

New Effective Population Size Findings for Atlantic and Shortnose Sturgeons

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Effective Population Size, N_e (N_{ef})

- The average number of individuals in a population that actually contribute genes to succeeding generations.
- Often considered as a hypothetical “idealized” population.
- In this ideal population there is random mating and no selection.
- *The effective size of a population is typically smaller than its actual size.*
- 50 / 500 rule

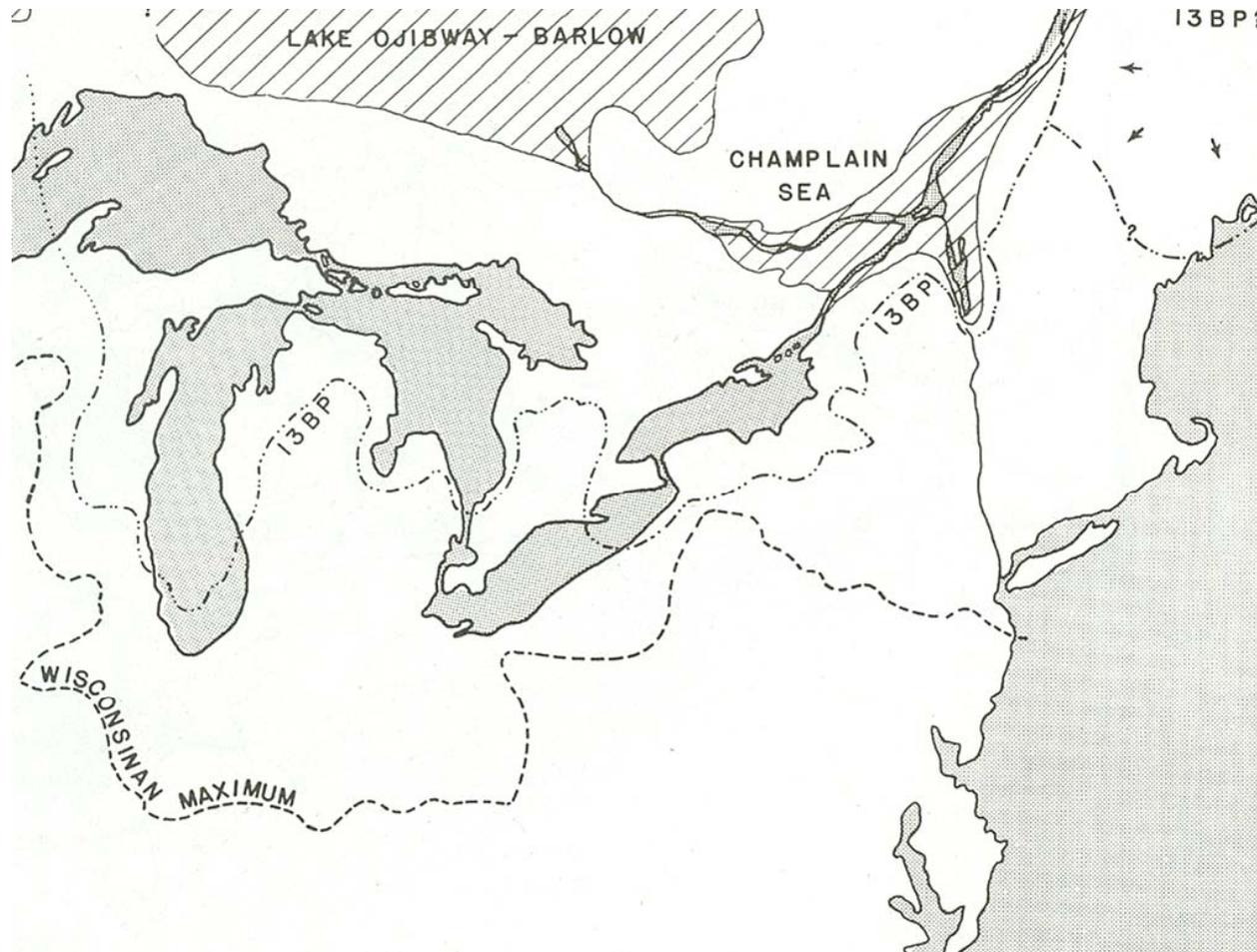
N_{ef} / N_f ratios

- Nunney & Elam (1994) – theory indicates *N_{ef} / N_f* should range between 0.25 and 0.75
- Red drum, Gulf of Mexico = 0.004
- Chinook salmon, Sacramento R. = 0.04
- Silvery minnow, Rio Grande = 0.005
- Frankham, 38 cases small pops = ~0.10

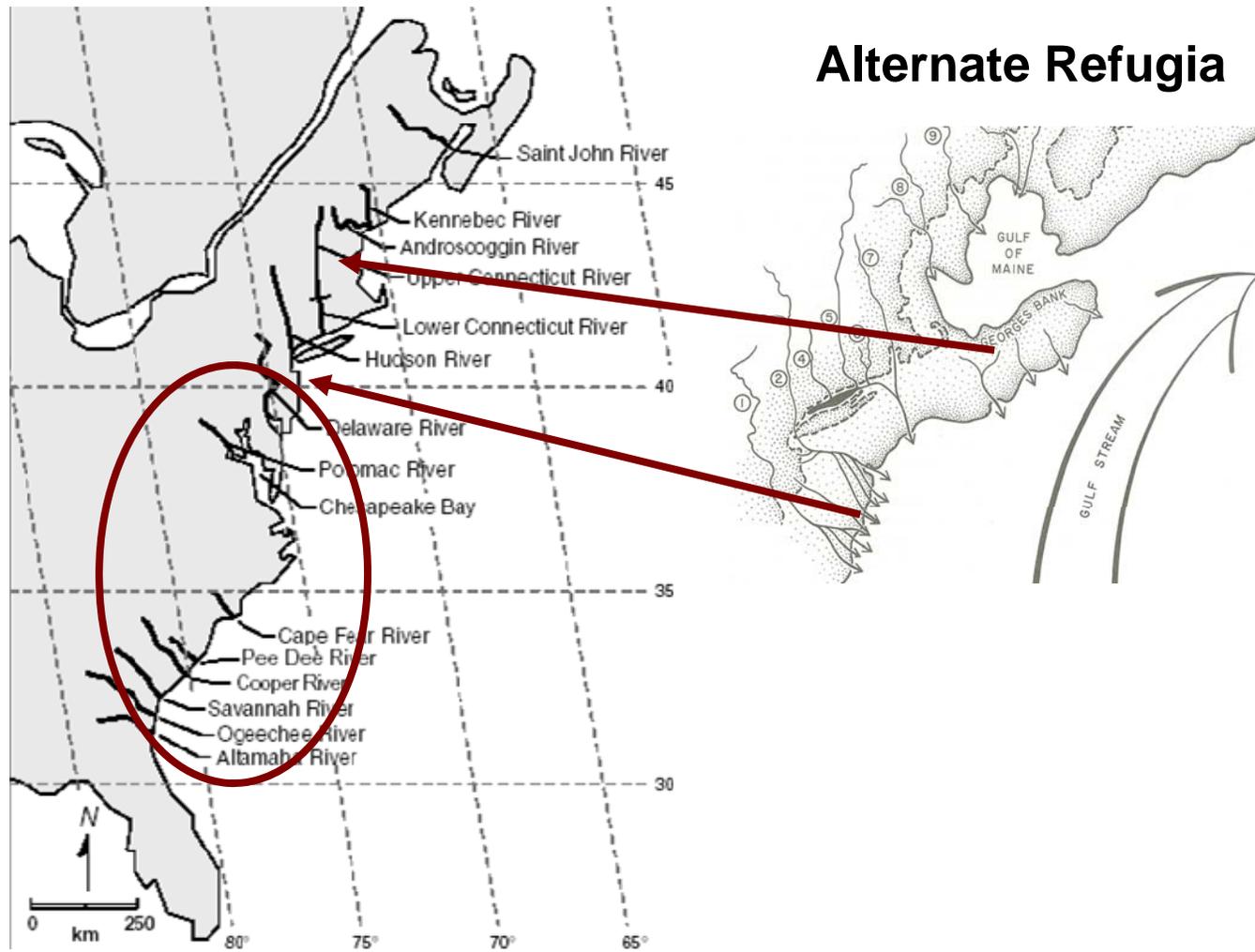
Methods

- MIGRATE software to estimate mutation-scaled effective population size parameter
- Bayesian inference (alternative of max likelihood did not stabilize)
- Independent runs: 4 Shortnose, 3 Atlantics
- More arcane details that I won't mention
- The conceit: two species, many pops, wide abundance ranges

“Northern” vs. “Southern” Rivers



Alternate Refugia



Atlantic Sturgeon

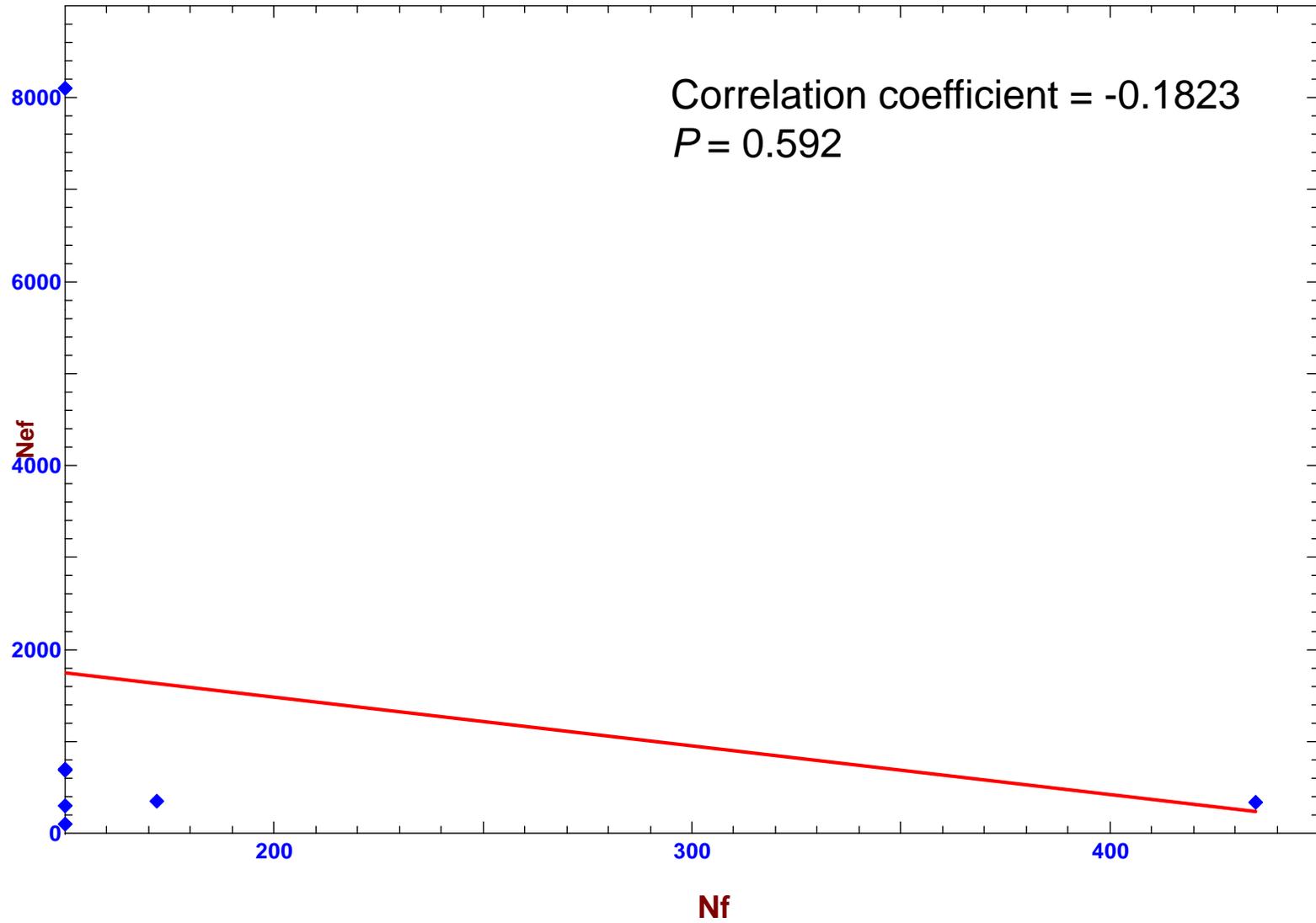
River	Sample size	# Haplotypes	<i>Nef</i>	<i>Nf (adults)</i>
St. Lawrence	46	1	82	150
St. John	76	1	97	150
Kennebec	19	2	2432**	150
Hudson	128	8	341	435
Delaware-YOY	19	6	106	150
James	65	4	294	150
Albemarle	48	6	680	150
ACE basin	72	11	5636	150
Savannah	58	18	8108	150
Ogeechee	47	9	697	150
Altamaha	140	5	352	172

(** estimates never stabilized)

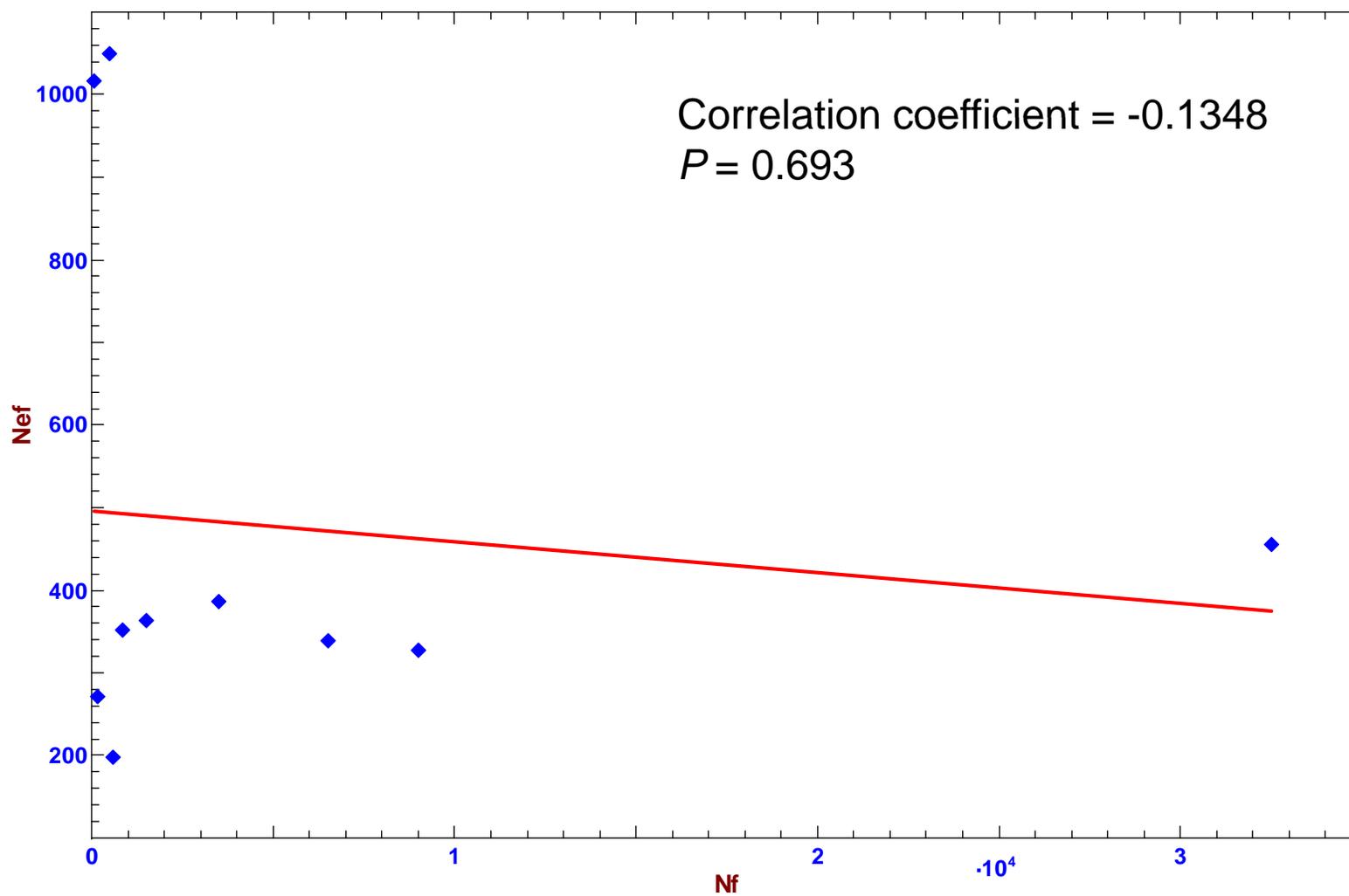
Shortnose Sturgeon

River	Sample size	# Haplotypes	<i>N_{ef}</i>	<i>N_f</i> (all ages)
St. John	42	8	327	9000
Kennebec	54	8	386	3500
Androscoggin	48	8	364	1500
Connecticut	46	4	198	600
Hudson	56	9	456	32,500
Delaware	57	8	339	6500
Winyah Bay	46	13	1099	500
Cooper	62	6	271	150
Savannah	25	7	352	837
Ogeechee	53	11	1016	74
Altamaha	69	10	583	3160

Atlantic Sturgeon



Shortnose Sturgeon



Two-Species Comparison

	Mean <i>Nef</i>	SD <i>Nef</i>	<i>Nef/Nf</i>	Range <i>Nef/Nf</i>
Atlantic	1711	2693	9.49	0.647 – 54.05
Shortnose	490	298	1.47	0.014 – 13.73

Note: Quattro et al. (2002) - shortnose sturgeon *Nef* in Southeast ~12,000

Zhang et al. (2003) – Chinese sturgeon in Yangtze, *Nef* ~2,800, *Nef/Nf* ~0.85

mtDNA vs. nDNA Microsatellites for Atlantic Sturgeon (Tim King)

- King used linkage-disequilibrium approach
- nDNA range $Ne = 7 - 169$
- nDNA Ne vs. $N = 0.02 - 0.30$
- mtDNA N_{ef} & nDNA Ne not correlated
($P > 0.05$)
- Confounding factors: molecular markers, genders, age classes in census estimates & statistical approaches

Preliminary Conclusions

- AS *Nef* higher but more variable than SNS
- *Nef* not correlated with *Nf* for either species
- AS *Nef* using mtDNA higher than with nDNA
- *Nef/Nf* much higher than for many other low abundance vertebrates
- 50/500 rule a concern depending on whether you trust mtDNA or nDNA results

Preliminary Conclusions

- No clear response either to range of current pop sizes nor to recent pop changes
- Benefits of “genetic storage effect”?
 - Long lived; skip spawning years; long time to maturity with sex differences – leading to considerable generation overlap that limits genetic drift