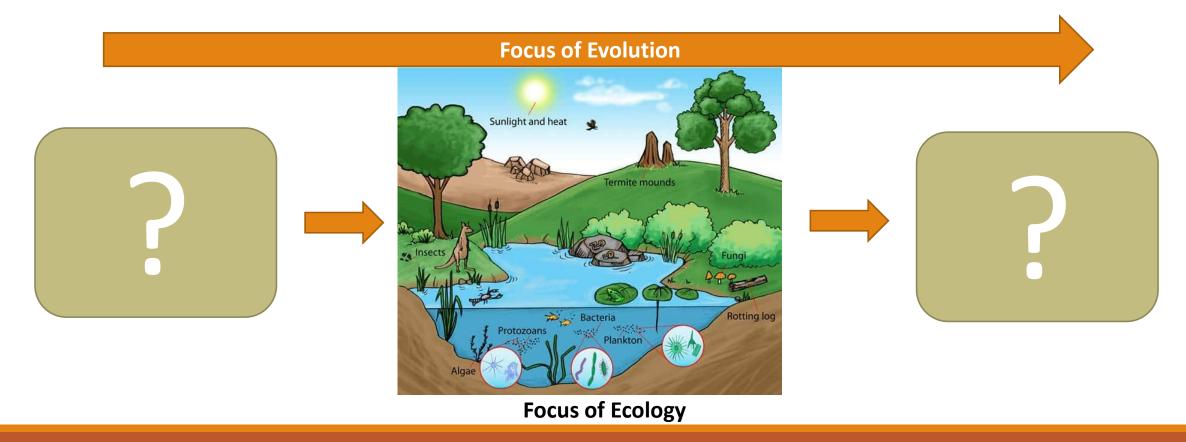
Using molecular approaches to understand the drivers of population divergence and speciation

2025 MMEE COURSE

MMEE; Ecology & Evolution

Do they always live like as they do now?

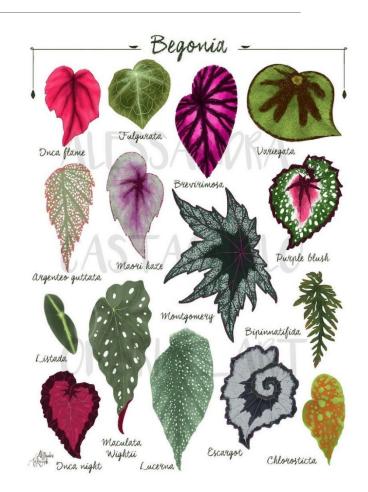


Classical ecology&evolution has limited resolution

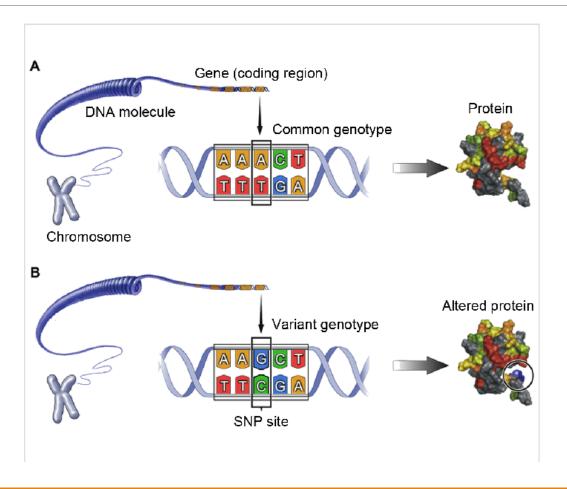
Nature can confuse us with organism morphology



Nikaido et al, 1999



SNPs can alter protein structure, so the function and ultimately phenotype



Iglesias et al, 2020

Targeted marker sequencing for identification

Good marker genes can reflect the difference among individual in molecular level

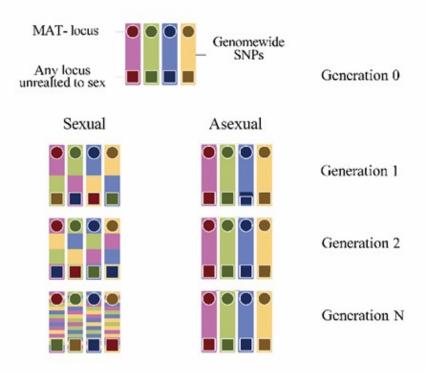


Genome encodes far higher number of genes than the number of markers we use

	Organism	# of protein- coding genes	# of genes naïve estimate: (genome size /1000)	BNID
viruses	HIV 1	9	10	105769
	Influenza A virus	10-11	14	105767
	Bacteriophage λ	66	49	105770
	Epstein Barr virus	80	170	103246
prokaryotes	Buchnera sp.	610	640	105757
	T. maritima	1,900	1,900	105766
	S. aureus	2,700	2,900	105500
	V. cholerae	3,900	4,000	105760
	B. subtilis	4,400	4,200	111448
	E. coli	4,300	4,600	105443
eukaryotes	S. cerevisiae	6,600	12,000	105444
	C. elegans	20,000	100,000	101364
	A. thaliana	27,000	140,000	111380
	D. melanogaster	14,000	140,000	111379
	F. rubripes	19,000	400,000	111375
	Z. mays	33,000	2,300,000	110565
	M. musculus	20,000	2,800,000	100308
	H. sapiens	21,000	3,200,000	100399, 111378
L	T. aestivum (hexaploid)	95,000	16,800,000	105448, 102713

However, loci of an organism do not coevolve together in most organisms





Lee et al, 2024

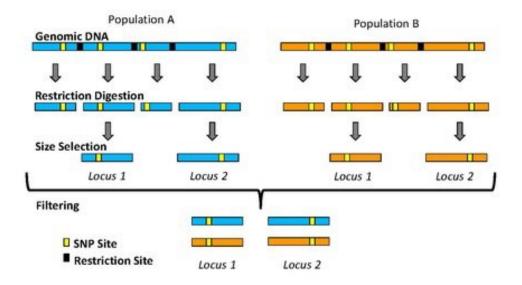
Cost is the main problem

Restriction enzyme associated DNA sequencing (RAD sequencing)

Double digested RAD sequencing

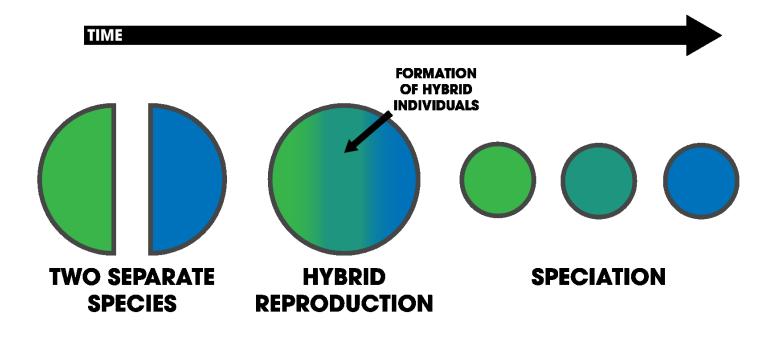
-> Sequence the subset of genome

Restriction-site Associate DNA Sequencing (RADseq)

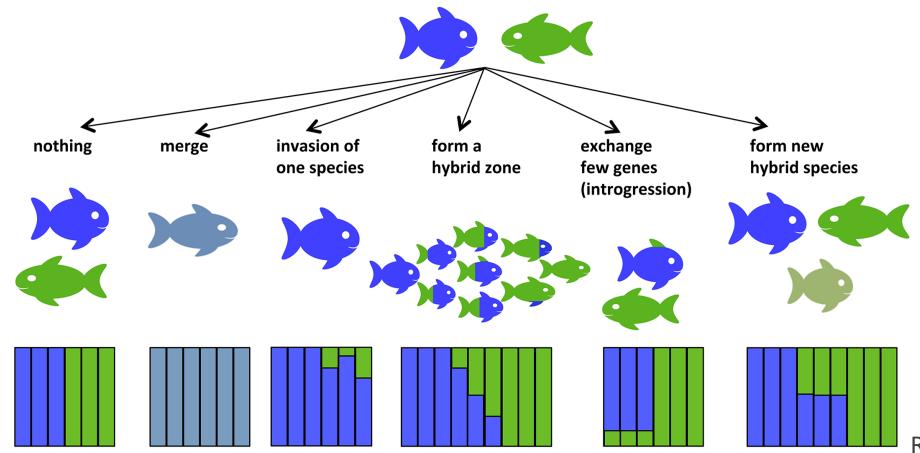


Speciation is continuum, and we cannot detect phenotypically until certain threshold

Genome investigation is a wonderful tool to detect this **even before it becomes obvious for phenotypes**

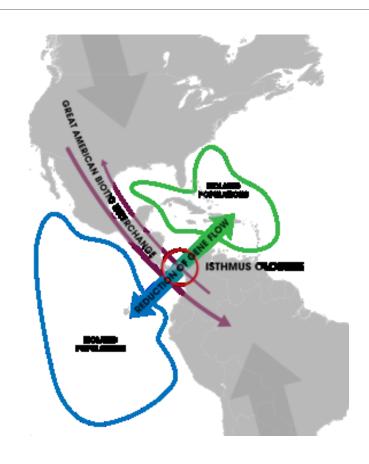


Theoretical models of hybridization



Runemark et al, 2019

During the practical parts of MMEE...



Types of Speciation

