

Slowing the Spread of Anti- Flu Shot Sentiment on Twitter

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James Chase @jason\_prosser · Jan 18

More out of curiosity, I hear people say that the flu shot makes them sick. My son & I both get the shot. We've been lucky, no flu yet.

#### Marni Hughes @marnihughesQ13

Health department isn't telling us yet @jason\_prosser But my son and I are recovering from the flu this week and we both got the shot.

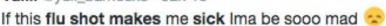


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Yuliii @yuli\_barriosx3 · Jan 18



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f#minor @wesleywyndam · Jan 13







I hate that getting a flu shot makes you sick







## **Experiment Question:**

Does the CDC's flu shot page slow the spread of misconceptions on Twitter?





# Designing the Administrator



- Friendly Avatar
- Healthy Living Enthusiast
- Non-bot Appearance
  - Unrelated tweets
  - Purchased followers



# Issues with Initial Experiment Design

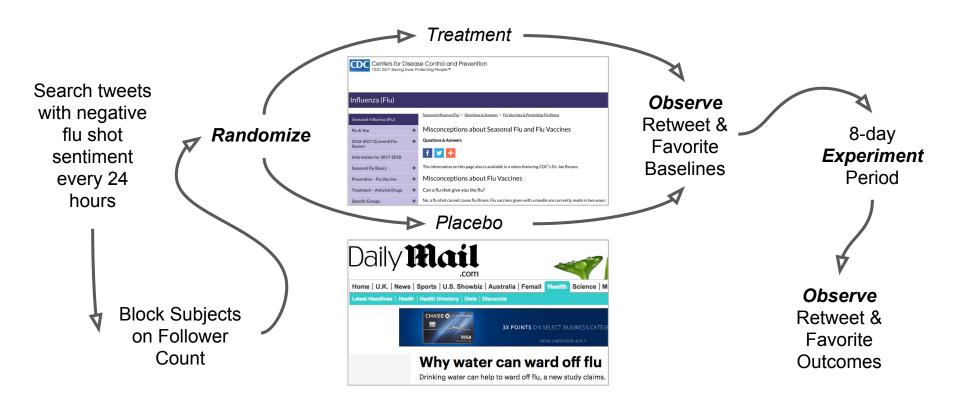


- Required Automation
  - NLP Classifier
  - Auto-replies
- Challenges
  - NLP package accuracy
  - Twitter spam blocker
  - US Flu season ending





# Final Experiment Design: ROXO

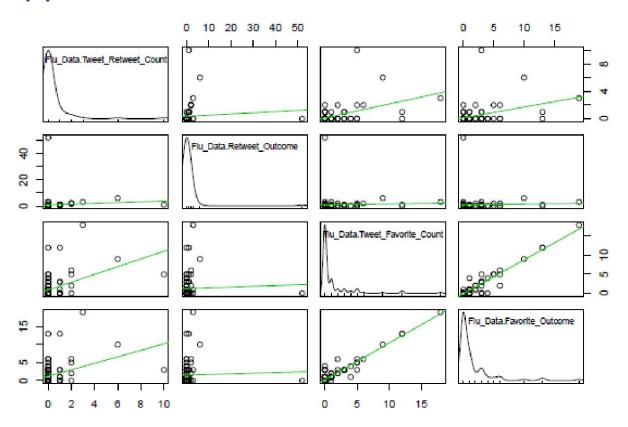




## **Collected Data**

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# EDA (I)





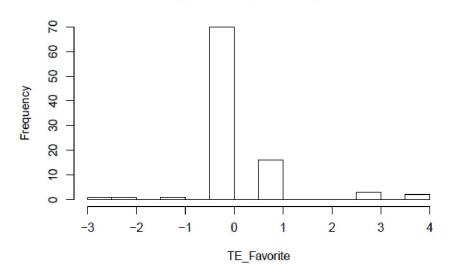
# EDA (II)

- Collected variables with numerical values have two characteristics:
  - right-skewed
  - outliers are far away from majority of data.
- Among variables User-Statuses-Count, User-Followers-Count, User-Friends-
- Count, User-Listed-Count, User-Followers-Count has most significant linear relationship with respect to Favorite-Outcome variable.
- By comparing data before and after experiments, Favorite-Outcome variable has much better trend with Tweet-Favorite-Count variable compared to Retweet-outcome and Tweet-Retweet-Count variable.



### **General ATE**

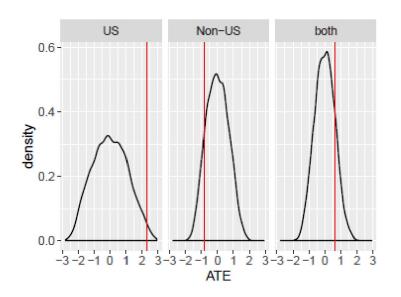
#### Histogram of Flu\_Data\$TE\_Favorite



- The individual treatment effect for each subject by favorite number
- For population treated with links
  - ATE = 0.53 (Retweet)
  - ATE = 0.29 (Favorite)
- Random assignment ATE for CDC treated links
  - ATE = 0 (Retweet)
  - ATE = 0.47 (Favorite)
- If we apply t-test, such difference is not statistically significant at the 0.05 level for Retweet measure, same for Favorite measure.



## Location Effect: U.S. v.s. Non-U.S.



- For CDC treatment:
  - ATE = 2.30 at U.S.
  - ATE = 0.85 at Non-U.S.
- We have to note that ATE value calculated from pool 0.6716826 different from ATE value obtained from combined one 0.6576287.
- This is a biased estimate because the probability of being assigned to the treatment group varies by block (area): in US this probability is 17 / 45 = 37 . 8%, while in Non-US the probability of being assigned to the treatment group is 26 / 49 = 53 . 1%.
- Besides, the number of Favorites is lower on average in Non-US, so the overall treatment effect calculated this way is larger than it actually is.
- Therefore, if outcomes were higher in the treatment group, it might reflect differences between US and Non-US rather than a treatment effect.



### **CACE Effect**

Treat. Ass.	Treated	No.#	Response
Baseline	NO	20	35%
Treatment	Yes	33	52%
Treatment	NO	10	34%
Placebo	Yes	40	33%
Placebo	No	11	36%

- Assume 80% of targets really get treated.
- For treatment and placebo groups, their compliance rate are not statistically different each other with p-value = 0.85 with t-test.
- The difference in the Never-Takers response rate between the treatment and placebo groups is not statistically significant since p = 0.77.
- CACE
  - 0.15 for receiving weblinks.
  - 0.19 among compilers.

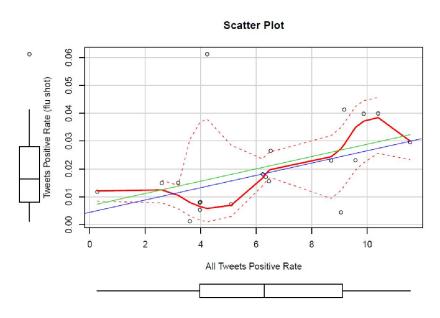


### Retweet v.s. Favorite

- Retweet\_Outcome ~ Assign\_Ind + Tweet\_Retweet\_Count +
   Tweet\_favorite\_Count (CDC = -1.0, p-value =0.4, negative effect with
   weaker statistical significance)
- Favorite\_Outcome ~ Assign\_Ind + Tweet\_Retweet\_Count +
   Tweet\_favorite\_Count (CDC = 0.3, p-value = 0.1, positive effect with
   stronger statistical significance)
- Favorite\_Outcome ~ Assign\_Ind\*Sex + Tweet\_Retweet\_Count + Tweet\_Favorite\_Count
  - the estimated effect of CDC is 0.4027 with p-value 0.1.
  - CDC has stronger causal effect for MALE.



# Gibbs Sampling for Missing New Tweets (I)



- During our experimental procedure, however, it is not easy to collect such data due to time constraints or other natural restrictions.
- The basic idea is that we first collect other available variable, the tweets positive rate (ALL) generated by a subject from his/her ALL previous Tweets averaged by days.
- The positive rate of new Tweets with flu shot topic is determined by linear model and tweets positive rate (ALL).

# Gibbs Sampling for Missing New Tweets (II)

• We assume we have paired data. We wish to find the posterior distributions of the coefficients \beta 0 (the intercept), \beta 1 (the gradient) and of the precision \tau, which is the reciprocal of the variance.

$$y_i \sim \mathcal{N}(eta_0 + eta_1 x_i, 1/ au)$$

- The massive advantage of Gibbs sampling over other MCMC methods is that no tuning parameters are required!
  - 1. Pick some initial  $\theta_2^{(i)}$ .
  - 2. Sample  $heta_1^{(i+1)}\sim p( heta_1| heta_2^{(i)},x)$ 3. Sample  $heta_2^{(i+1)}\sim p( heta_2| heta_1^{(i+1)},x)$

The estimated effect of CDC is -1.0 with p-value 0.38.



# **Future Improvements**



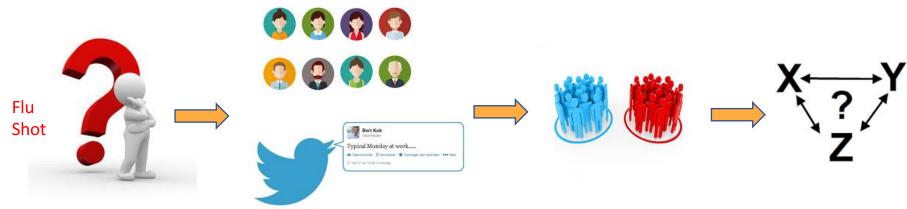




- Reply-bot circumventing Twitter spam rules
- Test different value engagement strategies
  - Authority
  - Objectivity
  - **□** Empathy
- Other public health outreach topics?
  - ☐ Childhood Immunizations
  - Antibiotics Use
  - Epidemics (Zika, Ebola, HIV, Malaria)







User	Tweets with Flu Shot	Control/Treatme nt	Read CDC/ No Read CDC	Increased Retweets number (outcome)	New tweets about flu shot after treatment (outcome)
api.search screen_name, user.name	api.search, tweets can do sentimental analysis	Group assignment for collected users.	Compliance or Non Compliance indicator	api.get_status	api.user_timeline