

Using Cluster Analysis to Develop a Tailored Contacting Strategy

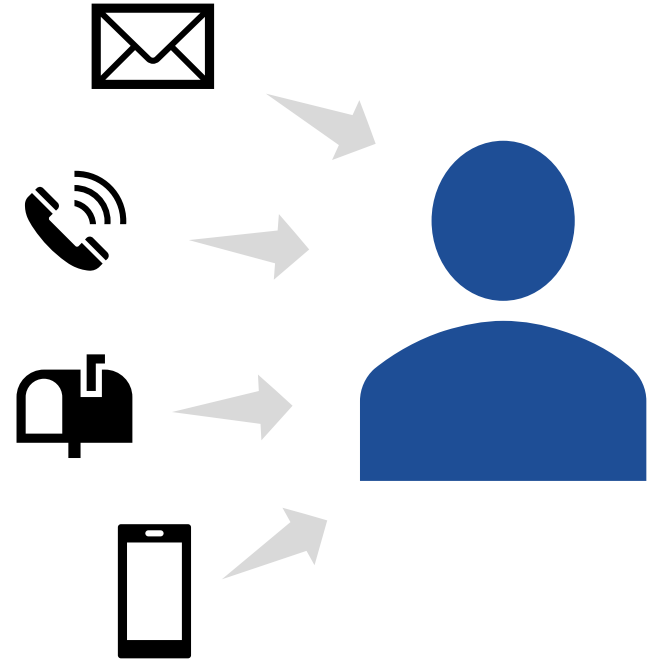
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Background

The Problem

- Researchers are challenged to find new methods for increasing survey response.
- Data collection effort cannot be increased indefinitely:
 - Limited project resources
 - Diminishing returns of additional contact attempts
 - Over-burdened sample members



Background

The Solution

- Find new ways to maximize the efficacy of each contact attempt.
- Contacting sample members using different modes (e.g., telephone, mail) is common practice, but all contact modes are not equally likely to result in response for all sample members (de Leeuw 2005).

Long term goal

Develop a **tailored contacting strategy** that identifies the optimal contacting approach for each SM:

- level of data collection effort (# of contacts)
- contact mode
- timing of contact attempts

Near term goal

Assess our ability to predict SMs' preferred contact times and whether these preferences are stable over time.

Research Questions

- Can clustering techniques be used to identify distinct groups of sample members that share similar characteristics?
- Does group membership have meaningful persistence across time? Or are they artifacts?
- Is there a simpler method to apply multiple interventions in succession (not remodeling each time)?
- Can cluster groups be used to inform individual data collection interventions?

About the Study

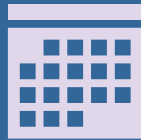
- Approach uses data from a nationally representative mixed-mode study of postsecondary students conducted in 2020 (base year).
- Pilot test of tailored contacting strategy conducted on follow-up study (subset of base year cohort) in 2022.



SAMPLE

Base year: ~170,000

Follow-up: ~37,000



DATA COLLECTION

Base year: 2020

Follow-up: 2022



SURVEY MODES

CATI & Web
(mobile-friendly)



CONTACT MODES

Phone, Email,
Hardcopy Mail, SMS

Sample Member Clustering Method

Our goal was to identify partitions among follow-up sample members that share similar characteristics (i.e., clusters).

Clustering Method

1. Subset the follow-up sample to base year respondents and cluster the cases using the k-prototypes algorithm.

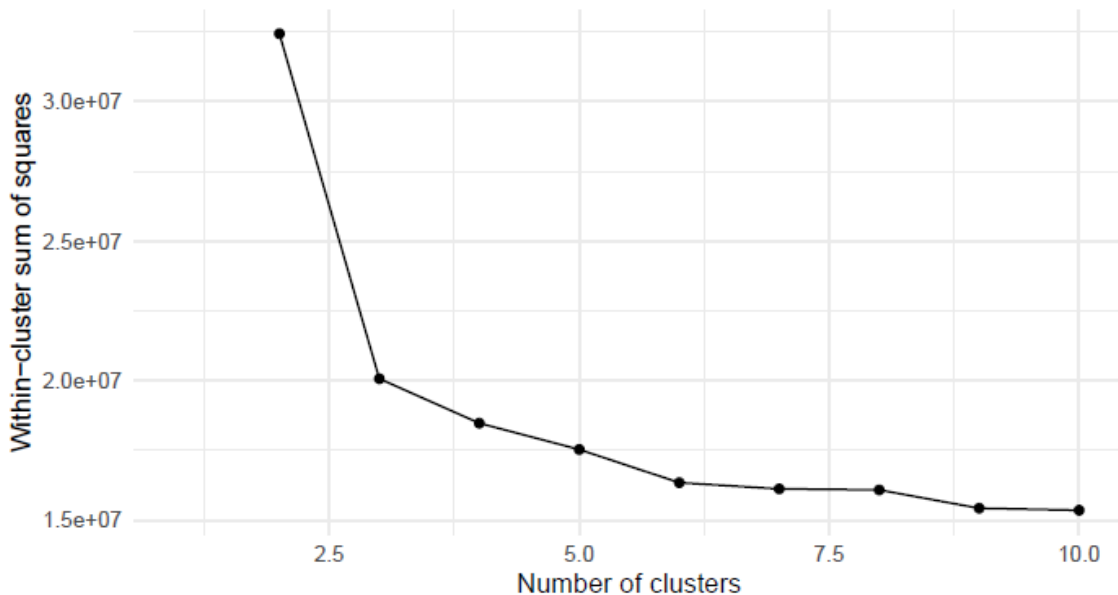
Used base year data (e.g., race, ethnicity, sex, institution sector) and paradata (e.g., completion mode and time, contact status)

2. Fit a support-vector machine (SVM) model to these clusters.
3. Predict cluster membership for the full follow-up sample, including base year respondents and nonrespondents.

Sample Member Clustering Method

Number of clusters

- The within-cluster sum of squares was evaluated for k-prototypes clusters solutions of 1 through 10.
- The 6-cluster solution was selected as optimal.



Sample Member Clustering Method

- Cluster analysis resulted in 6 sample clusters.
- Pilot experiment tested the preferred contact time predicted based on cluster.

Cluster	Predicted Preferred Contact Time	Sample Size
1	Morning	1,379
2	Evening	5,223
3	Afternoon	6,969
4	Afternoon	4,486
5	Afternoon	4,362
6	Afternoon	1,093
Total		23,512

Experiment Design

Contact tailoring pilot experiment:

- Vary timing of 1st reminder email
 - Experiment group: approximately 3/5 of full follow-up sample
 - Excludes cases that began data collection early or responded prior to reminder 1
- Send email in morning (10:00 EDT), afternoon (2:00 EDT), or evening (6:00 EDT) time slots based on cluster group.
- Email sent on April 19, 2022

Experiment Design

- Sample members were split into intervention and observation groups.
- Email was sent in the morning, afternoon, or evening.

Intervention group

Email time assigned based on cluster

Observation group

Email time assigned at random

	Intervention Group N = 11,932	Observation Group N = 11,930	
Email Time	Predicted & Actual Time	Predicted Time	Actual Time
Morning	678	701	3,923
Afternoon	8,454	8,456	3,914
Evening	2,630	2,593	3,913

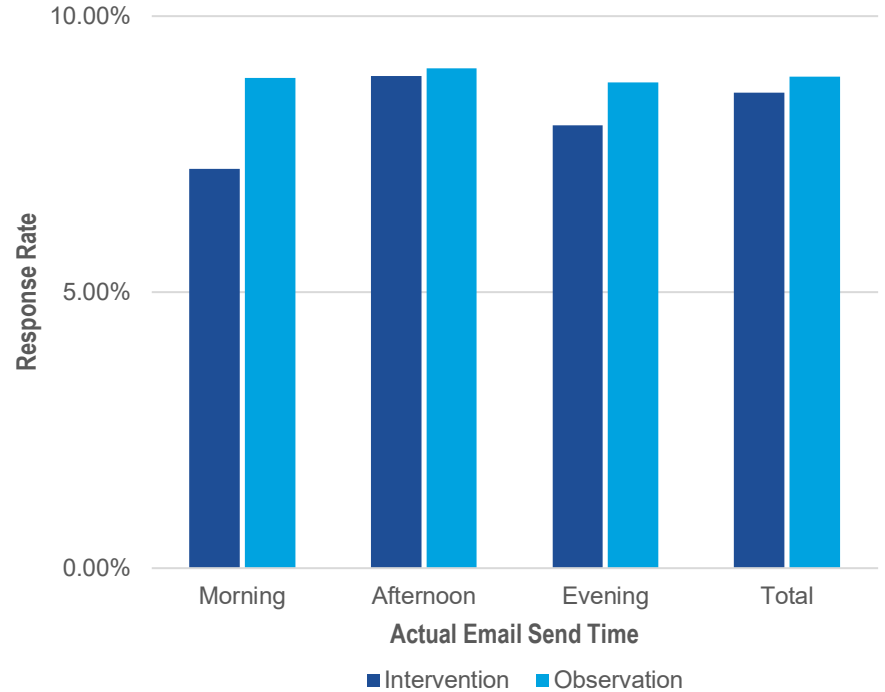
Results

- No significant difference in response rate between intervention and observation groups ($p < .05$).

Actual Email Time	Intervention Group RR (%)	Observation Group RR (%)
Morning	7.23	8.87
Afternoon	8.91	9.04
Evening	8.02	8.79
Total	8.61	8.90

Intervention vs. Observation Group by Email Send Time

As of 16 days post-intervention



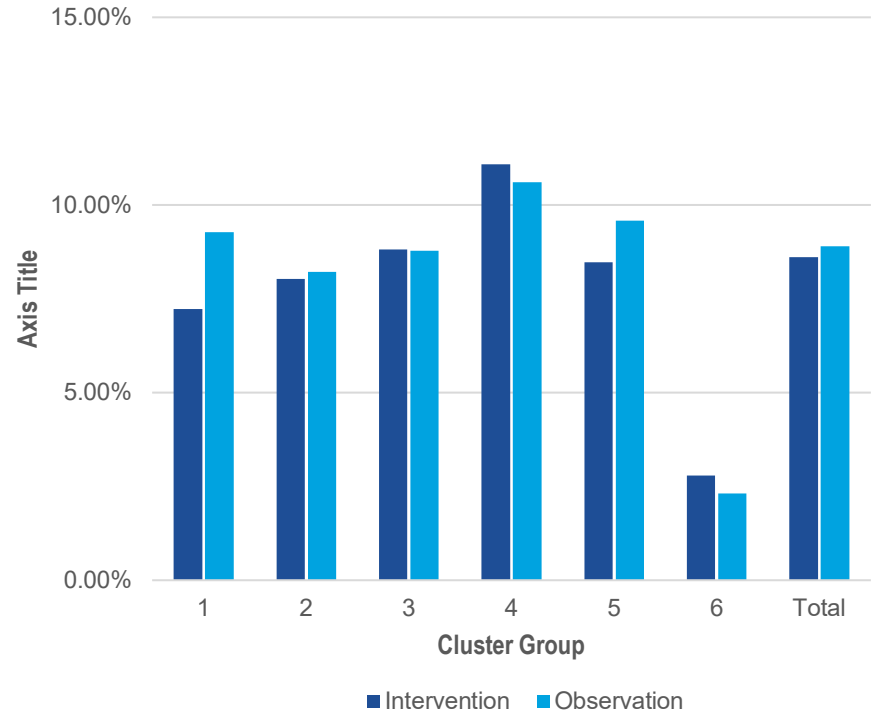
Results

- No significant difference in response rate between intervention and observation groups for any cluster ($p < .05$).

Cluster	Intervention Group RR (%)	Observation Group RR (%)
1	7.23	9.27
2	8.02	8.21
3	8.81	8.78
4	11.08	10.60
5	8.48	9.58
6	2.79	2.31
Total	8.61	8.90

Intervention vs. Observation Group by Cluster

As of 16 days post-intervention



Results

- Observation group email time was assigned at random.
- For approx. 1/3 of observation cases, actual and cluster-predicted time matched by chance (blue cells).
- No significant difference in response rate between matched and non-matched groups ($p < .05$).

Cluster & Predicted Preferred Contact Time	Observation Group RR (%) by Actual Email Time		
	Morning	Afternoon	Evening
1 (Morning)	7.91	10.13	9.64
2 (Evening)	9.06	8.69	6.87
3 (Afternoon)	7.73	9.37	9.23
4 (Afternoon)	10.94	10.05	10.85
5 (Afternoon)	10.01	9.22	9.49
6 (Afternoon)	1.91	1.73	3.16
Total	8.87	9.04	8.78
			RR (%)
Actual time MATCHES predicted time			8.52
Actual time DOES NOT MATCH predicted time			9.10

Conclusions

- Clustering analysis results indicate that it is possible to identify distinct groups of sample members that share similar characteristics.
- Our pilot attempt to convert these cluster groupings into an actionable contacting strategy based on time of day of contact was not successful.
- Possible explanations for null results:
 - Tailoring preferred time is not effective in isolation and must be combined with other interventions (e.g., day of the week, contact mode).
 - Time preference may not be stable over time or predictable a priori.



Thank you

Contact: Name | email: jwescott@rti.org

Appendix

k-prototypes algorithm

- Extends the more common k-means algorithm by defining a distance (pseudo) metric between a sample member vector X_i and prototypic cluster vector Q_l , or prototype.
 - x_{ij}^r and q_{lj}^r are values of numerical variables (as vector elements), and x_{ij}^c and q_{lj}^c are values of categorical variables for sample member i and prototypic cluster l .
- The variables m_r and m_c represent the number of numerical and categorical variables respectively, while γ_l serves as a weight for categorical variables for cluster l .
- Prototypes are analogous to k-means centroids.

$$d(X_i, Q_l) = \sum_{j=1}^{m_r} (x_{ij}^r - q_{lj}^r)^2 + \gamma_l \sum_{j=1}^{m_c} \delta(x_{ij}^c, q_{lj}^c) \quad \delta(p, q) = \begin{cases} 0 & p = q \\ 1 & p \neq q \end{cases}$$

Appendix

Predicting Cluster Membership for BY Non-Respondents

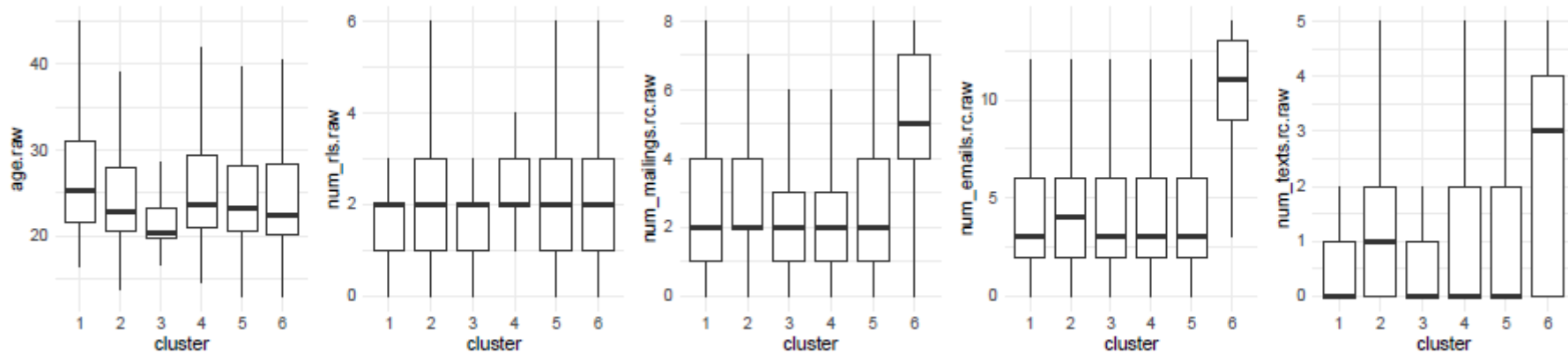
- Uses SVM supervised learning model, with a linear kernel.
 - Covariates were derived from sampling frame data that are known a priori for all students.
- To evaluate model performance, we selected a random 30% sample of clustered cases for testing and trained the SVM on the remaining 70% of cases.
 - Table 3 provides a confusion matrix contingency table for actual and predicted cluster membership using these cases. Entries on the main diagonal represent true-positives.
- Overall, the model performs well for all clusters, except cluster 6.

Predicted cluster	Actual cluster					
	1	2	3	4	5	6
1	2425	560	145	397	0	120
2	802	3473	173	845	10	185
3	433	90	3698	417	0	252
4	439	1270	38	6872	1	286
5	20	32	0	15	6832	347
6	1	2	0	2	2	0

Table 3: SVM confusion matrix

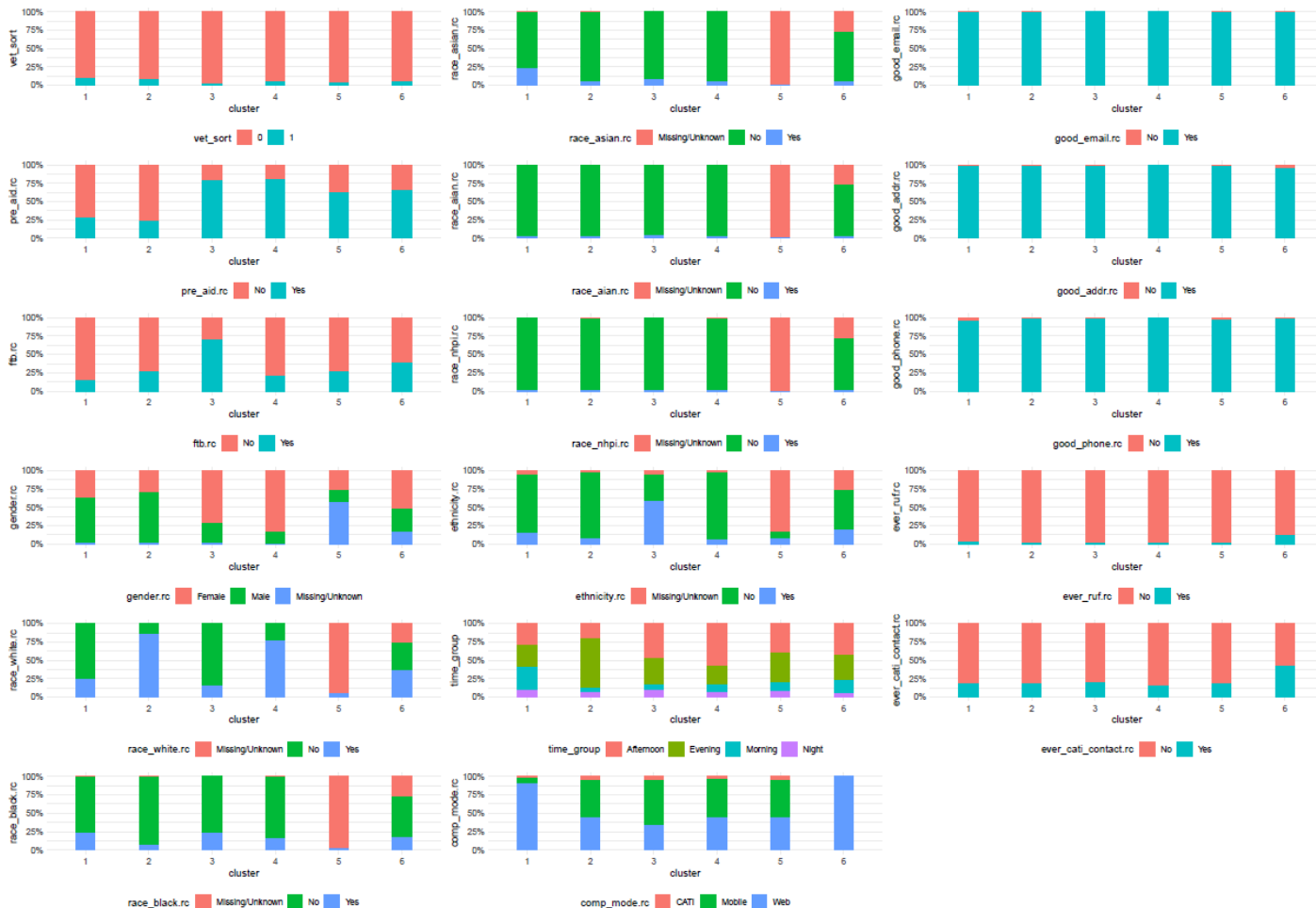
Appendix

Distribution of Numerical Variables by Cluster



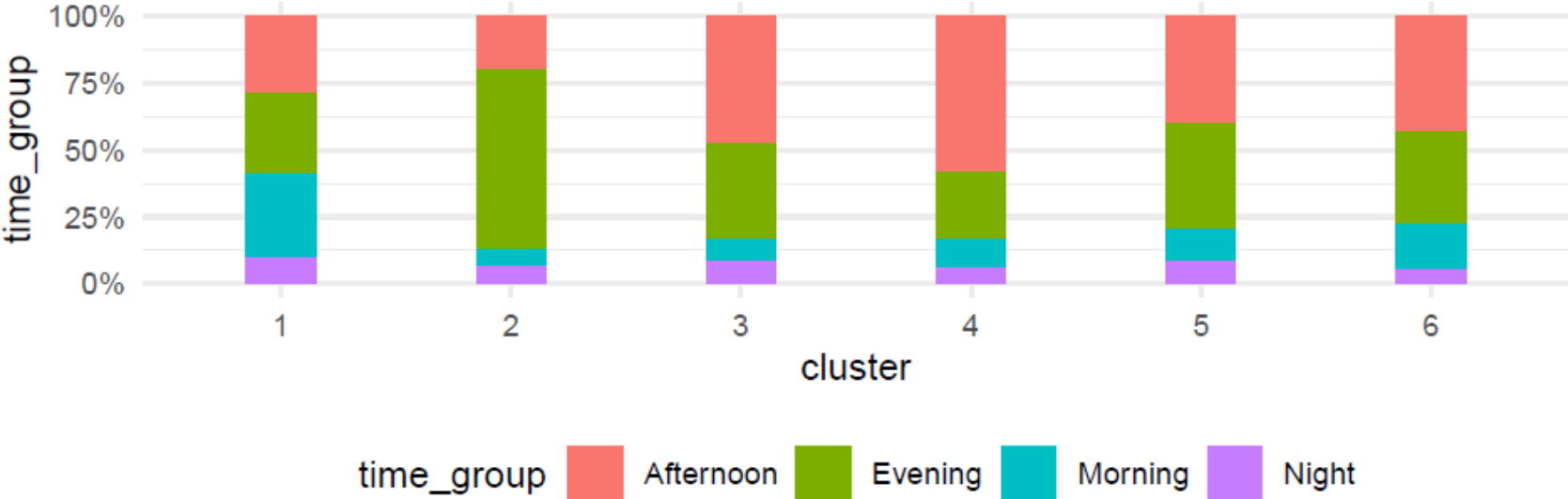
Appendix

Distribution of Categorical Variables by Cluster



Appendix

Distribution of Categorical Variables by Cluster



Appendix

Frame data

Variable	Description
<code>cip_orig1</code>	Classification of Instructional Programs (CIP) code for student's primary major
<code>ftb</code>	first-time beginner status
<code>vet_sort</code>	veteran status (a combination of <code>pre_vet</code> and <code>veteran</code> variables)
<code>sector11</code>	11-level sector
<code>gender</code>	gender
<code>ethnicity</code>	ethnicity (Hispanic)
<code>race_white</code>	race (White)
<code>race_black</code>	race (Black)
<code>race_asian</code>	race (Asian)
<code>race_aian</code>	race (American Indian and Alaska Native)
<code>race_nhpi</code>	race (Native Hawaiian and Pacific Islander)
<code>pre_aid</code>	student aid status

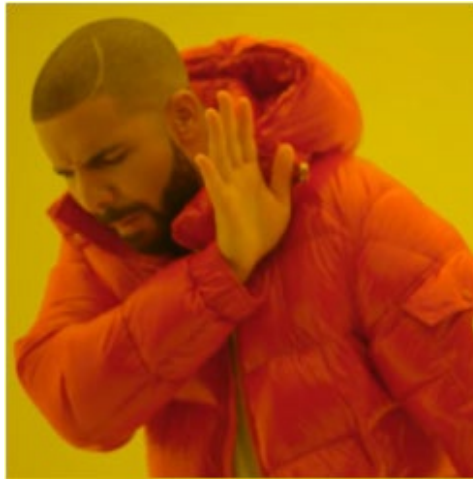
Table 1: Student sample file variables

Paradata

Variable	Description
<code>Num_RLs</code>	Number of roster lines
<code>COMP_MODE</code>	Completion mode (Web, CATI, Mobile)
<code>CATI.sumstat</code>	first-time beginner status
<code>SUMTIME</code>	Time final sumstat set
<code>GoodEmailFlag</code>	Good email indicator
<code>GoodAddrFlag</code>	Good address indicator
<code>GoodPhoneFlag</code>	Good phone number indicator
<code>Num_mailings</code>	Number of mailings sent
<code>Num_Emails</code>	Number of emails sent
<code>Num_Texts</code>	Number of texts sent
<code>EVER_ANY_REF</code>	Sample member ever refused
<code>Ever_CATI_contacted</code>	Ever CATI contact with sample member

Table 2: Case detail report variables

Just for fun...



Increasing
contact
attempts



Making
contact
attempts
more effective