

The genetic lottery goes to school: gene-environment interactions in the school context

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Motivation

- **Education** is a core determinant of life outcomes (Acemoglu and Autor, 2011; Hanushek and Woessmann, 2008; Krueger and Lindahl, 2001).
- **Equity** of education systems as a central policy goal:

Most fundamental, of course, is the question of how well schools reduce the inequity of birth by providing children an equitable foundation of mental skills and knowledge [...].

Coleman Report, p.36

- Effective education policies require understanding of the **production function**:

$$Y = f(\underbrace{G}_{\text{Nature}}, \underbrace{I^F, I^S}_{\text{Nurture}}).$$

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Why genes? Why now?

1. Genes account for 40% of variation in years of education (Branigan et al., 2013).
2. Genes matter for distributive justice (Koellinger and Harden, 2018).
3. Recent advances in molecular genetics now allow us to study the role of genes for education (Benjamin et al., 2024).

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A research agenda

Research question

Do better schools increase or decrease the effect of genes on educational attainment?

- **Evidence from the US:**

B. Arold, P. Hufe, and M. Stoeckli (in press). “Genetic Endowments, Educational Outcomes and the Moderating Influence of School Quality”. *Journal of Political Economy: Microeconomics*

- **Evidence from Norway:**

N. T. Borgen, R.G. Cheesman, P. Hufe, A.M.J. Sandso (2025). “The Genetic Lottery Goes to School: Better Schools Compensate for the Effects of Students Genetic Differences”. *Proceedings of the National Academy of the Sciences* 122 (43), e2511715122

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A a closely related research agenda (that I will not talk about)

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Outline

Measuring genetic factors

Evidence from the US

Evidence from Norway

Conclusion

Roadmap

Measuring genetic factors

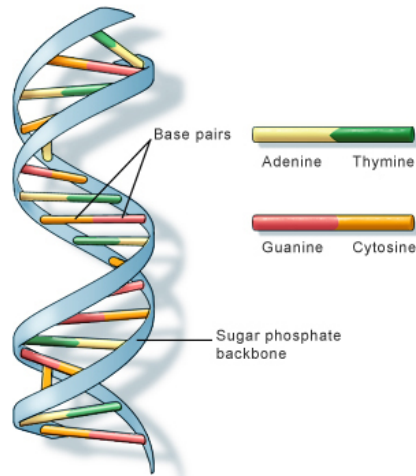
Evidence from the US

Evidence from Norway

Conclusion

Genetics 101

- Human genetic information stored in **23 chromosome pairs**.
- Each chromosome consists of a molecule called **DNA**.
- The “rungs of the ladder” of the DNA are **acid-base pairs**.
- **Genes** are sequences of acid-base pairs that are protein-coding.
 - There are 3.3 bn “rungs in the ladder.”
 - > 99.5% are the same for all human beings.



▶ Minor and major alleles

▶ Meiosis

▶ GWAS and PGI

Identification

- Estimation:

$$Y_i = \alpha PGI_i^{EA} + \beta Q_i + \kappa(PGI_i^{EA} \times Q_i) + \mathbf{X}_i\gamma + \epsilon_i$$

- Identification:

Requirement	Potential bias	Affected parameters	Potential solutions
Exogenous PGI^{EA}	indirect genetic effects	α, κ	genetic trios sibling design adoption design ...
Exogenous Q	selection into schools	β, κ	admission lotteries border discontinuities value-added estimates ...
Independent PGI^{EA}, Q	gene-environment correlation	κ	–

Roadmap

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Genetic Endowments, Educational Outcomes and the Moderating Influence of School Quality

B. Arold, P. Hufe, and M. Stoeckli

Journal of Political Economy: Microeconomics

National Longitudinal Study of Adolescent to Adult Health (Add Health)

- Initial information for a sample of adolescents ($N = 20,745$) collected in 1994/95.
- Nationally **representative** sample for students in grades 7-12.
- Follow up waves in **1996, 2001/02, 2008/09, 2016/18**.
- We restrict the sample to students of **European descent**.

Data inputs

- Recall our **estimation model**:

$$Y_i = \alpha PGI_i^{EA} + \beta Q_i + \kappa(PGI_i^{EA} \times Q_i) + \mathbf{X}_i\gamma + \epsilon_i$$

► Educational outcomes Y_i

► Genetic factors PGI_i^{EA}

► School quality Q

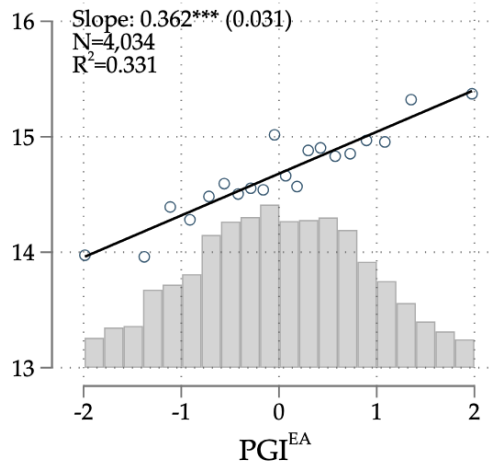
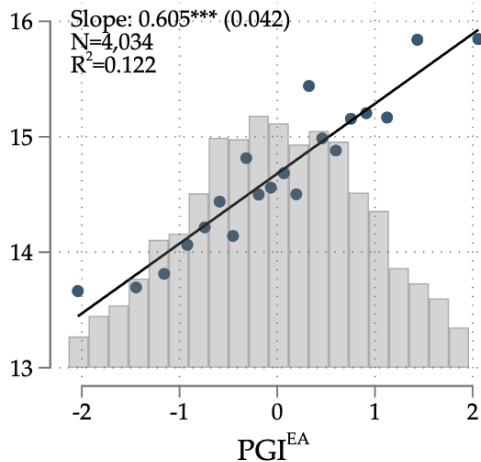
► Control \mathbf{X}_i

	N=4,034; High Schools=72			
	Mean	SD	Min	Max
Educational Attainment				
Years of Education	14.68	2.27	8.00	20.00
High School Degree	0.96	0.20	0.00	1.00
2-year College Degree	0.50	0.50	0.00	1.00
4-year College Degree	0.39	0.49	0.00	1.00
Post-Graduate Degree	0.14	0.35	0.00	1.00
Child and Family Characteristics				
PGI ^{EA}	0.00	1.00	-4.18	3.40
Female	0.54	0.50	0.00	1.00
Firstborn	0.48	0.50	0.00	1.00
Age in Months (Wave 1)	192.41	19.62	144.00	256.00
Maternal Age at Birth	25.33	4.83	16.00	46.08
Christian	0.82	0.38	0.00	1.00
Education Mother (in Years)	13.54	2.48	0.00	19.00
Education Father (in Years)	13.56	2.68	0.00	19.00
School Quality Indicators				
Q	0.00	1.00	-2.79	1.83
Teacher w/ MA (%)	51.20	24.11	0.00	95.00
Experienced Teacher (%)	66.65	23.43	0.00	98.00
New Teacher (%)	7.88	7.28	0.00	47.00
Class Size	24.40	4.50	12.00	38.00

Recap on identifying assumptions

1. No indirect genetic effects (α, κ).
2. No selection into schools (β, κ).
3. Independent variation in PGI^{EA} and Q (κ).

Identifying genetic effects



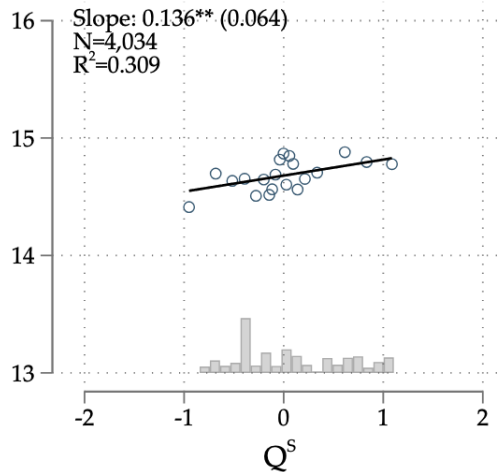
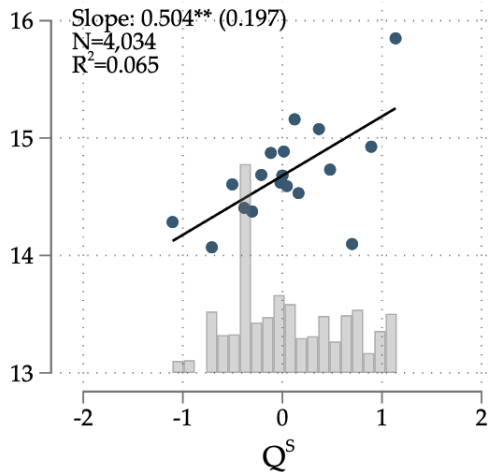
Identifying genetic effects

	Years of Education: Between- vs. Within-Family		Predicted Years of Education: w/o vs. w/ Control Function	
	(1)	(2)	(3)	(4)
PGI ^{EA}	0.415*** (0.085)	0.432*** (0.141)	–	–
Q	–	–	0.264** (0.129)	0.014 (0.050)
Difference in coefficients	-0.017 (0.129) [-0.269, 0.236]		0.250** (0.115) [0.023, 0.476]	
Child Controls	✓	✓	×	×
Family Controls	✓	✓	×	×
Control Function	✓	✓	×	✓
Sibling Fixed Effect	×	✓	×	×
N	677	677	4,034	4,034
R ²	0.420	0.795	0.084	0.184
Outcome Mean	14.722	14.722	14.681	14.681
Outcome SD	2.277	2.277	1.163	1.163

Recap on identifying assumptions

- ✓ No indirect genetic effects (α, κ).
- 2. No selection into schools (β, κ).
- 3. Independent variation in PGI^{EA} and Q (κ).

Identifying school effects



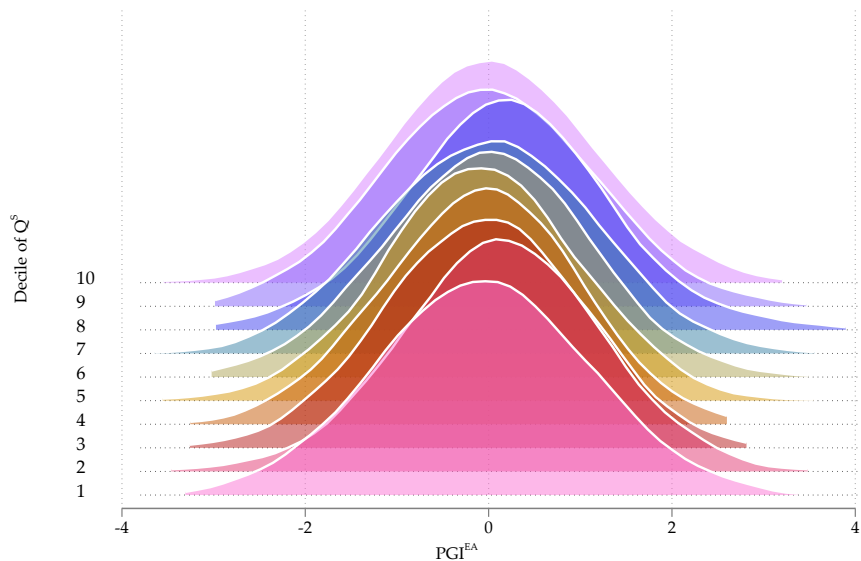
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Recap on identifying assumptions

- ✓ No **genetic nurture** (α, κ).
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- 3. **Independent variation** in PGI^{EA} and Q (κ).

Genes and school investments



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- ✓ **Independent variation** in PGI^{EA} and Q (κ).

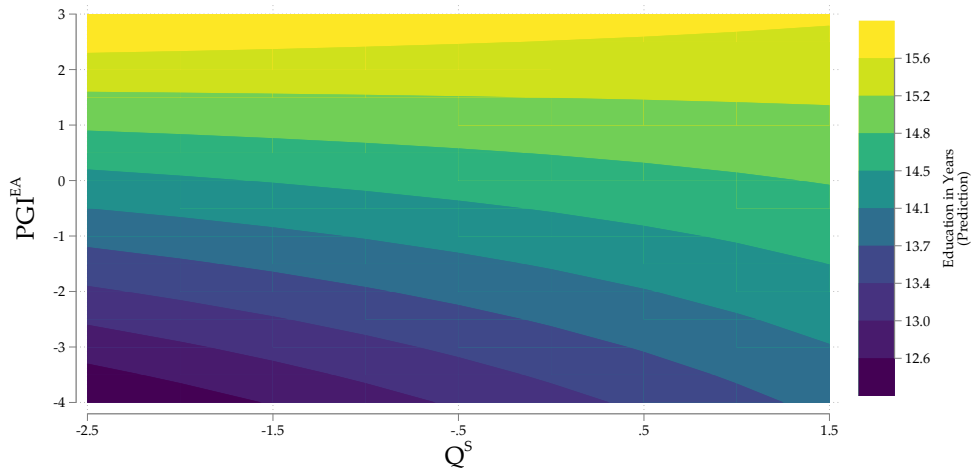
Gene-environment interaction

Outcome: Years of Education	Overall		Decomposition of Q			
	PCA (1)	Anderson (2008) (2)	(3)	(4)	(5)	(6)
PGI^{EA}	0.361*** (0.028)	0.361*** (0.029)	0.360*** (0.027)	0.362*** (0.029)	0.362*** (0.030)	0.362*** (0.030)
Q	0.124** (0.057)	0.098** (0.048)	-	-	-	-
$PGI^{EA} \times Q$	-0.068*** (0.026)	-0.064** (0.028)	-	-	-	-
Teacher w/ MA	-	-	0.166** (0.071)	-	-	-
$PGI^{EA} \times \text{Teacher w/ MA}$	-	-	-0.072*** (0.026)	-	-	-
Exp. Teacher	-	-	-	0.069 (0.059)	-	-
$PGI^{EA} \times \text{Exp. Teacher}$	-	-	-	-0.045* (0.026)	-	-
New Teacher	-	-	-	-	-0.020 (0.047)	-
$PGI^{EA} \times \text{New Teacher}$	-	-	-	-	0.038 (0.029)	-
Class Size	-	-	-	-	-	-0.008 (0.044)
$PGI^{EA} \times \text{Class Size}$	-	-	-	-	-	-0.004 (0.032)
Child Controls	✓	✓	✓	✓	✓	✓
Family Controls	✓	✓	✓	✓	✓	✓
Control Function	✓	✓	✓	✓	✓	✓
N	4,034	4,034	4,034	4,034	4,034	4,034
R ²	0.333	0.333	0.334	0.332	0.332	0.331
Outcome Mean	14.681	14.681	14.681	14.681	14.681	14.681
Outcome SD	2.268	2.268	2.268	2.268	2.268	2.268

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Exp. Teacher	-	-	-	0.069 (0.059)	-	-
$PGI^{EA} \times \text{Exp. Teacher}$	-	-	-	-0.045* (0.026)	-	-
New Teacher	-	-	-	-	-0.020 (0.047)	-
$PGI^{EA} \times \text{New Teacher}$	-	-	-	-	0.038 (0.029)	-
Class Size	-	-	-	-	-	-0.008 (0.044)
$PGI^{EA} \times \text{Class Size}$	-	-	-	-	-	-0.004 (0.032)
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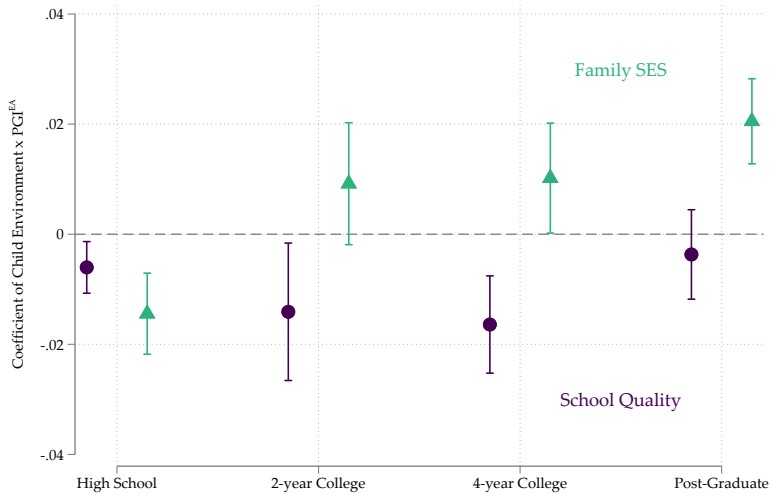
Gene-environment interaction



Robustness

- ✓ Inclusion of other school characteristics and policies, ▶ School characteristics
- ✓ Inclusion of other family/child characteristics, ▶ Family characteristics
- ✓ Inclusion of other PGI, ▶ Other PGI
- ✓ Placebo assignment to schools, ▶ Placebo
- ✓ Exclusion of outlier schools, ▶ Outlier
- ✓ No ceiling effects in educational attainment, ▶ Ceiling effects
- ✓ Sample selection and weighting criteria, ▶ Sample and weighting
- ✓ Correction for measurement error in PGI^{EA}. ▶ Becker et al. (2021)

Educational degrees



Roadmap

Measuring genetic factors

Evidence from the US

Evidence from Norway

Conclusion

The Genetic Lottery Goes to School: Better Schools Compensate for the Effects of Students Genetic Differences

N. T. Borgen, R.G. Cheesman, P. Hufe, A.M.J Sandsor

Proceedings of the National Academy of the Sciences

Data sources

- **MoBa:**
 - Initial information for a sample of mothers ($N > 114,000$) from 1999-2008.
 - 44,017 genotyped father-mother-child trios.
 - Linked to Norwegian register data.
 - We restrict the sample to birth cohorts 2002-2008 and students of European descent.
 - Effective sample size $N \approx 31,000$.
- **Norwegian registers:**
 - Population of students in Norway ($N \approx 60,000$ per cohort).
 - Information on standardized tests in reading and numeracy in grades 5, 8, and 9.
 - We restrict the sample to birth cohorts 1997-2007.
 - Effective sample size $N \approx 670,000$.

Data inputs

- Our **estimation model**:

$$Y_i = \alpha PGI_i^{EA} + \beta Q_i + \kappa(PGI_i^{EA} \times Q_i) + \mathbf{X}_i\gamma + \epsilon_i$$

► Educational outcomes Y_i

► Genetic factors PGI_i^{EA}

► School quality Q

► Controls \mathbf{X}_i

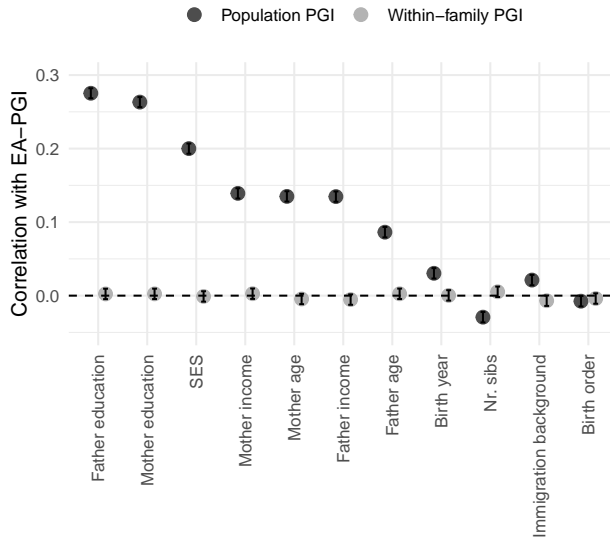
Summary statistics

	Analysis sample <i>N</i> = 30,939				MoBa (All) <i>N</i> = 56,533		Population <i>N</i> = 331,591	
	Mean	St. Dev.	Min	Max	Mean	St. Dev.	Mean	St. Dev.
Birth year	2004.9	1.6	2002	2008	2004.8	1.6	2004.5	1.7
Female	0.5	0.5	0.0	1.0	0.5	0.5	0.5	0.5
Migration background	0.1	0.3	0.0	1.0	0.1	0.3	0.2	0.4
Education (Father)	14.6	2.6	7.0	21.0	14.4	2.7	13.7	2.9
Education (Mother)	15.1	2.3	9.0	21.0	15.0	2.4	14.1	2.9
Inc. rank (Father)	58.5	25.6	0.0	99.0	57.1	26.2	50.9	28.3
Inc. rank (Mother)	61.0	25.4	0.0	99.0	59.9	25.7	51.5	27.6
Age (Father)	32.9	5.1	18.0	65.0	33.1	5.3	33.2	6.0
Age (Mother)	30.5	4.4	16.0	47.0	30.6	4.5	30.2	5.1
Reading (Grade 8)	0.3	0.9	-3.2	2.4	0.2	0.9	0.1	1.0
Numeracy (Grade 8)	0.3	0.9	-2.5	2.5	0.2	1.0	0.0	1.0
English (Grade 8)	0.2	1.0	-2.4	2.2	0.1	1.0	0.0	1.0

Recap on identifying assumptions

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3. Independent variation in PGI^{EA} and Q (κ).

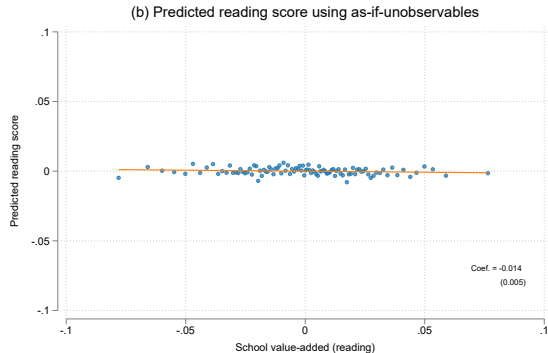
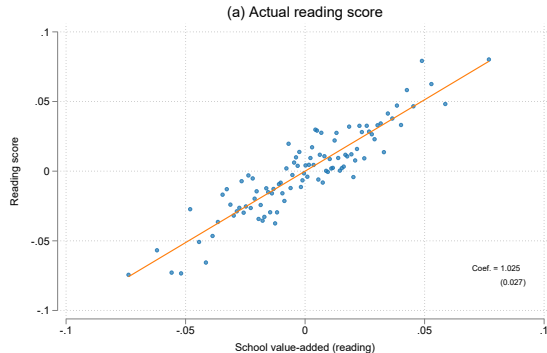
Identification of genetic effects



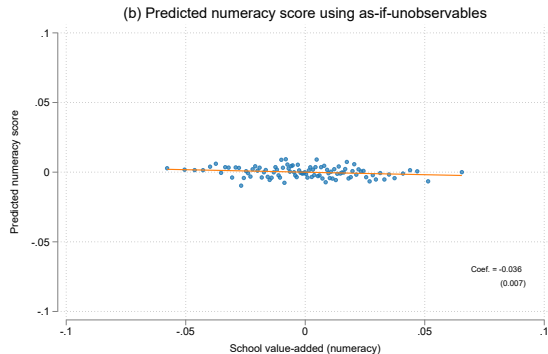
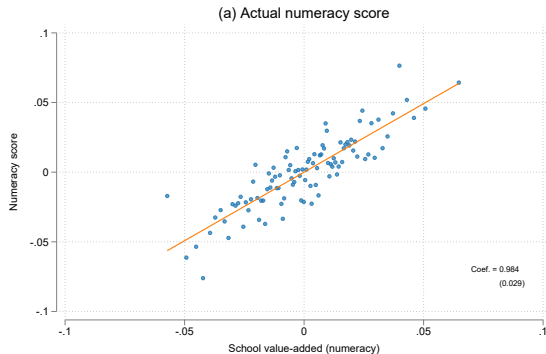
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Identification of school effects (Reading)



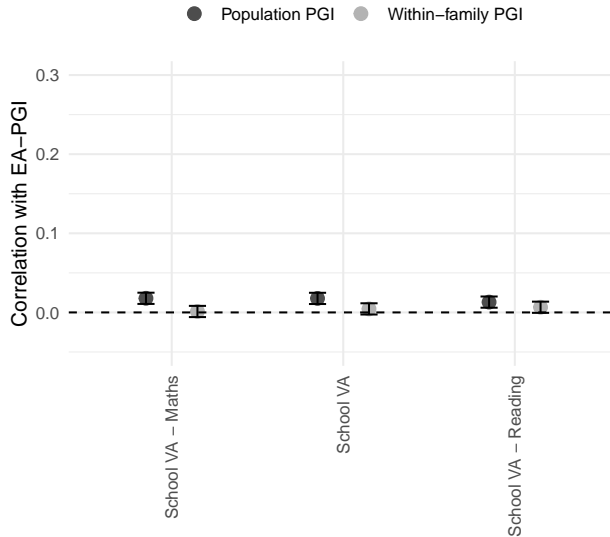
Identification of school effects (Numeracy)



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Gene-environment correlation



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Gene-environment interaction (Reading)

Outcome: Reading (Grade 9)	(1)	(2)	(3)	(4)
PGI^{EA}	0.304*** (0.006)	0.230*** (0.008)	0.231*** (0.005)	0.231*** (0.005)
$VA^{Reading}$	0.091*** (0.014)	0.090*** (0.013)	0.052*** (0.007)	0.050*** (0.007)
$PGI^{EA} \times VA^{Reading}$	-0.020* (0.008)	-0.020* (0.008)	-0.013* (0.005)	-0.013 (0.007)
Genetic controls	×	✓	✓	✓
School quality controls	×	×	✓	✓
2-way interactions (PGI^{EA} , $Q \times X$)	×	×	×	✓
R^2	0.096	0.104	0.654	0.657
N	30,939	30,939	30,939	30,939
Skill persistence ρ	–	–	0.462*** (0.006)	0.460*** (0.006)

Note: Own calculations. Standard errors (in parentheses) are clustered at the school level. Significance levels: * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$.

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PGI^{EA}	0.304*** (0.006)	0.230*** (0.008)	0.231*** (0.005)	0.231*** (0.005)
$VA^{Reading}$	0.091*** (0.014)	0.090*** (0.013)	0.052*** (0.007)	0.050*** (0.007)
$PGI^{EA} \times VA^{Reading}$	-0.020* (0.008)	-0.020* (0.008)	-0.013* (0.005)	-0.013 (0.007)
Genetic controls	×	✓	✓	✓
School quality controls	×	×	✓	✓
2-way interactions (PGI^{EA} , $Q \times X$)	×	×	×	✓
R^2	0.096	0.104	0.654	0.657
N	30,939	30,939	30,939	30,939
Skill persistence ρ	–	–	0.462*** (0.006)	0.460*** (0.006)

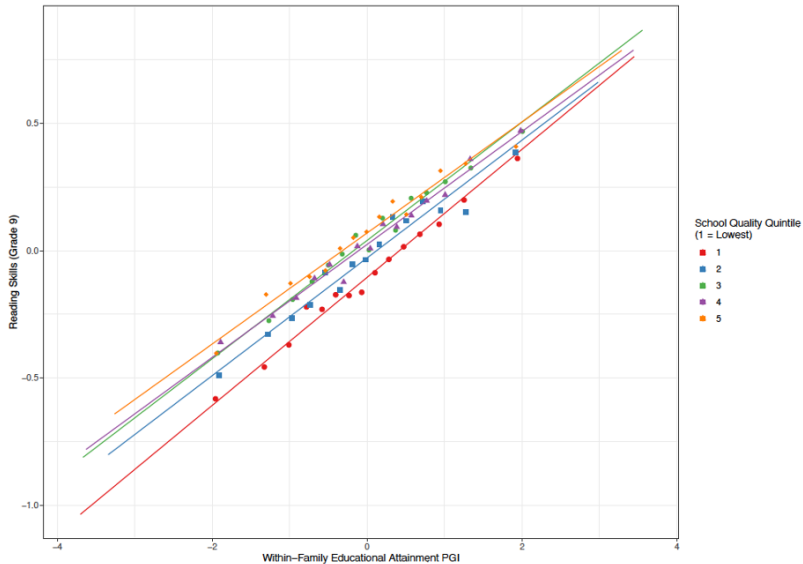
Note: Own calculations. Standard errors (in parentheses) are clustered at the school level. Significance levels: * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$.

Gene-environment interaction (Reading)

Outcome: Reading (Grade 9)	(1)	(2)	(3)	(4)
PGI^{EA}	0.304*** (0.006)	0.230*** (0.008)	0.231*** (0.005)	0.231*** (0.005)
$VA^{Reading}$	0.091*** (0.014)	0.090*** (0.013)	0.052*** (0.007)	0.050*** (0.007)
$PGI^{EA} \times VA^{Reading}$	-0.020* (0.008)	-0.020* (0.008)	-0.013* (0.005)	-0.013 (0.007)
Genetic controls	×	✓	✓	✓
School quality controls	×	×	✓	✓
2-way interactions (PGI^{EA} , Q_X)	×	×	×	✓
R^2	0.096	0.104	0.654	0.657
N	30,939	30,939	30,939	30,939
Skill persistence ρ	–	–	0.462*** (0.006)	0.460*** (0.006)

Note: Own calculations. Standard errors (in parentheses) are clustered at the school level. Significance levels: * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$.

Gene-environment interaction (Reading)

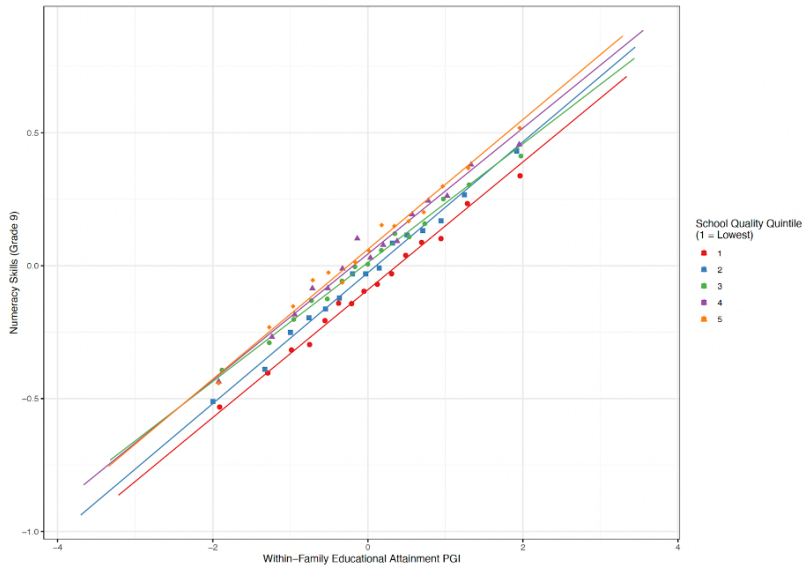


Gene-environment interaction (Numeracy)

Outcome: Numeracy (Grade 9)	(1)	(2)	(3)	(4)
PGI ^{EA}	0.314*** (0.006)	0.238*** (0.008)	0.239*** (0.004)	0.239*** (0.004)
VA ^{Numeracy}	0.076*** (0.013)	0.075*** (0.013)	0.039*** (0.005)	0.040*** (0.005)
PGI ^{EA} × VA ^{Numeracy}	-0.005 (0.007)	-0.006 (0.007)	-0.000 (0.004)	0.001 (0.005)
Genetic controls	×	✓	✓	✓
School quality controls	×	×	✓	✓
2-way interactions (PGI ^{EA} , Q, X)	×	×	×	✓
R ²	0.102	0.109	0.738	0.740
N	30,939	30,939	30,939	30,939
Skill persistence p	–	–	0.702*** (0.004)	0.703*** (0.004)

Note: Own calculations. Standard errors (in parentheses) are clustered at the school level. Significance levels: * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$.

Gene-environment interaction (Numeracy)



Contextualizing effect sizes

- Estimates pertain to a **low inequality country**. [▶ Inequality in VA](#)
 - Assuming cross-country portability of effects, substitutability would be 10% for grade 9 in Chicago high schools.
- Estimates pertain to **one year of schooling**.
 - Assuming linear additive effects, substitutability increases to 18% over the course of lower secondary school (grades 8-10) in Norway.
- Estimates can be compared to substitutability in **other dimensions of advantage**:
 - Latent family SES ($\Delta 1SD$): 2.87% (Jackson et al., 2024).

Roadmap

Measuring genetic factors

Evidence from the US

Evidence from Norway

Conclusion

Summary

- **What we do:**

- We study the (causal) interplay between PGL^{EA} and school quality.

- **What we find:**

- Students with lower PGIs benefit more from higher-quality schools.

- **Why it matters:**

- Investments in schools may help to students to (partially) overcome their draw in the genetic lottery and to reduce unequal opportunities in society.

Open questions

- **Generalizability:**

- Countries
- Learning domains
- Age groups
- ...

- **Mechanisms:**

- Features of good schools?
- Family responses as mediators?
- ...

Open questions

- **Generalizability:**

- Countries
- Learning domains
- Age groups
- ...

- **Mechanisms:**

- Features of good schools?
- Family responses as mediators?
- ...

Thank you for your attention! Questions?

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References I

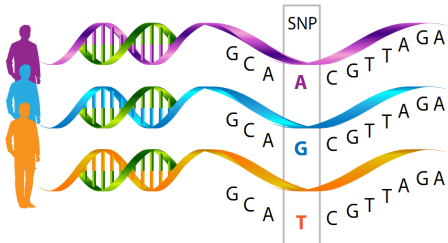
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Minor and major alleles

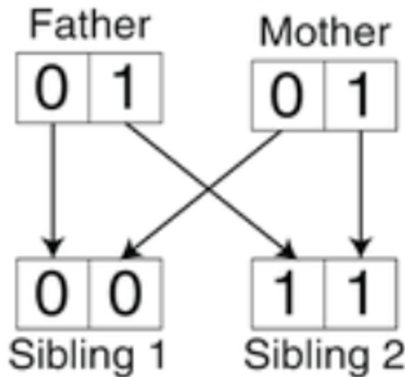
- Differences in base pairs across humans: **single-nucleotide polymorphism (SNP)**.
- Some of “rungs in the ladder” are more common than others. They are called **minor (major) alleles**.
- At each location individuals can have 0, 1, 2 minor (major) alleles.



Meiosis

- Parents pass one randomly selected allele to their offspring.
 - **Recombination**: Parental chromosome pairs cross a random number of times at random loci.
 - **Mendelian segregation**: For each parent, one of the recombined chromosome pairs is randomly transmitted to the germ cell.

- Siblings can end up having different alleles from both parents at a SNP.
- Conditional on the parental genotypes, offspring alleles at a SNP are random.



- Genetic discovery based on genome-wide association studies (**GWAS**):

$$y_i = \text{SNP}_{ij}\beta_j + \sum_1^n C_i^n \delta_j^n + u_i.$$

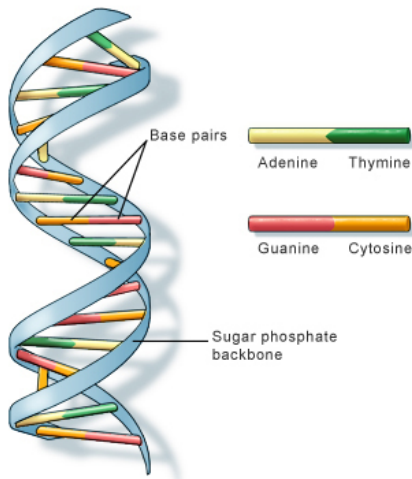
- Summary statistics via polygenic scores (**PGI**):

$$\text{PGI}_i = \sum_j \text{SNP}_{ij} \hat{\beta}_j.$$

Educational outcomes Y_i



- We use the **polygenic index (PGI)** for educational attainment from Lee et al. (2018):
 - Discovery sample of 1.1 mn people of European descent.
 - Explains 11% of variation in years of education.



School quality Q

- We use **survey and administrative information** on teachers:
 - Student-teacher ratio
 - Teacher w/ tenure < 1 year
 - Teacher w/ tenure > 5 years
 - Teacher w/ Master degree
- We aggregate information using PCA or through linear aggregation of standardized variables (Anderson, 2008; Kling et al., 2007).

- **Pre-determined characteristics** and **control function** (Altonji and Mansfield, 2018):

Family

- maternal age at birth
- years of education (m/f)
- non-US born (m/f)
- av. potential wage (m/f)
- SD potential wage (m/f)
- religion
- state FE

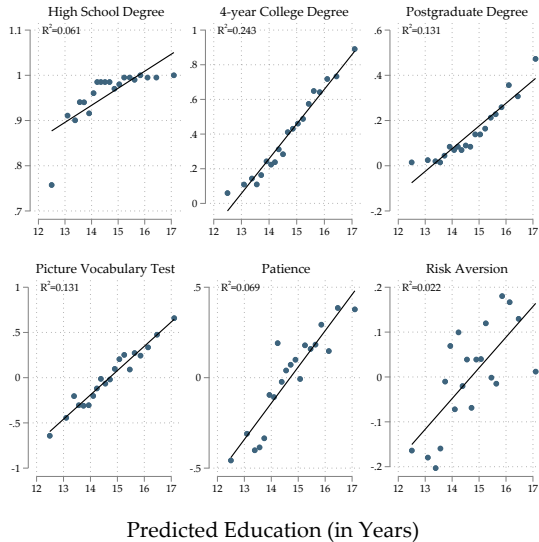
Child

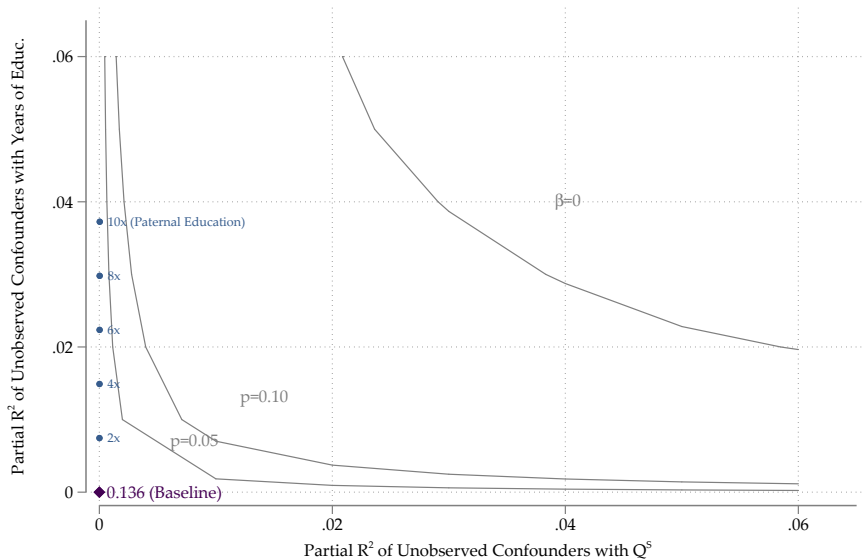
- firstborn
- gender x age in months
- 20 PC of full matrix of genetics data

Control function

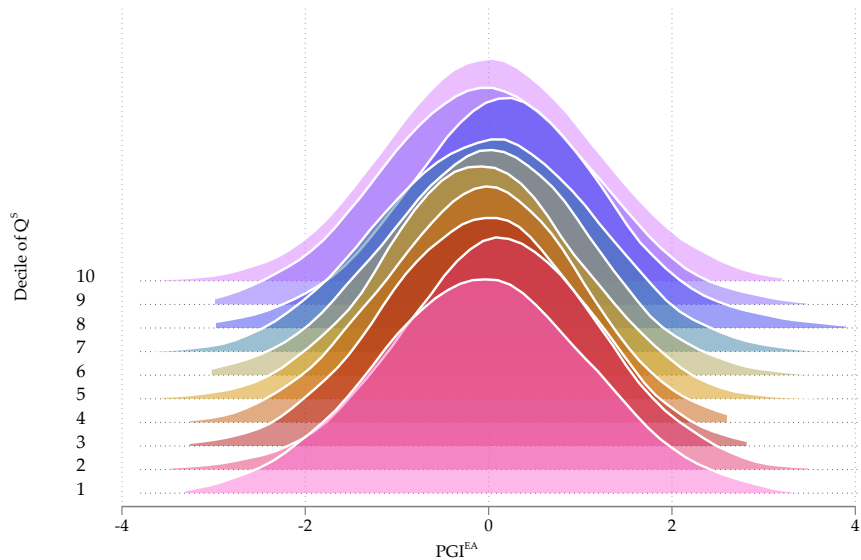
- white peers (%)
- single mothers (%)
- education mothers (av.)
- female peers (%)
- migrant peers (%)

Predictive power





Residualized distributions



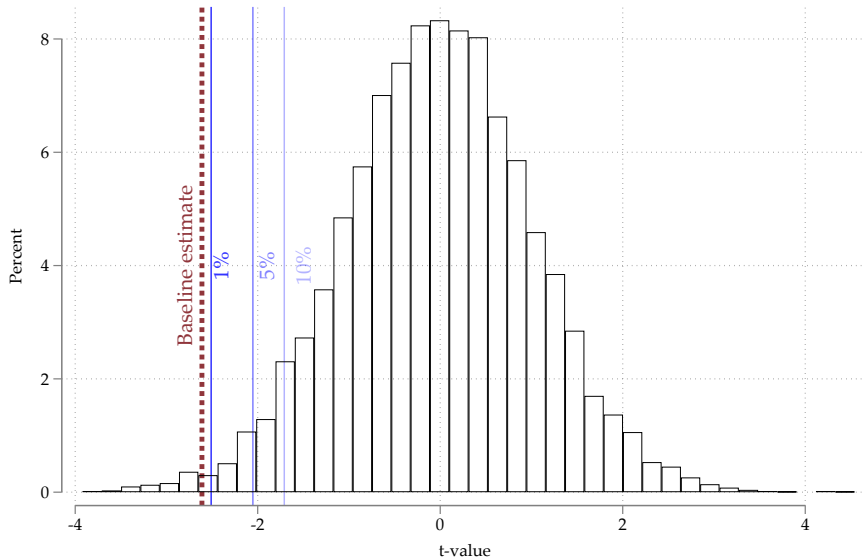
School characteristics

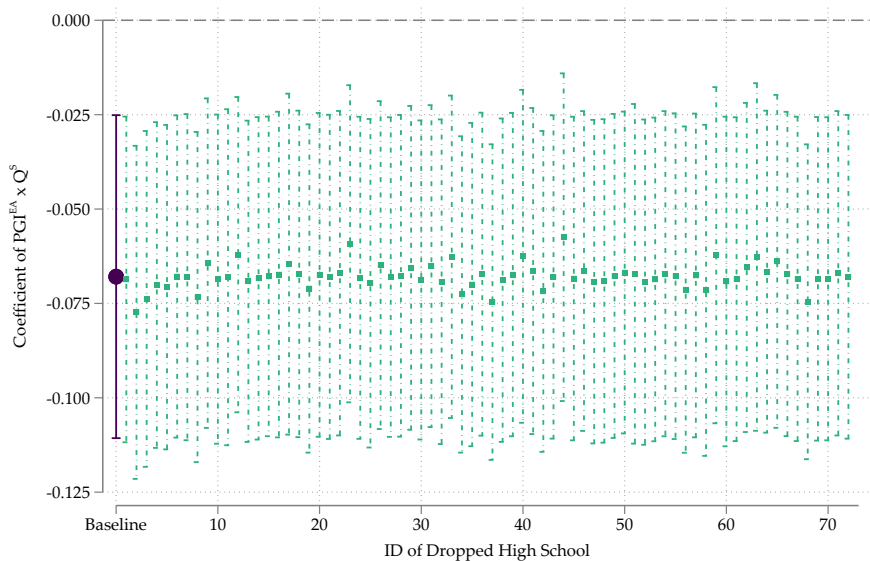
	Baseline	+ School Policies			+ Private School	+ Teacher Composition		+ School FE
Outcome: Years of Education	(1)	Retention Policy (2)	Ability Groups (3)	Strict. Index (4)	(5)	White Teacher (6)	Female Teacher (7)	(8)
PGI^{EA}	0.361*** (0.028)	0.366*** (0.029)	0.361*** (0.028)	0.362*** (0.029)	0.362*** (0.029)	0.361*** (0.028)	0.360*** (0.028)	0.350*** (0.029)
Q	0.124** (0.057)	0.116** (0.053)	0.127** (0.058)	0.135** (0.060)	0.144** (0.056)	0.121** (0.055)	0.135** (0.062)	-
$PGI^{EA} \times Q$	-0.068*** (0.026)	-0.066*** (0.025)	-0.068*** (0.025)	-0.064** (0.027)	-0.076*** (0.026)	-0.064** (0.026)	-0.065** (0.026)	-0.064** (0.027)
School Characteristic	-	-0.103* (0.060)	0.049 (0.034)	0.062* (0.036)	0.101** (0.043)	-0.013 (0.074)	-0.022 (0.048)	-
$PGI^{EA} \times$ School Characteristic	-	0.034 (0.030)	-0.016 (0.030)	0.019 (0.024)	-0.049** (0.023)	-0.029 (0.033)	0.045 (0.034)	-
Child Controls	✓	✓	✓	✓	✓	✓	✓	✓
Family Controls	✓	✓	✓	✓	✓	✓	✓	✓
Control Function	✓	✓	✓	✓	✓	✓	✓	✓
N	4,034	3,969	4,034	4,034	4,034	4,034	4,034	4,034
R ²	0.333	0.334	0.333	0.334	0.334	0.333	0.334	0.343

Family characteristics

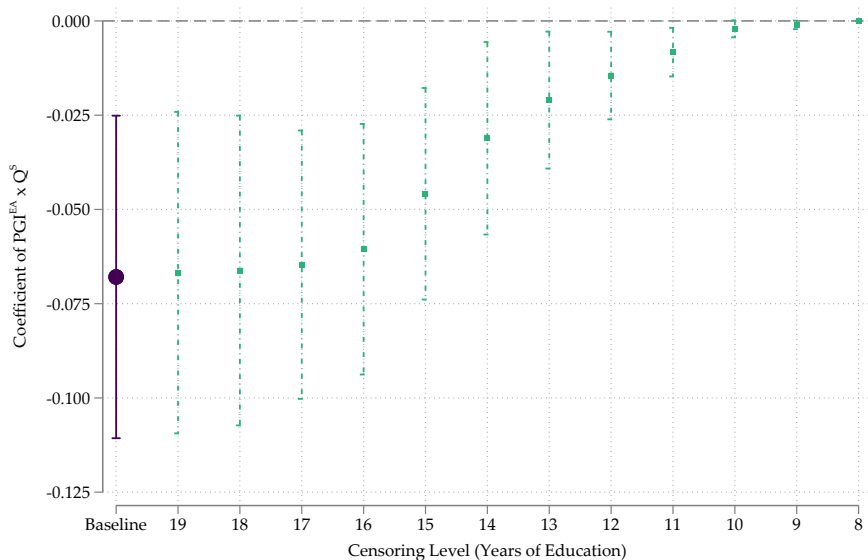
	Baseline	Interacted controls	Non-linearities	Subsample w/ lagged ability measures		
Outcome: Years of Education	(1)	(2)	(3)	(4)	(5)	(6)
PGI ^{EA}	0.361*** (0.028)	0.353*** (0.029)	0.389*** (0.036)	0.333*** (0.050)	0.334*** (0.048)	0.337*** (0.048)
Q	0.124** (0.057)	0.073 (0.059)	0.139** (0.066)	0.252* (0.144)	0.254* (0.132)	0.255* (0.133)
PGI ^{EA} × Q	-0.068*** (0.026)	-0.072** (0.030)	-0.075*** (0.026)	-0.088* (0.047)	-0.086** (0.042)	-0.089** (0.042)
PVT	-	-	-	-	0.223*** (0.047)	0.225*** (0.047)
PVT × Q	-	-	-	-	-	-0.003 (0.045)
GPA Science	-	-	-	-	0.381*** (0.060)	0.383*** (0.060)
GPA Science × Q	-	-	-	-	-	0.069 (0.059)
GPA Math	-	-	-	-	0.283*** (0.075)	0.281*** (0.074)
GPA Math × Q	-	-	-	-	-	-0.004 (0.080)
Child Controls	✓	✓	✓	✓	✓	✓
Family Controls	✓	✓	✓	✓	✓	✓
Control Function	✓	✓	✓	✓	✓	✓
All interactions (Q, PGI ^{EA} , X)	×	✓	×	×	×	×
2 nd Polynomial (Q, PGI ^{EA})	×	×	✓	×	×	×
N	4,034	4,034	4,034	1,039	1,039	1,039
R ²	0.333	0.345	0.334	0.437	0.510	0.511
Outcome Mean	14.681	14.681	14.681	14.520	14.520	14.520
Outcome SD	2.268	2.268	2.268	2.309	2.309	2.309

Outcome: Years of Education	Baseline	+ Controls for Other Polygenic Indexes					
	(1)	Body Mass Index (2)	ADHD (3)	Depressive Symptoms (4)	Intelligence (5)	Ever Smoker (6)	Sleep Duration (7)
PGI ^{EA}	0.361*** (0.028)	0.341*** (0.031)	0.330*** (0.028)	0.358*** (0.028)	0.349*** (0.031)	0.341*** (0.031)	0.360*** (0.028)
Q	0.124** (0.057)	0.121** (0.056)	0.120** (0.056)	0.120** (0.057)	0.124** (0.057)	0.122** (0.056)	0.124** (0.057)
PGI ^{EA} × Q	-0.068*** (0.026)	-0.076*** (0.029)	-0.071*** (0.026)	-0.065** (0.027)	-0.059** (0.028)	-0.067** (0.027)	-0.068*** (0.026)
Other PGI	–	-0.080*** (0.026)	-0.132*** (0.028)	-0.039 (0.030)	0.023 (0.030)	-0.097*** (0.036)	0.026 (0.028)
Other PGI × Q	–	-0.029 (0.028)	0.003 (0.028)	0.035 (0.029)	-0.018 (0.028)	0.017 (0.033)	-0.003 (0.029)
Child Controls	✓	✓	✓	✓	✓	✓	✓
Family Controls	✓	✓	✓	✓	✓	✓	✓
Control Function	✓	✓	✓	✓	✓	✓	✓
N	4,034	4,034	4,034	4,034	4,034	4,034	4,034
R ²	0.333	0.334	0.336	0.334	0.333	0.335	0.333





Ceiling effects



Sample and weighting

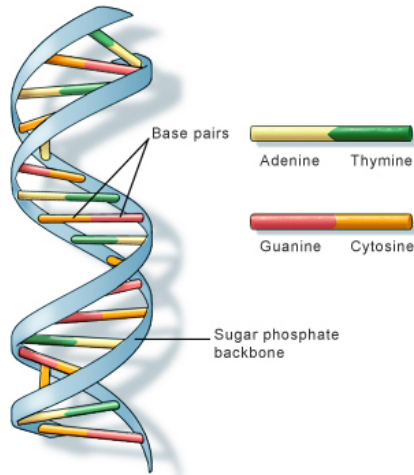
Outcome: Years of Education	Baseline	Alternative Sample Composition		
	(1)	Re-Weighted (2)	Excl. (Potential) Movers before High School (3)	Excl. (Potential) Movers during High School (4)
PGI ^{EA}	0.361*** (0.028)	0.347*** (0.031)	0.353*** (0.035)	0.343*** (0.040)
Q	0.124** (0.057)	0.115* (0.060)	0.159** (0.063)	0.101 (0.078)
PGI ^{EA} × Q	-0.068*** (0.026)	-0.061** (0.027)	-0.086*** (0.032)	-0.086** (0.038)
Child Controls	✓	✓	✓	✓
Family Controls	✓	✓	✓	✓
Control Function	✓	✓	✓	✓
N	4,034	3,968	2,962	2,439
R ²	0.333	0.313	0.350	0.344

	Coefficient	Standard Error	p-value	Substitutability
Baseline				
PGI^{EA}	0.361	0.032	0.000	
Q	0.124	0.064	0.052	
$PGI^{EA} \times Q$	-0.068	0.030	0.023	19%
Add Health ($\rho = 1.968$)				
PGI^{EA}	0.747	0.074	0.000	
Q	0.097	0.069	0.160	
$PGI^{EA} \times Q$	-0.108	0.061	0.075	15%
Health and Retirement Study ($\rho = 1.413$)				
PGI^{EA}	0.566	0.051	0.000	
Q	0.111	0.064	0.084	
$PGI^{EA} \times Q$	-0.093	0.043	0.030	16%
Wisconsin Longitudinal Study ($\rho = 1.649$)				
PGI^{EA}	0.718	0.068	0.000	
Q	0.099	0.068	0.145	
$PGI^{EA} \times Q$	-0.106	0.060	0.076	15%
UK Biobank ($\rho = 1.452$)				
PGI^{EA}	0.589	0.053	0.000	
Q	0.109	0.064	0.088	
$PGI^{EA} \times Q$	-0.096	0.044	0.030	16%

Standardized national tests in reading and numeracy (grade 9)

- **Low stakes**
 - Communicated to parents and teachers but mostly used to track student development.
- **Computer corrected**
 - Not affected by teacher biases.
- **Taken at beginning of the school year**
 - Measure skills accumulated until grade 9.
- **Same test as in grade 8**
 - Allow mapping for VA calculation.
- **Highly predictive of later life-outcomes**
 - 1 SD \uparrow in numeracy, increases high school graduation at age 21 by 9.5 p.p.

- We use the **polygenic index (PGI)** for educational attainment from Okbay et al. (2022):
 - Discovery sample of 3 mn people of European descent.
 - Explains 16% of variation in years of education.
 - \approx 56% of explanatory power due to direct genetic effects.



School quality Q

1. We construct **school VA for reading and numeracy** in grade 8 (Angrist et al., 2023).
2. We model educational outcomes Y of student i attending school j in cohort c for subject d :

$$Y_{ijc}^d = \beta^d Z_{ijc} + \underbrace{VA_{jc}^d + \epsilon_{ijc}^d}_{=e_{ijc}^d}$$

3. We estimate VA in subject d by averaging over residuals in school-cohort cells:

$$VA_{jc}^d = \sum e_{ijc}^d / N_{jc}$$

4. We apply the **Bayesian Shrinkage estimator** à la Chetty et al. (2014).
5. **Highly predictive of later life-outcomes**
 - 1 SD ↑ in VA, increases years of schooling by 0.5-0.8 years (Kirkebøen, 2022).

Child controls

- Lagged test scores in numeracy, reading, English
- Parental years of education
- Migration status
- Age of arrival in Norway
- # of siblings
- Gender
- Year of birth
- Birth order

School controls

- School-cohort averages of all child background variables

Parental PGI

- PGI^{EA} mother
- PGI^{EA} father

Genotyping controls

- Genotyping center
- Genotyping batch
- Genotyping plate
- Imputation batch

Saturation controls

- Interaction of child background controls, school controls, and parental PGIs with PGI^{EA} and Q

Inequality in VA

