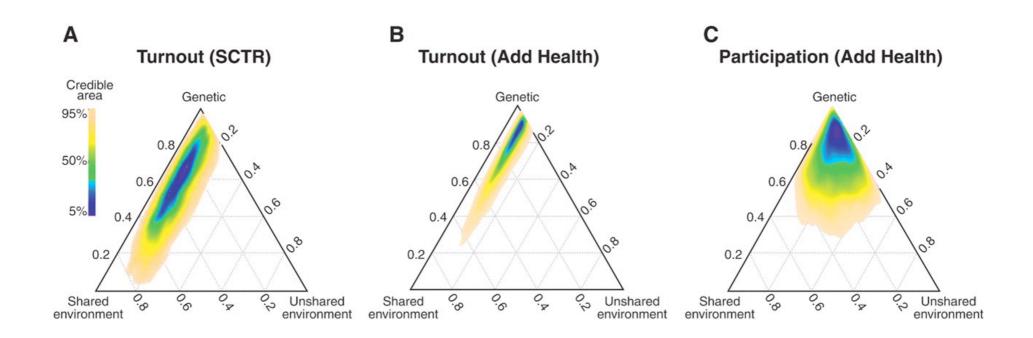
The CHRNA6 gene, patience, and voter turnout

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Road Map

- Behaviour genetics, genopolitics
- Patience and voter turnout
- CHRNB3 and CHRNA6
- Results



Fowler, Baker and Dawes. 2008. "Genetic Variation in Political Participation." APSR.

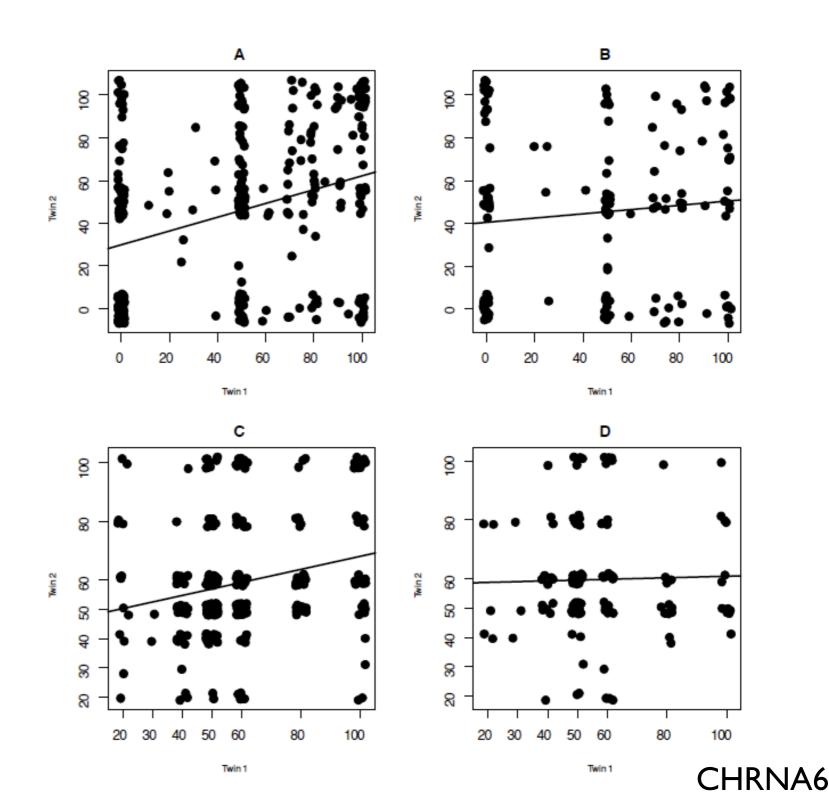
Cesarini et al. 2008.

"Genetic Variation in Preferences for Giving and Risk-Taking." Quarterly Journal of Economics.

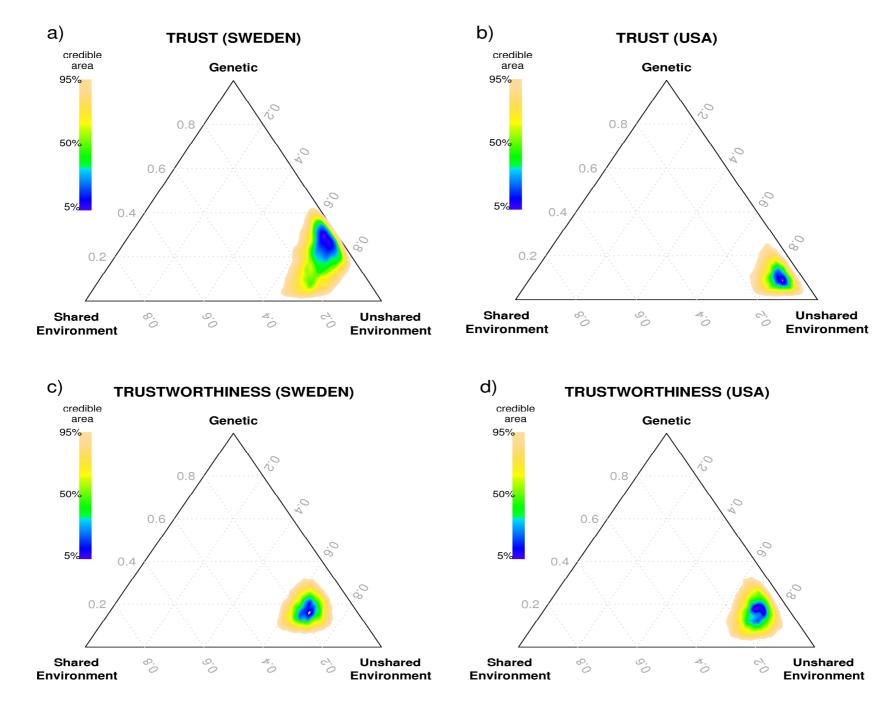
Panel A: DG, MZ. Panel B: DG, DZ.

Panel C: Risk Aver., MZ

Panel D: Risk Aver., DZ

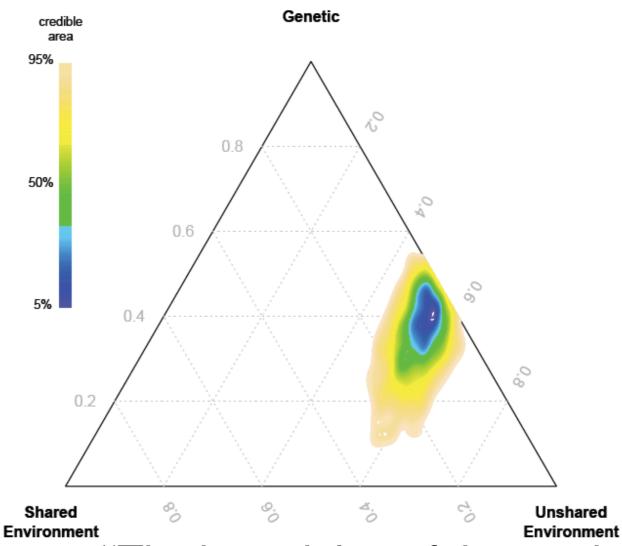


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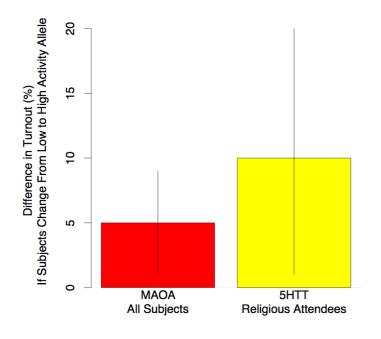
Cesarini et al. 2008. "Heritability of cooperative behavior in the trust game." PNAS.

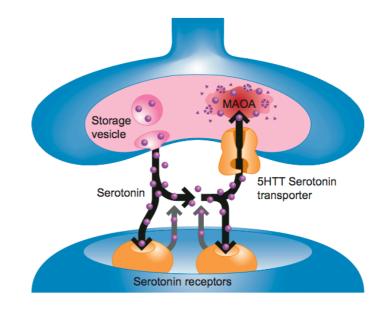
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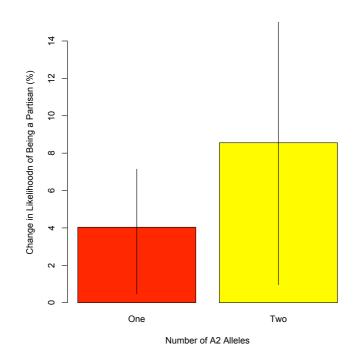


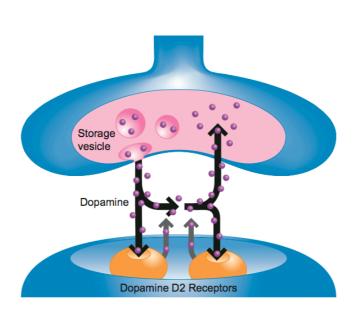
Loewen and Dawes. "The heritability of duty and voter turnout." R and R, Political Psychology

Evidence from gene association studies









Fowler and Dawes. 2008. "Two Genes Predict Voter Turnout." *JOP*. Dawes and Fowler. 2009. "Partisanship, Voting, & the Dopamine D2 Receptor Gene." *JOP*.

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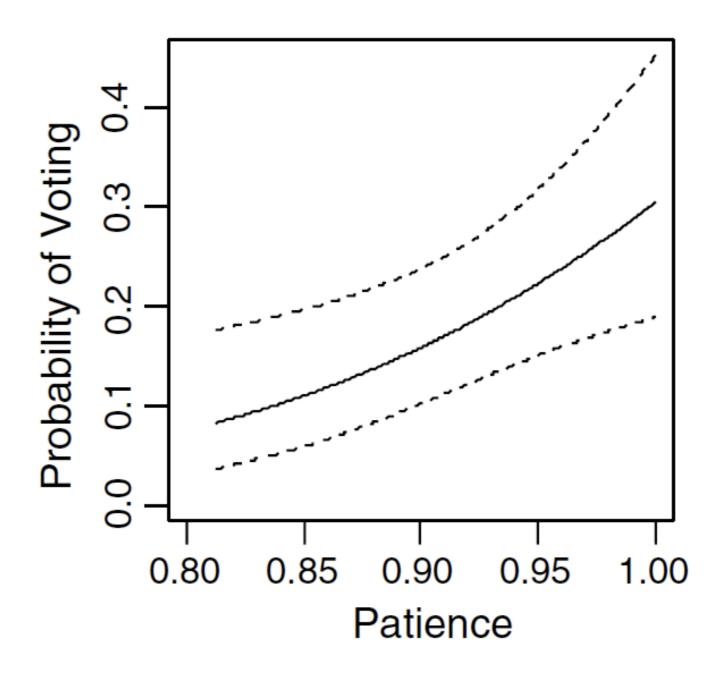
CHRNA, Patience and Turnout

- Patience and turnout
- CHRNB3 and CHRNA6
- Results

The Role of Patience

- The traditional calculus of voting does not take account of time.
- We vote today, but the benefits come later.
- Smaller discount factors should lead to higher turnout.
- This applies to expressive voting with equal felicity.

The Role of Patience

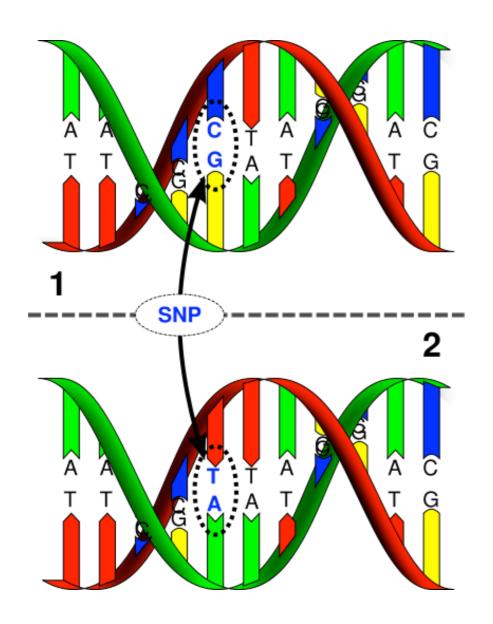


Fowler and Kam. 2006. "Patience as a Political Virtue: Delayed Gratification and Turnout." *Political Behavior.*

CHRNA

- Nicotine acetycholine receptors (nAChRs)
- Implicated in neurotransmission on the sending side.
- Two subunits (in CHRNA6 and CHRNB3) influence the release of dopamine.
- Dopamine has been related to behaviour in delay discounting tasks (Cui et al 2008)
- CHRNA6 and CHRNB3 have been shown to be related to nicotine initiation and dependence (e.g. Lie et al 2005) and alcohol abuse (Hoft et al 2009).
- In turn, these behaviours are related to delay discounting behaviour.
- Note, finally, the central role of dopamine in regulating (social) behaviour.

CHRNA



Four SNPs

CHRNA6: rs2304297 (G) rs892413 (G)

CHRNB3: rs4950 (C) rs13280604 (G)

	0	1	2
SNP1	0.46	0.39	0.15
SNP2	0.51	0.36	0.13
SNP3	0.48	0.38	0.14
SNP4	0.49	0.35	0.16

Data

- Add Health
- Limited genetic sample
- 2574 respondents in the third wave of the study.
- 18-26 years old.
- Surveyed in 2001 or 2002.
- DV: Voting in the 2000 Presidential election.

Results

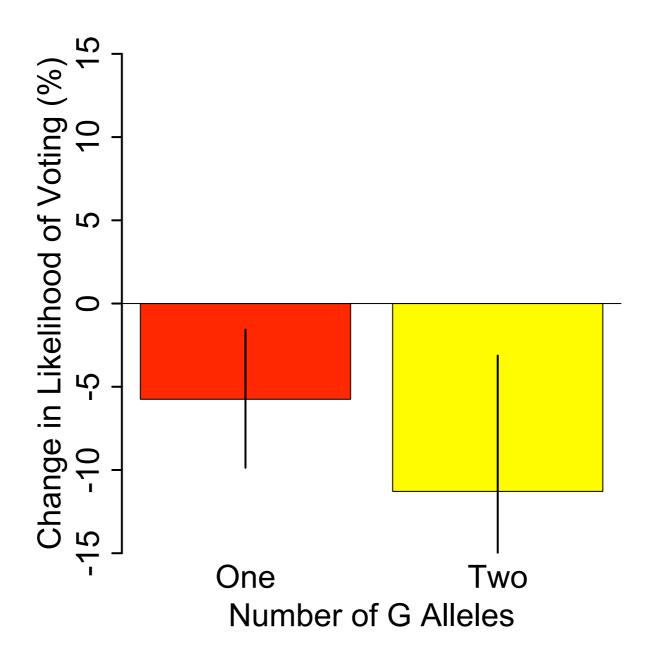
$$logit(P[Y_{ij} = 1 | Z_{ij}, U_j]) = \beta_0 + \beta_X X_{ij} + \beta_Z Z_{ij} + U_j$$

Table 2: Association between CHRNA6 and CHRNB3 SNPs and Turnout

	SN	IP1	S٨	IP2	S٨	IP3	SNP4		SN (whi	
	OR	р	OR	р	OR	р	OR	р	ÓR	p
SNP1: G	0.79	0.01		•		•		•	0.73	0.01
SNP2: C			0.84	0.04						
SNP3: C					0.88	0.11	0.89	0.15		
SNP4: G	1.13	0.00	1.12	0.00	1.11	0.00	1.13	0.15		
Age	1.07	0.51	1.03	0.78	1.06	0.55	1.08	0.46	1.10	0.01
Male	1.07	0.51	1.03	0.78	1.06	0.55	1.08	0.46	1.20	0.16
Black	2.06	0.00	1.86	0.00	1.66	0.00	1.77	0.00		
Asian	0.73	0.16	0.82	0.37	0.86	0.48	0.78	0.27		
Native	0.87	0.71	0.84	0.64	0.78	0.49	0.90	0.77		
American										
Hispanic	0.83	0.30	0.81	0.22	0.78	0.15	0.80	0.19		
N	21	17	21	93	21	75	20	68	13	40
LL	-14	25	-14	78	-14	72	-13	99	-9	04
LL	-14	43	-14	193	-14	84	-14	15	-9	11
(constant)										

Table 2: Models of association between CHRNA6 and CHRNB3 SNPs and voter turnout. All results are expressed in odds ratios (OR). The model is a mixed-effects logit which estimates a random intercept for each family (not shown).

Results



Replication

- 400 adults, principally in California
- Imputed SNP, where the measure is the "dosage" of CHRNA6.
- Control for age, gender, and ethnicity.

Replication results

Table 4: Model of association between CHRNA6 and voting in replication sample

	S	SNP1		
	OR	p value		
rs2304297	0.54	0.00		
MDS	0.01	0.38		
Age	1.08	0.00		
Male	0.61	0.05		
N	400			
LL	-661.57			
LL(constant)	-725.41			

Table 4: Model of association between CHRNA6 and voting in replication sample. The model is a mixed-effects ordered logit regression which estimates a random intercept for each family (not shown).

Conclusions

- Voting may be an act of the patient.
- We can link genes to political behaviour through behavioural mediators/mechanisms.
- Next steps include risk, trust, inequality aversion, and strategic inference.
- Genes are the institutions of the human body.

Thanks



Chris Dawes



James Fowler



Jaime Settle

Results

	Retrospective Logit		
	OR	P value	
Vote	0.62	0.03	
Age	1.07	0.34	
Male	0.85	0.48	
Intercept	0.23	0.33	
\overline{N}		328	
LL		441	
LL(constant)		446	

Table 3: Family-based test of association between CHRNA6 and turnout. The model is a mixed-effects logit which estimates a random intercept for each family (not shown). The dichotomous dependent variable is whether or not the kth allele (k = 1, 2) is an G allele [G allele = 1, G allele = 0]. Odds ratios (OR) are presented. Detailed variable descriptions are provided in the appendix.