions_medicare
q.good_health_provider_ı	medicare
q.mortality_4conditions_r	medicare
q.CLABSI_infection_ratio	
q.info_recovery_hospitaliz	zed
q.info_recovery_hospitalia	zed
q.info_recovery_hospitalia q.good_hospital_staff_ho	
q.good_hospital_staff_ho	spitalized
	spitalized
q.good_hospital_staff_ho	spitalized ehealth
q.good_hospital_staff_ho q.improve_mobility_home q.improved_wounds_hom	spitalized ehealth
q.good_hospital_staff_ho	spitalized ehealth
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q.good_hospital_staff_ho q.improve_mobility_home q.improved_wounds_hom q.sores_NHres q.antipsychotic_med_NHr	spitalized ehealth nehealth
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q.good_hospital_staff_hode. q.improve_mobility_home. q.improved_wounds_home. q.sores_NHres q.antipsychotic_med_NHr.u.summary_avoidable_hode. u.hosp_asthma_child	spitalized ehealth nehealth res sp_cost
q.good_hospital_staff_hode. q.improve_mobility_home. q.improved_wounds_home. q.sores_NHres q.antipsychotic_med_NHr.u.summary_avoidable_hode. u.hosp_asthma_child	spitalized ehealth nehealth res sp_cost
q.good_hospital_staff_hode. q.improve_mobility_home. q.improved_wounds_home. q.sores_NHres q.antipsychotic_med_NHru.summary_avoidable_hode. u.hosp_asthma_child	spitalized ehealth nehealth res sp_cost
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q.good_hospital_staff_hodenimprove_mobility_homenimproved_wounds_homenimproved_wounds_homenimproved_wounds_homenimproved_wounds_homenimproved_NHres q.antipsychotic_med_NHresusummary_avoidable_hodenimproved_woundsp_asthma_child u.hosp_asthma_child u.hosp_ambulatory_65toouhosp_ambulatory_aboveu.30day_hosp_readmit_m	spitalized ehealth nehealth res sp_cost 74yrs e75yrs
q.good_hospital_staff_hosq.improve_mobility_homeq.improved_wounds_homq.sores_NHres q.antipsychotic_med_NHru.summary_avoidable_hou.hosp_asthma_child u.hosp_ambulatory_65toou.hosp_ambulatory_abovu.30day_hosp_readmit_m	spitalized ehealth nehealth res sp_cost 74yrs e75yrs nedicare
q.good_hospital_staff_hodenimprove_mobility_homenimproved_wounds_homenimproved_wounds_homenimproved_wounds_homenimproved_wounds_homenimproved_nantipsychotic_med_NHru.summary_avoidable_hodenimproved_woundsp_asthma_child u.hosp_ambulatory_65toouhosp_ambulatory_abovenimproved_wooday_hosp_readmit_mu.30day_hosp_readmit_nantiproved_wooday_hosp_readmit_N	spitalized ehealth nehealth res sp_cost 74yrs e75yrs

OR	IGI	NAL	LABEL	/ EXP	LANATI	ON

a001: Adults ages 19‰ÛÒ64 uninsured

a002: Children ages 0‰ÛÒ18 uninsured

a004: Adults who went without care because of cost in past year

a005: Individuals under age 65 with high out-of-pocket medical costs relative to their annual household income

a006: At-risk adults without a routine doctor visit in past two years

a003: Adults without a dental visit in past year

q100: Prevention and Treatment Dimension Summary

q002: Adults with a usual source of care

q001a: Adults with age- and gender-appropriate cancer screenings

q001b: Adults with age-appropriate vaccines

q003: Children with a medical home

q004: Children with a medical and dental preventive care visit in the past year

q005: Children with emotional, behavioral, or developmental problems who received needed mental health care in the past year

q006: Children ages 19‰ûò35 months who received all recommended doses of seven key vaccines

q014: Medicare beneficiaries who received at least one drug that should be avoided in the elderly

q015: Medicare beneficiaries with dementia, hip/pelvic fracture, or chronic renal failure who received a prescription drug that is contraindicated for that condition

q016: Medicare fee-for-service patients whose health provider always listens, explains, shows respect, and spends enough time with them

q009: Risk-adjusted 30-day mortality among Medicare beneficiaries hospitalized for heart attack, heart failure, pneumonia, or stroke

q022: Central line-associated bloodstream infections (CLABSI),

Standardized Infection Ratio

beneficiaries

q010: Hospitalized patients given information about what to do during their recovery at home

q011: Hospitalized patients who reported hospital staff always managed pain well, responded when needed help to get to bathroom or pressed call button, and explained medicines and side effects

q017: Home health patients who get better at walking or moving around

 $\ensuremath{\mathsf{q018}}\xspace$. Home health patients whose wounds improved or healed after an operation

q019: High-risk nursing home residents with pressure sores

q020: Long-stay nursing home residents with an antipsychotic medication

u100: Avoidable Hospital Use and Cost Dimension Summary

u001: Hospital admissions for pediatric asthma, per 100,000 children u002a: Hospital admissions among Medicare beneficiaries for ambulatory care‰ÛÒsensitive conditions, ages 65‰ÛÒ74, per 1,000

u002b: Hospital admissions among Medicare beneficiaries for ambulatory care‰ÛÒsensitive conditions, age 75 and older, per 1,000 beneficiaries

u004: Medicare 30-day hospital readmissions, rate per 1,000 beneficiaries

u005: Short-stay nursing home residents readmitted within 30 days of hospital discharge to nursing home

u006: Long-stay nursing home residents hospitalized within a six-month period

u007: Home health patients also enrolled in Medicare with a hospital admission

u003: Potentially avoidable emergency department visits among Medicare beneficiaries, per 1,000 beneficiaries u010: Total reimbursements per enrollee (ages 18‰0ò64) with employer-sponsored insurance u009: Total Medicare (Parts A & B) reimbursements per enrollee h100: Healthy Lives Dimension Summary h001: Mortality amenable to health care, deaths per 100,000 population h015: Years of potential life lost before age 75 h009: Breast cancer deaths per 100,000 female population h010: Colorectal cancer deaths per 100,000 population h007: Suicide deaths per 100,000 population h007: Suicide deaths per 100,000 population h008: Infant mortality, deaths per 1,000 live births h005: Adults ages 18‰0ò64 who report fair/poor health or activity limitations because of physical, mental, or emotional problems h003: Adults who smoke h004: Adults ages 18‰0ò64 who are obese (BMI >= 30) h011: Children ages 10‰0ò17 who are overweight or obese (BMI >= 85th percentile) h008: Adults ages 18‰0ò64 who have lost six or more teeth because of tooth decay, infection, or gum disease u008: Total single premium per enrolled employee at private-sector establishments that offer health insurance u908: Total single premium per enrolled employee at private-sector establishments that offer health insurance, unadjusted u909: Total Medicare (Parts A & B) reimbursements per enrollee, unadjusted h901b: Mortality amenable to health care, Black race, deaths per 100,000 population h901w: Mortality amenable to health care, White race, deaths per 100,000 population q001: Adults ages 50 and older who received recommended screening	
u010: Total reimbursements per enrollee (ages 18‰ÛÒ64) with employer-sponsored insurance u009: Total Medicare (Parts A & B) reimbursements per enrollee h100: Healthy Lives Dimension Summary h001: Mortality amenable to health care, deaths per 100,000 population h015: Years of potential life lost before age 75 h009: Breast cancer deaths per 100,000 female population h010: Colorectal cancer deaths per 100,000 population h007: Suicide deaths per 100,000 population h007: Suicide deaths per 100,000 population h008: Adults ages 18‰ÛÒ64 who report fair/poor health or activity limitations because of physical, mental, or emotional problems h003: Adults who smoke h004: Adults ages 18‰ÛÒ64 who are obese (BMI >= 30) h011: Children ages 10‰ÛÒ17 who are overweight or obese (BMI >= 85th percentile) h008: Adults ages 18‰ÛÒ64 who have lost six or more teeth because of tooth decay, infection, or gum disease u008: Total single premium per enrolled employee at private-sector establishments that offer health insurance u908: Total single premium per enrolled employee at private-sector establishments that offer health insurance, unadjusted u909: Total Medicare (Parts A & B) reimbursements per enrollee, unadjusted h901b: Mortality amenable to health care, Black race, deaths per 100,000 population h901w: Mortality amenable to health care, White race, deaths per 100,000 population	u003: Potentially avoidable emergency department visits among
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unadjusted h901b: Mortality amenable to health care, Black race, deaths per 100,000 population h901w: Mortality amenable to health care, White race, deaths per 100,000 population	establishments that offer health insurance, unadjusted
h901b: Mortality amenable to health care, Black race, deaths per 100,000 population h901w: Mortality amenable to health care, White race, deaths per 100,000 population	u909: Total Medicare (Parts A & B) reimbursements per enrollee,
100,000 population h901w: Mortality amenable to health care, White race, deaths per 100,000 population	unadjusted
h901w: Mortality amenable to health care, White race, deaths per 100,000 population	h901b: Mortality amenable to health care, Black race, deaths per
100,000 population	100,000 population
	h901w: Mortality amenable to health care, White race, deaths per
q001: Adults ages 50 and older who received recommended screening	100,000 population
-	q001: Adults ages 50 and older who received recommended screening
and preventive care	and preventive care

u.avoidable_ER_medicare
u.tot_reimb_employer_ins
u.tot_reimb_medicare
h.summary_healthy_lives
h deaths accepte
h.deaths_amenable
h.yrs_lost_potential_life_before75
h.deaths_breast_cancer_F
h.deaths_colorectal_cancer
h.deaths_suicide
h.deaths_infant_mortality
h.poor health adult
h.smoke_adult
h.obese_adult
in.obcsc_dudit
h.obese_child
h.poor_dental_adult
u.premium_emp_private
u.premium_emp_private_unadj
u.reimb_medicare_unadj
u.deaths_amenable_black
u.deaths_amenable_white
q.with_prev_screening_above50yrs

Code - data cleaning

Our full data cleaning code is below.

Data cleaning - states health trends

```
rm(list = ls())
library(leaflet)
library(maps)
library(rgdal)
## Loading required package: sp
## rgdal: version: 1.2-5, (SVN revision 648)
## Geospatial Data Abstraction Library extensions to R successfully loaded
## Loaded GDAL runtime: GDAL 2.1.2, released 2016/10/24
## Path to GDAL shared files: /Library/Frameworks/R.framework/Versions/3.3/Resources/library/rgdal/gda
## Loaded PROJ.4 runtime: Rel. 4.9.1, 04 March 2015, [PJ_VERSION: 491]
## Path to PROJ.4 shared files: /Library/Frameworks/R.framework/Versions/3.3/Resources/library/rgdal/p
## Linking to sp version: 1.2-4
library(DT)
library(ggplot2)
library(ggthemes)
library(plotly)
## Attaching package: 'plotly'
## The following object is masked from 'package:ggplot2':
##
##
       last_plot
## The following object is masked from 'package:stats':
##
##
       filter
## The following object is masked from 'package:graphics':
##
##
       layout
library(magrittr)
library(readxl)
library(plyr)
##
## Attaching package: 'plyr'
## The following objects are masked from 'package:plotly':
##
##
       arrange, mutate, rename, summarise
## The following object is masked from 'package:maps':
##
##
       ozone
library(dplyr)
```

##

```
## Attaching package: 'dplyr'
## The following objects are masked from 'package:plyr':
##
##
       arrange, count, desc, failwith, id, mutate, rename, summarise,
##
       summarize
## The following objects are masked from 'package:stats':
##
##
       filter, lag
## The following objects are masked from 'package:base':
##
##
       intersect, setdiff, setequal, union
library(tidyr)
##
## Attaching package: 'tidyr'
## The following object is masked from 'package:magrittr':
##
##
       extract
library(readr)
library(stringi)
library(RColorBrewer)
library(countrycode)
require(gridExtra)
## Loading required package: gridExtra
##
## Attaching package: 'gridExtra'
## The following object is masked from 'package:dplyr':
##
##
       combine
## IMPORT DATA
# General ACA information
aca_general_raw <- read_excel("Data/states_aca_general.xlsx", sheet = 2, col_names = FALSE)
# States health outcomes
health_ind_raw <- read_excel("Data/states_health_ind.xlsx", sheet = 4, col_names = FALSE)
# Insurance coverage data Long format
insurance_long <- read_excel("Data/insurance-clean.xlsx", sheet = 1, col_names = TRUE)</pre>
# Wide format
insurance <- read_excel("Data/insurance-clean.xlsx", sheet = 2, col_names = TRUE)</pre>
# Demographic information
demographics1 <- read.csv("Data/Population_Age_Income.csv", header = TRUE)</pre>
# Provided with Assignment 4 (want the yearly population data)
demographics2 <- read_excel("Data/PopulationEstimates.xls")</pre>
# Election Results
```

```
election <- read_excel("Data/US_Presidential_Results_by_State_1828-2016.xlsx",</pre>
    sheet = 2, col_names = FALSE)
## CLEAN DATA
# List of states abbreviations
state abbreviations <- c("AL", "AK", "AZ", "AR", "CA", "CO", "CT", "DE", "FL",
    "GA", "HI", "ID", "IL", "IN", "IA", "KS", "KY", "LA", "ME", "MD", "MA",
    "MI", "MN", "MS", "MO", "MT", "NE", "NV", "NH", "NJ", "NM", "NY", "NC", "ND", "OH", "OK", "OR", "PA", "RI", "SC", "SD", "TN", "TX", "UT", "VT",
    "VA", "WA", "WV", "WI", "WY")
## CLEAN INSURANCE RATES DATA (WIDE)
# Drop unnecessary columns
insurance <- insurance[1:50, -1]</pre>
# Rename variables
colnames(insurance) <- c("state_abb", "state", "uninsured_num_2008", "uninsured_pct_2008",</pre>
    "uninsured_num_2009", "uninsured_pct_2009", "uninsured_num_2010", "uninsured_pct_2010",
    "uninsured_num_2011", "uninsured_pct_2011", "uninsured_num_2012", "uninsured_pct_2012",
    "uninsured_num_2013", "uninsured_pct_2013", "uninsured_num_2014", "uninsured_pct_2014",
    "uninsured_num_2015", "uninsured_pct_2015")
# Create variables for insurance rates
insurance$insured_pct_2008 <- 100 - insurance$uninsured_pct_2008</pre>
insurance$insured_pct_2009 <- 100 - insurance$uninsured_pct_2009</pre>
insurance$insured_pct_2010 <- 100 - insurance$uninsured_pct_2010</pre>
insurance$insured pct 2011 <- 100 - insurance$uninsured pct 2011
insurance$insured_pct_2012 <- 100 - insurance$uninsured_pct_2012</pre>
insurance$insured_pct_2013 <- 100 - insurance$uninsured_pct_2013</pre>
insurance$insured_pct_2014 <- 100 - insurance$uninsured_pct_2014</pre>
insurance$insured_pct_2015 <- 100 - insurance$uninsured_pct_2015</pre>
# Reorder variables
insurance <- insurance[, c("state_abb", "state", "uninsured_num_2008", "uninsured_pct_2008",</pre>
    "insured_pct_2008", "uninsured_num_2009", "uninsured_pct_2009", "insured_pct_2009",
    "uninsured_num_2010", "uninsured_pct_2010", "insured_pct_2010", "uninsured_num_2011",
    "uninsured_pct_2011", "insured_pct_2011", "uninsured_num_2012", "uninsured_pct_2012",
    "insured_pct_2012", "uninsured_num_2013", "uninsured_pct_2013", "insured_pct_2013",
    "uninsured_num_2014", "uninsured_pct_2014", "insured_pct_2014", "uninsured_num_2015",
    "uninsured_pct_2015", "insured_pct_2015")]
## CLEAN INSURANCE RATES DATA (LONG)
# Rename variables
colnames(insurance_long) <- c("year", "state", "population", "uninsured_num",</pre>
    "uninsured_pct", "insured_num", "insured_pct")
# Add state abbreviations First create a look-up table (based on wide data)
state_abb_lookup <- insurance[, c("state", "state_abb")]</pre>
# Then do the matching
```

```
insurance_long$state_abb <- state_abb_lookup$state_abb[match(insurance_long$state,</pre>
    state_abb_lookup$state)]
# Reorder variable
insurance_long <- insurance_long[, c(1:2, ncol(insurance_long), 3:(ncol(insurance_long) -</pre>
    1))]
# Order observations (by state, then by year)
insurance long <- insurance long[order(insurance long$state, insurance long$year),
## CLEAN ACA GENERAL DATA
# Keep only relevant rows and columns
aca_general <- aca_general_raw[6:57, 1:73]</pre>
# Label variables for ACA general data
names(aca_general) <- c("state", "unins_all_pct_10", "unins_all_pct_15", "unins_all_decr_pct",</pre>
    "unins_all_decr", "cov_emp", "cov_parents_plan", "lifetime_lim_preACA_tot",
    "lifetime_lim_preACA_child", "lifetime_lim_preACA_adultM", "lifetime_lim_preACA_adultF",
    "cov_private_tot", "cov_private_child", "cov_private_adultM", "cov_private_adultF",
    "premium_emp_avg_growth_pct_00to10", "premium_emp_avg_growth_pct_10to15",
    "premium_emp_savings_15", "premium_emp_savings_16", "MLR_rebate_beneficiaries_12",
    "MLR_rebate_amt_12", "MLR_rebate_beneficiaries_13", "MLR_rebate_amt_13",
    "MLR_rebate_beneficiaries_14", "MLR_rebate_amt_14", "MLR_rebate_beneficiaries_15",
    "MLR_rebate_amt_15", "MLR_rebate_amt_12to15", "medicaid_enroll_13", "medicaid_enroll_16",
    "medicaid enroll child 13", "medicaid enroll incr 13to16", "medicaid full duals",
    "medicaid_partial_duals", "medicaid_full_or_partial_duals", "state_has_expanded",
    "insurance_incr_medicaid", "cholest_scr_incr_medicaid", "mammogram_incr_medicaid",
    "papsmear_incr_medicaid", "clinic_care_incr_medicaid", "all_care_incr_medicaid",
    "phycisian_visit_ann_incr_medicaid", "depression_decr_medicaid", "good_health_incr_medicaid",
    "deaths_ann_decr_medicaid", "catastrophic_oop_ann_decr_medicaid", "indebted_ppl_decr_medicaid",
    "fed_spending_net_incr_inMil", "uncompensated_care_decr_inMil", "mental_substance_elig_medicaid_sha
    "mental_substance_elig_medicaid", "preexisting_condition_09", "cov_mkt_plan_16",
    "cov_mkt_16", "receive_tax_credit", "avg_tax_credit", "receive_cost_sharing",
    "avg_num_mkt_plans_avail_17", "cov_mkt_under75D_pct", "cov_mkt_under100D_pct",
    "cov_offmkt_elig_tax_credit", "rate_review_funds_to_state", "HIECP_grant_award_to_state",
    "medicare_enroll_16", "medicare_benef_donuthole", "medicare_benef_donuthole_savings_tot",
    "medicare_benef_donuthole_savings_avg", "medicare_partB_free_prev_services",
    "medicare_partB_free_prev_services_share", "medicare_incr_readmit_rate",
    "medicare_avoided_readmit", "accountable_care_org_num")
# Trim white space on state variable
aca general$state <- stri trim both(aca general$state)</pre>
# Change format of variables Apart from T/F variable for 'state has
# expanded'
aca_general$state_has_expanded[aca_general$state_has_expanded == "yes"] <- TRUE
aca_general$state_has_expanded[aca_general$state_has_expanded == "no"] <- FALSE
# Convert others to numeric form
for (j in c(2:35, 37:73)) {
    aca_general[[j]] <- as.numeric(aca_general[[j]])</pre>
```

```
# Save USA data separately
aca_general_USA <- aca_general[aca_general$state == "United States", ]</pre>
# Drop USA and DC data
aca_general <- aca_general[!(aca_general$state %in% c("United States", "District of Columbia")),
# Add state abbreviations
aca_general$state_abb <- state_abbreviations</pre>
aca_general <- aca_general[, c(1, ncol(aca_general), 2:(ncol(aca_general) -
    1))]
# Drop raw data
rm(aca_general_raw)
## CLEAN HEALTH INDICATOR DATA
# Keep only relevant rows and columns
health_ind <- health_ind_raw[4:6703, c(1, 3:6)]
# Label variables
names(health_ind) <- c("state", "measure", "year", "rate", "rank")</pre>
# Simplify date (years) - only keep last year if range was given
health_ind\$year[health_ind\$year == "07/2009 - 06/2012"] <- "2012"
health_ind\$year[health_ind\$year == "07/2010 - 06/2013"] <- "2013"
health_ind\$year[health_ind\$year == "07/2012 - 06/2015"] <- "2015"
\label{eq:health_ind} $$ health_ind\$year == "10/2013-9/2014"] <- "2014" 
health_ind\$year[health_ind\$year == "2008-09"] <- "2009"
health_ind\$year[health_ind\$year == "2010-11"] <- "2011"
health_ind\$year[health_ind\$year == "2011-12"] <- "2012"
health_ind\$year[health_ind\$year == "2011/12"] <- "2012"
health_ind\$year[health_ind\$year == "2012-13"] <- "2013"
health_ind\$year[health_ind\$year == "2013-14"] <- "2014"
health_ind\$year[health_ind\$year == "2013(Q2-Q4)"] <- "2013"
health_ind\$year[health_ind\$year == "2014-15"] <- "2015"
health_ind\$year[health_ind\$year == "2015(Q2-Q4)"] <- "2015"
# Export to excel to manually label variable names
unique_measures <- unique(as.factor(health_ind$measure))</pre>
# Copied in from excel sheet
unique_measures_relabelled <- c("a.summary_access", "a.unins_adult", "a.unins_child",
    "a.no_care_bc_cost_adult", "a.high_00P_relative_under65", "a.at_risk_no_routine_doc_adult",
    "a.no_dental_adult", "q.summary_prev_treat", "q.with_usual_care_adult",
    "q.with_cancer_screening_adult", "q.with_vaccines_adult", "q.with_medical_home_child",
    "q.with_prev_medical_dental_child", "q.with_mental_healthcare_child", "q.with_vaccines_infant",
    "q.drug_should_avoid_medicare", "q.drug_should_avoid_3conditions_medicare",
    "q.good_health_provider_medicare", "q.mortality_4conditions_medicare", "q.CLABSI_infection_ratio",
    "q.info_recovery_hospitalized", "q.good_hospital_staff_hospitalized", "q.improve_mobility_homehealt
    "q.improved_wounds_homehealth", "q.sores_NHres", "q.antipsychotic_med_NHres", "u.summary_avoidable_hosp_cost", "u.hosp_asthma_child", "u.hosp_ambulatory_65to74yrs",
    "u.hosp_ambulatory_above75yrs", "u.30day_hosp_readmit_medicare", "u.30day_hosp_readmit_NHres",
```

```
"u.hosp_6mos_NHres", "u.hosp_medicare_homehealth", "u.avoidable_ER_medicare",
    "u.tot_reimb_employer_ins", "u.tot_reimb_medicare", "h.summary_healthy_lives",
    "h.deaths amenable", "h.yrs lost potential life before75", "h.deaths breast cancer F",
    "h.deaths_colorectal_cancer", "h.deaths_suicide", "h.deaths_infant_mortality",
    "h.poor_health_adult", "h.smoke_adult", "h.obese_adult", "h.obese_child",
    "h.poor_dental_adult", "u.premium_emp_private", "u.premium_emp_private_unadj",
    "u.reimb_medicare_unadj", "u.deaths_amenable_black", "u.deaths_amenable_white",
   NA, "q.with_prev_screening_above50yrs", NA)
# Replace variable names with abbreviated version
for (i in 1:57) {
   health_ind$measure <- replace(health_ind$measure, health_ind$measure ==</pre>
        unique_measures[i], unique_measures_relabelled[i])
}
# Remove the NA's
health_ind <- health_ind[!is.na(health_ind$measure), ]</pre>
# Remove duplicates By state, measure.year
health_ind <- health_ind[!duplicated(health_ind[, 1:3]), ]
# Interact measure and year
health_ind$measure_year <- interaction(health_ind$measure, health_ind$year)
# Convert to character variable
health_ind$measure_year <- as.character(health_ind$measure_year)
# Pre-reshape: Store existing data as 'long'
health_ind_long <- health_ind
# Reshape wide
health_ind <- as.data.frame(cbind(health_ind$state, health_ind$measure_year,
   health_ind$rate))
# Drop rank data for now
names(health_ind) <- c("state", "measure_year", "rate")</pre>
health_ind <- reshape(health_ind, idvar = "state", timevar = "measure_year",
    direction = "wide")
# Drop 'rate' from the name
names(health_ind)[2:ncol(health_ind)] <- substr(names(health_ind), 6, nchar(names(health_ind)))[2:ncol()</pre>
# Drop variables who have all entries as NA Due to 'unbalanced panel' in
# reshaping First create a vector to record if variable is all NA
variable_all_NA <- vector(mode = "logical", length = ncol(health_ind))</pre>
for (j in 1:ncol(health_ind)) {
    if (sum(is.na(health_ind[[j]])) == 52) {
        variable_all_NA[j] <- TRUE</pre>
   }
# Only keep the variables that are NOT all NA
health_ind <- health_ind[, !variable_all_NA]
# Change format of variables To character variable
```

```
health_ind$state <- as.character(health_ind$state)</pre>
# To numeric variable
for (j in c(2:164)) {
    health_ind[[j]] <- as.character(health_ind[[j]])
    health_ind[[j]] <- as.numeric(health_ind[[j]])</pre>
}
# Save USA data separately
health_ind_USA <- health_ind[health_ind$state == "United States", ]
# Drop USA and DC data
health_ind <- health_ind[!(health_ind$state %in% c("United States", "District of Columbia")),
# Label state abbreviations
health_ind$state_abb <- state_abbreviations
health_ind <- health_ind[, c(1, ncol(health_ind), 2:(ncol(health_ind) - 1))]
# Drop raw data
rm(health_ind_raw)
## CLEAN DEMOGRAPHIC (1) DATA
# Drop unnecessary row and column
demographics1 <- demographics1[-1, -1]</pre>
# Rename variables
colnames(demographics1) <- c("state", "income", "population", "ppl_age0to4",</pre>
    "ppl_age5to9", "ppl_age10to14", "ppl_age15to19", "ppl_age20to24", "ppl_age25to29",
    "ppl_age30to34", "ppl_age35to39", "ppl_age40to44", "ppl_age45to49", "ppl_age50to54",
    "ppl_age55to59", "ppl_age60to64", "ppl_age65to69", "ppl_age70to74", "ppl_age75to79",
    "ppl_age80to84", "ppl_age85plus")
# Drop observations DC and PR
demographics1 <- demographics1[!demographics1$state == "District of Columbia" &</pre>
    !demographics1$state == "Puerto Rico", ]
# Label state abbreviations
demographics1$state_abb <- state_abbreviations</pre>
demographics1 <- demographics1[, c(1, ncol(demographics1), 2:(ncol(demographics1) -</pre>
    1))]
# Change format of variables To character variable
demographics1$state <- as.character(demographics1$state)</pre>
# To numeric variable
for (j in c(3:ncol(demographics1))) {
    demographics1[[j]] <- as.character(demographics1[[j]])</pre>
    demographics1[[j]] <- as.numeric(demographics1[[j]])</pre>
}
# BEA regions
demographics1$BEA_region <- ""</pre>
```

```
demographics1$BEA_region[demographics1$state_abb %in% c("ME", "NH", "VT", "MA",
    "CT", "RI")] <- "New England"
demographics1$BEA_region[demographics1$state_abb %in% c("NY", "NJ", "PA", "MD",
    "DE")] <- "Mideast"
demographics1$BEA_region[demographics1$state_abb %in% c("WI", "IL", "MI", "IN",
    "OH")] <- "Great Lakes"
demographics1$BEA_region[demographics1$state_abb %in% c("WV", "KY", "VA", "TN",
    "NC", "SC", "AR", "LA", "MS", "AL", "GA", "FL")] <- "Southeast"
demographics1$BEA region[demographics1$state abb %in% c("ND", "SD", "NE", "KS",
    "MN", "IA", "MO")] <- "Plains"
demographics1$BEA_region[demographics1$state_abb %in% c("MT", "ID", "WY", "UT",
    "CO")] <- "Rocky Mountains"</pre>
demographics1$BEA region[demographics1$state abb %in% c("AZ", "NM", "TX", "OK")] <- "Southwest"
demographics1$BEA_region[demographics1$state_abb %in% c("WA", "OR", "CA", "NV",
    "AK", "HI")] <- "Far West"
# Tag income level (category)
demographics1$income_level <- ""</pre>
demographics1$income_level[demographics1$income >= 60000] <- "High"</pre>
demographics1$income_level[demographics1$income >= 50000 & demographics1$income <
    60000] <- "Upper Middle"
demographics1$income_level[demographics1$income >= 45000 & demographics1$income <
    50000] <- "Lower Middle"
demographics1$income level[demographics1$income < 45000] <- "Low"
# Convert to factor variable
demographics1$income_level <- factor(demographics1$income_level, levels = c("Low",</pre>
    "Lower Middle", "Upper Middle", "High"))
## CLEAN DEMOGRAPHIC (2) DATA
colnames(demographics2) <- demographics2[2, ]</pre>
# Drop first few rows (non-observations)
demographics2 <- demographics2[-(1:2), ]</pre>
# Drop county-level data, keep state-level data only
demographics2 <- demographics2[demographics2$Area Name == "United States" |</pre>
    demographics2$State != lag(demographics2$State), ]
# Keep only columns of population from 2010 to 2015
demographics2 <- demographics2[, c("State", "Area Name", "POP ESTIMATE 2010",</pre>
    "POP_ESTIMATE_2011", "POP_ESTIMATE_2012", "POP_ESTIMATE_2013", "POP_ESTIMATE_2014",
    "POP_ESTIMATE_2015")]
# Change format of variables To character variable
for (j in c(3:8)) {
    demographics2[[j]] <- as.numeric(demographics2[[j]])</pre>
}
# Rename variables
```

```
colnames(demographics2) <- c("state_abb", "state", "population_2010", "population_2011",</pre>
    "population_2012", "population_2013", "population_2014", "population_2015")
## CLEAN ELECTION DATA
# Keep only 2012 and 2008 data, and data for the 50 states
election <- election[4:53, c("X0", "X5", "X6", "X9", "X10")]
# Rename variables
colnames(election) <- c("state", "Dem_pct_2012", "Rep_pct_2012", "Dem_pct_2008",
    "Rep_pct_2008")
# Add state abbreviations Match with look-up table
election$state_abb <- state_abb_lookup$state_abb[match(election$state, state_abb_lookup$state)]</pre>
# Change format of variables To numeric variable
for (j in c(4:6)) {
    election[[j]] <- as.numeric(election[[j]])</pre>
}
# Calculate winning party for each state
election$party_2012 <- ifelse(election$Dem_pct_2012 > election$Rep_pct_2012,
    "Democratic", "Republican")
election$party_2008 <- ifelse(election$Dem_pct_2008 > election$Rep_pct_2008,
    "Democratic", "Republican")
# Reorder variables
election <- election[, c("state", "state_abb", "Dem_pct_2012", "Rep_pct_2012",</pre>
    "party_2012", "Dem_pct_2008", "Rep_pct_2008", "party_2008")]
Data cleaning - NY Times articles
library(tm)
## Loading required package: NLP
##
## Attaching package: 'NLP'
## The following object is masked from 'package:ggplot2':
##
##
       annotate
library(tm.plugin.lexisnexis)
library(readxl)
library(gtools) # for smartbind
library(dplyr) # for data_frame
library(lubridate) # for date formatting
## Attaching package: 'lubridate'
## The following object is masked from 'package:plyr':
##
##
       here
```

```
## The following object is masked from 'package:base':
##
##
       date
library(stringr)
library(tools) # Title case
library(quanteda)
## quanteda version 0.9.9.24
## Using 3 of 4 cores for parallel computing
##
## Attaching package: 'quanteda'
## The following objects are masked from 'package:tm':
##
##
       as.DocumentTermMatrix, stopwords
## The following object is masked from 'package:NLP':
##
##
       ngrams
## The following object is masked from 'package:readr':
##
       tokenize
## The following object is masked from 'package:utils':
##
##
       View
## The following object is masked from 'package:base':
##
##
       sample
# library(ggplot2)
library(quanteda)
library(stringr)
library(tm)
library(qdap)
## Loading required package: qdapDictionaries
## Loading required package: qdapRegex
##
## Attaching package: 'qdapRegex'
## The following objects are masked from 'package:dplyr':
##
##
       escape, explain
## The following object is masked from 'package:ggplot2':
##
       %+%
##
## Loading required package: qdapTools
##
## Attaching package: 'qdapTools'
## The following object is masked from 'package:dplyr':
```

```
##
##
       id
## The following object is masked from 'package:plyr':
##
##
       id
##
## Attaching package: 'qdap'
  The following objects are masked from 'package:quanteda':
##
##
       as.DocumentTermMatrix, as.wfm, ngrams, weight
  The following object is masked from 'package:stringr':
##
##
##
       %>%
  The following objects are masked from 'package:tm':
##
##
##
       as.DocumentTermMatrix, as.TermDocumentMatrix
  The following object is masked from 'package:NLP':
##
##
##
       ngrams
## The following object is masked from 'package:tidyr':
##
       %>%
##
##
  The following object is masked from 'package:dplyr':
##
##
       %>%
  The following object is masked from 'package:magrittr':
##
##
       %>%
  The following object is masked from 'package:plotly':
##
##
       %>%
##
## The following object is masked from 'package:DT':
##
       %>%
##
## The following object is masked from 'package:leaflet':
##
##
       %>%
## The following object is masked from 'package:base':
##
##
       Filter
library(SnowballC)
library(dplyr)
library(tidytext)
library(wordcloud)
# Combine CSV and HTML Files
data <- read_excel("LexisNexis/NYTimes_Metadata.xlsx")</pre>
```

```
colnames(data) <- tolower(colnames(data))</pre>
# Correct data
data$date <- substr(parse_date_time(data$date, c("mdy")), 1, 10)</pre>
data$author <- toTitleCase(tolower(data$byline))</pre>
data$byline <- NULL
## Get Text files
source1 <- LexisNexisSource("LexisNexis/The_New_York1.html")</pre>
source2 <- LexisNexisSource("LexisNexis/The_New_York2.html")</pre>
corpus1 <- Corpus(source1, readerControl = list(language = NA))</pre>
corpus2 <- Corpus(source2, readerControl = list(language = NA))</pre>
corpus <- c(corpus1, corpus2)</pre>
# Convert to quanteda corpus
corpus <- quanteda::corpus(corpus)</pre>
## Add Metadata Check: match(data$headline, corpus$documents$heading)
corpus$documents$datetimestamp <- substring(corpus$documents$datetimestamp,</pre>
corpus$documents$date <- corpus$documents$datetimestamp</pre>
corpus$documents$description <- corpus$documents$id</pre>
# options(width = 200) kwic(corpus, 'Trump')
save(corpus, file = "nytimes.rda")
```

Data cleaning - Twitter API

```
counts = table(tweets.df$screenName)
barplot(counts)
# Let's do something hacky: Limit the data set to show only folk who tweeted
# twice or more in the sample cc=subset(counts,counts>1)
barplot(cc, las = 2, cex.names = 0.3)
```

Design Evolution and Planning Process

We started from drawing the graphs on paper and whiteboard sessions. As we already have a general idea of our datasets, we were able to create some graph that can visualize information from different datasets. We divided our project into three sections.

The following was our original plan of the plots, though it had changed significantly since then.

The first section focus on the general insurance coverage in the US.

Graph 1: The first graph will show the general domestic information and also an insurance coverage rate.

Section 1

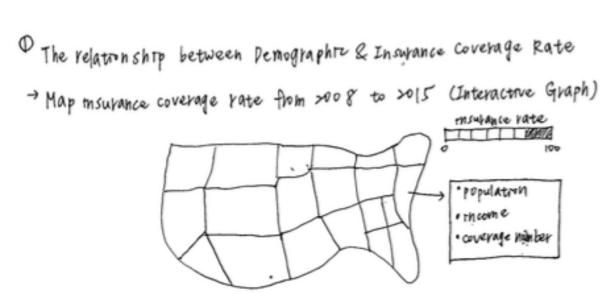


Figure 2:

Graph 2 and 3 focus on insurance rate. Graph 2 will show the trend of uninsured rate while graph 3 is a bar graph that will show the source of insurance by region

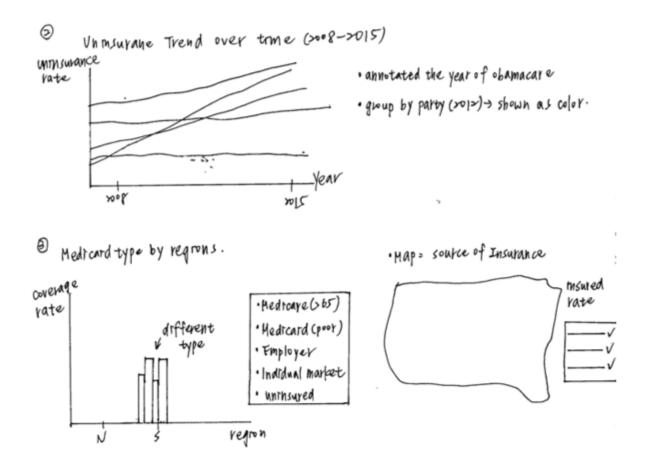


Figure 3:

Section 2 tries to see the insurance coverage in depth. We want to control some variables and see their relationship with insurance coverage rate.

The graph 4 is supposed to show the relationship between mortality rate and insurance coverage rate. We want to control some omitted variable like income, and expansion of Obamacare,

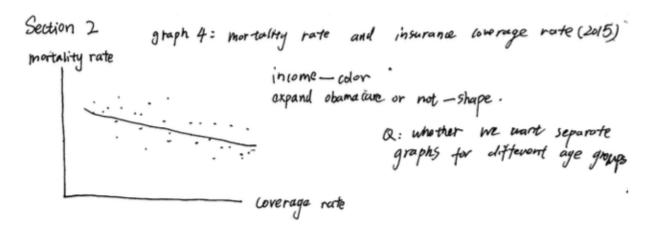
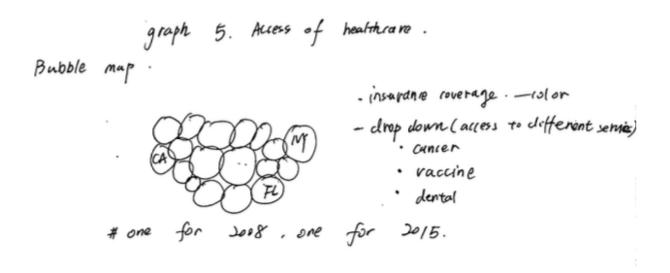


Figure 4:

Graph 5 will try to evaluate the relationship between the access of healthcare and insurance coverage. We divided health services into several categories based on our dataset like vaccine, dental and so on.



 $Figure\ 5:$

The section 3 is the text analysis. We planned to first have a look of the change of numbers of tweets mentioned ACA and then sentiment analysis of tweets mentioned ACA over years.

Code for data exploration and refinement of plots

Shiny App

Figure 1

We attempted to create a simply Shiny App and it was pretty challenging! We took quite a while to figure out reactivity when it came to updating the map. Our shiny app allowed users to use a slider to control the year, and have the US map of uninsured rates shift in response.

Shiny Code - Data cleaning

```
# Import insurance data
insurance_shiny <- read.csv("shiny_insurance.csv")</pre>
# insurance_shiny <- read.csv('/Users/gracekongyx/Documents/*6_Data
# Science/QMSS G4063 Data Visualization/Final Project/R
# code/Health_Insurance_Shiny/shiny_insurance.csv') insurance_shiny <-</pre>
# read.csv('/Users/gracekongyx/Documents/*6_Data Science/QMSS G4063 Data
# Visualization/Final Project/Data/shiny_insurance.csv')
## DATA PROCESSING
# Save list of state abbreviations
state_abb_lookup <- insurance_shiny[, c("state", "state_abb")]</pre>
# Obtain shape files of US states
map states = map("state", fill = TRUE, plot = FALSE)
# Standardize name format with rest of assignment Function to conver to
# proper case
properCase <- function(x) {</pre>
    s <- strsplit(x, " ")[[1]]
    paste(toupper(substring(s, 1, 1)), substring(s, 2), sep = "", collapse = " ")
}
# Convert to proper case
map_states$state <- sapply(map_states$names, properCase)</pre>
# Rename the main component of states with multiple parts in the map
map_states$state[map_states$state == "Massachusetts:main"] <- "Massachusetts"
map_states$state[map_states$state == "Michigan:south"] <- "Michigan"</pre>
map_states$state[map_states$state == "New York:main"] <- "New York"</pre>
map_states$state[map_states$state == "North Carolina:main"] <- "North Carolina"
map states$state[map states$state == "Virginia:main"] <- "Virginia"</pre>
map_states$state[map_states$state == "Washington:main"] <- "Washington"</pre>
# Match with state abbreviations
map_states$state_abb <- state_abb_lookup$state_abb[match(map_states$state, state_abb_lookup$state)]
# Then strip off all the extra location information for states with multiple
# parts (leaving just the state name)
for (i in 1:length(map states$state)) {
    map_states$state[i] <- unlist(strsplit(map_states$state[i], ":"))[1]</pre>
}
# Import key indicators data, matched by state Remember to later match
# information based on state, not state abbreviation (due to how we labelled
# above)
map_states_ins <- map_states</pre>
map_states_ins$uninsured_pct_2008 <- insurance_shiny$uninsured_pct_2008[match(map_states$state,
```

```
insurance_shiny$state)]
map_states_ins$uninsured_pct_2009 <- insurance_shiny$uninsured_pct_2009[match(map_states$state,
    insurance_shiny$state)]
map_states_ins$uninsured_pct_2010 <- insurance_shiny$uninsured_pct_2010[match(map_states$state,
    insurance_shiny$state)]
map_states_ins$uninsured_pct_2011 <- insurance_shiny$uninsured_pct_2011[match(map_states$state,
    insurance_shiny$state)]
map states ins$uninsured pct 2012 <- insurance shiny$uninsured pct 2012[match(map states$state,
    insurance shiny$state)]
map_states_ins$uninsured_pct_2013 <- insurance_shiny$uninsured_pct_2013[match(map_states$state,
    insurance_shiny$state)]
map_states_ins$uninsured_pct_2014 <- insurance_shiny$uninsured_pct_2014[match(map_states$state,
    insurance_shiny$state)]
map_states_ins$uninsured_pct_2015 <- insurance_shiny$uninsured_pct_2015[match(map_states$state,
    insurance_shiny$state)]
# Get coordinates of state centers
states_centers <- state.center</pre>
state.center <- cbind(states_centers, state_abb_lookup)</pre>
Shiny Code - Actual
## THE SHINY APP
# User interface
ui <- fluidPage(titlePanel("Uninsured Rate in the United States from 2008-2015"),
    sidebarLayout(sidebarPanel(sliderInput(inputId = "map_year", label = "Year",
        value = 2008, min = 2008, max = 2015, sep = "", round = TRUE)), mainPanel(leafletOutput("uninsu
# Server
server <- function(input, output) {</pre>
    variable_name <- reactive({</pre>
        paste("uninsured_pct_", as.character(input$map_year), sep = "")
   })
    # variable <- reactive({map_states_ins[[paste('uninsured_pct_',</pre>
    # input$map_year, sep = '')]]})
    output$uninsured_map <- renderLeaflet({</pre>
        map <- (leaflet(map_states_ins) %% setView(lat = 39.8282, lng = -96,
            zoom = 3.5))
       map
   })
    observe({
        classes <- 6
        pal <- colorNumeric(palette = "Oranges", domain = c(0:30), n = classes)</pre>
        map <- leafletProxy("uninsured_map") %>% clearShapes() %>% clearControls() %>%
            clearMarkers() %>% addPolygons(data = map_states_ins, fillColor = ~pal(map_states_ins[[vari
            smoothFactor = 0.5, fillOpacity = 0.6, color = "#333333", weight = 1,
            popup = paste("<b>State: </b>", map_states$state, "<br/>", "<b>Uninsured Rate: </b>",
                map_states_ins[[variable_name()]], "%")) %% addLabelOnlyMarkers(data = filter(state.ce
            state_abb != "AK" & state_abb != "HI"), lng = ~x, lat = ~y, label = ~state_abb,
            labelOptions = labelOptions(textsize = "9px", noHide = T, direction = "top",
                textOnly = T)) %>% addLegend(pal = pal, values = c(0:30), bins = classes,
            position = "bottomright", title = paste("Uninsured", "<br/>", "Rate ",
                "(", as.character(input$map_year), ")", "<br/>", "(%)", sep = ""),
```

```
opacity = 0.7)
})

# Run the application
shinyApp(ui = ui, server = server)
```

One of our challenges with Shiny was slightly silly, but we took a while to figure out the scale to use for map. At first, taking direction from the class code, we set the scale / legend of each map to be defined by the variable being plotted (i.e. uninsured rate of a certain year). However, the colors of resulting maps would then show the relative difference between the states, and did not have the effect of showing the decreasing trend across the board.

So we realized we needed the same scale across the maps. We started out choosing the first insurance data year as the anchor but ran into problems of out-of-range values for the other years (see Texas below):

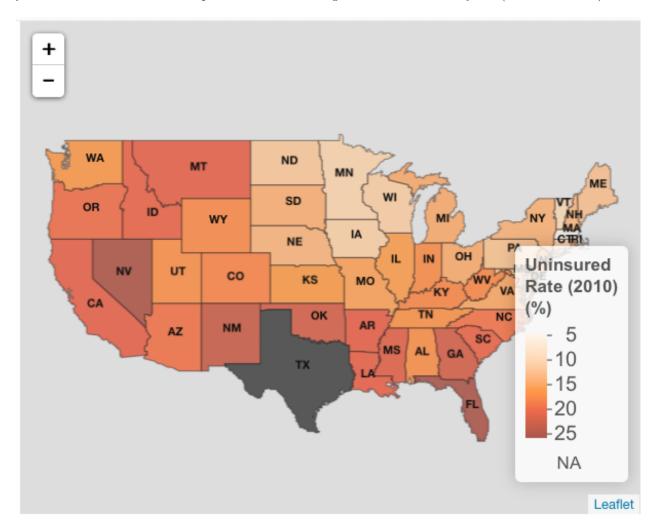


Figure 6:

Then we realized we could just manually input a numerical scale, one that encompassed all the possible values (between 0 and 30 percent). Our final app thus looked like this.

Uninsured Rate in the United States from 2008-2015

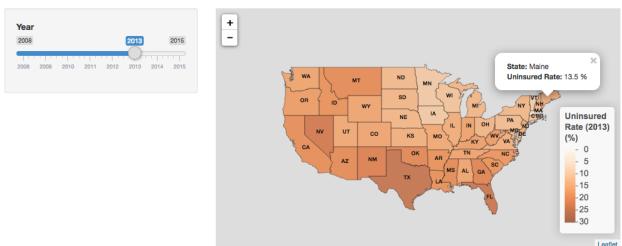


Figure 7:

Uninsured Rate in the United States from 2008-2015



Figure 8:

Link to Shiny App: https://gracekongyx.shinyapps.io/health_insurance_shiny/

States trends

```
## SOME DATA PREPARATION
# Look-up information into insurance and insurance_long, on: Party (2012)
insurance_long$party_2012 <- election$party_2012[match(insurance_long$state,</pre>
    election$state)]
insurance$party_2012 <- election$party_2012[match(insurance$state, election$state)]</pre>
# BEA region
insurance_long$BEA_region <- demographics1$BEA_region[match(insurance_long$state,</pre>
    demographics1$state)]
insurance$BEA_region <- demographics1$BEA_region[match(insurance$state, demographics1$state)]</pre>
# Income level
insurance_long$income_level <- demographics1$income_level[match(insurance_long$state,</pre>
    demographics1$state)]
insurance$income_level <- demographics1$income_level[match(insurance$state,</pre>
    demographics1$state)]
# Also find nation-wide trend for insurance
insurance_long_US <- data.frame(year = c(2008, 2009, 2010, 2011, 2012, 2013,
    2014, 2015), uninsured_pct = c(16.8, 17.5, 18.2, 17.2, 16.9, 16.6, 13.3,
    10.5))
```

Figure 2A

Final figure 2A. This was very close to our original plan. For the interactivity part, we decided to group the legend in terms of region instead of just list all the states, as it would take very long to 'unclick' 49 states to get to the state you want.

This plot is interactive on the website.

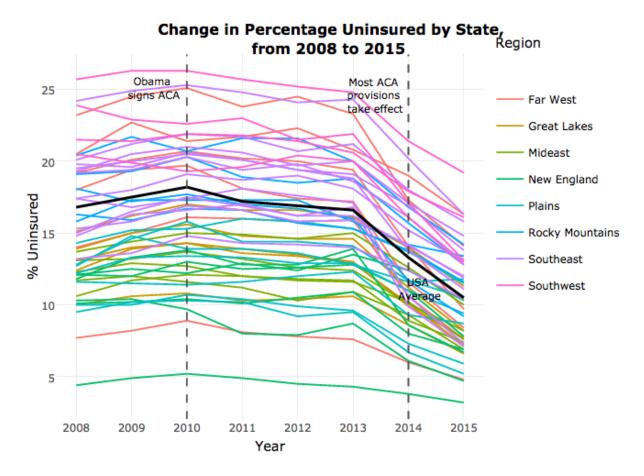
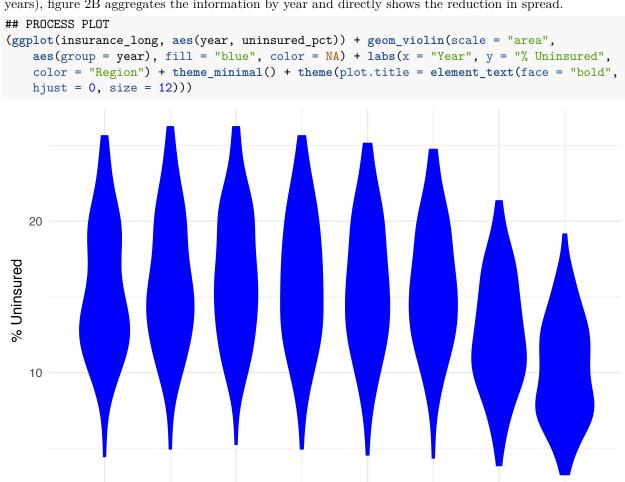


Figure 9:

Figure 2B

We realised that there was less spread in the uninsured percentage data points as the years passed. Therefore, figure 2B is a good contrast to figure 2A. While figure 2A grouped information to each state (through the years), figure 2B aggregates the information by year and directly shows the reduction in spread.



Year

Final figure 2B:

```
## PLOT 2B
plot2b <- (ggplot(insurance_long, aes(year, uninsured_pct)) + geom_violin(scale = "area",
    aes(group = year), fill = "dark blue", color = NA) + scale_x_continuous(breaks = c(2008,
    2009, 2010, 2011, 2012, 2013, 2014, 2015)) + scale_y_continuous(breaks = c(5,
    10, 15, 20, 25, 30)) + labs(x = "Year", y = "% Uninsured") + ggtitle("Narrowing of Uninsured Rates theme_minimal() + theme(plot.title = element_text(face = "bold", hjust = 0.5,
    size = 12)))
plot2b</pre>
```

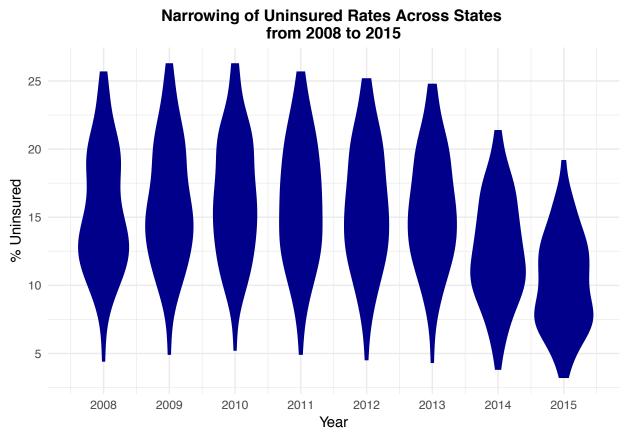


Figure 3

Final figure 3. This was close to our original plan as well.

This plot is interactive on the website.

```
## PI.OT 3
plot3 <- (ggplot(insurance_long, aes(year, uninsured_pct)) + geom_line(aes(color = party_2012,</pre>
    group = state, label = uninsured_num)) + scale_x_continuous(breaks = c(2008,
    2009, 2010, 2011, 2012, 2013, 2014, 2015)) + scale_y_continuous(breaks = c(5,
    10, 15, 20, 25, 30)) + scale_color_manual(values = c("steel blue", "firebrick1")) +
    labs(x = "Year", y = "% Uninsured", color = "Party \n(2012 Winner)") + geom_line(data = insurance_l
   lwd = 1) + geom_vline(xintercept = 2010, linetype = 2, color = "gray30") +
    geom_vline(xintercept = 2014, linetype = 2, color = "gray30") + annotate("text",
   x = 2009.4, y = 25, label = "Obama \nsigns ACA", color = "black", fontface = 2,
    size = 3) + annotate("text", x = 2013.4, y = 24.5, label = "Most ACA \nprovisions \ntake effect",
    color = "black", fontface = 2, size = 3) + annotate("text", x = 2014.2,
   y = 11, label = "USA \nAverage", color = "black", fontface = 2, size = 3) +
    ggtitle("Change in Percentage Uninsured by Party, \nfrom 2008 to 2015") +
    theme_minimal() + theme(plot.title = element_text(face = "bold", hjust = 0,
    size = 12)))
ggplotly(plot3, dynamicTicks = FALSE, tooltip = c("state", "year", "uninsured_pct",
    "uninsured num"))
```

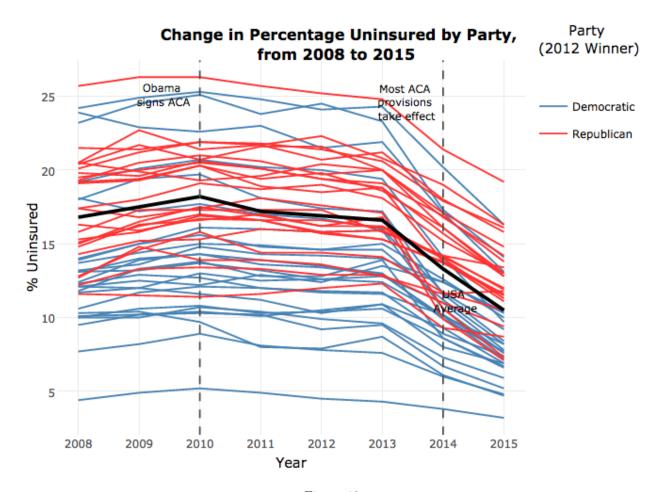


Figure 10:

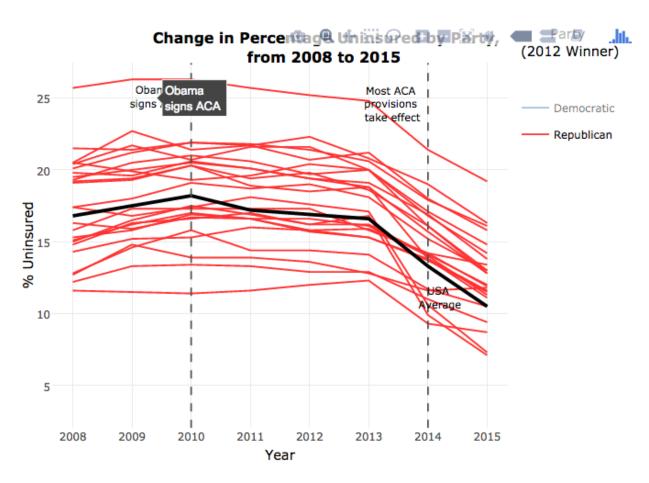
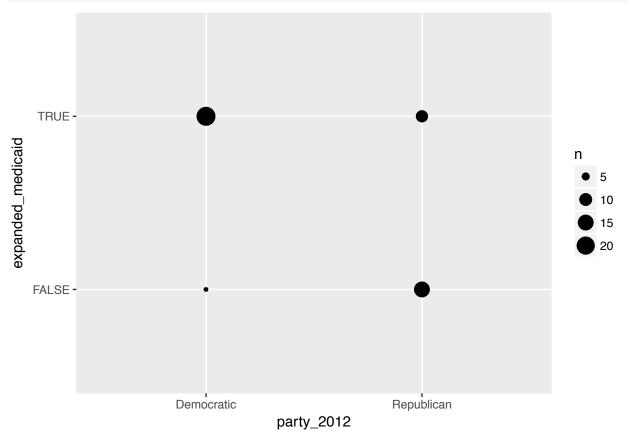


Figure 11:

Figure 4

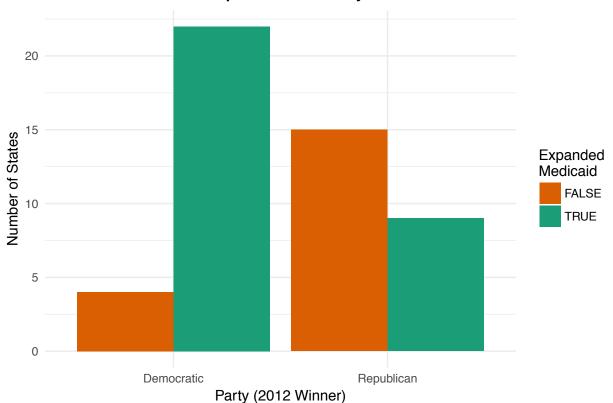
We decided to add an additional plot correlating medicaid expansion with state party, as we hypothesized that one of the reasons for party differences was their differential adoption of medicaid expansion. Thus this plot was important for confirming our hypothesis.

We tried a geom count at first since it was 2 discrete variables. But we felt that the graph looked too sparese and not aesthetically appealing.



Final figure 4:

Medicaid Expansion and Party of State



In response to student comments that we should present information comparing the states in the decrease in uninsured rates, and not just on the static rates at a point in time, we created another plot that ranks states on the gains in insured rate from 2010 to 2015. This also helped us gain insight as to whether the republican states are catching up or not (answer: to some extent, as they are in the middle of the group).

Final figure 5. This plot is interactive on the website.

```
insurance$unins_pct_decr_10_15 <- insurance$uninsured_pct_2010 - insurance$uninsured_pct_2015
insurance$unins_num_decr_10_15 <- insurance$uninsured_num_2010 - insurance$uninsured_num_2015

## PLOT 5
plot5 <- (ggplot(insurance, aes(x = reorder(state, unins_pct_decr_10_15), y = unins_pct_decr_10_15,
    fill = party_2012, label = unins_num_decr_10_15)) + geom_col() + coord_flip() +
    labs(x = "", y = "Decrease in % Uninsured", fill = "Party \n(2012 Winner)") +
    scale_fill_manual(values = c("steel blue", "firebrick1")) + ggtitle("Decrease in in Percentage Unin
    theme_minimal() + theme(plot.title = element_text(face = "bold", hjust = 0,
    size = 12), axis.text.y = element_text(size = 6.5)))
plot5</pre>
```

Decrease in in Percentage Uninsured from 2010 to 2015

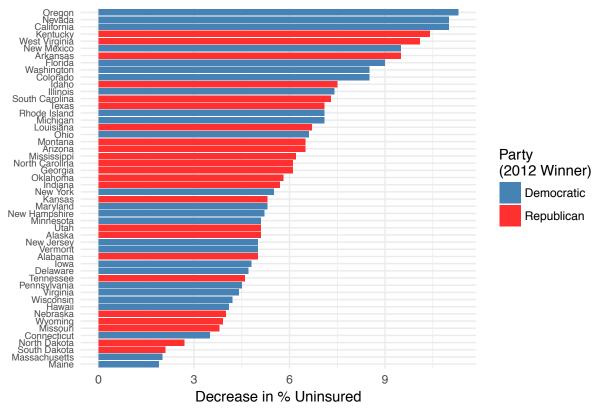


Figure 6A

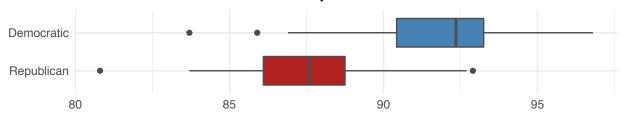
We wanted to present a snapshot of how states are doing in 2015, compared across different dimensions.

Final figure 6A:

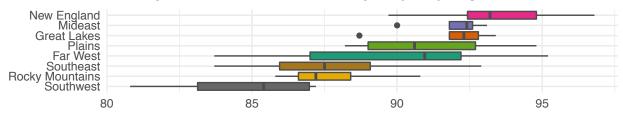
```
## PLOT 6A
# Subset relevant variables
insurance_2010_2015 <- insurance[, c("state", "insured_pct_2010", "unins_pct_decr_10_15",</pre>
    "insured_pct_2015", "party_2012", "BEA_region", "income_level")]
names(insurance_2010_2015) [names(insurance_2010_2015) == "unins_pct_decr_10_15"] <- "ins_pct_incr_10_15"</pre>
# Boxplots by party
plot6a1 <- (ggplot(insurance_2010_2015, aes(x = reorder(party_2012, insured_pct_2015),</pre>
    y = insured_pct_2015)) + geom_boxplot(aes(fill = party_2012, label = insured_pct_2015),
    color = "gray30") + coord_flip() + scale_fill_manual(values = c("steel blue",
    "firebrick")) + ggtitle("Democratic vs. Republican States") + theme_minimal() +
    theme(plot.title = element_text(face = "bold", hjust = 0.37, size = 12),
        legend.position = "none", axis.title.x = element_blank(), axis.title.y = element_blank()))
# Boxplots by BEA region
plot6a2 <- (ggplot(insurance_2010_2015, aes(x = reorder(BEA_region, insured_pct_2015),</pre>
    y = insured_pct_2015)) + geom_boxplot(aes(fill = BEA_region, label = insured_pct_2015),
    color = "gray30") + coord_flip() + scale_fill_manual(values = brewer.pal(n = 8,
   name = "Dark2")) + ggtitle("By Bureau of Economic Analysis (BEA) Region") +
    theme_minimal() + theme(plot.title = element_text(face = "bold", hjust = 0.24,
    size = 12), legend.position = "none", axis.title.x = element_blank(), axis.title.y = element_blank()
# Boxplots by income level
plot6a3 <- (ggplot(insurance_2010_2015, aes(x = income_level, y = insured_pct_2015)) +</pre>
    geom_boxplot(aes(fill = income_level, label = insured_pct_2015), color = "gray30") +
    coord_flip() + scale_fill_manual(values = brewer.pal(n = 4, name = "Dark2")) +
   labs(x = "", y = "% Insured (2015)") + ggtitle("By State's Income Level") +
    theme_minimal() + theme(plot.title = element_text(face = "bold", hjust = 0.39,
    size = 12), legend.position = "none", axis.title.y = element_blank()))
# Combine plots vertically
grid.arrange(plot6a1, plot6a2, plot6a3, nrow = 3, top = "2015 Insurance Rates at a Glance")
```

2015 Insurance Rates at a Glance

Democratic vs. Republican States



By Bureau of Economic Analysis (BEA) Region



By State's Income Level

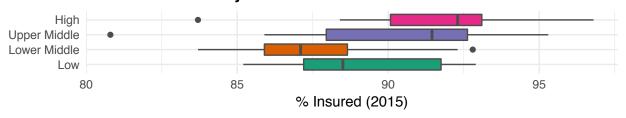


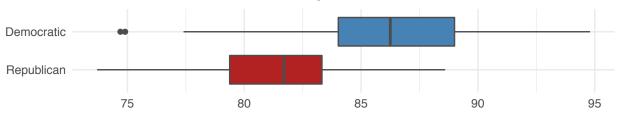
Figure 6B

In response to student comments to show how states did both before and after the ACA was rolled out, we initially wanted to put in a similar plot for the 2010 (pre-ACA) rates, as below.

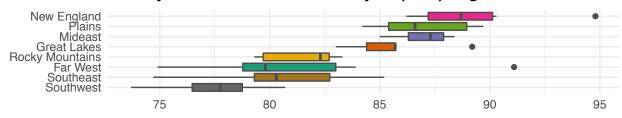
```
# PROCESS PLOT
# Boxplots by party
plot3a1 <- (ggplot(insurance_2010_2015, aes(x = reorder(party_2012, insured_pct_2010),</pre>
    y = insured pct 2010)) + geom boxplot(aes(fill = party 2012, label = insured pct 2010),
    color = "gray30") + coord_flip() + scale_fill_manual(values = c("steel blue",
    "firebrick")) + ggtitle("Democratic vs. Republican States") + theme_minimal() +
    theme(plot.title = element_text(face = "bold", hjust = 0.37, size = 12),
        legend.position = "none", axis.title.x = element_blank(), axis.title.y = element_blank()))
# Boxplots by BEA region
plot3b1 <- (ggplot(insurance_2010_2015, aes(x = reorder(BEA_region, insured_pct_2010),</pre>
    y = insured_pct_2010)) + geom_boxplot(aes(fill = BEA_region, label = insured_pct_2010),
    color = "gray30") + coord_flip() + scale_fill_manual(values = brewer.pal(n = 8,
   name = "Dark2")) + ggtitle("By Bureau of Economic Analysis (BEA) Region") +
   theme_minimal() + theme(plot.title = element_text(face = "bold", hjust = 0.24,
    size = 12), legend.position = "none", axis.title.x = element blank(), axis.title.y = element blank(
# Boxplots by income level
plot3c1 <- (ggplot(insurance_2010_2015, aes(x = income_level, y = insured_pct_2010)) +</pre>
    geom_boxplot(aes(fill = income_level, label = insured_pct_2010), color = "gray30") +
    coord flip() + scale fill manual(values = brewer.pal(n = 4, name = "Dark2")) +
   labs(x = "", y = "% Insured (2015)") + ggtitle("By State's Income Level") +
    theme_minimal() + theme(plot.title = element_text(face = "bold", hjust = 0.39,
    size = 12), legend.position = "none", axis.title.y = element_blank()))
# Combine plots vertically
grid.arrange(plot3a1, plot3b1, plot3c1, nrow = 3, top = "2010 Insurance Rates at a Glance")
```

2010 Insurance Rates at a Glance

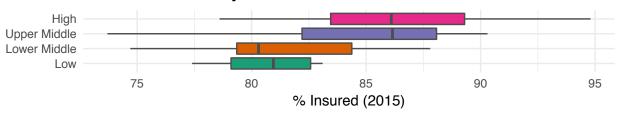
Democratic vs. Republican States



By Bureau of Economic Analysis (BEA) Region



By State's Income Level



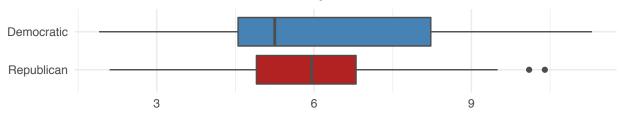
However, we though this didn't really give us much new information, the rankings were roughly the same. Yet, when tried a new plot that compared *change in insured rate from 2010 to 2015* instead, we found the interesting result that the rankings were often flipped, indicating that there was some 'catch-up' (but not enough).

Final figure 6B:

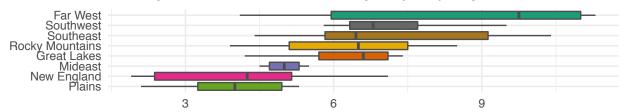
```
## PLOT 6B
# Boxplots by party
plot6b1 <- (ggplot(insurance_2010_2015, aes(x = reorder(party_2012, ins_pct_incr_10_15),</pre>
    y = ins_pct_incr_10_15)) + geom_boxplot(aes(fill = party_2012, label = ins_pct_incr_10_15),
    color = "gray30") + coord_flip() + scale_fill_manual(values = c("steel blue",
    "firebrick")) + ggtitle("Democratic vs. Republican States") + theme_minimal() +
    theme(plot.title = element_text(face = "bold", hjust = 0.37, size = 12),
        legend.position = "none", axis.title.x = element_blank(), axis.title.y = element_blank()))
# Boxplots by BEA region
plot6b2 <- (ggplot(insurance_2010_2015, aes(x = reorder(BEA_region, ins_pct_incr_10_15),</pre>
    y = ins_pct_incr_10_15)) + geom_boxplot(aes(fill = BEA_region, label = ins_pct_incr_10_15),
    color = "gray30") + coord_flip() + scale_fill_manual(values = brewer.pal(n = 8,
   name = "Dark2")) + ggtitle("By Bureau of Economic Analysis (BEA) Region") +
    theme_minimal() + theme(plot.title = element_text(face = "bold", hjust = 0.24,
    size = 12), legend.position = "none", axis.title.x = element_blank(), axis.title.y = element_blank(
# Boxplots by income level
plot6b3 <- (ggplot(insurance_2010_2015, aes(x = income_level, y = ins_pct_incr_10_15)) +</pre>
    geom_boxplot(aes(fill = income_level, label = ins_pct_incr_10_15), color = "gray30") +
    coord_flip() + scale_fill_manual(values = brewer.pal(n = 4, name = "Dark2")) +
   labs(x = "", y = "Increase in % Insured") + ggtitle("By State's Income Level") +
    theme_minimal() + theme(plot.title = element_text(face = "bold", hjust = 0.39,
    size = 12), legend.position = "none", axis.title.y = element_blank()))
# Combine plots vertically
grid.arrange(plot6b1, plot6b2, plot6b3, nrow = 3, top = "Increase in Insurance Rates (2010-2015) at a G
```

Increase in Insurance Rates (2010-2015) at a Glance

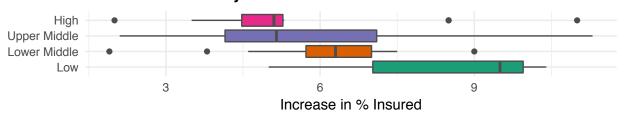
Democratic vs. Republican States



By Bureau of Economic Analysis (BEA) Region



By State's Income Level

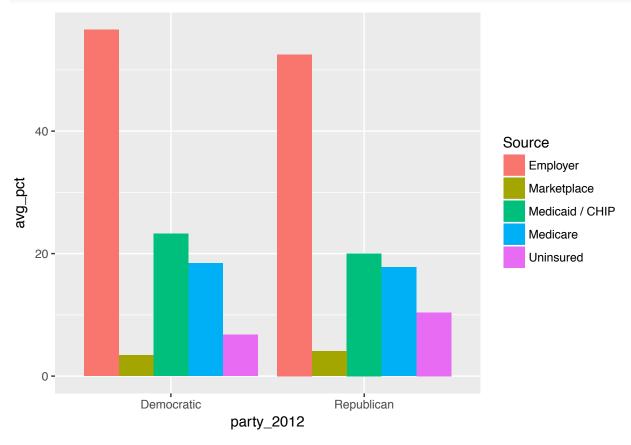


```
## SOME DATA RESHAPING
# Reshape aca_general (long), keeping the interesting quantities Import
# uninsured number (2015)
aca_general$unins_all_15 <- insurance$uninsured_num_2015[match(aca_general$state,
    insurance$state)]
# Keep only select variables
aca_general_long <- aca_general[, c("state", "state_abb", "cov_emp", "cov_mkt_plan_16",
    "medicaid_enroll_16", "medicare_enroll_16", "unins_all_15")]
# The actual reshaping
aca_general_long <- reshape(aca_general_long, varying = c("cov_emp", "cov_mkt_plan_16",</pre>
    "medicaid_enroll_16", "medicare_enroll_16", "unins_all_15"), v.names = "Number",
    timevar = "Source", times = c("Employer", "Marketplace", "Medicaid / CHIP",
        "Medicare", "Uninsured"), new.row.names = 1:1000, direction = "long")
# Look-up information into aca_general_long, on: Party (2012)
aca_general_long$party_2012 <- election$party_2012[match(aca_general_long$state,
    election$state)]
# BEA region
aca_general_long$BEA_region <- demographics1$BEA_region[match(aca_general_long$state,
    demographics1$state)]
# Income level
aca_general_long$income_level <- demographics1$income_level[match(aca_general_long$state,
    demographics1$state)]
# Population
aca_general_long$population_2015 <- demographics2$population_2015[match(aca_general_long$state,
    demographics2$state)]
# Calculate percentage from number and population
aca_general_long$percentage <- aca_general_long$population_2015 *</pre>
   100
```

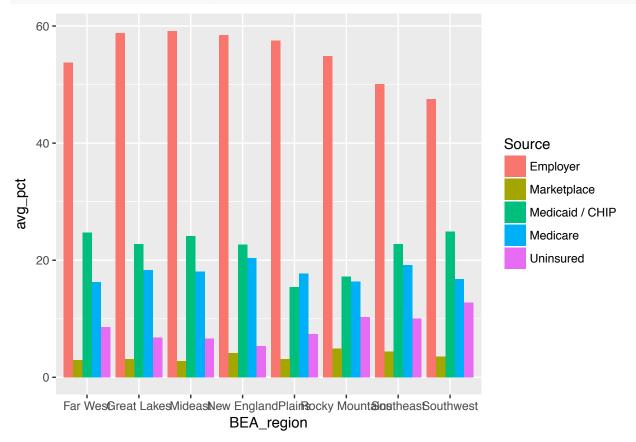
As we had planned from the start, we wanted to do a graph on the different sources where people got their insurance. And we wanted to do some sort of correlation with state characteristics, including party, BEA region and income level.

Therefore, we made exploratory plots for each of these dimensions.

By party



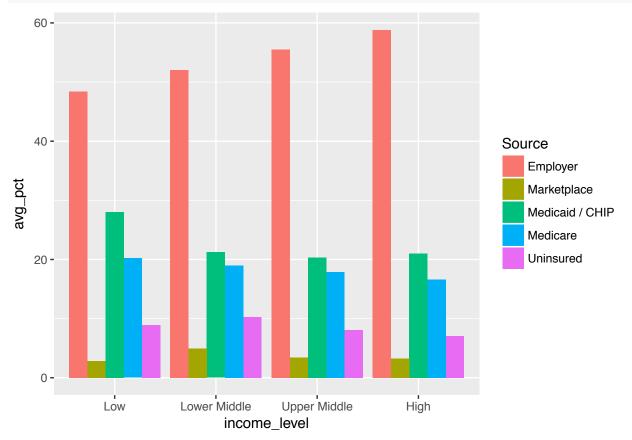
By region



By income level

```
## PROCESS PLOT
aca_general_long_by_income <- ddply(aca_general_long, c("income_level", "Source"),
    summarise, avg_pct = mean(percentage))

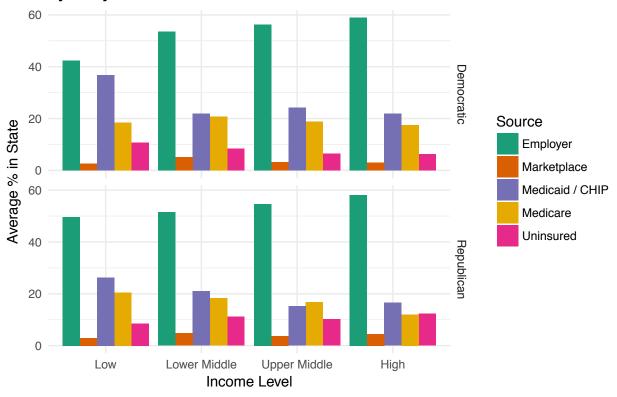
(ggplot(aca_general_long_by_income, aes(income_level, avg_pct, fill = Source)) +
    geom_col(position = "dodge"))</pre>
```



We felt that the trends in the party and the income level plots were most significant adn meaningful. So we dropped regional differences and presented trends by party and income level in the final faceted plot.

Final figure 7. This plot is interactive on the website.

Sources of Health Insurance, by Party and Income Level



States outcomes - health and financial

To prepare for comparing the states across the different health and financial outcomes, we create a dataframe of (10+) key indicators in these two dimensions. This required going through our 100+ outcome variables and selecting those that were relevant to our question, while still keeping enough breadth.

For example, we asked what represented a good health outcome? We wanted to capture aspects of mortality itself, access to health care (routine), and access to preventative healthcare like cancer screenings and vaccines.

And how about what represented a good financial outcome? That was trickier as there were many trade offs in terms of who bore the most financial burden in each state. Thus, we tried to include information on all these financial metrics, like OOP (private, incidental cost), premium (private, pooled cost), and federal spending (public cost).

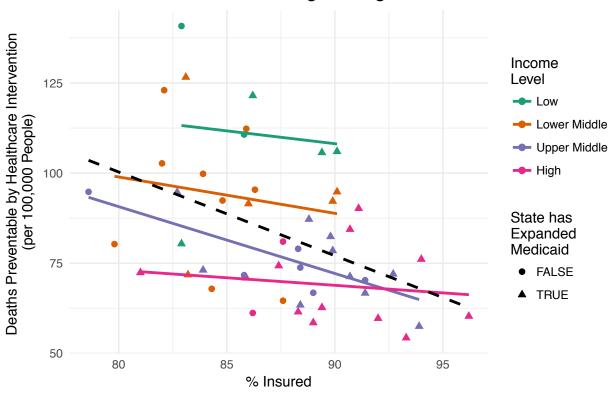
```
## PREPARATION FOR FURTHER PLOTS (INCLUDING DATA TABLE)
# Want to combine information on insurance coverage with key information on
# health outcomes/access and affordability
key_indicators <- state_abb_lookup</pre>
## Merge in information to the data table on the following
# Demographic Information Income level
key_indicators$income_level <- demographics1$income_level[match(key_indicators$state_abb,
    demographics1$state_abb)]
# Insurance Coverage Percentage insured (2015)
key_indicators$insured_pct_2015 <- insurance$insured_pct_2015[match(key_indicators$state_abb,
    insurance$state_abb)]
# Percentage insured (2014)
key_indicators$insured_pct_2014 <- insurance$insured_pct_2014[match(key_indicators$state_abb,
    insurance$state abb)]
# Indicator: Whether state has expanded medicare
key_indicators$state_has_expanded <- aca_general$state_has_expanded[match(key_indicators$state_abb,
    aca_general$state_abb)]
# Health outcomes and access Mortality amenable to health care, deaths per
# 10000 population
key_indicators$deaths_amenable_2014 <- health_ind$h.deaths_amenable.2014[match(key_indicators$state_abb
   health_ind$state_abb)]
# Years of Potential Life Lost before 75
key_indicators$years_life_lost <- health_ind$h.yrs_lost_potential_life_before75.2014[match(key_indicators)]
   health_ind$state_abb)]
# Adults with a usual source of care (%)
key_indicators$usual_care_2015 <- health_ind$q.with_usual_care_adult.2015[match(key_indicators$state_ab
   health_ind$state_abb)]
# Adults with age/gender-appropriate cancer screenings (%)
key_indicators$with_cancer_screening <- health_ind$q.with_cancer_screening_adult.2014[match(key_indicat
   health_ind$state_abb)]
# Adults with age-appropriate vaccines (%)
key_indicators$with_vaccines <- health_ind$q.with_vaccines_adult.2015[match(key_indicators$state_abb,
   health_ind$state_abb)]
# Affordability and cost-efficiency Individuals under age 65 with high OOP
```

medical costs relative to annual household income

```
key_indicators$high_00P_relative <- health_ind$a.high_00P_relative_under65.2015[match(key_indicators$st
    health_ind$state_abb)]
# Average annual growth in family premiums for employer coverage (between
# 2010 and 2015)
key_indicators$premium_ann_growth_10_15 <- aca_general$premium_emp_avg_growth_pct_10to15[match(key_indi
    aca_general$state_abb)]
# Marketplace consumers who could select a plan for less than $100
key_indicators$IM_plan_under_100 <- aca_general$cov_mkt_under100D_pct[match(key_indicators$state_abb,
    aca_general$state_abb)]
# Net increase in federal spending (millions)
key_indicators$incr_fed_spending_mil <- aca_general$fed_spending_net_incr_inMil[match(key_indicators$state_abb)]</pre>
```

We tried correlating mortality measures with insurance rates, as planned in our very early discussions.

Relationship Between Avoidable Mortality and Health Insurance Coverage Among States



Ideally, we wanted to make the plot interactive so that viewers could find out which point represented which state. However, ggplotly messed up our legend so we abandoned that thought.

PROCESS PLOT
ggplotly(plot8_old)

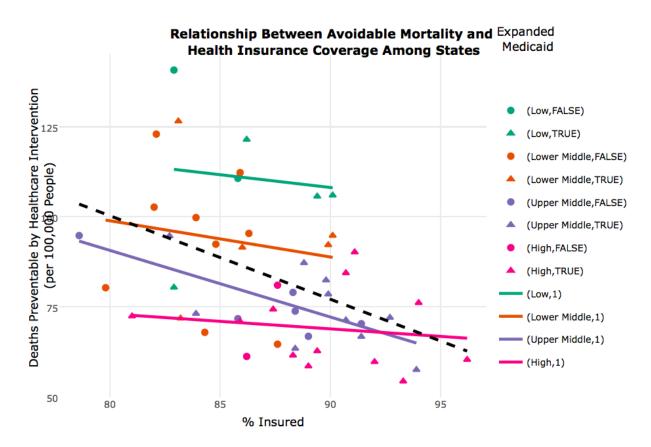


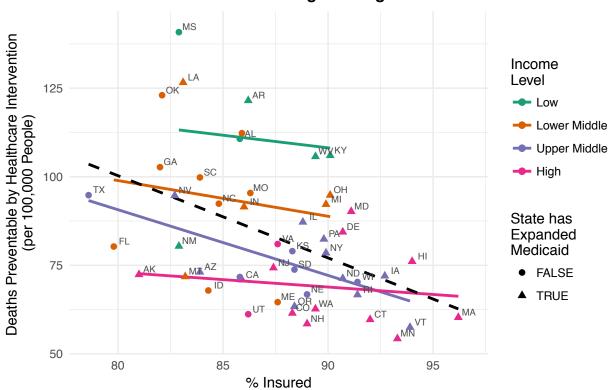
Figure 12:

Ultimately, we went with labeling the state abbreviations on the plot itself, ensuring that it was small enough not to be too distracting and that it didn't overlap with the points.

Final figure 8:

```
## PLOT 8
plot8 <- (ggplot(key_indicators, aes(insured_pct_2014, deaths_amenable_2014)) +
    geom_smooth(aes(color = income_level, group = income_level), method = "lm",
        lwd = 1, se = FALSE) + geom_smooth(color = "black", method = "lm", linetype = 2,
    lwd = 1, se = FALSE) + geom_point(aes(color = income_level, shape = state_has_expanded),
    size = 2) + geom_text(aes(label = state_abb), nudge_x = 0.5, nudge_y = 1.5,
    check_overlap = TRUE, size = 2.5, color = "gray30") + labs(x = "% Insured",
    y = "Deaths Preventable by Healthcare Intervention \n(per 100,000 People)",
    color = "Income \nLevel", shape = "State has \nExpanded \nMedicaid") + scale_color_manual(values = name = "Dark2")) + ggtitle("Relationship Between Avoidable Mortality and \nHealth Insurance Coverag theme_minimal() + theme(plot.title = element_text(face = "bold", hjust = 0.5,
    size = 12)))
plot8</pre>
```

Relationship Between Avoidable Mortality and Health Insurance Coverage Among States



Next, to prepare our data for the leaflet maps sections, we did a lot of pre-processing of our map_states spatial object. Things we had to deal with included the fact that some states had multiple polygons (e.g. Michigan which is split up), yet when evaluating variables we had to match the correct colors to all the polygons within the state. The following code was how we figured out the process.

```
## DATA PROCESSING FOR MAPS
# Obtain shape files of US states
map_states = map("state", fill = TRUE, plot = FALSE)
# Standardize name format with rest of assignment Function to conver to
# proper case
properCase <- function(x) {</pre>
    s <- strsplit(x, " ")[[1]]
    paste(toupper(substring(s, 1, 1)), substring(s, 2), sep = "", collapse = " ")
}
# Convert to proper case
map_states$state <- sapply(map_states$names, properCase)</pre>
# Rename the main component of states with multiple parts in the map
map_states$state[map_states$state == "Massachusetts:main"] <- "Massachusetts"</pre>
map_states$state[map_states$state == "Michigan:south"] <- "Michigan"</pre>
map states$state[map states$state == "New York:main"] <- "New York"</pre>
map_states$state[map_states$state == "North Carolina:main"] <- "North Carolina"
map_states$state[map_states$state == "Virginia:main"] <- "Virginia"</pre>
map_states$state[map_states$state == "Washington:main"] <- "Washington"</pre>
# Match with state abbreviations
map states$state abb <- state abb lookup$state abb[match(map states$state, state abb lookup$state)]
# Then strip off all the extra location information for states with multiple
# parts (leaving just the state name)
for (i in 1:length(map_states$state)) {
    map_states$state[i] <- unlist(strsplit(map_states$state[i], ":"))[1]</pre>
}
# Get coordinates of state centers
states_centers <- state.center</pre>
state.center <- cbind(states_centers, state_abb_lookup)</pre>
# Health outcomes: Import key indicators data, matched by state Remember to
# later match information based on state, not state abbreviation (due to how
# we labelled above)
map_states$deaths_amenable_2014 <- key_indicators$deaths_amenable_2014[match(map_states$state,
    key indicators$state)]
map_states$usual_care_2015 <- key_indicators$usual_care_2015[match(map_states$state,
    key_indicators$state)]
map_states$with_cancer_screening <- key_indicators$with_cancer_screening[match(map_states$state,
    key_indicators$state)]
map_states$with_vaccines <- key_indicators$with_vaccines[match(map_states$state,
    key_indicators$state)]
# Cost outcomes: Import key indicators data, matched by state Remember to
# later match information based on state, not state abbreviation (due to how
# we labelled above)
map_states$high_00P_relative <- key_indicators$high_00P_relative[match(map_states$state,
    key_indicators$state)]
```

We deliberated with what kind of scales to use for our leaflet map. We tried out a numeric scale and then a 6-color quantile scale but found that the color was actually too much detail and distraction. We decided that 4 bins was actually ideal, and so stuck with the default.

Numeric scale

```
# PROCESS PLOT
(leaflet(map_states) %>% setView(lat = 39.8282, lng = -96, zoom = 4) %>% addPolygons(color = "#333333",
   weight = 1, smoothFactor = 0.5, fillOpacity = 0) %>% addPolygons(group = "Preventable Deaths",
   fillColor = ~colorNumeric("RdYlGn", -deaths_amenable_2014)(-deaths_amenable_2014),
    smoothFactor = 0.5, stroke = FALSE, fillOpacity = 0.6, popup = paste("<b>State: </b>",
       map_states$state, "<br/>", "<b>Preventable Mortality</b>", "<br/>",
        "<b>per 100,000: </b>", map_states$deaths_amenable_2014)) %% addPolygons(group = "Access to Us"
   fillColor = ~colorNumeric("RdYlGn", usual_care_2015)(usual_care_2015), smoothFactor = 0.5,
    stroke = FALSE, fillOpacity = 0.6, popup = paste("<b>State: </b>", map_states$state,
        "<br/>", "<b>Access to Usual</b>", "<br/>", "<b>Source of Care: </b>",
       map_states$usual_care_2015, "%")) %>% addPolygons(group = "Cancer Screenings Rate",
   fillColor = ~colorNumeric("RdYlGn", with_cancer_screening)(with_cancer_screening),
   smoothFactor = 0.5, stroke = FALSE, fillOpacity = 0.6, popup = paste("<b>State: </b>",
       map_states$state, "<br/>", "<b>Cancer Screenings</b>", "<br/>", "<b>Rate: </b>",
       map_states$with_cancer_screening, "%")) %>% addPolygons(group = "Adult Vaccination Rate",
   fillColor = ~colorNumeric("RdYlGn", with_vaccines)(with_vaccines), smoothFactor = 0.5,
   stroke = FALSE, fillOpacity = 0.6, popup = paste("<b>State: </b>", map_states$state,
        "<br/>", "<b>Vaccination</b>", "<br/>", "<b>Rate: </b>", map_states$with_vaccines,
       "%")) %>% addLabelOnlyMarkers(data = filter(state.center, state_abb !=
   "AK" & state_abb != "HI"), lng = ~x, lat = ~y, label = ~state_abb, labelOptions = labelOptions(noHi
   direction = "top", textOnly = T)) %>% addLayersControl(baseGroups = c("Preventable Deaths",
   "Access to Usual Care", "Cancer Screenings Rate", "Adult Vaccination Rate"),
   options = layersControlOptions(collapsed = FALSE)))
```

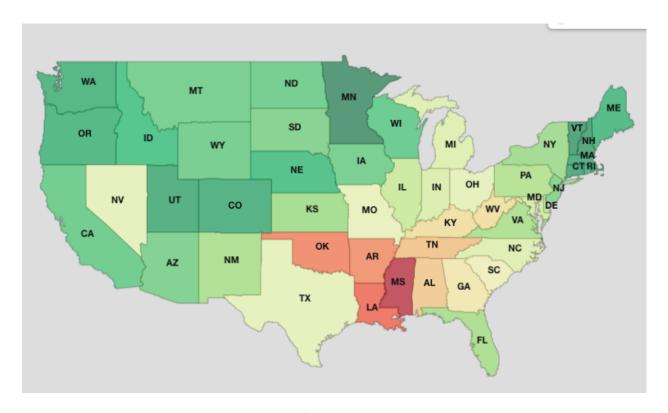


Figure 13:

6-level quantile scale

```
## PROCESS PLOT
(leaflet(map_states) %>% setView(lat = 39.8282, lng = -96, zoom = 4) %% addPolygons(color = "#333333",
    weight = 1, smoothFactor = 0.5, fillOpacity = 0) %>% addPolygons(group = "Preventable Deaths",
   fillColor = ~colorQuantile("RdYlGn", -deaths_amenable_2014, n = 6)(-deaths_amenable_2014),
    smoothFactor = 0.5, stroke = FALSE, fillOpacity = 0.6, popup = paste("<b>State: </b>",
        map_states$state, "<br/>", "<b>Preventable Mortality</b>", "<br/>",
        "<b>per 100,000: </b>", map_states$deaths_amenable_2014)) %>% addPolygons(group = "Access to Us
   fillColor = ~colorQuantile("RdYlGn", usual_care_2015, n = 6)(usual_care_2015),
    smoothFactor = 0.5, stroke = FALSE, fillOpacity = 0.6, popup = paste("<b>State: </b>",
        map_states$state, "<br/>", "<b>Access to Usual</b>", "<br/>", "<b>Source of Care: </b>",
        map_states$usual_care_2015, "%")) %>% addPolygons(group = "Cancer Screenings Rate",
    fillColor = ~colorQuantile("RdYlGn", with_cancer_screening, n = 6)(with_cancer_screening),
    smoothFactor = 0.5, stroke = FALSE, fillOpacity = 0.6, popup = paste("<b>State: </b>",
        map_states$state, "<br/>", "<b>Cancer Screenings</b>", "<br/>", "<b>Rate: </b>",
        map_states$with_cancer_screening, "%")) %% addPolygons(group = "Adult Vaccination Rate",
    fillColor = ~colorQuantile("RdYlGn", with_vaccines, n = 6)(with_vaccines),
    smoothFactor = 0.5, stroke = FALSE, fillOpacity = 0.6, popup = paste("<b>State: </b>",
        map_states$state, "<br/>", "<b>Vaccination</b>", "<br/>", "<b>Rate: </b>",
        map_states$with_vaccines, "%")) %>% addLabelOnlyMarkers(data = filter(state.center,
    state_abb != "AK" & state_abb != "HI"), lng = ~x, lat = ~y, label = ~state_abb,
   labelOptions = labelOptions(noHide = T, direction = "top", textOnly = T)) %>%
    addLayersControl(baseGroups = c("Preventable Deaths", "Access to Usual Care",
        "Cancer Screenings Rate", "Adult Vaccination Rate"), options = layersControlOptions(collapsed =
```

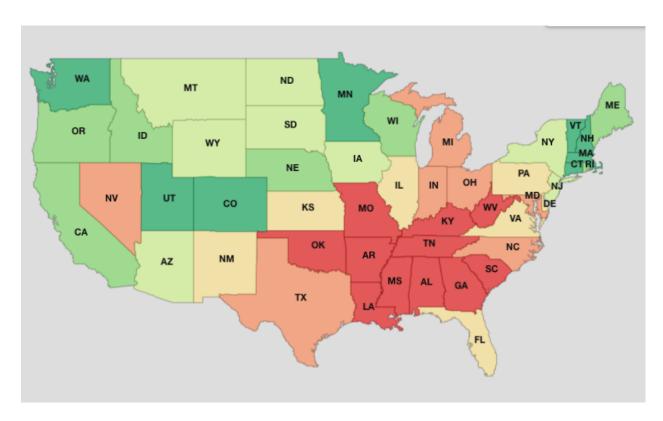


Figure 14:

Final figure 9:

4-level quantile scale

```
## PLOT 9: LEAFLET MAP - HEALTH OUTCOMES
(leaflet(map_states) %>% setView(lat = 39.8282, lng = -96, zoom = 4) %>% addPolygons(color = "#333333",
    weight = 1, smoothFactor = 0.5, fillOpacity = 0) %% addPolygons(group = "Preventable Deaths (2014)
    fillColor = ~colorQuantile("RdYIGn", -deaths_amenable_2014)(-deaths_amenable_2014),
    smoothFactor = 0.5, stroke = FALSE, fillOpacity = 0.6, popup = paste("<b>State: </b>",
        map_states$state, "<br/>", "<b>Preventable Mortality</b>", "<br/>",
        "<b>per 100,000: </b>", map_states$deaths_amenable_2014)) %% addPolygons(group = "Access to Us"
    fillColor = ~colorQuantile("RdYlGn", usual_care_2015)(usual_care_2015),
    smoothFactor = 0.5, stroke = FALSE, fillOpacity = 0.6, popup = paste("<b>State: </b>",
        map_states$state, "<br/>", "<b>Access to Usual</b>", "<br/>", "<b>Source of Care: </b>",
        map_states$usual_care_2015, "%")) %>% addPolygons(group = "Cancer Screenings Rate (2014)",
   fillColor = ~colorQuantile("RdYlGn", with_cancer_screening)(with_cancer_screening),
    smoothFactor = 0.5, stroke = FALSE, fillOpacity = 0.6, popup = paste("<b>State: </b>",
       map_states$state, "<br/>", "<b>Cancer Screenings</b>", "<br/>", "<b>Rate: </b>",
        map_states$with_cancer_screening, "%")) %>% addPolygons(group = "Adult Vaccination Rate (2015)"
   fillColor = ~colorQuantile("RdYlGn", with vaccines)(with vaccines), smoothFactor = 0.5,
    stroke = FALSE, fillOpacity = 0.6, popup = paste("<b>State: </b>", map_states$state,
        "<br/>", "<b>Vaccination</b>", "<br/>", "<b>Rate: </b>", map_states$with_vaccines,
        "%")) %>% addLabelOnlyMarkers(data = filter(state.center, state_abb !=
    "AK" & state_abb != "HI"), lng = ~x, lat = ~y, label = ~state_abb, labelOptions = labelOptions(noHi
   direction = "top", textOnly = T)) %>% addLayersControl(baseGroups = c("Preventable Deaths (2014)",
    "Access to Usual Care (2015)", "Cancer Screenings Rate (2014)", "Adult Vaccination Rate (2015)"),
    options = layersControlOptions(collapsed = FALSE)))
```

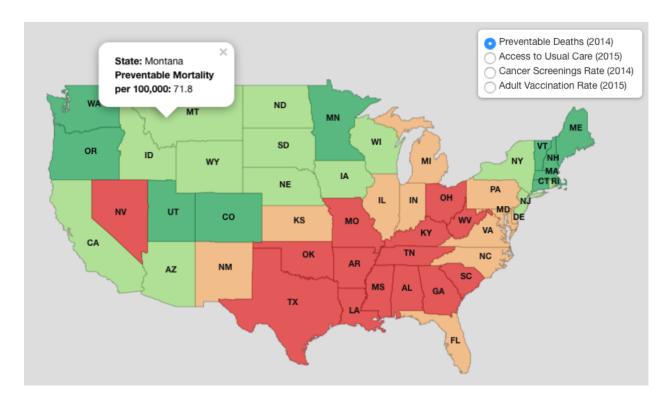


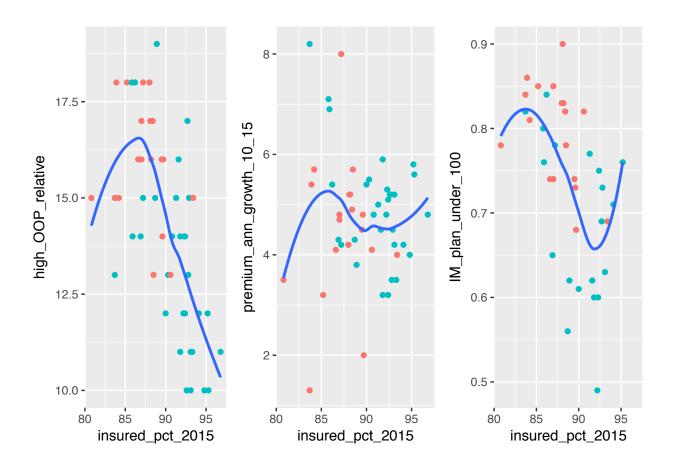
Figure 15:

This was a new plot from after our presentation, but we realized we did not have a plot explicitly correlating financial outcomes to insurance rates (like what we had for our health outcomes section).

However, there were 3-4 key financial variables that correlated differently with insured rates. We thus explored correlating each of them separately, to see which had a significant trend we could report.

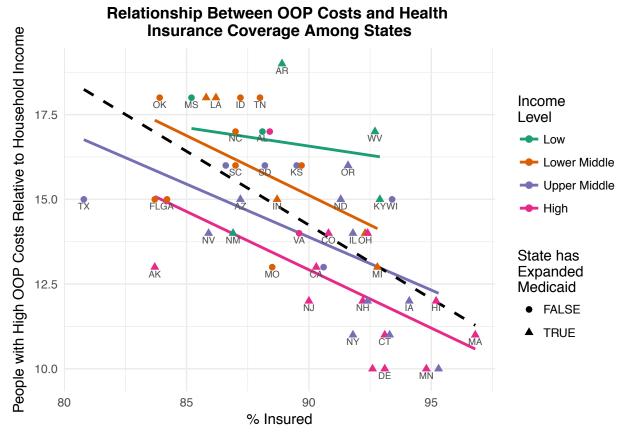
```
# PROCESS PLOTS
plot10a <- (ggplot(key_indicators, aes(insured_pct_2015, high_00P_relative)) +
    geom_point(aes(color = state_has_expanded)) + geom_smooth(se = FALSE) +
    theme(legend.position = "none"))
plot10b <- (ggplot(key_indicators, aes(insured_pct_2015, premium_ann_growth_10_15)) +
    geom_point(aes(color = state_has_expanded)) + geom_smooth(se = FALSE) +
    theme(legend.position = "none"))
plot10c <- (ggplot(key_indicators, aes(insured_pct_2015, IM_plan_under_100)) +
    geom_point(aes(color = state_has_expanded)) + geom_smooth(se = FALSE) +
    theme(legend.position = "none"))
grid.arrange(plot10a, plot10b, plot10c, ncol = 3)

## `geom_smooth()` using method = 'loess' and formula 'y ~ x'
## `geom_smooth()` using method = 'loess' and formula 'y ~ x'
## `geom_smooth()` using method = 'loess' and formula 'y ~ x'</pre>
```



The only variable with a significant correlation trend with insured percentage was the percentage of high OOP people. Therefore, we used that as our y-variable, and create a graph that looked analogous to our health outcomes plot.

Final figure 10:



Final figure 11 - leaflet plot of cost outcomes:

```
## PLOT 11: LEAFLET MAP - AFFORDABILITY / COST OUTCOMES
marketplace_pct <- map_states$IM_plan_under_100</pre>
marketplace_pct <- marketplace_pct * 100</pre>
marketplace_pct[is.na(marketplace_pct)] <- "No State Marketplace"</pre>
marketplace_pct_symbol <- ifelse(marketplace_pct == "No State Marketplace",</pre>
    "", "%")
(leaflet(map_states) %>% setView(lat = 39.8282, lng = -96, zoom = 4) %% addPolygons(color = "#333333",
    weight = 1, smoothFactor = 0.5, fillOpacity = 0) %% addPolygons(group = "Contained OOP Costs (2015
   fillColor = ~colorQuantile("RdYlGn", -high_00P_relative)(-high_00P_relative),
    smoothFactor = 0.5, stroke = FALSE, fillOpacity = 0.6, popup = paste("<b>State: </b>",
        map_states$state, "<br/>", "<b>People with High OOP</b>", "<br/>", "<b>Relative to Income: </b>
        map_states$high_OOP_relative, "%")) %>% addPolygons(group = "Premium Growth Rate (2010-2015)",
    fillColor = ~colorQuantile("RdYlGn", -premium_ann_growth_10_15)(-premium_ann_growth_10_15),
    smoothFactor = 0.5, stroke = FALSE, fillOpacity = 0.6, popup = paste("<b>State: </b>",
        map_states$state, "<br/>", "<b>Annual Premium Growth</b>", "<br/>",
        "<b>Rate (2010-2015): </b>", map_states$premium_ann_growth_10_15, "%")) %%
    addPolygons(group = "Affordable Marketplace Plan (2017)", fillColor = ~colorQuantile("RdYlGn",
        IM_plan_under_100)(IM_plan_under_100), smoothFactor = 0.5, stroke = FALSE,
        fillOpacity = 0.6, popup = paste("<b>State: </b>", map_states$state,
            "<br/>", "<b>Marketplace Consumers</b>", "<br/>", "<b>Who Can Select Plan</b>",
            "<br/>", "<b>Under $100: </b>", marketplace_pct, marketplace_pct_symbol)) %>%
    addPolygons(group = "Increase in Federal Spending (2016)", fillColor = ~colorQuantile("RdYlGn",
        -incr fed spending mil)(-incr fed spending mil), smoothFactor = 0.5,
        stroke = FALSE, fillOpacity = 0.6, popup = paste("<b>State: </b>", map_states$state,
            "<br/>", "<b>Net Increase in Federal</b>", "<br/>", "<b>Spending</b>: $",
            map_states$incr_fed_spending_mil, "million")) %% addLabelOnlyMarkers(data = filter(state.c
    state abb != "AK" & state abb != "HI"), lng = ~x, lat = ~y, label = ~state abb,
   labelOptions = labelOptions(noHide = T, direction = "top", textOnly = T)) %>%
    addLayersControl(baseGroups = c("Contained OOP Costs (2015)", "Premium Growth Rate (2010-2015)",
        "Affordable Marketplace Plan (2017)", "Increase in Federal Spending (2016)"),
        options = layersControlOptions(collapsed = FALSE)))
```

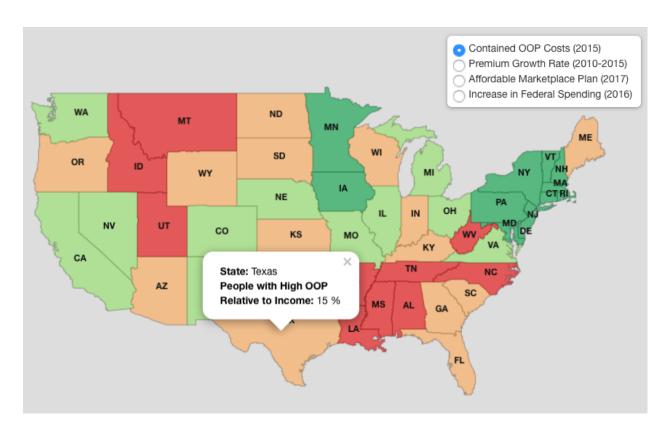


Figure 16:

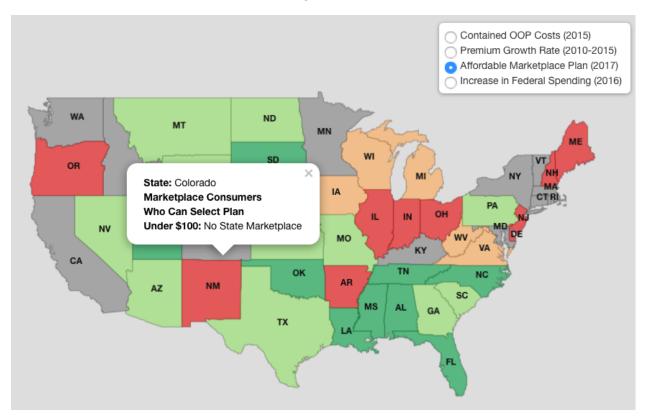


Figure 17:

Finally, we prepared to rank states on various measures using a data table. We distilled form our 10+ key indicators to 6 dimensions to rank states on. We made a version of the data table with values, and a version with the explicit ranks.

```
# Create data table reporting values
data_table_values <- key_indicators[, c("state", "insured_pct_2015", "deaths_amenable_2014",
    "usual_care_2015", "high_00P_relative", "premium_ann_growth_10_15", "incr_fed_spending_mil")]
# Create data table reporting rank
data_table_rank <- data_table_values[, c("state", "insured_pct_2015", "deaths_amenable_2014",
    "usual_care_2015", "high_00P_relative", "premium_ann_growth_10_15")]
# Convert each function to rank Rank 1 is always more favorable (regardless
# of definition of variable)
data_table_rank$r_insured_pct_2015 <- rank(-data_table_values$insured_pct_2015,
   na.last = "keep", ties.method = "min")
data_table_rank$r_deaths_amenable_2014 <- rank(data_table_values$deaths_amenable_2014,
   na.last = "keep", ties.method = "min")
data_table_rank$r_usual_care_2015 <- rank(-data_table_values$usual_care_2015,
   na.last = "keep", ties.method = "min")
data_table_rank$r_high_00P_relative <- rank(data_table_values$high_00P_relative,
   na.last = "keep", ties.method = "min")
data_table_rank$r_premium_ann_growth_10_15 <- rank(data_table_values$premium_ann_growth_10_15,
   na.last = "keep", ties.method = "min")
data_table_rank$r_incr_fed_spending_mil <- rank(data_table_values$incr_fed_spending_mil,
   na.last = "keep", ties.method = "min")
# Remove value variables
data_table_rank$insured_pct_2015 <- NULL</pre>
data_table_rank$deaths_amenable_2014 <- NULL
data_table_rank$usual_care_2015 <- NULL</pre>
data_table_rank$high_00P_relative <- NULL</pre>
data_table_rank$premium_ann_growth_10_15 <- NULL
data_table_rank$incr_fed_spending_mil <- NULL</pre>
```

We intially thought that putting the values into the data table would add additional useful information. And we thought that the rank would be apparent just by looking at the relative position in the data table. But because there were 50 rows, it was unfeasible for a viewer to count which row the state belonged to. Thus, we decided to report the explicit ranks instead, since ranking was the objective of this section. The detailed values could be seen in our leaflet maps anyway.

Showing 1 to 10 of 50 entries

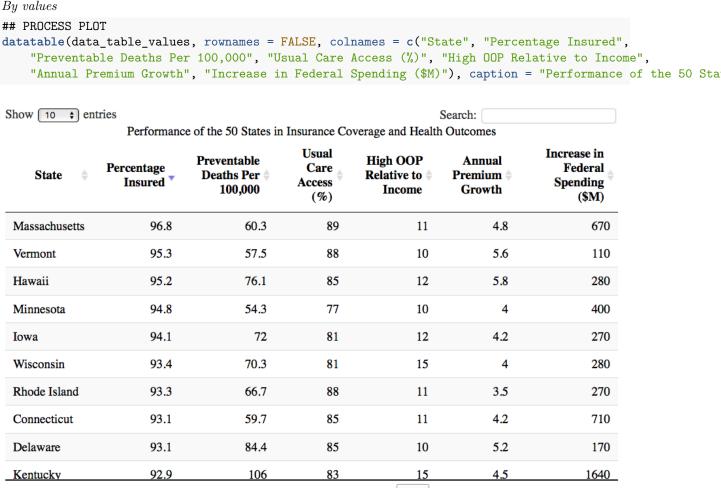


Figure 18:

Previous

2

3

5

Next

Final figure 12 (data table):

By ranks

Show 10 \$\display\$ entries Rank of the 50 States in Insurance Coverage				Search: [ge, Health and Cost Outcomes		
State \$	Percentage _ Insured	Fewer Preventable † Deaths	Usual Care \$ Access	OOP Contained	Lower Annual Premium Growth	Lower Increase in Federal Spending
Massachusetts	1	5	1	5	24	22
Vermont	2	2	2	1	42	2
Hawaii	3	24	7	9	45	14
Minnesota	4	1	34	1	10	18
Iowa	5	19	18	9	14	12
Wisconsin	6	14	18	26	10	14
Rhode Island	7	11	2	5	6	12
Connecticut	8	4	7	5	14	24
Delaware	8	31	7	1	32	5
Kentucky	10	44	12	26	20	40

Figure 19:

After the presentation, where we tried to verbally assess the rankings of the states, we realized that what we needed was an overall rank. Yes, it is a simplification in many regards, as we try to combine outcomes in three dimensions and reflect it in an overall rank. But as a journalistic piece, often the viewers would wonder... so which state is the best in the end?

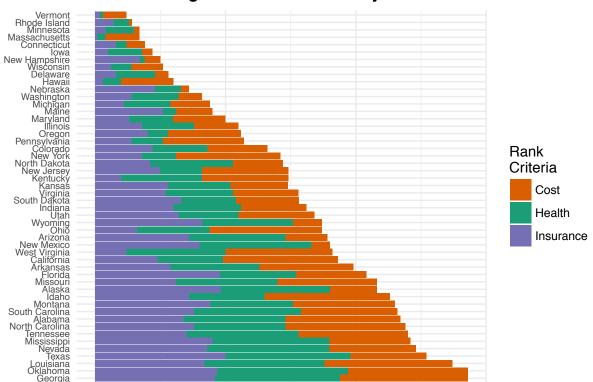
Thus, we used a simple method of aggregating the rankings of the states in the above six health measures into an overall rank. We rank each state on each of three dimensions, insurance coverage, health outcomes, and cost control outcomes. For the health and cost control outcomes, since they have multiple subcomponents (2 and 3 respectively, as in the data table), we take an average over these subcomponent ranks. We then add up the ranks in the three dimensions to find an overall rank to order the states.

```
# Compute overall ranking of states by each of three measures
data_table_rank_cat <- data_table_rank</pre>
# Insurance coverage
data_table_rank_cat$insurance_rank <- data_table_rank$r_insured_pct_2015
# Health
data_table_rank_cat$health_rank_sum <- data_table_rank$r_deaths_amenable_2014 +
    data_table_rank$r_usual_care_2015
data_table_rank_cat$health_rank <- rank(data_table_rank_cat$health_rank_sum,
    na.last = "keep", ties.method = "min")
# Finance
data_table_rank_cat$cost_rank_sum <- data_table_rank$r_high_00P_relative + data_table_rank$r_premium_an
    data table rank$r incr fed spending mil
data_table_rank_cat$cost_rank <- rank(data_table_rank_cat$cost_rank_sum, na.last = "keep",</pre>
    ties.method = "min")
data_table_rank_cat <- data_table_rank_cat[, c("state", "insurance_rank", "health_rank",</pre>
    "cost_rank")]
data_table_rank_cat$sum_of_ranks <- data_table_rank_cat$insurance_rank + data_table_rank_cat$health_ran
    data_table_rank_cat$cost_rank
data_table_rank_cat$overall_rank <- rank(data_table_rank_cat$sum_of_ranks, na.last = "keep",
    ties.method = "min")
data_table_rank_cat_long <- data_table_rank_cat</pre>
data_table_rank_cat_long <- reshape(data_table_rank_cat, varying = c("insurance_rank",</pre>
    "health_rank", "cost_rank"), v.names = "Rank", timevar = "Category", times = c("Insurance",
    "Health", "Cost"), new.row.names = 1:1000, direction = "long")
```

Final figure 13. Our actual plot is an interactive one, where you can toggle the legend to chose which ranks to 'add up' in the figure.

```
plot13 <- (ggplot(data_table_rank_cat_long, aes(x = reorder(state, -overall_rank),
    y = Rank, fill = Category, label = overall_rank)) + geom_col(position = "stack") +
    coord_flip() + labs(x = "", y = "", fill = "Rank \nCriteria") + scale_fill_manual(values = brewer.p
    name = "Dark2")[c(2, 1, 3)]) + ggtitle("Overall Ranking of States Healthcare Systems") +
    theme_minimal() + theme(plot.title = element_text(face = "bold", hjust = 0,
    size = 12), axis.text.y = element_text(size = 6.5), axis.ticks.x = element_blank(),
    axis.text.x = element_blank()))
plot13</pre>
```

Overall Ranking of States Healthcare Systems



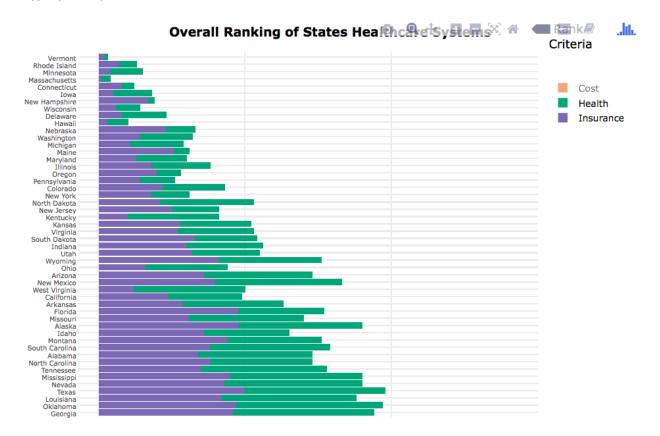


Figure 20:

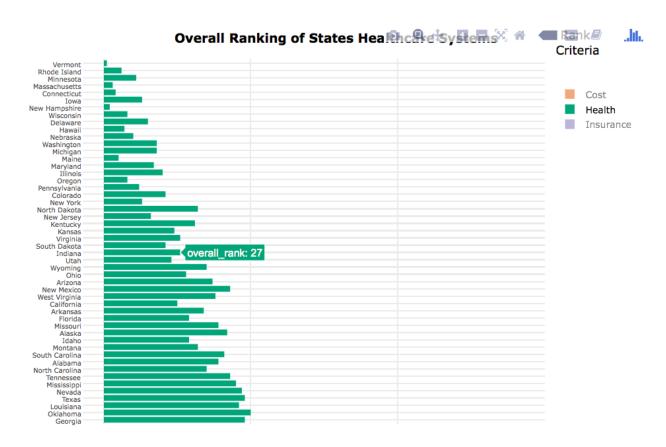


Figure 21:

Perceptions of ACA - NY Times Analaysis

We are also interested in how people think about Affordable care act(ACA). We want to have a look of the tweets and newspaper related to ACA. To realize this, we use API to download articles realted to ACA on New york times from 2011-2017 and transform them into corpus to do the text analysis.

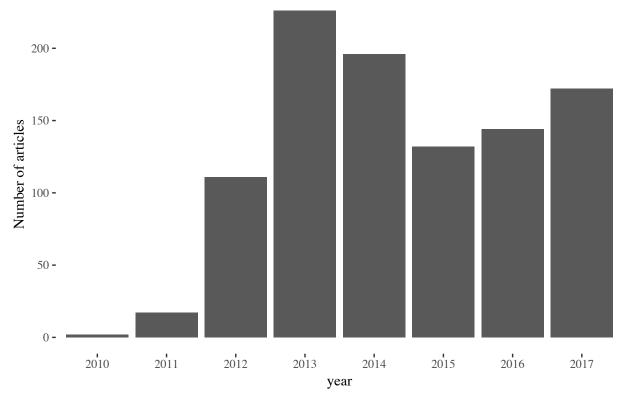
Research Questions: 1. The overll trend of public attention, which can be shown as the number of articles related to ACA in different years. 2. What people discuss about when they discuss ACA, which can be shown as the word frequencies. We could see that people discussed ACA a lot when obama first signed it, and the election in 2017 made it a hot topic again. And not suprisingly, people always talk about Trump when they talk about ACA

Figure 14

We could see that people discussed ACA a lot when Obama first signed it, and the election in 2017 made it a hot topic again. And not suprisingly, people always talk about Trump when they talk about ACA.

Draft plot:

Number of Articles Mentioning ACA by Year



Final figure 14:

Number of Articles Mentioning ACA by Year

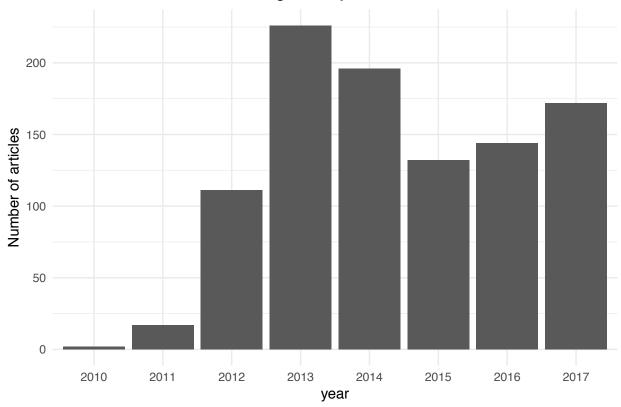


Figure 15

```
load("nytimes.rda")
dfmtotal <- dfm(corpus, remove = stopwords("english"), stem = TRUE, removePunct = TRUE,</pre>
    removeNumbers = TRUE, tolower = TRUE, verbose = TRUE)
## Creating a dfm from a corpus ...
##
      ... lowercasing
##
      ... tokenizing
      ... found 1,000 documents, 22,386 features
##
## ...
## removed 169 features, from 174 supplied (glob) feature types
## ... stemming features (English)
\#\# , trimmed 7834 feature variants
      ... created a 1,000 x 14,383 sparse dfm
      ... complete.
##
## Elapsed time: 1.89 seconds.
dfmtotal[, 1:5]
## Document-feature matrix of: 1,000 documents, 5 features (88.9% sparse).
head(stopwords("english"), 20)
## [1] "i"
                                   "my"
                     "me"
                                                "myself"
                                                              "we"
```

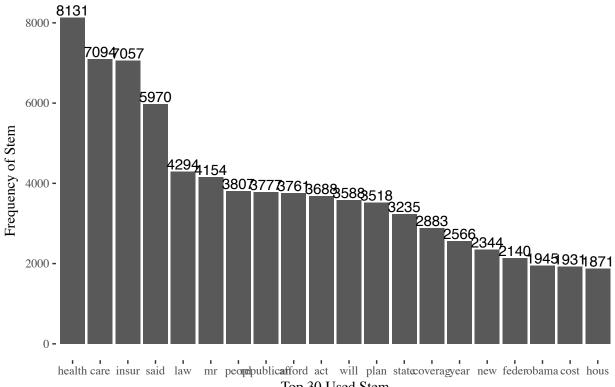
```
[6] "our"
                        "ours"
                                      "ourselves"
                                                                    "your"
##
   [11] "yours"
                        "yourself"
                                      "yourselves" "he"
                                                                    "him"
  [16] "his"
                       "himself"
                                      "she"
                                                                    "hers"
freq <- topfeatures(dfmtotal, 20)</pre>
wf <- data.frame(word = names(freq), freq = freq)</pre>
```

The graph shows that the frequently use words are health, insur and care. And other words like law, people and republican are also frequently used. It is not surprising that the articles most 40 features on the health care issues, and yet the topics of people, policy and politics are also highly concerned in New York Times.

Draft plot:

```
p <- ggplot(subset(wf, freq > 50), aes(word, freq))
p <- p + geom_bar(stat = "identity")</pre>
p <- p + geom_text(aes(label = freq), vjust = -0.2) + scale_x_discrete(limits = wf$word)</pre>
p <- p + theme(axis.text.x = element_text(angle = 45, hjust = 1), plot.title = element_text(face = "bol
    hjust = 0.5, size = 12)) + ggtitle("Word Frequency from Articles on ACA, 2011-2017") +
    theme tufte() + ylab("Frequency of Stem") + xlab("Top 30 Used Stem")
p
```

Word Frequency from Articles on ACA, 2011–2017



Top 30 Used Stem

Final figure 15 - included interactivity with plotly.

Word Frequency from Articles on ACA, 2011-2017

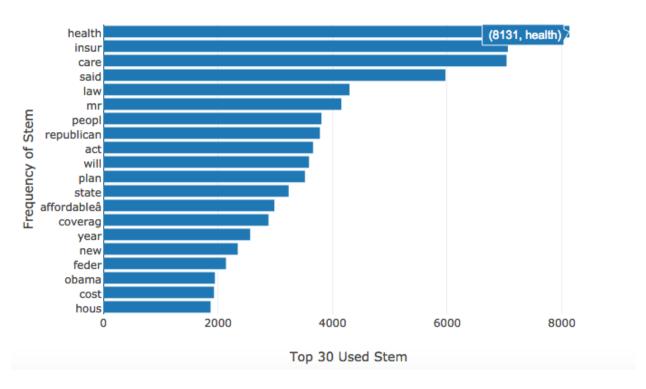


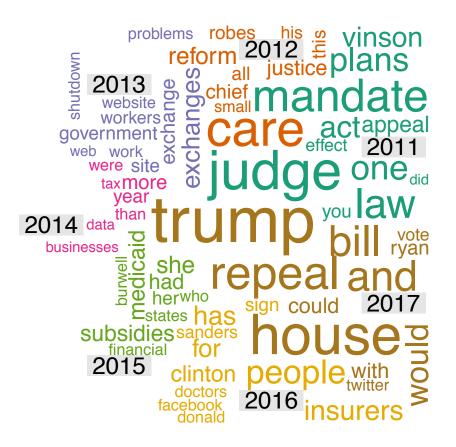
Figure 22:

```
load("nytimes.rda")
NYT_source <- VectorSource(corpus$documents[, 1])</pre>
NYT <- VCorpus(NYT_source)</pre>
documents <- corpus$documents
removeNumPunct <- function(x) {</pre>
    gsub("[^[:alpha:][:space:]]*", "", x)
clean corpus <- function(corpus) {</pre>
    corpus <- tm_map(corpus, removePunctuation)</pre>
    corpus <- tm_map(corpus, content_transformer(tolower))</pre>
    corpus <- tm_map(corpus, content_transformer(replace_symbol))</pre>
    corpus <- tm_map(corpus, removeWords, c(stopwords("english"), "will", "can"))</pre>
    # We could add more stop words as above
    corpus <- tm_map(corpus, stripWhitespace)</pre>
    corpus <- tm_map(corpus, removeNumbers)</pre>
    corpus <- tm_map(corpus, content_transformer(removeNumPunct))</pre>
    return(corpus)
}
NYT_clean <- clean_corpus(NYT)</pre>
NYT_stem <- tm_map(NYT_clean, stemDocument)</pre>
meta(NYT_stem, type = "local", tag = "author") <- documents$author</pre>
NYT dtm <- DocumentTermMatrix(NYT stem)</pre>
NYT_tdm <- TermDocumentMatrix(NYT_stem)</pre>
NYT tdm2 <- tidy(NYT tdm)</pre>
NYT dtm2 <- tidy(NYT dtm)
NYT_tidy <- tidy(NYT_stem)</pre>
NYT_tdm2 <- merge(NYT_tdm2, NYT_tidy, by.x = "document", by.y = "id", all.x = TRUE)
NYT_dtm2 <- merge(NYT_dtm2, NYT_tidy, by.x = "document", by.y = "id", all.x = TRUE)
atxt <- documents$texts[documents$datetimestamp == "2011"]</pre>
btxt <- documents$texts[documents$datetimestamp == "2012"]</pre>
ctxt <- documents$texts[documents$datetimestamp == "2013"]</pre>
dtxt <- documents$texts[documents$datetimestamp == "2014"]</pre>
etxt <- documents$texts[documents$datetimestamp == "2015"]</pre>
ftxt <- documents$texts[documents$datetimestamp == "2016"]</pre>
gtxt <- documents$texts[documents$datetimestamp == "2017"]</pre>
clean.text <- function(x) {</pre>
    # tolower
    x = tolower(x)
    # remove rt
    x = gsub("rt", "", x)
    # remove at
    x = gsub("@/\w+", "", x)
    # remove punctuation
    x = gsub("[[:punct:]]", "", x)
    # remove numbers
    x = gsub("[[:digit:]]", "", x)
    # remove links http
    x = gsub("http\\w+", "", x)
    # remove tabs
  x = gsub("[ | t]{2,}", "", x)
```

```
# remove blank spaces at the beginning
    x = gsub("^", "", x)
    # remove blank spaces at the end
    x = gsub(" $", "", x)
    return(x)
}
aclean <- clean.text(atxt)</pre>
bclean <- clean.text(btxt)</pre>
cclean <- clean.text(ctxt)</pre>
dclean <- clean.text(dtxt)</pre>
eclean <- clean.text(etxt)</pre>
fclean <- clean.text(ftxt)</pre>
gclean <- clean.text(gtxt)</pre>
a <- paste(aclean, collapse = " ")
b <- paste(bclean, collapse = " ")</pre>
c <- paste(cclean, collapse = " ")</pre>
d <- paste(dclean, collapse = " ")</pre>
e <- paste(eclean, collapse = " ")
f <- paste(fclean, collapse = " ")</pre>
g <- paste(gclean, collapse = " ")
# put everything in a single vector
all \leftarrow c(a, b, c, d, e, f, g)
all <- removeWords(all, c("will", "can", "the", "that", "are", "mrs", "not",
    "said", "cou"))
```

Wordcloud by Year: In the beginning, the articles mainly focused on the law issues as we can see from the frequently used words such as federal, act, mandate, judge, reform, etc. As time passed, the topics mainly focused on financial issues, the words that articles frequently mentioned are workers, business, tax, financial, subsidies, medicaid, etc. In the latest two years, due to the presidential election, the topics are changed to political concerns. For example, the words like people, trump, republicans, house, and repeal are frequently used in the articles.

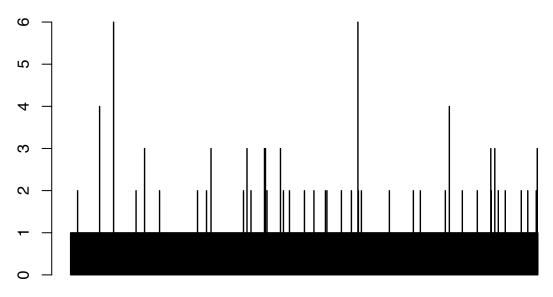
Final figure 16:



Perceptions of ACA - Twitter API Analaysis

```
library(httr)
##
## Attaching package: 'httr'
## The following object is masked from 'package:NLP':
##
##
       content
## The following object is masked from 'package:plotly':
##
##
       config
# library(oauth)
library(ROAuth)
library(twitteR)
##
## Attaching package: 'twitteR'
## The following object is masked from 'package:qdapTools':
##
##
## The following objects are masked from 'package:dplyr':
##
##
       id, location
## The following object is masked from 'package:plyr':
##
##
       id
library(RCurl)
## Loading required package: bitops
##
## Attaching package: 'RCurl'
## The following object is masked from 'package:tidyr':
##
##
       complete
library(RJSONIO)
library(stringr)
# secretkey myapp <- oauth_app('twitter', key = 'liLn6XJFenGjtvWFwi5LnDS1M',
# secret = 'dsCBm9Kyaeu9GMKlM9xwKl7eKmDn6qsjP31LQtwMGkF60QdLh6') Get OAuth
# credentials twitter_token <- oauth1.0_token(oauth_endpoints('twitter'),</pre>
# myapp)
# Declare Twitter API Credentials
api key <- "liLn6XJFenGjtvWFwi5LnDS1M" # From dev.twitter.com
api_secret <- "dsCBm9Kyaeu9GMK1M9xwK17eKmDn6qsjP31LQtwMGkF60QdLh6" # From dev.twitter.com
token <- "772176811455381505-PYuNAEqhHFc02r83WS9Y5dnsZciIY5v" # From dev.twitter.com
token_secret <- "mgRPwKeHZEw9Y486h2GMtCBxDztPfQLX1Msd5vog1hiwv" # From dev.twitter.com
# Create Twitter Connection
```

```
library("base64enc")
setup_twitter_oauth(api_key, api_secret, token, token_secret)
## [1] "Using direct authentication"
# Run Twitter Search. Format is searchTwitter('Search Terms', n=100,
# lang='en', geocode='lat,lng', also accepts since and until).
tweets <- searchTwitter("Obamacare OR ACA OR 'Affordable Care Act' OR #ACA",
   n = 1000, lang = "en", since = "2014-08-20")
# Transform tweets list into a data frame
tweets.df <- twListToDF(tweets)</pre>
head(tweets.df, 3)
##
## 1 Republicans wasted eight yrs. complaining & whipping frenzi over what 76% Majority of American
          I'm grateful my pre-existing condition (SIN) was covered by the Affordable Care Act of JESUS
## 3
                                                                    How the Affordable Care Act Drove Do
##
    favorited favoriteCount replyToSN
                                                    created truncated
## 1
         FALSE
                           0
                                  <NA> 2017-05-09 17:14:00
                                                                 TRUE
## 2
         FALSE
                           0
                                  <NA> 2017-05-09 17:13:20
                                                                FALSE
## 3
         FALSE
                           Λ
                                  <NA> 2017-05-09 17:13:00
                                                                FALSE
                                id replyToUID
    replyToSID
## 1
           <NA> 861992660822704128
                                          <NA>
## 2
           <NA> 861992493352669186
                                          <NA>
## 3
           <NA> 861992411777683456
                                          <NA>
##
                                                               statusSource
## 1
        <a href="http://twitter.com" rel="nofollow">Twitter Web Client</a>
## 2 <a href="http://www.facebook.com/twitter" rel="nofollow">Facebook</a>
## 3
                  <a href="http://bufferapp.com" rel="nofollow">Buffer</a>
##
         screenName retweetCount isRetweet retweeted longitude latitude
## 1
           etecbill
                               0
                                     FALSE
                                                FALSE
                                                                      NA
## 2 wildlyfejblood
                               0
                                     FALSE
                                                FALSE
                                                             NA
                                                                      NA
        spediatrics
                               0
                                     FALSE
                                                FALSE
                                                             NA
                                                                      NA
counts = table(tweets.df$screenName)
barplot(counts)
```



_MelissaR Corevity H3RJay kayfey mikev58 rbole tayinuk

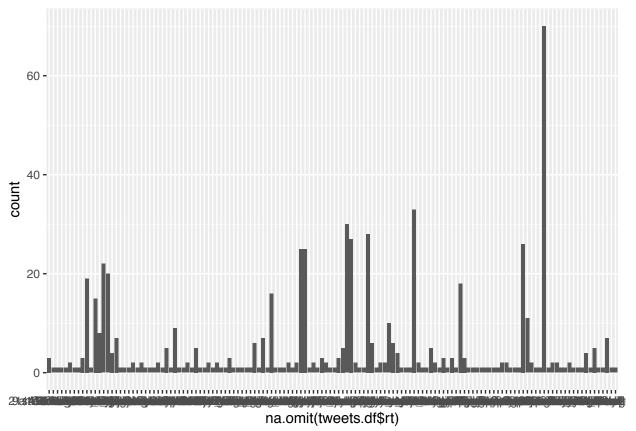
Some data exploration

library(stringr)

Let's do something hacky: Limit the data set to show only folk who tweeted twice or more in the sample.

Now we can plot a chart showing how often a particular person was RT'd in our sample.

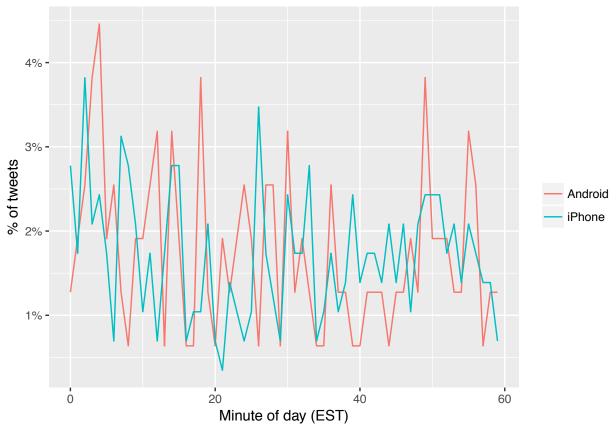
```
library(ggplot2)
ggplot() + geom_bar(aes(x = na.omit(tweets.df$rt)))
```

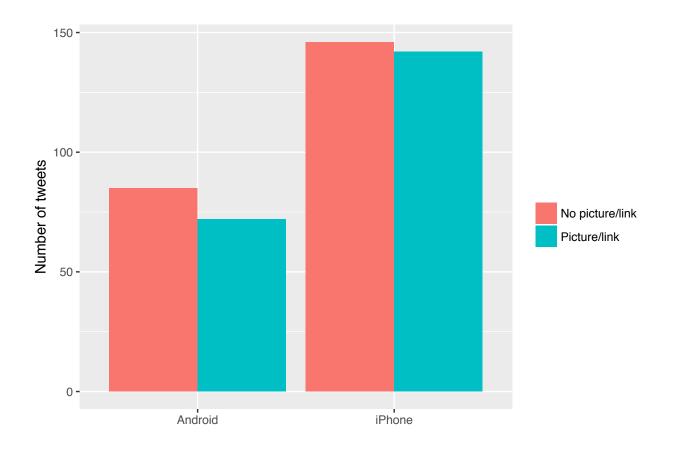


```
library(tidyr)
library(dplyr)
library(purrr)
```

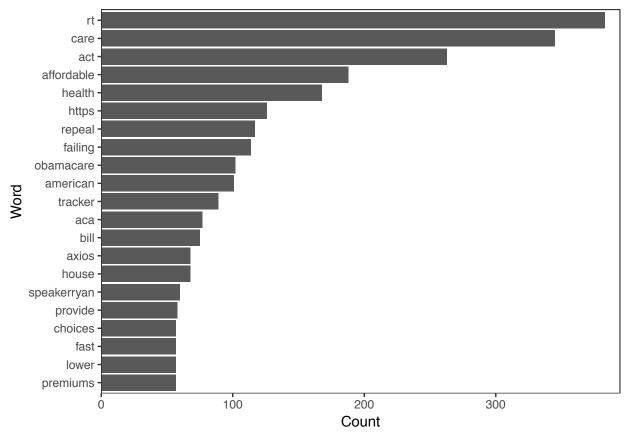
```
##
## Attaching package: 'purrr'
## The following object is masked from 'package:qdap':
##
## %>%
## The following objects are masked from 'package:dplyr':
##
## contains, order_by
## The following object is masked from 'package:plyr':
```

```
##
##
       compact
## The following object is masked from 'package:magrittr':
##
       set_names
## The following object is masked from 'package:maps':
##
       map
tweets <- tweets.df %>% select(id, statusSource, text, created) %>% extract(statusSource,
    "source", "Twitter for (.*?)<") %>% filter(source %in% c("iPhone", "Android"))
table(tweets$source)
##
## Android iPhone
##
       157
               288
library(lubridate)
library(scales)
##
## Attaching package: 'scales'
## The following object is masked from 'package:purrr':
##
##
       discard
## The following objects are masked from 'package:readr':
##
       col_factor, col_numeric
##
tweets %>% count(source, minute = minute(with_tz(created, "EST"))) %>% mutate(percent = n/sum(n)) %>%
    ggplot(aes(minute, percent, color = source)) + geom_line() + scale_y_continuous(labels = percent_formula)
    labs(x = "Minute of day (EST)", y = "% of tweets", color = "")
```



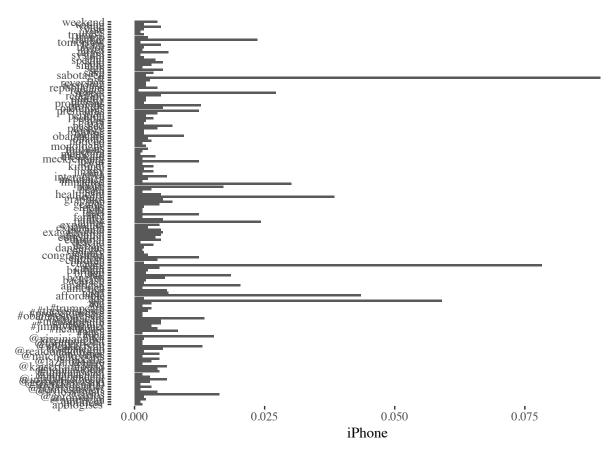


Comparison of words Now that we're sure there's a difference, what can we say about the difference in the content? We'll use the tidytext package. We start by dividing into individual words using the unnest_tokens function, and removing some common stopwords. As we can see the most frequently used word when discussing ACA on Twitter is rt (Republican Party), care, act, afforadable, health, obama, and so on. From the table, we found that most people discussed about political issues (i.e., the topics of republican and democratic parties), affordable care act itself, and financial concerns (i.e., bill, pay, etc).



```
android_iphone_ratios <- tweet_words %>% count(word, source) %>% filter(sum(n) >=
    5) %>% spread(source, n, fill = 0) %>% ungroup() %>% mutate_each(funs((. +
    1)/sum(. + 1)), -word) %>% mutate(logratio = log2(Android/iPhone)) %>% arrange(desc(logratio))

ggplot(data = android_iphone_ratios, aes(x = word, y = iPhone)) + geom_bar(stat = "identity") +
    # geom_text(label= Android, color='red') +
xlab(NULL) + coord_flip() + theme_tufte()
```



Sentiment analysis: Since we've observed a difference in sentiment between the Android and iPhone tweets, let's try quantifying it. We'll work with the NRC Word-Emotion Association lexicon, available from the tidytext package, which associates words with 10 sentiments: positive, negative, anger, anticipation, disgust, fear, joy, sadness, surprise, and trust.

To measure the sentiment of the Android and iPhone tweets, we can count the number of words in each category. (For example, we see that 41 of the 2331 words in the Android tweets were associated with "anger"). We then want to measure how much more likely the Android account is to use an emotionally-charged term relative to the iPhone account. Since this is count data, we can use a Poisson test to measure the difference:

```
nrc <- sentiments %>% filter(lexicon == "nrc") %>% dplyr::select(word, sentiment)
sources <- tweet_words %>% group_by(source) %>% mutate(total_words = n()) %>%
    ungroup() %>% distinct(id, source, total_words)
by_source_sentiment <- tweet_words %>% inner_join(nrc, by = "word") %>% count(sentiment,
    id) %>% ungroup() %>% # complete(sentiment, id, fill = list(n = 0)) %>%
inner_join(sources) %% group_by(source, sentiment, total_words) %>% summarize(words = sum(n)) %>%
    ungroup()
## Joining, by = "id"
head(by_source_sentiment)
## # A tibble: 6 × 4
##
                sentiment total_words words
      source
##
       <chr>
                                <int> <int>
                    <chr>
## 1 Android
                    anger
                                 1812
                                         66
```

89

1812

2 Android anticipation

```
## 3 Android
                   disgust
                                   1812
                                            8
## 4 Android
                                   1812
                                            81
                      fear
## 5 Android
                       joy
                                   1812
                                            30
## 6 Android
                                   1812
                  negative
                                           113
```

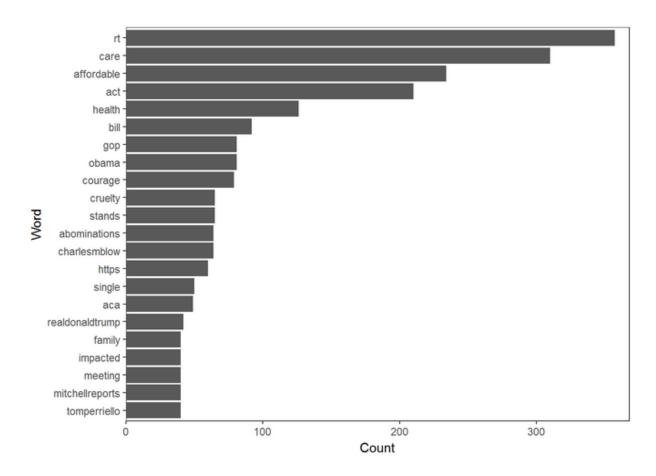
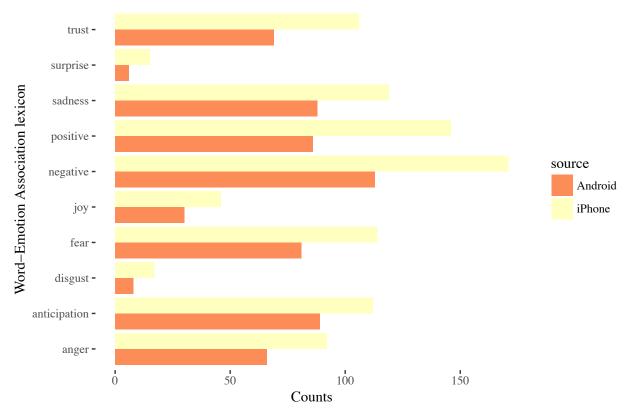


Figure 23:

From the below table, we could see that most people feel quite positive about ACA, the words they used are expressed their trust, anticipation, joy and positive feeling.

Sentiment Analysis from Tweets on ACA



Data Preparation using Twitter: The Twitter search API does not return an exhaustive list of tweets that match your search criteria, as Twitter only makes available a sample of recent tweets. For a more comprehensive search, we will need to use the Twitter streaming API, creating a database of results and regularly updating them, or use an online service that can do this. Now that we have tweet texts, we need to clean them up before doing any analysis. This involves removing content, such as punctuation, that has no emotional content, and removing any content that causes errors.

```
#text cleaning
library(tm)
# build a corpus, and specify the source to be character vectors
myCorpus <- Corpus(VectorSource(tweets.df$text))</pre>
# convert to lower case
# tm v0.6
myCorpus <- tm_map(myCorpus, content_transformer(tolower))</pre>
# tm v0.5-10
# myCorpus <- tm_map(myCorpus, tolower)</pre>
# remove URLs
removeURL <- function(x) gsub("http[^[:space:]]*", "", x)</pre>
# tm v0.6
myCorpus <- tm_map(myCorpus, content_transformer(removeURL))</pre>
# tm v0.5-10
# myCorpus <- tm map(myCorpus, removeURL)</pre>
# remove anything other than English letters or space
removeNumPunct <- function(x) gsub("[^[:alpha:][:space:]]*", "", x)</pre>
myCorpus <- tm_map(myCorpus, content_transformer(removeNumPunct))</pre>
# remove punctuation
# myCorpus <- tm map(myCorpus, removePunctuation)</pre>
# remove numbers
# myCorpus <- tm_map(myCorpus, removeNumbers)</pre>
# add two extra stop words: "available" and "via"
myStopwords <- c(stopwords('english'), "available", "via")</pre>
# remove "r" and "biq" from stopwords
myStopwords <- setdiff(myStopwords, c("r", "big"))</pre>
# remove stopwords from corpus
myCorpus <- tm_map(myCorpus, removeWords, myStopwords)</pre>
# remove extra whitespace
myCorpus <- tm_map(myCorpus, stripWhitespace)</pre>
# keep a copy of corpus to use later as a dictionary for stem completion
myCorpusCopy <- myCorpus</pre>
# stem words
myCorpus <- tm_map(myCorpus, stemDocument)</pre>
# inspect the first 5 documents (tweets)
# inspect(myCorpus[1:5])
# The code below is used for to make text fit for paper width
for (i in c(1:2, 320)) {
cat(paste0("[", i, "] ")) writeLines(strwrap(as.character(myCorpus[[i]]), 60))
}
## [1] exampl call java code r
## [2] simul mapreduc r big data analysi use flight data rblogger
## [320] r refer card data mine now cran list mani use r function
## packag data mine applic
# tm v0.5-10
# myCorpus <- tm map(myCorpus, stemCompletion)</pre>
# tm v0.6
```

```
stemCompletion2 <- function(x, dictionary) {</pre>
x <- unlist(strsplit(as.character(x), " "))</pre>
# Unexpectedly, stemCompletion completes an empty string to
# a word in dictionary. Remove empty string to avoid above issue.
x <- x[x != ""]
x <- stemCompletion(x, dictionary=dictionary)</pre>
x <- paste(x, sep="", collapse=" ") PlainTextDocument(stripWhitespace(x))
#myCorpus <- lapply(myCorpus, stemCompletion2, dictionary=myCorpusCopy) myCorpus <- Corpus(VectorSource
# count frequency of "mining"
miningCases <- lapply(myCorpusCopy,</pre>
function(x) { grep(as.character(x), pattern = "\\<mining")} )</pre>
sum(unlist(miningCases))
## [1] 82
# count frequency of "miner"
minerCases <- lapply(myCorpusCopy,</pre>
function(x) {grep(as.character(x), pattern = "\\miner")} )
sum(unlist(minerCases))
## [1] 5
# replace "miner" with "mining"
myCorpus <- tm_map(myCorpus, content_transformer(gsub),</pre>
pattern = "miner", replacement = "mining")
tdm <- TermDocumentMatrix(myCorpus,</pre>
control = list(wordLengths = c(1, Inf)))
## <<TermDocumentMatrix (terms: 822, documents: 320)>>
## Non-/sparse entries: 2460/260580
## Sparsity: 99%
## Maximal term length: 27
## Weighting : term frequency (tf)
#__figure 17___
#Frequent Words and Associations
term.freq <- rowSums(as.matrix(tdm))</pre>
term.freq <- subset(term.freq, term.freq >= 15)
df <- data.frame(term = names(term.freq), freq = term.freq)</pre>
ggplot(df, aes(x = term, y = freq)) + geom_bar(stat = "identity") +
xlab("Terms") + ylab("Count") + coord_flip()
```

Unsurprisingly, Word Cloud shows that most frequently mentioned terms are health, care, affordable, aca, and act. Except the discussion of insurance policy itself, people do talk frequently about political-related topics, such as Republican Party (rt), Grand Old Party (gop), and Obama. Also, some terms related to push forward or hold back the policy, such as courage, urge and repeal. Financial words are used, like pay, bill and save. Finally, specific names of people and place were mentioned in the tweets (i.e., Larry Levitt and Mecklenburg).

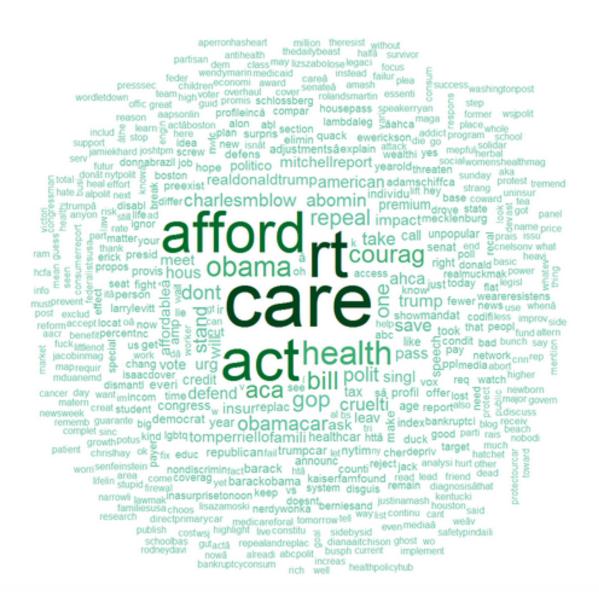


Figure 24:

Topic Modelling

n text mining, we often have collections of documents, such as social media posts or news articles, that we'd like to divide into natural groups so that we can understand them separately. Topic modeling is a method for unsupervised classification of such documents, similar to clustering on numeric data, which finds natural groups of items even when we're not sure what we're looking for.

Latent Dirichlet allocation (LDA) is a particularly popular method for fitting a topic model. It treats each document as a mixture of topics, and each topic as a mixture of words. This allows documents to "overlap" each other in terms of content, rather than being separated into discrete groups, in a way that mirrors typical use of natural language.

Latent Dirichlet allocation is one of the most common algorithms for topic modeling. Without diving into the math behind the model, we can understand it as being guided by two principles: Every document is a mixture of topics and Every topic is a mixture of words.

This visualization lets us understand the eight topics that were extracted from the tweets. The most common words in topic 1 include "rt", "care", and "health", which suggests it may represent health insurance and republican issues. Those most common in topic 2 include "obama", "act", and "bill", suggeting that this topic represents issues related to obamacare. One important observation about the words in each topic is that some words, such as "act" are common within both topics. This is an advantage of topic modeling as opposed to "hard clustering" methods: topics used in natural language could have some overlap in terms of words.

```
library(ggplot2)
top_terms %>% mutate(term = reorder(term, beta)) %>% ggplot(aes(term, beta,
    fill = factor(topic))) + geom_col(show.legend = FALSE) + facet_wrap(~topic,
    scales = "free") + coord_flip()
```

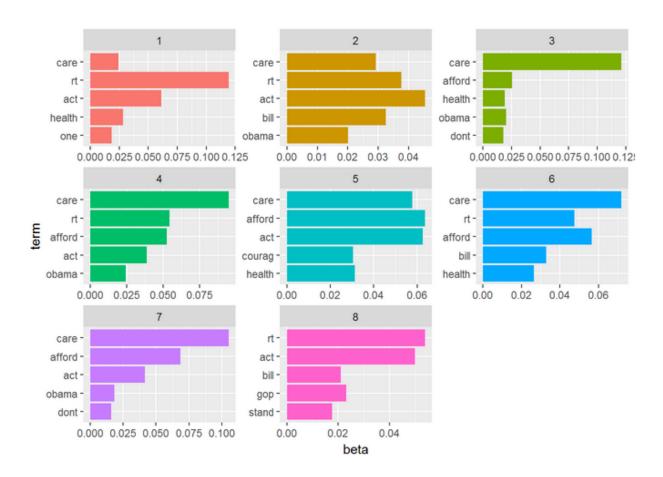


Figure 25:

Next Steps

We want to do more for the text analysis part, maybe include some sentiment analysis. And for the New York Times analysis, when we convert the articles into metadata, we abandoned many variables, which also limit the possibility to explore some possible relationship, so we might try to do some text analysis while control other variables. As we were not able to download all the tweets we didn't finish the text analysis based on tweets.

One further thing we would like to explore if we had full access to the tweets during the years, is to find out how perceptions of the ACA differed by state. In particular, we can investigate if there are party and regional differences, which would link up well with the earlier section of our project. Given tweets in this full time range, we would have liked to explore how these sentiments changed over time, especially in response to significant policy and political events.