

# QUANTITATIVE MEDICINE

TRANSFORMING DRUG DISCOVERY

Technology Evaluation Consortium  
Cambridge Healthtech Associates

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# Agenda

- ToxCast Study Design
- Simulation Results
- Prospective Uses
- Next Study

# ToxCast Study Design

**Purpose:** Develop a Model to Actively Learn and Accurately Predict All Observations in ToxCast

# ToxCast Data

- Total Assays: 1197
  - ACEA – change in cell growth kinetics
  - AttaGene – reporter gene assay
  - BioSeek - ELISA
  - Cellumen – High Content Screening
  - CellzDirect – change in expression levels
  - GenTronix – *in vitro* measure of growth arrest
  - NCGC - reporter gene assay
  - NovaScreen – enzyme inhibition assays
  - Solidus – Microarray?
  - ToxRefDB – Many toxicity measures

# ToxCast Data

- Compounds: 309
- Experimental results:
  - Total possible results: 369,873
  - Available results: 321,033 (87% coverage)
  - Continuous measures from database
  - Discrete inferred from Continuous measures

# How do we determine what is Ground Truth?

- Threshold data based on ToxCast results

CASRN	NAME	CLM_CellLoss_24hr	CLM_Hepat_Apoptosis_24hr	CLM_Hepat_DNADamage_24hr
50594-66-6	Acifluorfen	1000000	1000000	1000000
15972-60-8	Alachlor	61.46	1000000	1000000
116-06-3	Aldicarb	1000000	1000000	1000000
834-12-8	Ametryn	1000000	1000000	1000000
33089-61-1	Amitraz	123.9	144	NA
101-05-3	Anilazine	1000000	1000000	1000000
3337-71-1	Asulam	1000000	1000000	1000000
1912-24-9	Atrazine	1000000	1000000	1000000
35575-96-3	Azamethiphos	102.4	1000000	52.1
131860-33-8	Azoxystrobin	1000000	1000000	1000000

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# Receiver Operating Characteristic

- What are we predicting?
  - Assay results for all “unobserved” experiments in ToxCast
- How do we measure accuracy of those predictions?
  - Calculate a ROC curve by sweeping across prediction thresholds and comparing to hidden ground truth in ToxCast

# Compound Diversity-Based Selection

**Purpose:** Develop a Model to Actively Learn and Accurately Predict All Observations in ToxCast

**Standard Approach (single pass):**

**Selection method:** Cluster all compounds by their fingerprints into K clusters. Choose one compound from each cluster.

**Prediction method:** Learn a QSAR model using Lasso regression or RandomForest.

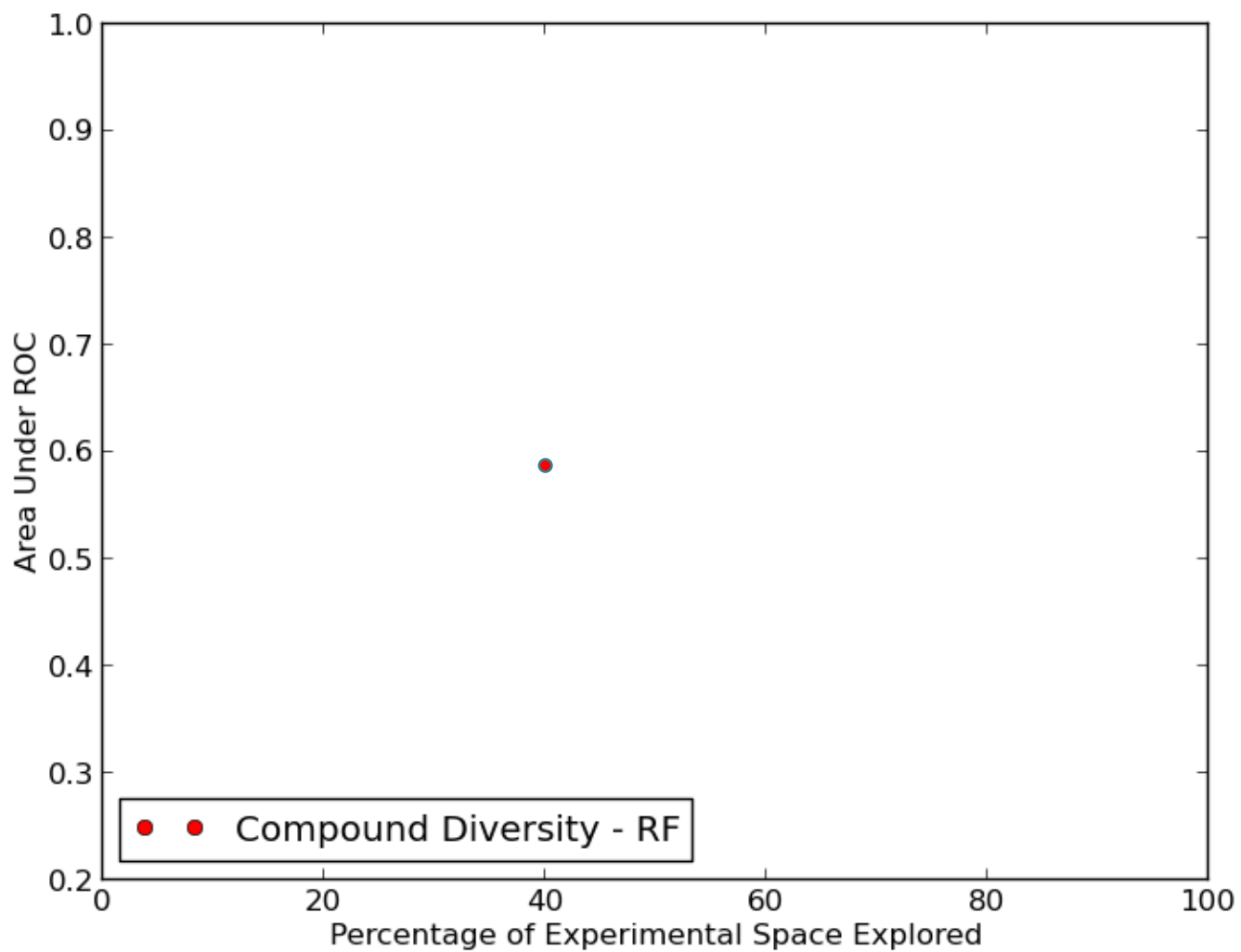
# AFRS-Based Experiment Selection

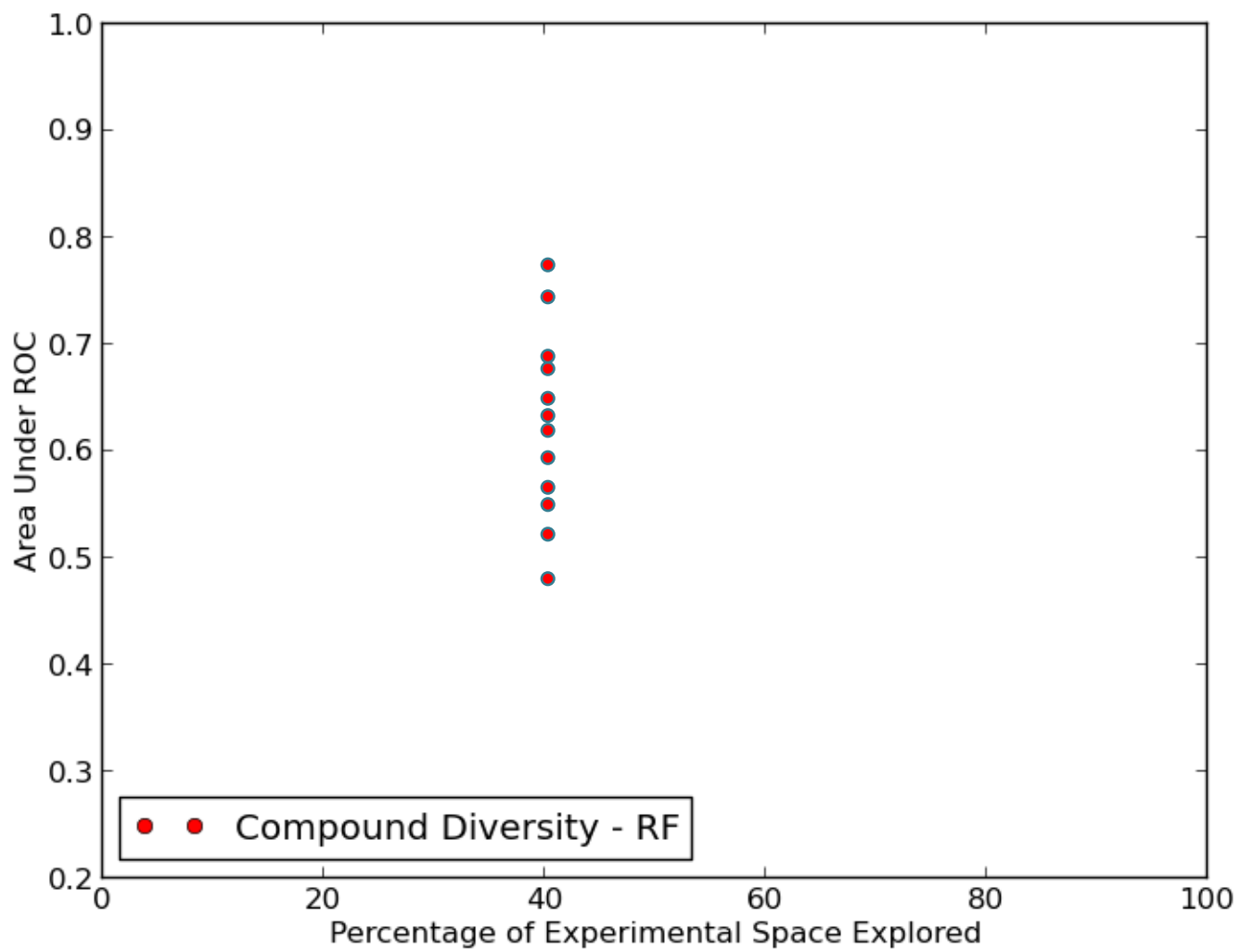
**Purpose:** Develop a Model to Actively Learn and Accurately Predict All Observations in ToxCast

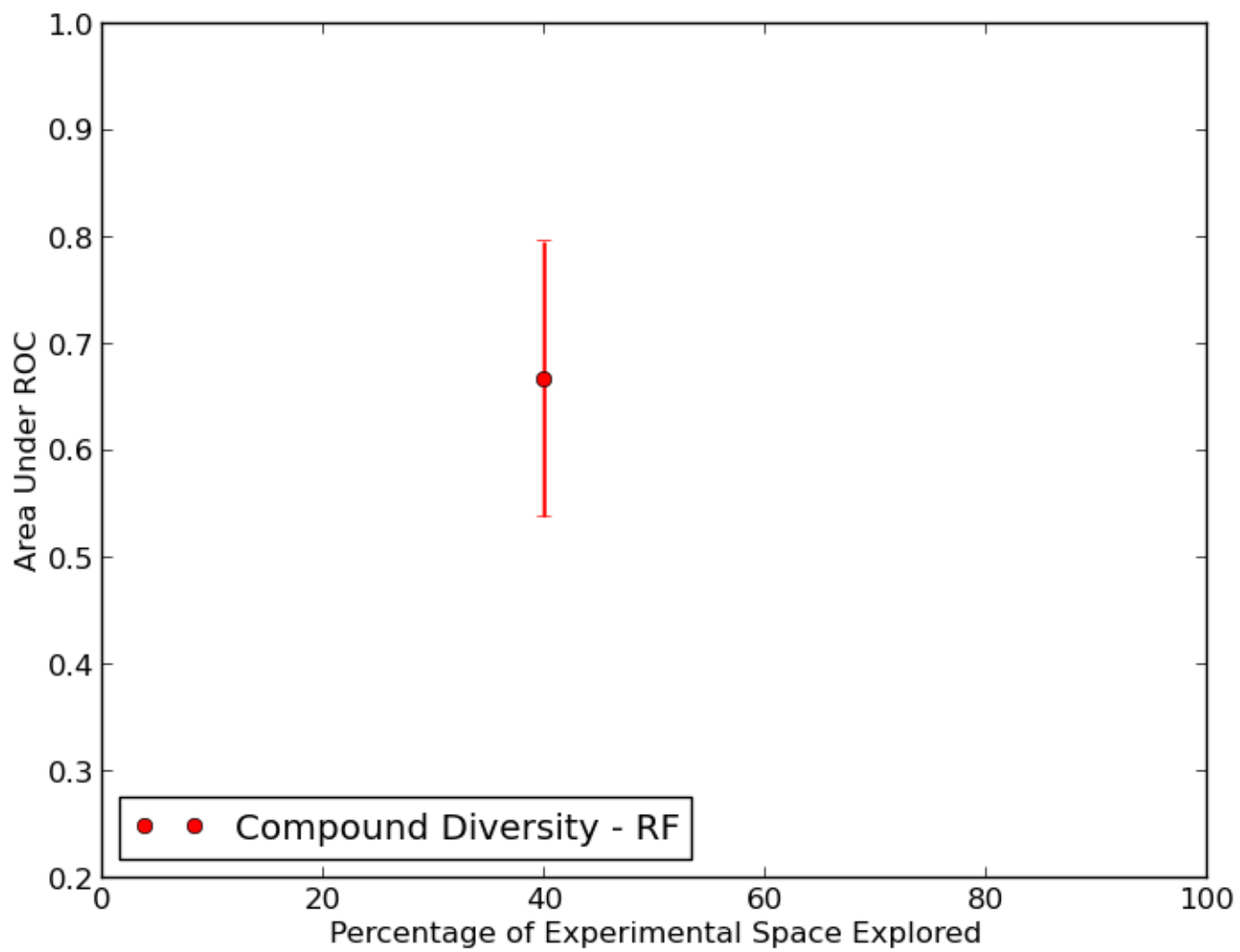
**AFRS Approach (iterative):**

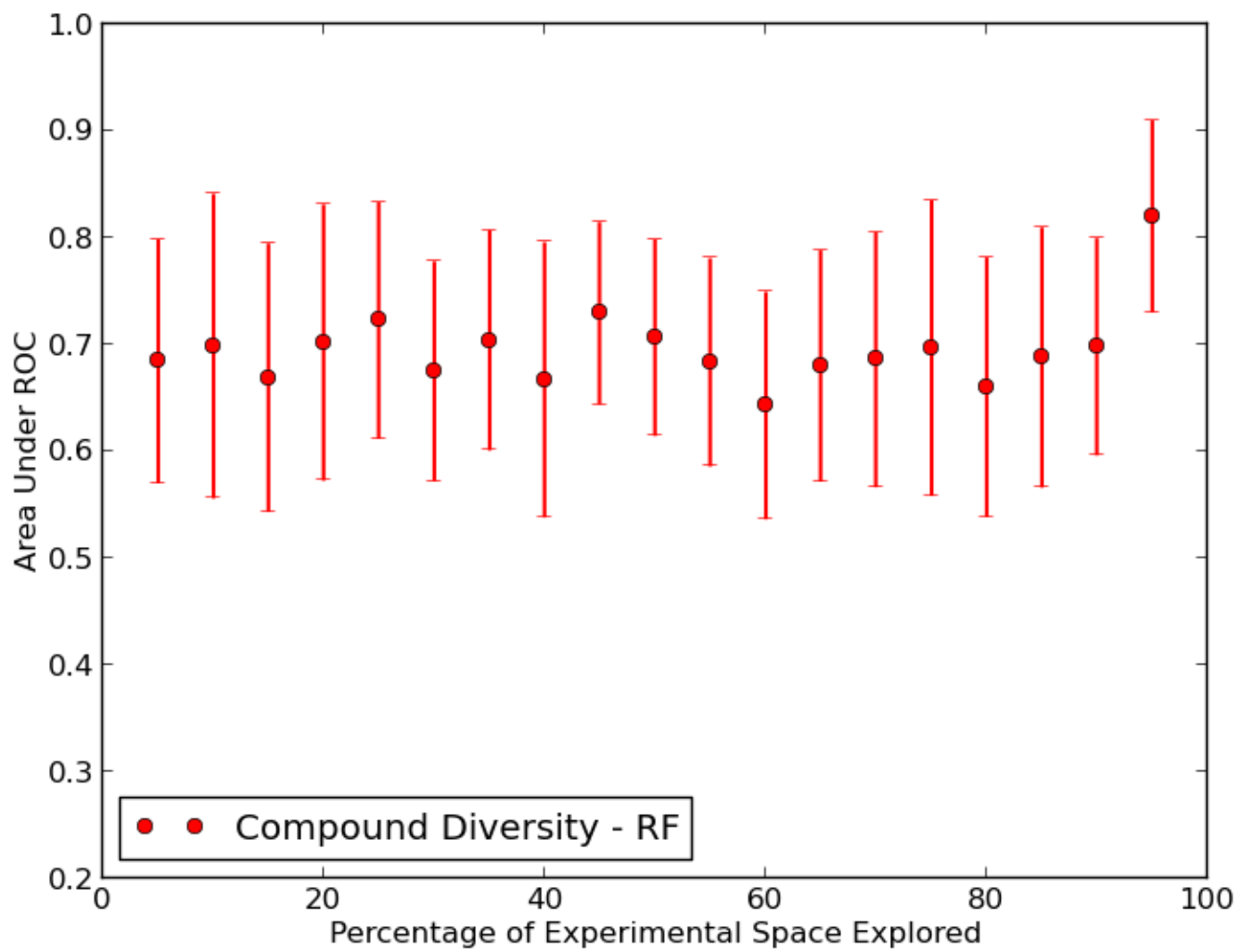
**AFRS Selection Method:** Choose batches of 1% of experiments believed to be the most informative.

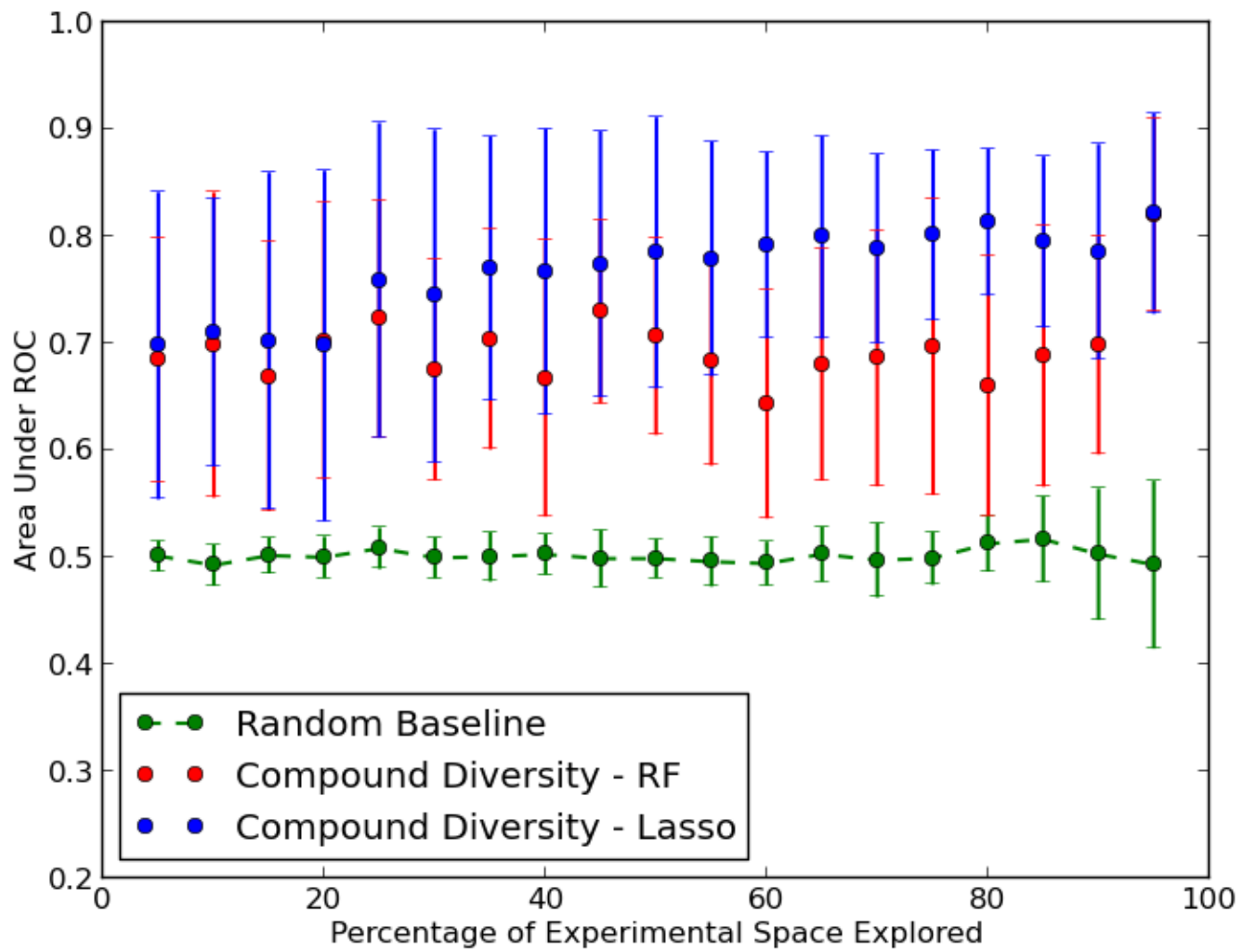
**AFRS Prediction Method:** Learn a model using the AFRS for to predict the results of all untested experiments.



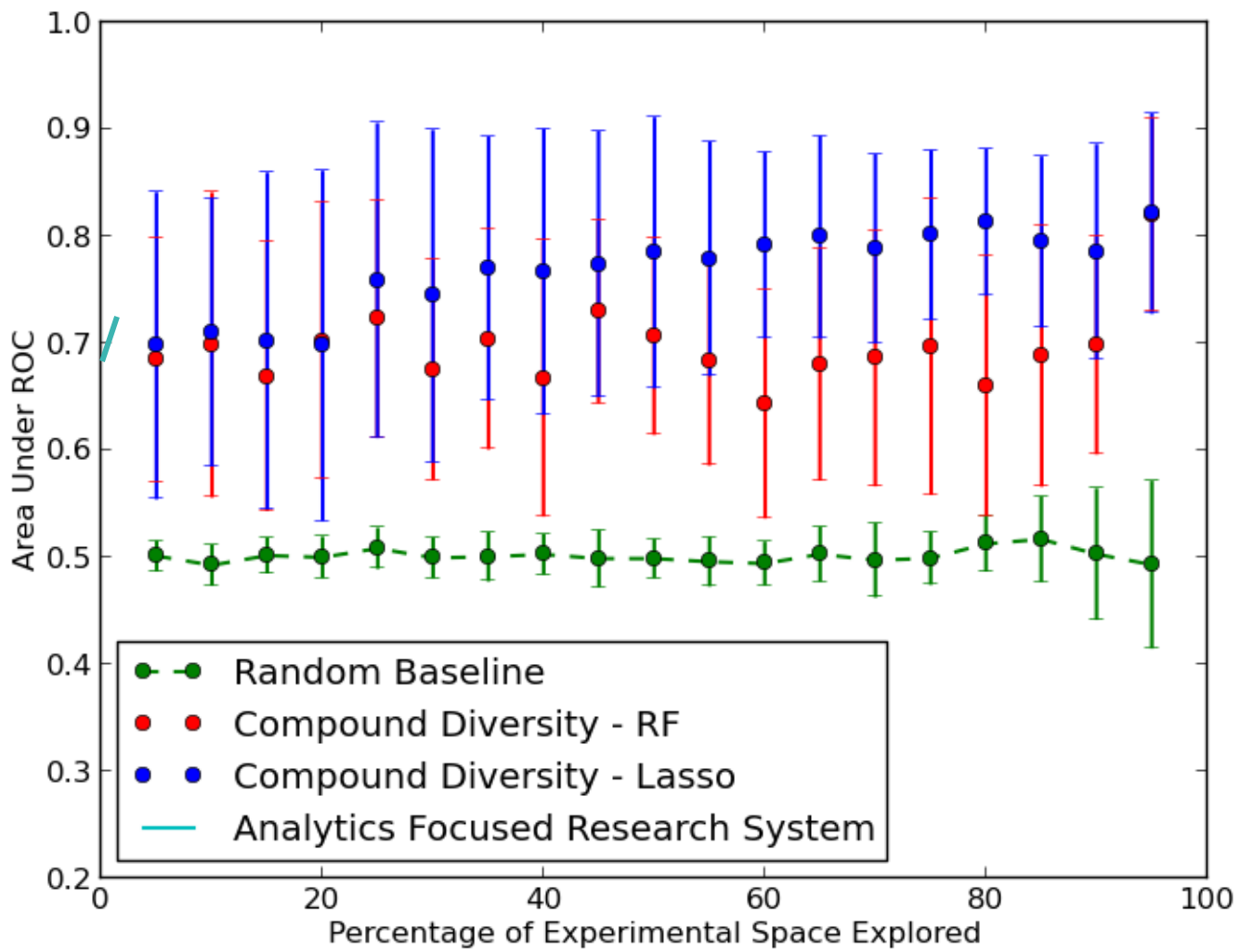




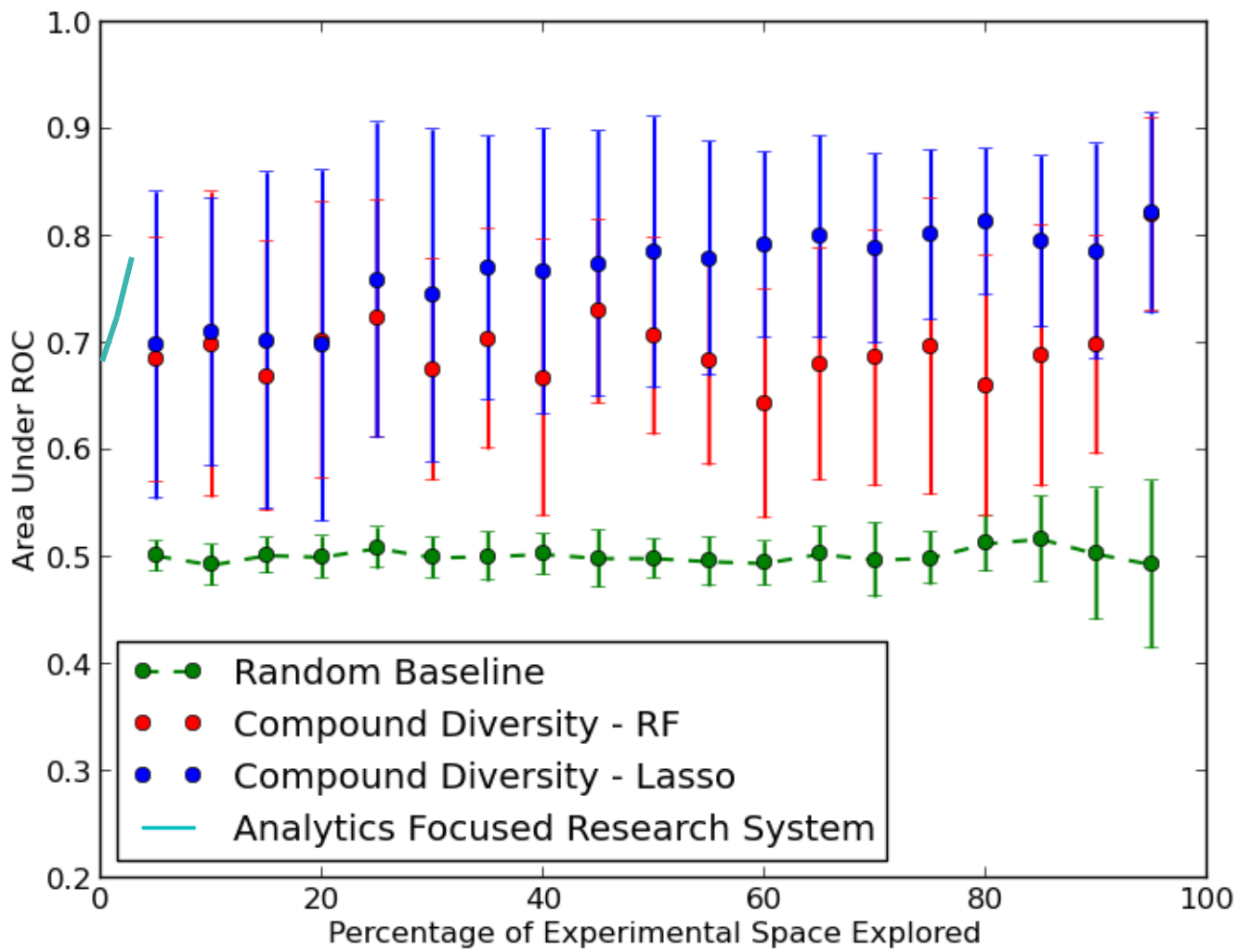




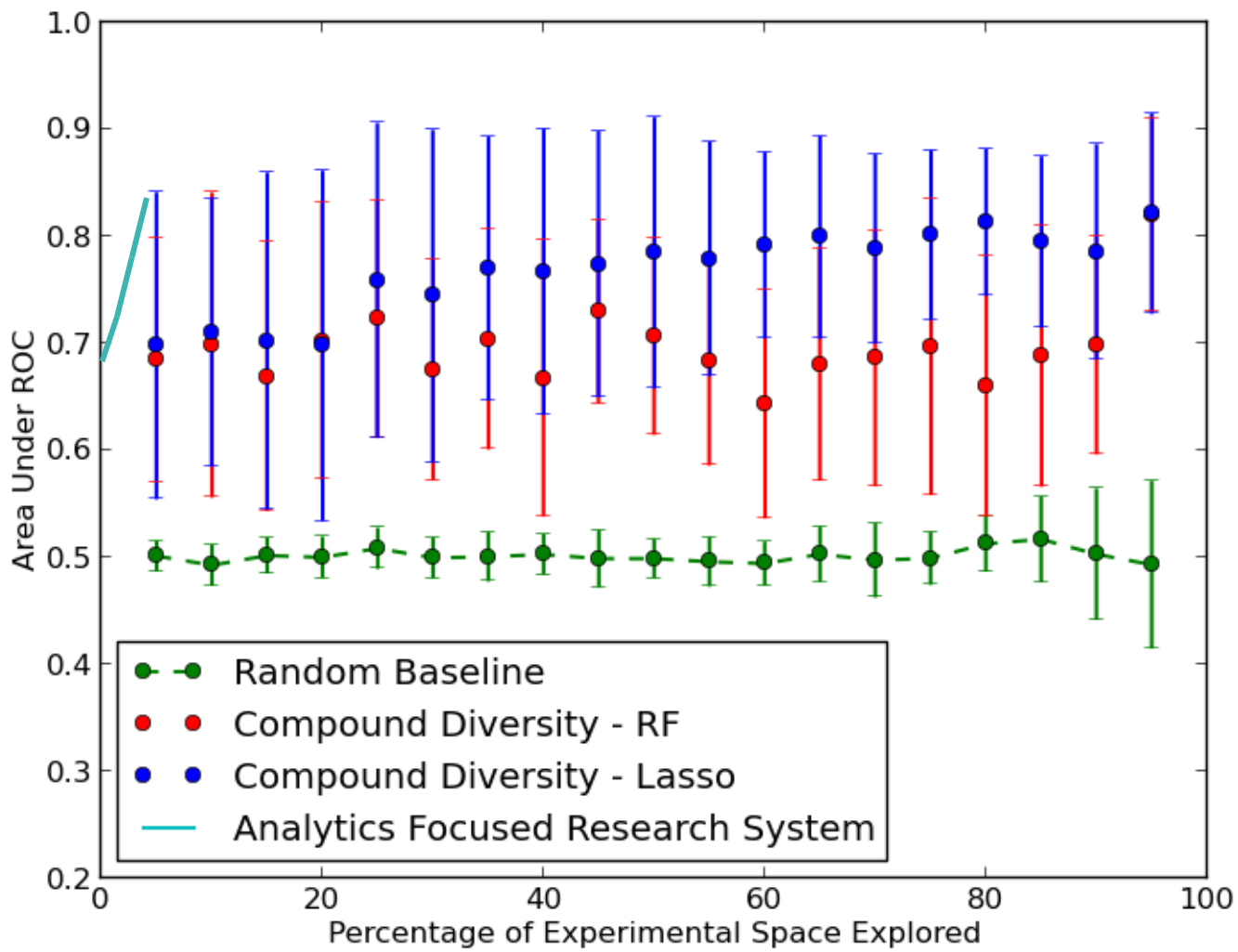




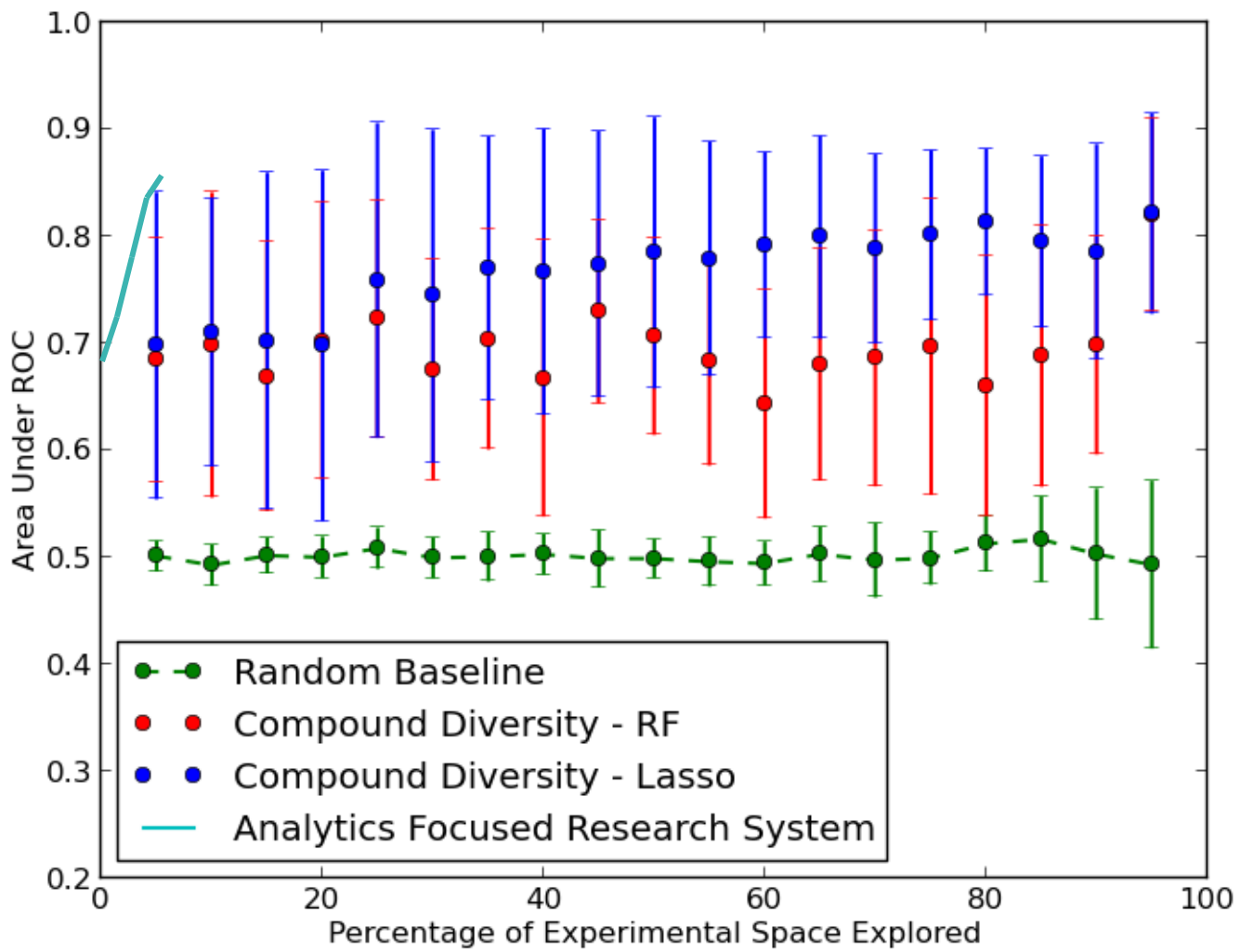
Active Learning Cycle Iteration: 1



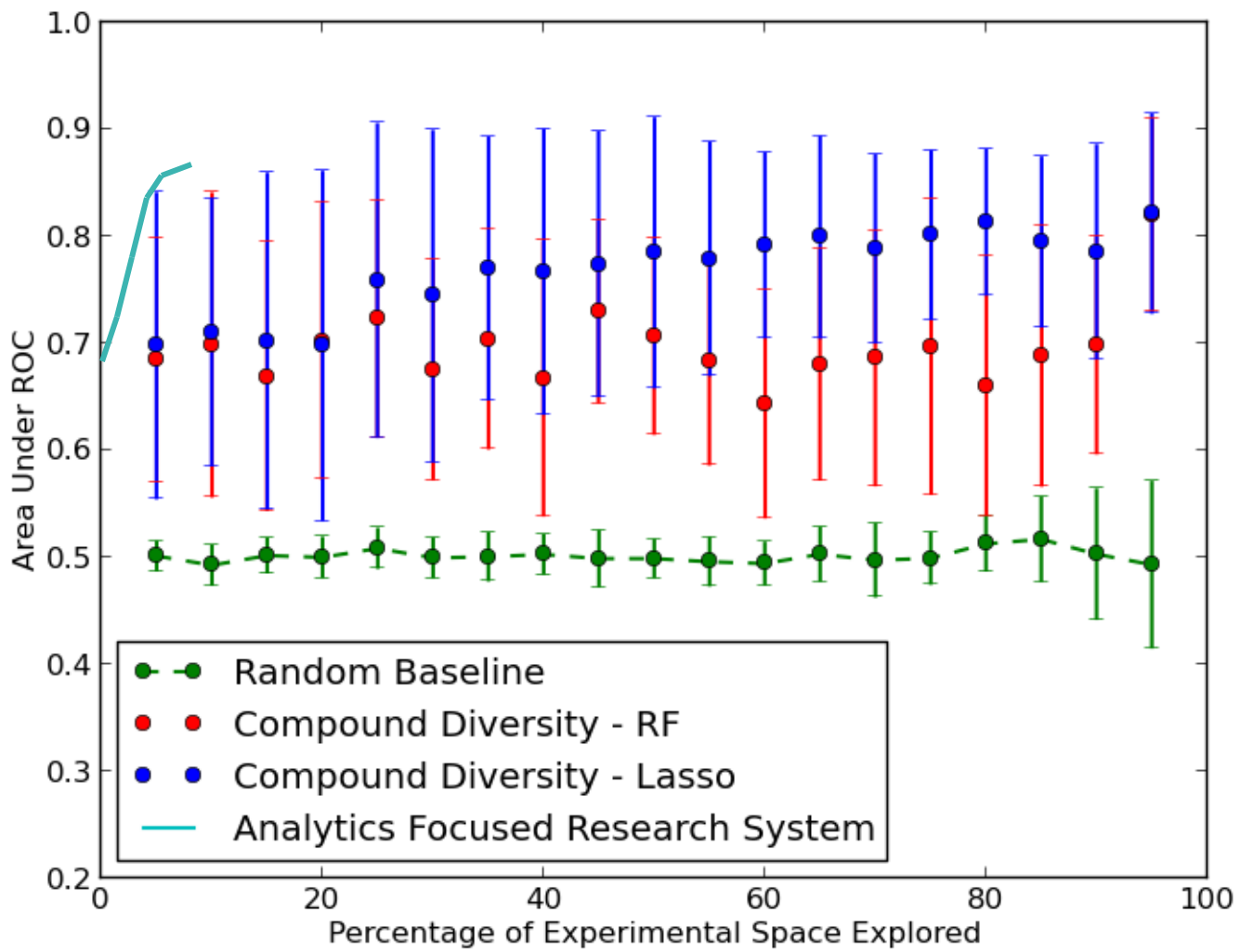
Active Learning Cycle Iteration: 3



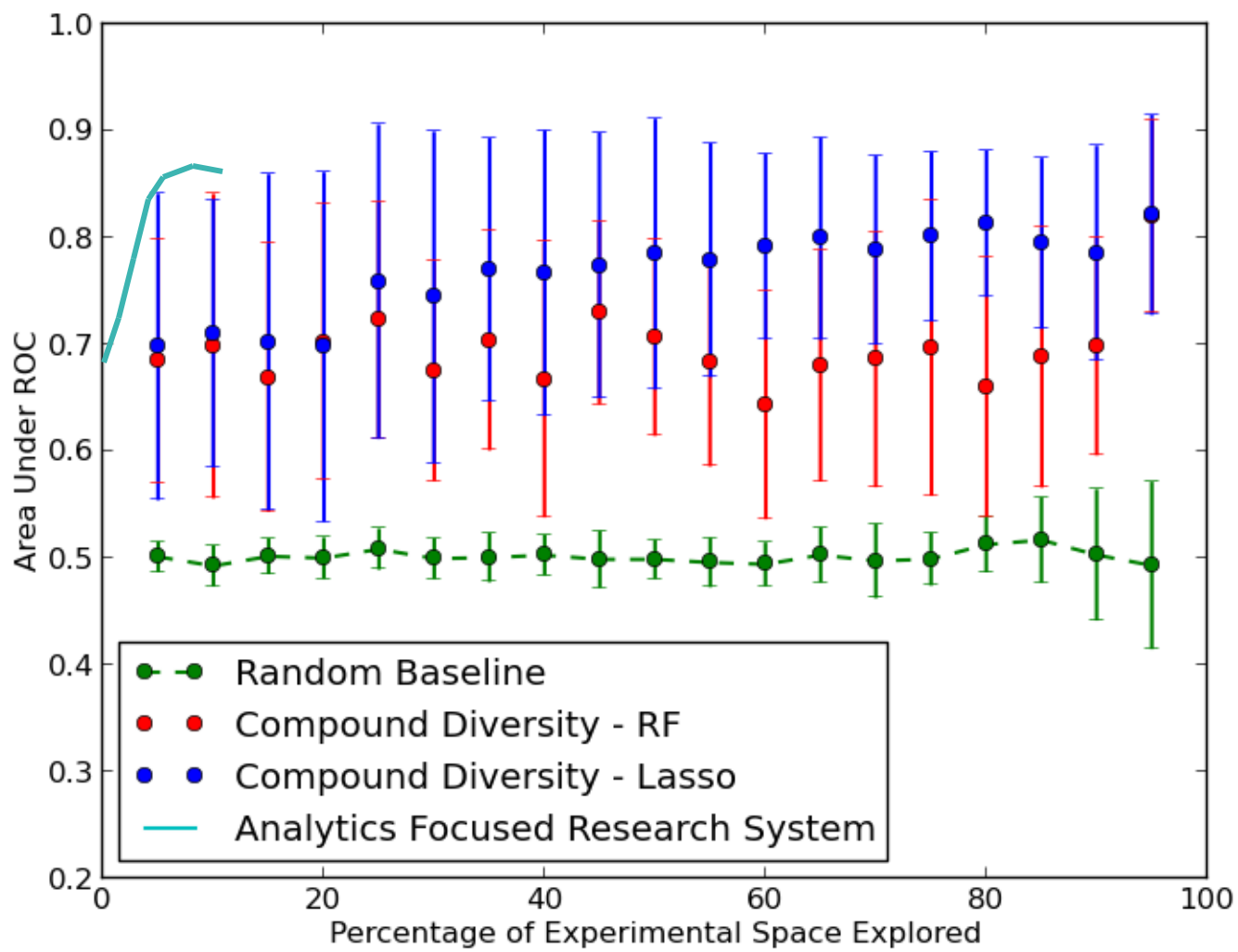
Active Learning Cycle Iteration: 4



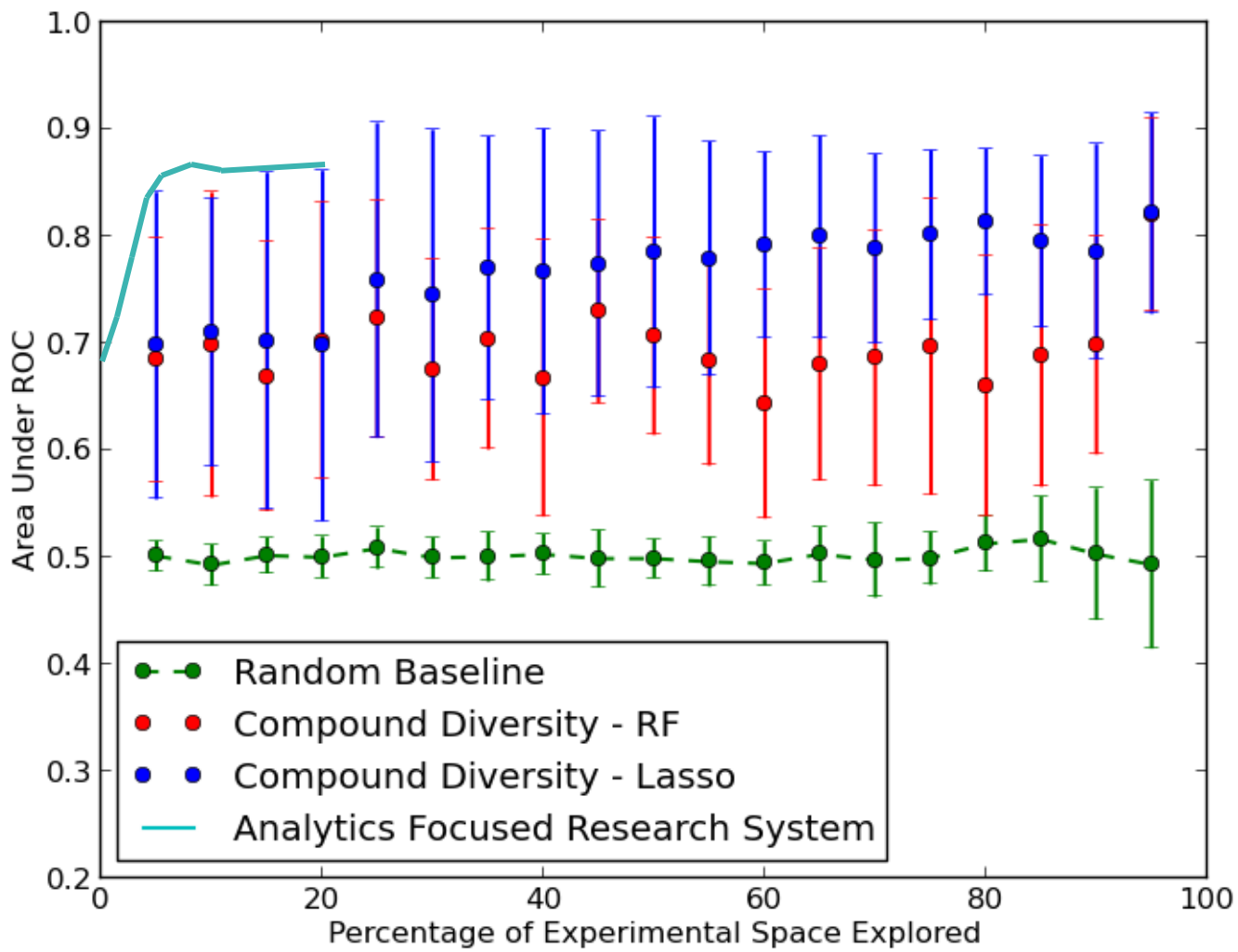
Active Learning Cycle Iteration: 6



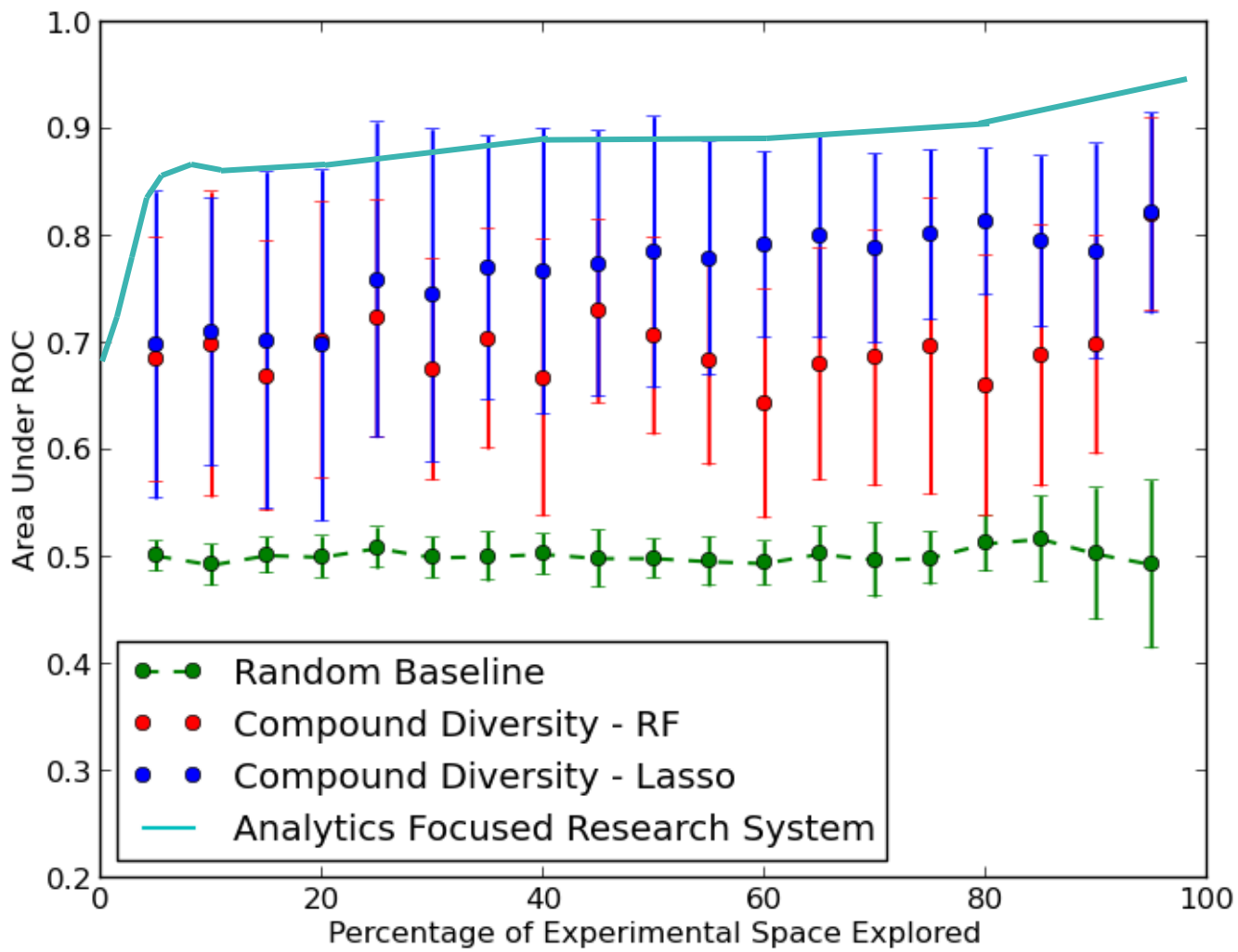
Active Learning Cycle Iteration: 8



Active Learning Cycle Iteration: 11

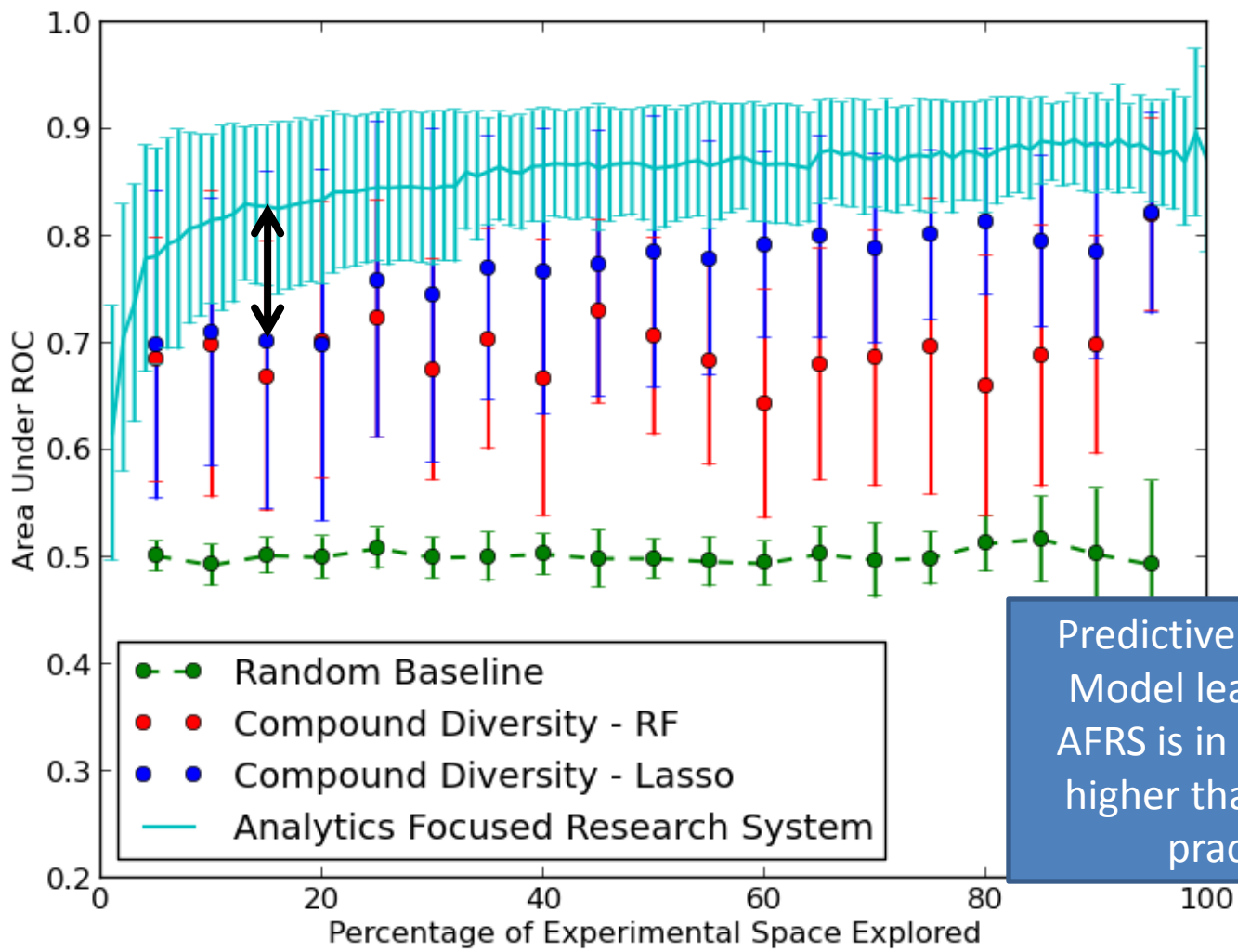


Active Learning Cycle Iteration: 20

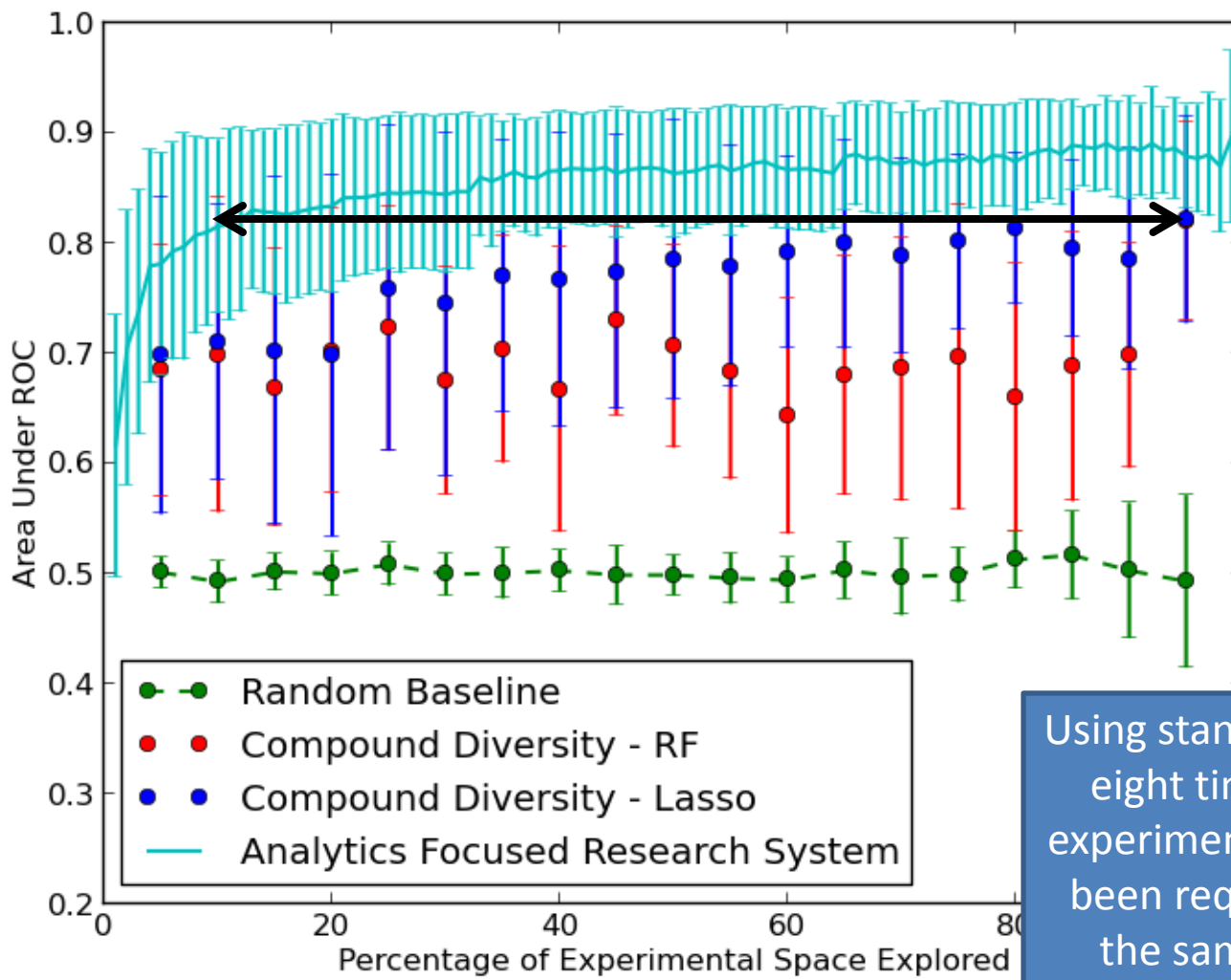


Active Learning Cycle Iteration: 99

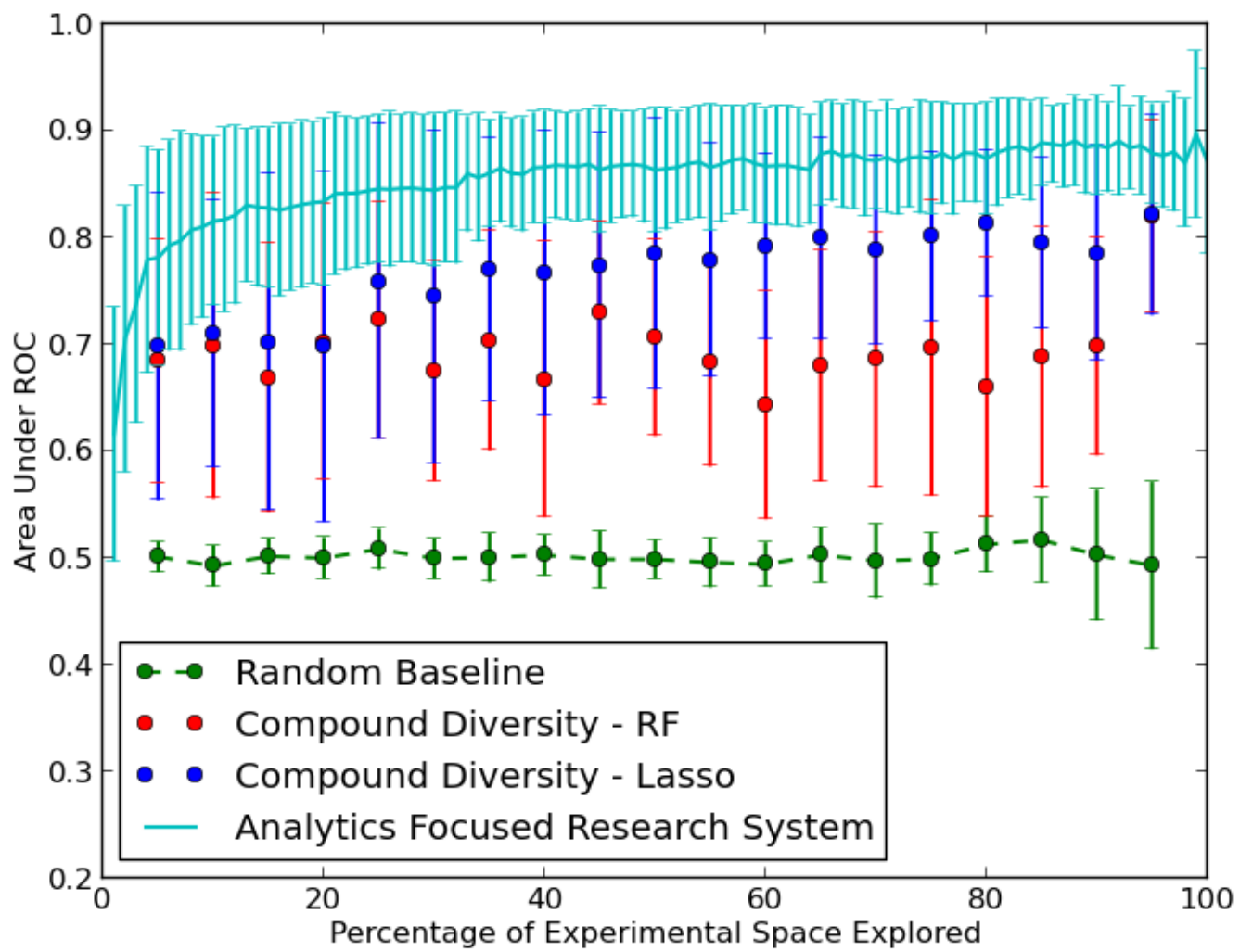




Predictive accuracy of Model learned using AFRS is in expectation higher than standard practices.



Using standard methods, eight times as many experiments would have been required to have the same accuracy.



# Example From a Single Simulation

- When predicting experimental results for Atrazine:

AFRS			Standard Approach		
Total Experiments Executed	Estimated Experimental Cost	Accuracy of Prediction for Atrazine (AUC)	Total Experiments Executed	Estimated Experimental Cost	Accuracy of Prediction for Atrazine (AUC)
32,000	\$600,000	85%	260,000	\$4,800,000	85%

ToxCast Phase 1 Estimated Total Cost: \$6 million

# Prospective Uses of AFRS Process

- For a small set of compounds:
  - Make predictions for toxicity
  - Prioritize toxicology assays to most effectively eliminate compounds (faster-to-fail)
  - Prioritize assays for improving predictive model

	<b>emamectin benzoate</b>	<b>Aldicarb</b>
Round 3	Novascreen - cytochrome P450, family 19, subfamily A, polypeptide 1  ELISA – chemokine (C-C motif) ligand 2	None
Round 7	Novascreen - ADME_hCYP4F12_Activator	Novascreen - cytochrome P450, family 19, subfamily A, polypeptide 1

# Prospective Uses of AFRS Process

- For a new series of synthesized compounds:
  - Select a small number of experiments to effectively expand the predictions to new areas of chemical space as needed
  - Prioritize compounds based on predicted toxicity

# Prospective Uses of AFRS Process

- For multiple assays:
  - Prioritize assays based on human toxicity predictivity
- For a large digital library of ***unsynthesized*** compounds
  - Select a small number of experiments from available ***synthesized*** compounds which give the most information about ***unsynthesized*** compounds
  - Prioritize compounds for synthesis
    - Based on predicted lack of toxicity
    - Based on potential information gain from testing synthesized compounds

# Next studies with other data?

- Using your data, expand experimental space to include results from other areas of your company?
- Direct prediction of human toxicity with your data or alternative public source?
- Prospective study?



# QUANTITATIVE MEDICINE

## TRANSFORMING DRUG DISCOVERY

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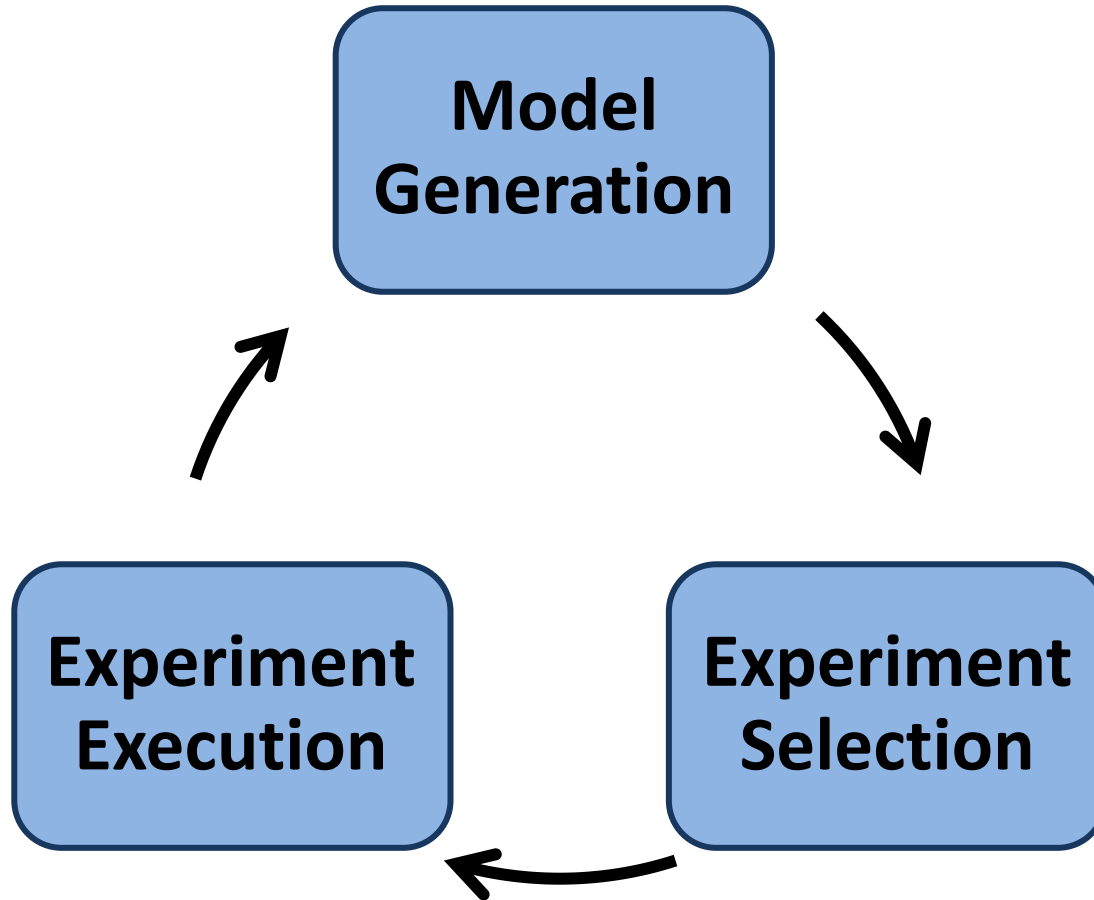
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CHIEF SCIENCE OFFICER

# Active Learning



# AFRS Directed Study Types

## Retrospective

- Experiment Direction Simulated
- Show *potential* for improved **accuracy** and **efficiency**
- May discover new useful knowledge through AFRS analytics

## Prospective

- Experiments Directed as Selected
- Yield *actual* improvements in accuracy and efficiency
- Likely to discover new useful knowledge through directed experimentation **and** AFRS analytics