	Conventional wisdom			
Are we still evolving? Alan R. Rogers	Something must have happened to weaken the selective pressure drastically. We cannot escape the conclusion that man's evolution towards manness suddenly came to a halt. —Ernst Mayr 1963 Natural selection has almost become irrelevant in human evolution. There's been no biological change in humans in 40,000 or 50,000 years. Everything we call culture and civilization we've built with			
September 24, 2012	the same body and brain. —Stephen Jay Gould 2000			
September 24, 2013	Certainly, human nature is fixed. It's universal and unchanging —common to every baby that's born, down through the history of our species. —Helena Cronin. 2000 Getting Human Nature Right.			
If humans had been evolving lately, how would we know?	Look at some DNA sequence data			
How can recent selection be detected? Let's examine some human DNA.	RefT G C A T G T A A T G C TA A G A A .B A G AC T . G CDC G G CDC			
What happens when a favorable mutation strikes some lucky chromosome?	Cross-overs shuffle DNA			
	Image: Constraint of the second secon			

- ► Better-than-average chance of spreading through population.
- What is it that spreads?
 - ► The entire chromosome?
 - Just the mutated nucleotide site?

Neither: the answer is in between.



- shuffle parental chromosomes.
- sites far apart shuffled more.
- result: "recombinant" chromosomes

The chromosome you pass on to your child

What happens to lucky chromosome?

- ▶ Not like either of those you inherited from Mom or Dad.
- A mosaic, made of pieces of Mom's and Dad's.
- ► Technical terms: *recombination*, *recombinant chromosomes*
- Favorable mutation surrounded by region from original chromosome.
- \blacktriangleright Recombination gradually reduces size of that region.
- The older the mutation, the smaller the region.
- ► Technical term: *linkage disequilibrium* (LD)

How a selective sweep generates LD



Common neutral mutations

- May accidentally drift to high frequency, but this takes a long time.
- Plenty of time for recombination.
- Sit on *short* stretches of original chromosome.

Common favorable mutations

DNA sequences from region of human lactase gene

	cgcttcaggcattcctatctaaacagaccaacgtaAgggtacaatgcctaacccagacgtttcaactct
20	
21	
22	
23	
24	
25	
26	
27	t
28	tt
29	c
37	Ga.gttgac.c.tgtct.
38	ccggagatatggctc.gGaaa.gccttttgct.t
39	ccggagatatggctc.gGaaa.gccttttgct.t
40	tccagtag.t.catgtttccgGa.gttgac.c.tgtct.
41	tccagtag.t.catgt.gttccgGa.gttgac.c.tgtct.
42	tccagtag.t.catgt.gttccgGa.gttgac.c.tgtct.
43	tccagtag.t.catgt.g.tc.gGa.gttgac.c.tgtct.
44	tccagtag.t.catgtttc.gGacgttgac.c.tgtct.
45	tccagtag.t.catgt.gttc.gGa.gttgac.c.tgtct.
46	ccggagatatggctc.gGaaa.gccttttgcg.gt.tc
47	tccagtag.t.catgt.gttccgGa.gttgac.c.tgtct.
48	tccagtag.t.catgt.gttccgGa.gttgac.c.tgtct.
49	tccagtag.t.catgt.gttccgGa.gttgac.c.tgtct.
50	tatccggag.tc.atcgg.tc.g.tg.tc.gGa.g.gtgggtcg.gt.tc
51	ta.ccggag.tatcgg.tc.g.tg.tc.gGa.g.gtgggtcg.gt.tc
52	ta.ccggag.tatc.g.tc.g.tg.tc.gGa.g.gtgggtcg.gt.tc
53	$\texttt{ta.ccgga} \dots \texttt{.g.t.atcgg.tc.g.tg.tc.gG} \dots \texttt{.a.g.g} \dots \texttt{tgggt} \dots \texttt{cg.gt.t.c}$

- Increase rapidly in frequency
- ► Little time for recombination.
- Sit on *long* stretches of original chromosome.

Evidence for natural selection at lactase gene

Lucky chromosomes nearly identical.

▶ Region of LD covers nearly a million nucleotides in European

Unlucky chromosomes vary.

Absent in Africa and Asia.

population.

LD on human chromosome 2 (Voigt et al 2006)





GO Nesting	GO Category	ASN	CEU	YRI
	5,			
21–1	Chemosensory perception	-	0.0006	0.0004
21-1-1	Olfaction	-	0.0006	0.0008
22–2	Gametogenesis	0.008	-	-
22-2-2	Spermatogenesis and motility	0.02	0.03	-
22–3	Fertilization	0.004	0.003	-
1–11	Other carbohydrate metabolism	0.0002	-	-
6	Electron transport	-	0.0002	-
4–13	Chromatin packaging/remodeling	< 0.0001	0.01	-
16–1–1	MHC-I-mediated immunity	-	<0.0001	0.02
3–2	Steroid metabolism	-	-	< 0.0001
3–5	Lipid and fatty acid binding	0.001	-	-
4-4-2	mRNA transcription initiation	-	0.002	-
5–3	Protein modification	0.002	-	-
7–5	Vitamin/cofactor transport	0.002	-	-
9	Phosphate metabolism	0.002	0.03	-
13–4	Peroxisome transport	-	-	0.002

All *p*-values are one-sided, testing for enrichment of categories in each population; "-" indicates that the *p*-value is >0.05.

What evolutionary changes might we expect?

Cause	\longrightarrow Expected response	
diet	\longrightarrow metabolism (protein, carbs, etc)	
longevity	\longrightarrow DNA repair	
intelligence	\longrightarrow neural function	
population density \longrightarrow immune response		





Why the recent burst of evolution?

Summary

Hawks et al (2007) argue that we have changed our own environment:

- population size
- ► agriculture
- complex society
- Iong-distance trade

To say nothing of the more recent changes.

- Recent positive selection on many human genes.
- ► Favored alleles originated within past 40 kyr.
- A burst of adaptive evolution.
- Several functional categories.
- Suggests responses to increased population size, to shifts in diet, and to longevity.