Substructured population growth in the Ashkenazi Jews inferred with Approximate Bayesian Computation Ariella L. Gladstein^{1*} and Michael F. Hammer²

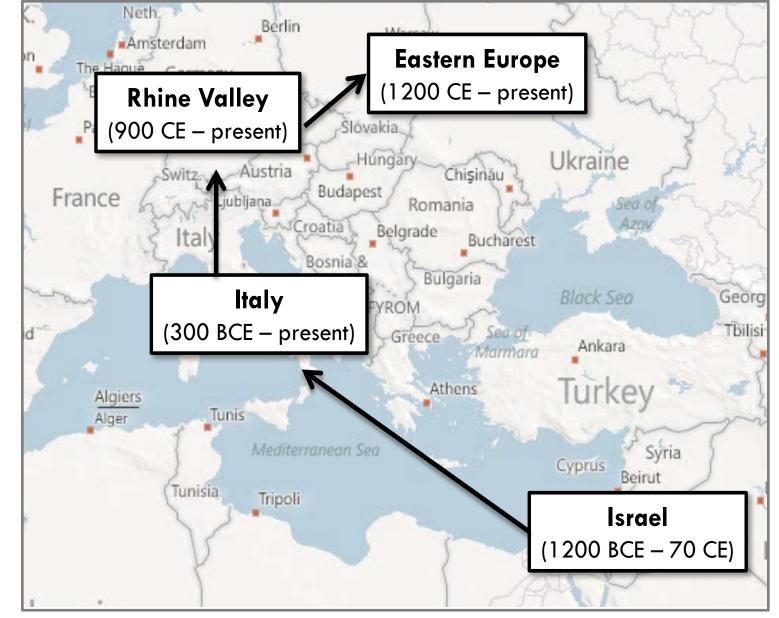
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Introduction

Who are the Ashkenazi Jews (AJ)?

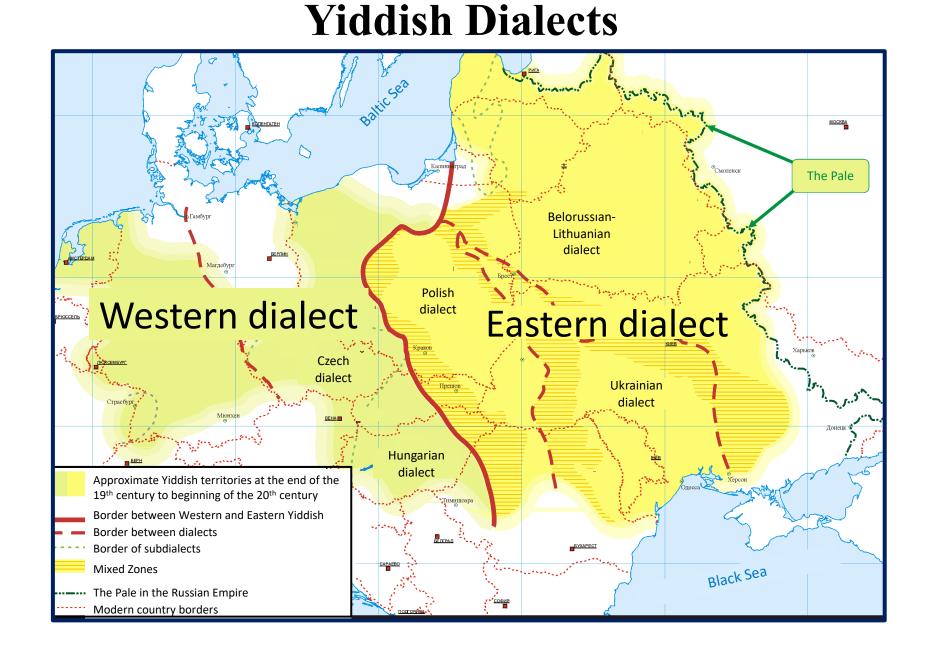
AJ culturally, religiously, and linguistically identify as Jews whose ancestors came from the Rhine Valley.

Hypothesis of AJ origins



Western vs. Eastern AJ

- 11th 16th centuries AJ expelled from Central Europe and invited to Poland by nobility.
- From 16th century culturally distinct communities developed.



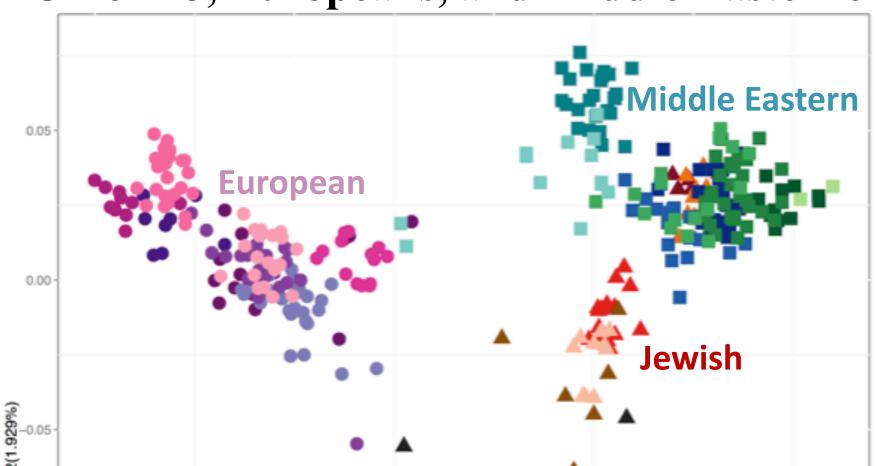
Results

- Eastern and Western AJ are separated on PCA.
- ABC chosen model is M2 (see Model Choice table below).
- Eastern and Western growth rate calculated from estimated Ne's are on the order of census growth rates.

Model Choice

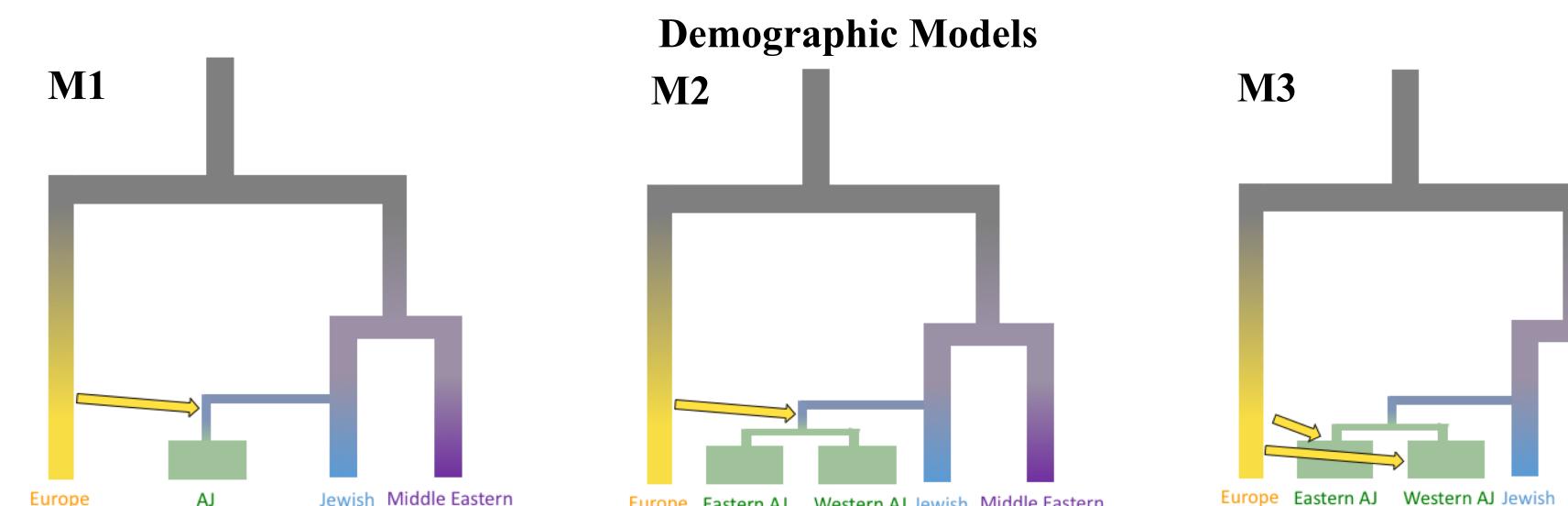
											_		
		M1	M2	M3			• •			A	Jew i	ish	
Bayes Factor		0.0005	2.01	0.50	(%6 6 -0.05 -						*		
P(false negative M2)		0.08		0.14	PC2(1.929%)				•	4			
Exponen	tial Growth R	ate (per	gene	ration)	-0.10 -						Eastern		
	Eastern AJ	West	ern A	J							Ash	kenazi	
ABC	0.25	0.069			-0.15 -					^	Weste	ern	
Census	0.18 +- 0.01	7 0.081	+- 0.0	026							4		
	Bes	t Model				-0.10		-0.05	PC1(4.197	0.00	0.05	6	
									(Census	5 ⁵ Grov	vth	
							ion size	6e+06			Ea	astern	•
		Tem	$n \approx 12 \text{ ky}$	a			Census population	+06				لحر	'
							od sn	- 3e+					
							Censi	0e+00		===	Western	/	-
									1200	1400	1600	1800	
			Tr	nj \approx 5 kya	,			-			Year	_	•
								E	lack line	es are est	imated E	astern Jew	V1

PCA of AJ, Europeans, and Middle Easterners



Questions:

- 1) Is there substructure within AJ?
- 2) If so, is this substructure purely due to genetic isolation, or does differential migration from European populations play a role?

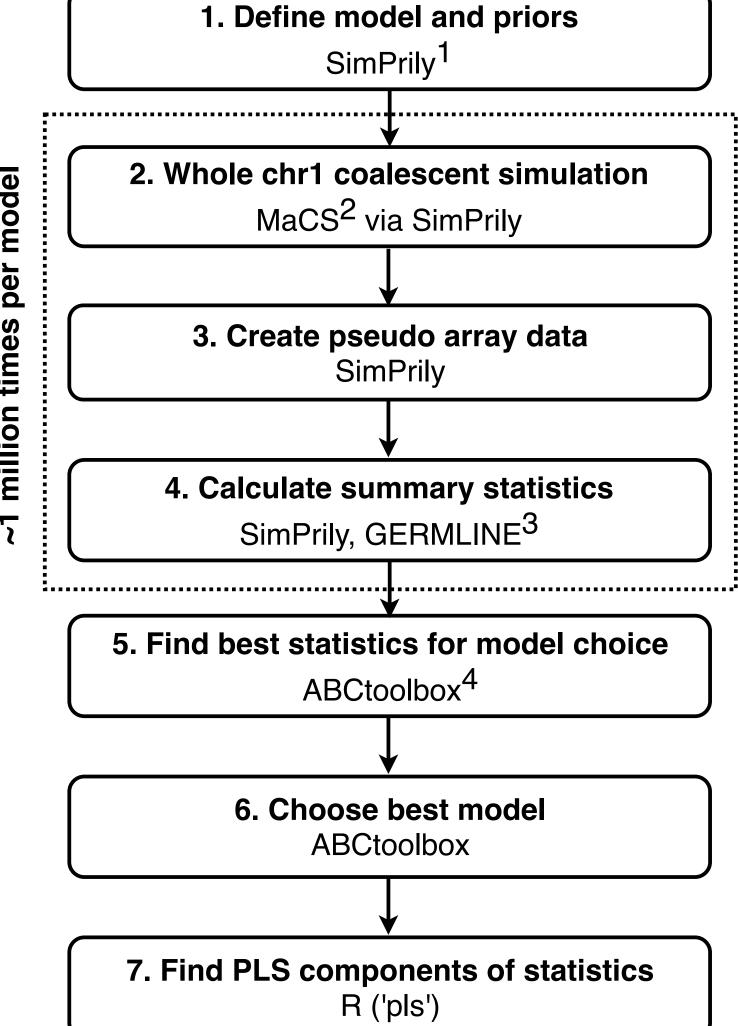


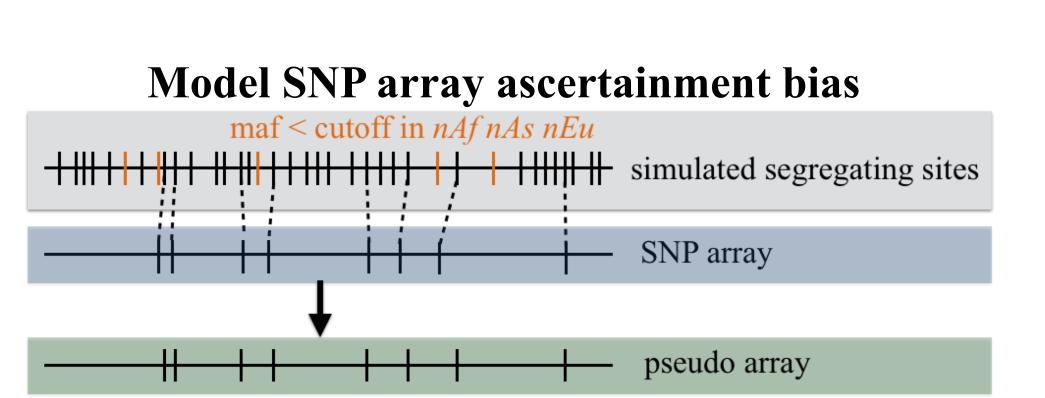
Europe

Europe Eastern AJ Western AJ Jewish Middle Eastern

Europe Eastern AJ Western AJ Jewish Middle Eastern

Methods



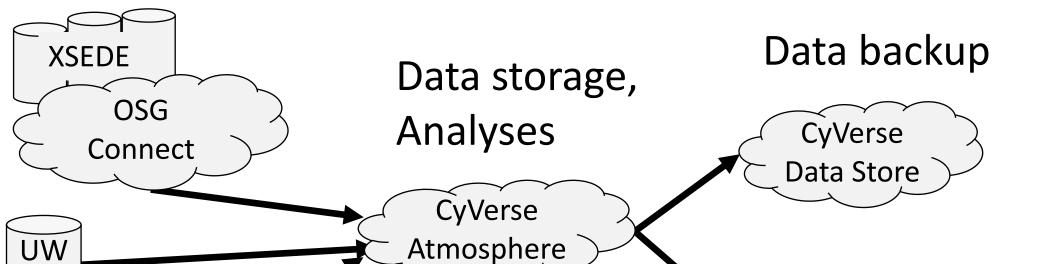


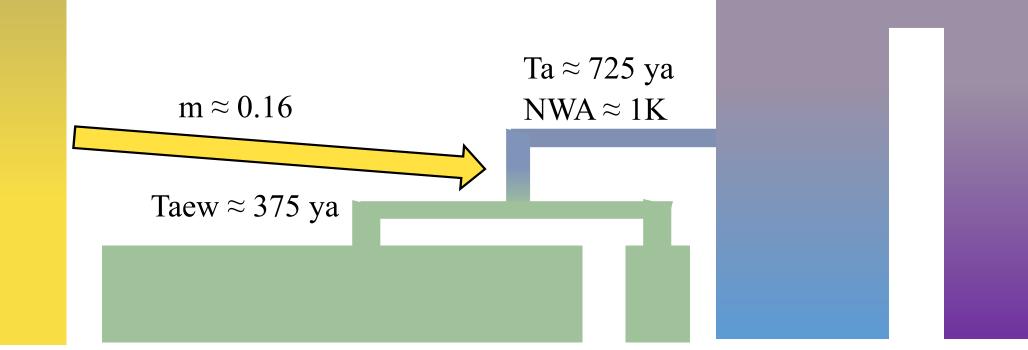
181 summary statistics

- Site frequency spectrum-based stats for older history
- Identity by descent-based stats for recent history

Simulations on HTC clusters, analyses on cloud

Simulations





Middle Eastern Eastern AJ Europe Western AJ Jewish $NM \approx 470K$ $NJ \approx 400K$ NWA $\approx 8K$ $NEA \approx 1.5M$

Conclusions:

- 1) There is substructure within AJ
- 2) Differentiation between Eastern and Western AJ is best explained by isolation followed by differences in population growth rates (higher in Eastern AJ).

 \checkmark

Historical differences that may have contributed to greater growth in Eastern AJ

	Eastern	Central
	Europe	Europe
Restricted movement and living area.		\checkmark
Limitation on number of families.		\checkmark

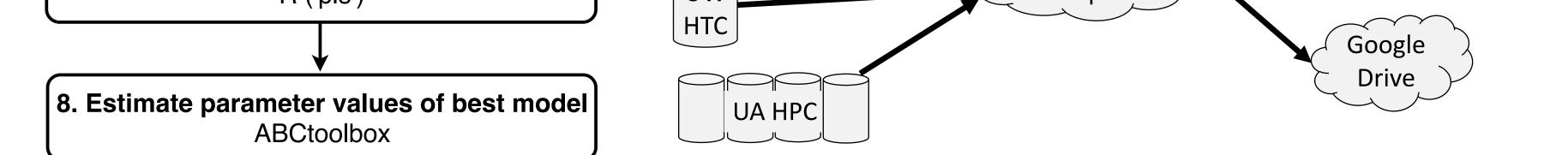
References

1) Gladstein et al., 2018; 2) Chen et al. 2009; 3) Gusev et al. 2009; 4) Wegmann et al. 2009; 5) DellaPergola 2001

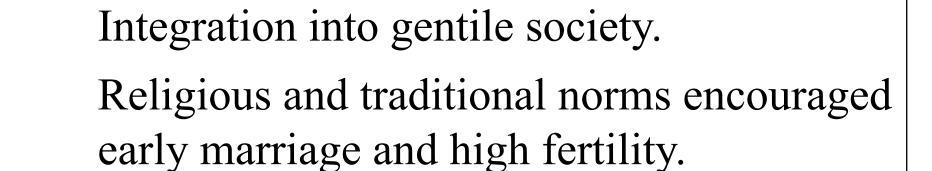
Acknowledgments

We would like to thank Mats Rynge for his help setting up the Pegasus workflow and running it on the Open Science Grid. We would also like to thank Research Bazaar Arizona for programming help.

census sizes and grey lines are estimated Western/Southern Jewish census sizes. Dashed lines are exponential growth models fitted to the census data.



Full pipeline available at https://bitbucket.org/agladstein/macsswig_simsaj/wiki/Home



Compute Resources University of Arizona Research Computing HPC; Open Science Grid; CyVerse; Extreme Science and Discovery Environment (Comet, Jetstream, Bridges); University of Wisconsin Center for HTC

