

Advances in Survey Estimation with Imperfectly Matched Auxiliary Data

Jay Breidt

Colorado State University

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*Joint work with Chien-Min Huang and Jean Opsomer
Colorado State University and Westat*

- About 450 charter boats and 15,000 boat trips along the Atlantic Coast of South Carolina each year
- How many black sea bass were caught in 2018?

$$\begin{aligned} U &= \{1, 2, \dots, N\} \\ &= \{\text{all SC charter fishing boat trips in 2018}\} \end{aligned}$$

- number of black sea bass caught on k th trip: y_k
- total black sea bass caught: $T = \sum_{k \in U} y_k$
- Infeasible to obtain data on all $N \simeq 15,000$ boat trips: instead, use a **probability sample** $s \subset U$

Two sources of information on the charter boat fishery 3

Sample with angler interviews: Monthly logbook records:



Revised 4-2012

SOUTH CAROLINA CHARTERBOAT LOGBOOK

Vessel (Please Print): _____ Date: _____ Permit No.: _____

Number of Anglers: _____ Trip Start Time: _____ Actual Hours Fished: _____ Location: _____ Example: 32-78-C1 (see map)

Trip Start _____ Artificial _____ Target _____
Location: _____ Reef Name: _____ Species: _____

Locale: ☐ Estuarine Method: ☐ Trawl ☐ Cast / Fly Water Depth: _____ feet
☐ 0 - 3 miles ☐ Bottom ☐ Dive ☐ Gig Shallowest: _____ feet
☐ Offshore ☐ Offshore Deepest: _____ feet

AGENCY USE ONLY

MAIL OR FAX REPORT BY:
THE 10TH OF THE MONTH TO:
SCDNR Fisheries Statistics Section, P.O.
Box 12559, Charleston, SC 29422-2559
FAX: (843) 953-9362 Phone: (843) 953-9313

Yr	Mo	Day	Permit #	Location	Locale	Ang#	Meth
Target Sp.	Hrs.	Reef	Trip Start	Shallowest	Deepest		

Species	# Kept	Lbs Kept	# Released Alive	# Released Dead	Species	# Kept	Lbs Kept	# Released Alive	# Released Dead
1050 Dolphin					1423 Gag				
4710 Wahoo					1424 Scamp				
4655 Yellowfin Tuna					1414 Snowy Grouper				
4658 Blackfin Tuna					1416 Red Grouper				
3026 Sailfish					1410 Other Grouper				
2177 White Marlin					(Specify)				
2179 Blue Marlin					3202 Red Porgy (Pink)				
1940 King Mackerel					3295 Other Porgies				
3840 Spanish Mackerel					(Specify)				
4653 Little Tunny					3764 Red Snapper				
0330 Bonita					3765 Vermilion Snapper				
4654 Skip Jack					3560 Black Sea Bass				
0180 Barracuda					3514 Spottail Pinfish				
3810 Spadefish					1441 White Grunt				
0030 Amberjack					1440 Other Grunts				
0870 Crevalle Jack					(Specify)				
0230 Bluefish					4560 Triggerfish				
0570 Cobia					1082 Red Drum				
4350 Tarpon					1081 Black Drum				
Other Fish					3447 Spotted Seatrout				
(Specify)					3446 Weakfish				
					1209 Flounder				
					3560 Sheephead				
					4410 Ladyfish				
					1970 Whiting				
					2670 Inshore Pinfish				
					3518 Sharpnose Shark				
					3495 Blacktip Shark				
					3483 Bonnethead Shark				
					3521 Spiny Dogfish				
					3511 Smooth Dogfish				
					3508 Other Sharks				
					(Specify)				
					2860 Stingrays				

Captain's Notes: _____

Signature: _____

Print Name: _____

- Design-based difference estimators
- Extension to multiple frames
- Require matching of sampled elements to auxiliary records
 - most theory and methods assume matching is done without error
- Some results on estimation under imperfect matching
 - properties of difference estimators
 - simulation results based on South Carolina charter boat fishing

- Draw probability sample $s \subset U$ via design with known, positive **inclusion probabilities** $\Pr[k \in s] = \pi_k > 0$
- Sample membership indicator $I_k = 1$ if $k \in s$, $I_k = 0$ otherwise

$E[I_k] = \pi_k$, averaging over all possible samples

- Since $E[I_k/\pi_k] = 1$ under repeated sampling, unbiased **Horvitz-Thompson estimator** of T is

$$\hat{T} = \sum_{k \in s} \frac{y_k}{\pi_k} = \sum_{k \in U} y_k \frac{I_k}{\pi_k}$$

Now suppose we have the following:

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- **Auxiliary data** $\mathbf{x}_\ell$ for all ℓ in some database $\mathcal{A}$
- Perfect, known **matching** from $\mathcal{A}$ to population U :

$$M_{k\ell} = \begin{cases} 1, & \text{if } \ell \in \mathcal{A} \text{ matches } k \in U, \\ 0, & \text{otherwise} \end{cases}$$

- A “**method**” $\mu(\cdot)$ for predicting y_k from $\mathbf{x}_\ell$:

$$\sum_{\ell \in \mathcal{A}} M_{k\ell} \mu(\mathbf{x}_\ell) = \tilde{y}_k \text{ predicts } y_k$$

- for each element k , look up the correct $\mathbf{x}_\ell$
- apply $\mu(\cdot)$, which does not depend on the sample

Difference estimator combines sample and auxiliary data 7

- Difference estimator of T is then

$$\begin{aligned}\tilde{T} &= \sum_{k \in U} \sum_{\ell \in \mathcal{A}} M_{k\ell} \mu(\mathbf{x}_\ell) + \sum_{k \in s} \frac{y_k - \sum_{\ell \in \mathcal{A}} M_{k\ell} \mu(\mathbf{x}_\ell)}{\pi_k} \\ &= \sum_{k \in U} \tilde{y}_k + \sum_{k \in U} (y_k - \tilde{y}_k) \frac{I_k}{\pi_k} \\ &= (\text{auxiliary-based prediction}) + (\text{bias adjustment})\end{aligned}$$

where $\tilde{y}_k$ is not random

- Expectation is

$$\mathbb{E} [\tilde{T}] = \sum_{k \in U} \tilde{y}_k + \sum_{k \in U} (y_k - \tilde{y}_k) \mathbb{E} \left[\frac{I_k}{\pi_k} \right] = T$$

$$\begin{aligned} \text{Var} \left(\sum_{k \in U} \tilde{y}_k + \sum_{k \in U} (y_k - \tilde{y}_k) \frac{I_k}{\pi_k} \right) \\ = \sum_{j, k \in U} \Delta_{jk} \frac{(y_j - \tilde{y}_j)}{\pi_j} \frac{(y_k - \tilde{y}_k)}{\pi_k} \end{aligned}$$

- Compare to Horvitz-Thompson estimator:

$$\text{Var} \left(\sum_{k \in U} y_k \frac{I_k}{\pi_k} \right) = \sum_{j, k \in U} \Delta_{jk} \frac{y_j}{\pi_j} \frac{y_k}{\pi_k}$$

- Difference estimator is **exactly** unbiased, regardless of the quality of the method $\mu(\cdot)$
- Has smaller variance than HT provided “residuals”

$$y_k - \tilde{y}_k$$

have smaller variation than “raw values” y_k

– (If $M_{k\ell} \equiv 0$, we get back HT)

- Have an **exactly** unbiased variance estimator
- Above results assume (1) one frame covers the universe and (2) matching is perfect

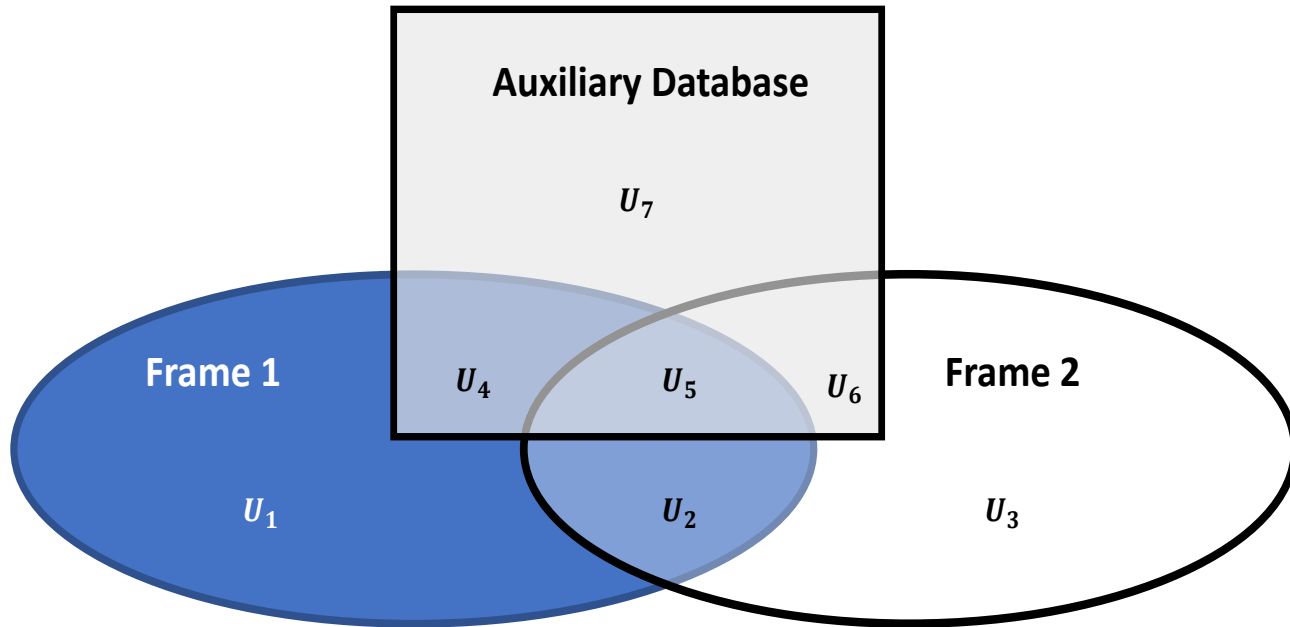
- Assume that the universe U is completely covered by disjoint “overlap domains”:

$$U = \left\{ \bigcup_{g \in G_1} U_g \right\} \cup \left\{ \bigcup_{g \in G_2} U_g \right\} \cup \left\{ \bigcup_{g \in G_3} U_g \right\}$$

- If $g \in G_1$, overlap domain U_g is covered by one or more frames, **but not** the database
- If $g \in G_2$, overlap domain U_g is covered by one or more frames **and** the database
- If $g \in G_3$, overlap domain U_g is covered **only** by the database

$$U = \{U_1 \cup U_2 \cup U_3\} \cup \{U_4 \cup U_5 \cup U_6\} \cup \{U_7\}$$

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Overall estimation approach

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		In Auxiliary Database?	
		No	Yes
In Sampling Frame(s)?	No		• G_3 • Synthetic predictor • Biased • Zero sampling variance
	Yes	• G_1 • Mecatti estimator • Unbiased • Potentially large variance	• G_2 • Difference estimator • Unbiased • Small variance if auxiliary information is good

- From frame f , draw a sample s_{fg} to represent U_g
- Compute Horvitz-Thompson estimator

$$\hat{T}_{fg} = \sum_{k \in s_{fg}} \frac{y_k}{\pi_k^{(f)}}, \quad \text{where } \mathbb{E} [\hat{T}_{fg}] = T_g$$

- Define the coverage indicator

$$F_{fg} = \begin{cases} 1, & \text{if overlap domain } g \text{ is covered by frame } f \\ 0, & \text{otherwise} \end{cases}$$

- Adjust for multiplicity by constructing weights

$$\psi_{fg} = \frac{F_{fg}}{\left(\sum_f F_{fg}\right)}$$

($\psi_{fg} = 1$ if domain covered by only one frame; $1/2$ if two frames, etc.)

- Unbiased Mecatti/multiplicity estimator for $\sum_{g \in G_1} T_g$ is

$$\sum_{g \in G_1} \sum_{f=1}^F \psi_{fg} \hat{T}_{fg}$$

- Multiplicity-adjusted difference estimator for $g \in G_2$:

$$\begin{aligned}\tilde{T}_g^* &= \sum_{k \in U_g} \sum_{\ell \in \mathcal{A}} M_{k\ell} \mu(\mathbf{x}_\ell) + \sum_{f=1}^F \psi_{fg} \sum_{k \in s_{fg}} \frac{y_k - \sum_{\ell \in \mathcal{A}} M_{k\ell} \mu(\mathbf{x}_\ell)}{\pi_k^{(f)}} \\ &= \sum_{k \in U_g} \tilde{y}_k + \sum_{f=1}^F \psi_{fg} \sum_{k \in U_g} (y_k - \tilde{y}_k) \frac{I_k^{(f)}}{\pi_k^{(f)}}\end{aligned}$$

- Unbiased difference estimator for $\sum_{g \in G_2} T_g$ is then

$$\sum_{g \in G_2} \tilde{T}_g^*$$

- G_3 has no sampling frame coverage
- Can only predict with the auxiliary data,

$$\tilde{T}_g = \sum_{k \in U_g} \sum_{\ell \in \mathcal{A}} M_{k\ell} \mu(\mathbf{x}_\ell) = \sum_{k \in U_g} \tilde{y}_k$$

- Synthetic predictor for $\sum_{g \in G_3} T_g$ is then

$$\sum_{g \in G_3} \tilde{T}_g = \sum_{g \in G_3} \sum_{k \in U_g} \tilde{y}_k$$

- Zero sampling variance, unknown bias

- Replace $M_{k\ell} = 0$ or 1 by **match metrics** $m_{k\ell} \in [0, 1]$
 - known only for sampled k
- Produced by deterministic algorithm
- Could involve formal **probabilistic record linkage** (Fellegi and Sunter 1969, Winkler 2009) or other methods
 - conditional probabilities, likelihood ratios, ...
- Whatever their origin, treat $m_{k\ell}$ as fixed in what follows

- Under perfect matching, multi-frame estimator is

$$\begin{aligned} & \sum_{g \in G_1} \sum_{f=1}^F \psi_{fg} \hat{T}_{fg} + \sum_{\ell \in \mathcal{A}} \left(\sum_{g \in G_2 \cup G_3} \sum_{k \in U_g} M_{k\ell} \right) \mu(\mathbf{x}_\ell) \\ & + \sum_{g \in G_2} \sum_{f=1}^F \psi_{fg} \sum_{k \in s_{fg}} \frac{y_k - \sum_{\ell \in \mathcal{A}} M_{k\ell} \mu(\mathbf{x}_\ell)}{\pi_k^{(f)}} \end{aligned}$$

- Under imperfect, $m_{k\ell}$ is known only for $k \in s_{fg}$
- Cannot just substitute $m_{k\ell}$ for $M_{k\ell}$ in second term, but ok in third

- Second term under perfect matching is

$$\sum_{\ell \in \mathcal{A}} \left(\sum_{g \in G_2 \cup G_3} \sum_{k \in U_g} M_{k\ell} \right) \mu(\mathbf{x}_\ell)$$

- If ℓ th record matches some element in $\cup_{g \in G_2 \cup G_3} U_g$, then
(parenthetical term) = 1
- Under imperfect matching, estimate parenthetical term
as equal to 1
— (or construct a complicated, and biased, estimator)

- Analogue of perfect-match multi-frame estimator

$$\sum_{g \in G_1} \sum_{f=1}^F \psi_{fg} \hat{T}_{fg} + \sum_{\ell \in \mathcal{A}} \left(\sum_{g \in G_2 \cup G_3} \sum_{k \in U_g} M_{k\ell} \right) \mu(\mathbf{x}_\ell) \\ + \sum_{g \in G_2} \sum_{f=1}^F \psi_{fg} \sum_{k \in s_{fg}} \frac{y_k - \sum_{\ell \in \mathcal{A}} M_{k\ell} \mu(\mathbf{x}_\ell)}{\pi_k^{(f)}}$$

is then

$$\tilde{T}_{diff} = \sum_{g \in G_1} \sum_{f=1}^F \psi_{fg} \hat{T}_{fg} + \sum_{\ell \in \mathcal{A}} (1) \mu(\mathbf{x}_\ell) \\ + \sum_{g \in G_2} \sum_{f=1}^F \psi_{fg} \sum_{k \in s_{fg}} \frac{y_k - \sum_{\ell \in \mathcal{A}} m_{k\ell} \mu(\mathbf{x}_\ell)}{\pi_k^{(f)}}$$

- Bias depends on matching and prediction error:

$$\begin{aligned} \mathbb{E} \left[\tilde{T}_{diff} \right] - T &= - \sum_{g \in G_3} T_g + \sum_{\ell \in \mathcal{A}} \mu(\mathbf{x}_\ell) - \sum_{\ell \in \mathcal{A}} \left(\sum_{g \in G_2} \sum_{k \in U_g} m_{k\ell} \right) \mu(\mathbf{x}_\ell) \\ &= -(\text{total uncovered}) + (\text{database}) - (\text{overlap}) \end{aligned}$$

- Sufficient conditions for unbiased estimation are

$$G_3 = \emptyset \text{ and } \sum_{g \in G_2} \sum_{k \in U_g} m_{k\ell} = 1 \text{ for all } \ell \in \mathcal{A}$$

- Asymptotic unbiasedness and mean square consistency requires some combination of “not too much” matching error or undercoverage, and “good” prediction of the uncovered population

- Variance of the estimator is (setting $m_{k\ell} \equiv 0$ for $k \in \cup_{g \in G_1} U_g$):

$$\sum_{f=1}^F \sum_{g \in G_1 \cup G_2} \sum_{g' \in G_1 \cup G_2} \psi_{fg} \psi_{fg'} \sum_{j \in U_g} \sum_{k \in U_{g'}} \Delta_{jk}^{(f)} \frac{d_j}{\pi_j^{(f)}} \frac{d_k}{\pi_k^{(f)}}$$

$$\text{with } d_j = \begin{cases} y_j - \sum_{\ell \in \mathcal{A}} M_{j\ell} \mu(\mathbf{x}_\ell), & \text{perfect matching} \\ \text{prediction error} \\ y_j - \sum_{\ell \in \mathcal{A}} m_{j\ell} \mu(\mathbf{x}_\ell), & \text{imperfect matching} \\ \text{matching and/or prediction error} \end{cases}$$

- $\text{Var} \left(\tilde{T}_{diff} \right) = O \left(\frac{N^2}{\min_f n_f} \right)$ and $N^{-1} \tilde{T}_{diff} \xrightarrow{\text{m.s.}} N^{-1} \mathbb{E} \left[\tilde{T}_{diff} \right]$
- Unbiased variance estimation provided all $\pi_{jk}^{(f)} > 0$ in each frame

Use SC recreational fishery to devise a simulation study ²³

- About 450 charter boats and 15,000 boat trips along the Atlantic Coast each year
- Survey data from sampled angler on boat trip on the actual date
 - coverage error: not all sites and times are in-frame
 - lots of sampling error
- Logbook data from captain's report, later that month
 - nonresponse
 - measurement error
- Lots of matching error!

- **Perfect match:** $m_{k\ell} = 1$ for $\ell = \ell_1$ and 0 otherwise
- **High-quality match:** $m_{k\bullet} = \sum_{\ell \in \mathcal{A}} m_{k\ell} = 1$

$$m_{k\ell} = \begin{cases} 1/3, & \text{if } \ell = \ell_1, \ell = \ell_2 \text{ or } \ell = \ell_3, \\ 0, & \text{otherwise} \end{cases}$$

- **Low-quality match:** $m_{k\bullet} = \sum_{\ell \in \mathcal{A}} m_{k\ell} < 1$

$$m_{k\ell} = \begin{cases} 1/6, & \text{if } \ell = \ell_1, \ell = \ell_2 \text{ or } \ell = \ell_3, \\ 0, & \text{otherwise} \end{cases}$$

- **No match:** $m_{k\ell} = 0$ for all $\ell \in \mathcal{A}$

- Match metrics $\{m_{kl}\}_{k \in s, l \in \mathcal{A}}$ developed by South Carolina Department of Natural Resources staff

Interview variables	Logbook variables
Date of interview	Date of reported trip
Time of interview	Estimated trip end time
License number of vessel	License number of vessel
Name of vessel given	Name of vessel reporting
Interview site	Reported start site

- Large fraction of unmatched trips and low-quality matches

	No Match	LQ	HQ	Perfect
Empirical	11.0%	52.5%	36.5%	0.0%

- Use real logbook data to create artificial population with $|U| = 10,647$ boat trips, sorted in space and time
- Use Markov chain to assign (unobservable) states to groups of population boat trips:

	state from Markov chain			
	no match	LQ	HQ	perfect
size of group of elements:	1	10	5	1
logbook records created:	0	5	5	1
metric sum:	0	1/2	1	1

- If an **LQ** element is selected, metrics (correctly) indicate it might match one of five records, or none of them

- Set Markov chain parameters to simulate match metrics $\{m_{k\ell}\}$ and logbook database $\mathcal{A}$ under two scenarios:

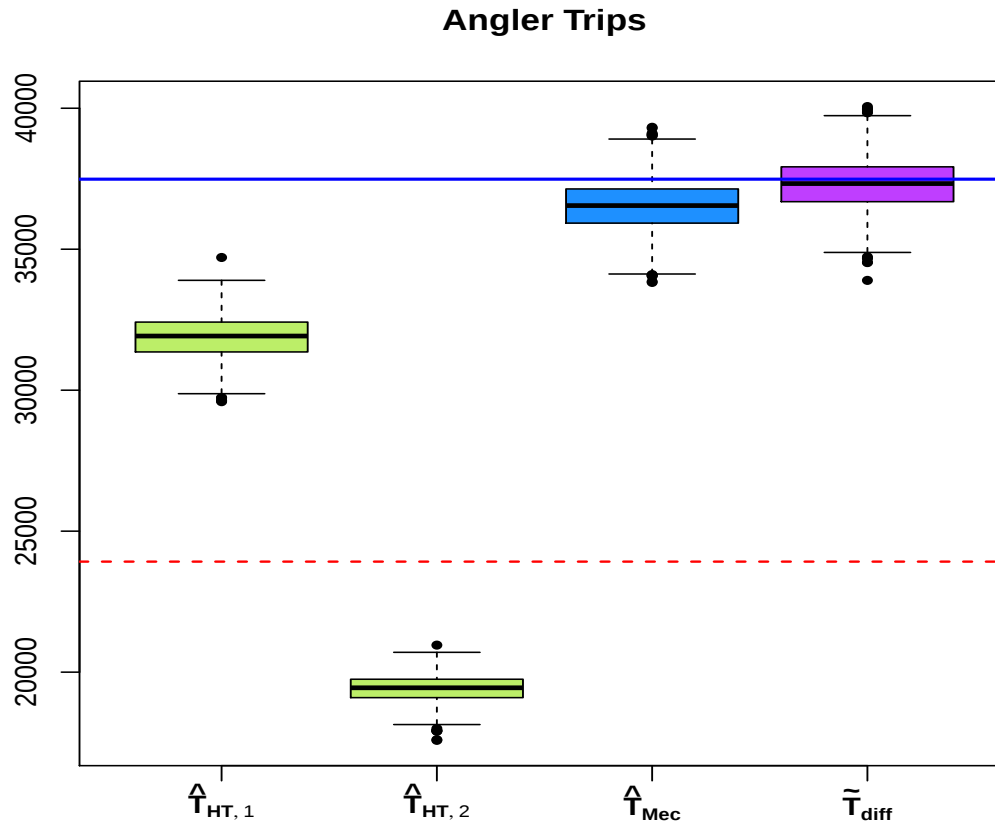
	$ \mathcal{A} $	No Match	LQ	HQ	Perfect
Poor Match	6,836	8.6%	54.4%	31.7%	5.3%
Better Match	9,031	2.3%	23.3%	69.8%	4.7%
Empirical		11.0%	52.5%	36.5%	0.0%

- Population of boat-trips and database of logbook records is then fixed
- Create two incomplete frames, partially overlapping
- Sample repeatedly from this finite population

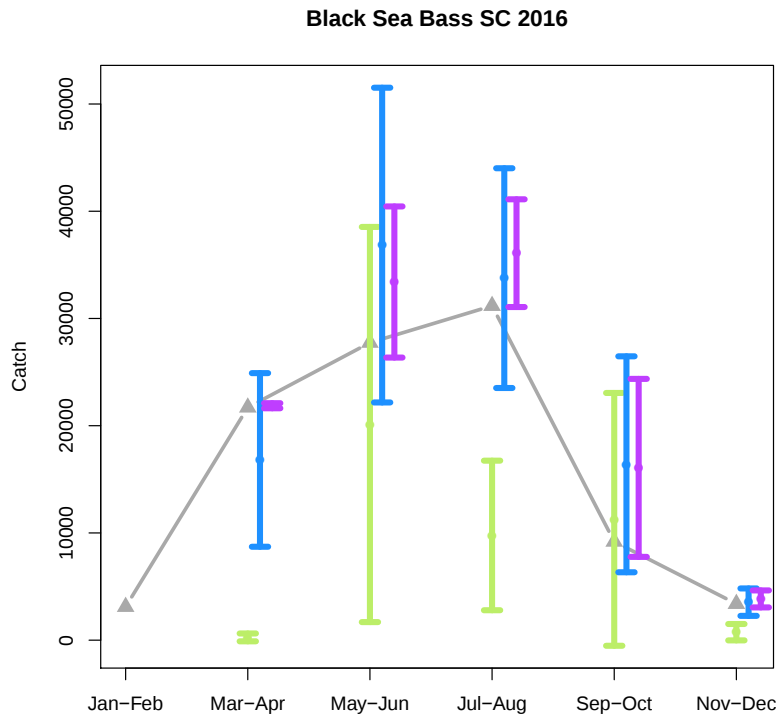
- Simplifications:
 - no “differential matching”: quality of $m_{k\ell}$ does not depend on y_k
 - no measurement error: $\mu(\mathbf{x}_\ell) = y_k$ for perfect match
- Draw 1000 repeated samples from simulated population
 - stratified, two-stage, unequal-probability selection
- Compute $\hat{T}_{HT,1}$, $\hat{T}_{HT,2}$, $\hat{T}_{Mec}$, $\tilde{T}_{diff}$ for number of angler trips and several species in each simulated sample
- Assess bias, variance, and MSE for each estimator

Even with poor match, difference dominates Mecatti

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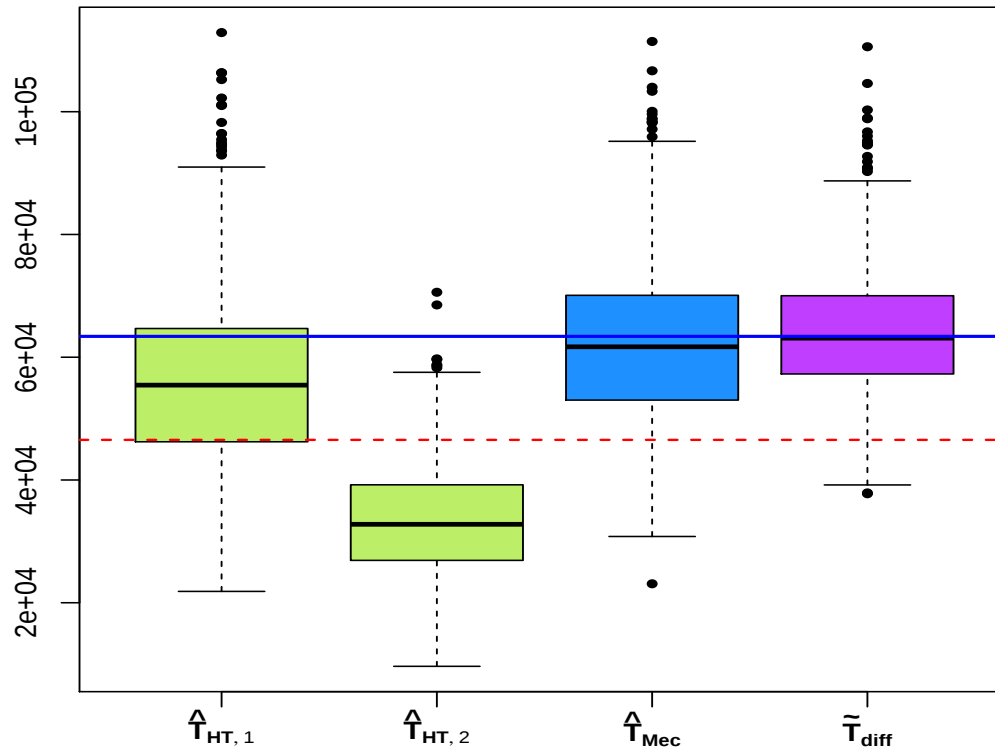
- Frequently targeted and caught; appears regularly in both sources



Difference estimator dominates Mecatti

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Black Sea Bass Catch



- Auxiliary information is useful even with imperfect matching
 - naive difference estimator improves accuracy and precision of multiplicity estimator
 - variance estimators and confidence intervals (not shown) work well
- Matching across frames or matching across auxiliary databases adds challenges
- Grazie mille!
Contact info: FJay.Breidt@colostate.edu