## Supplemental Material

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Table S1: results from training education polygenic scores in 50,000 non-adopted individuals in the UK Biobank. Note: this p-value  $(10^{-50})$  held for all of the standard set-values tested in PRSice (0.001, 0.05, 0.1, 0.2, 0.3, 0.4, 0.5, 1).

Threshold	PRS.R <sup>2</sup>	Full.R <sup>2</sup>	Null.R <sup>2</sup>	Coefficient	SE	Р	Num_SNP
1	0.071066	0.11043	0.039365	196340	3110.32	p < 10 <sup>-50</sup>	87372

Table S2: full polygenic prediction results.  $PRS.SE = standard error around the R^2 value, derived from bootstrapping with 1000 replications. Note that the samples comprised 6500 non-adopted individuals and 6311 adoptees.$ 

Group	PRS.R <sup>2</sup>	PRS.SE	Coefficient	SE	Р	N-SNPs	Empirical P
Adopted	0.037	0.0044	27581	1759.93	2.47x10 <sup>-54</sup>	87372	9.99x10 <sup>-05</sup>
Non-adopted	0.074	0.0063	38359.5	1681.36	6.78x10 <sup>-111</sup>	87372	9.99x10 <sup>-05</sup>

Table S3: full results from GxE (education polygenic score x adoption status) model. The outcome variable was years of education. All quantitative covariates were standardised. The table shows results for a linear model testing for additive interaction. The interaction term of interest is highlighted in yellow.

	Estimate	SE	t	Pr(> t )
(Intercept)	13.42	0.4	33.91	<2x10 <sup>-16</sup>
PRS_std	1.2	0.28	4.27	2x10 <sup>-05</sup>
adopted	0.12	0.53	0.23	0.82
Sex.0.0	0.63	0.12	5.24	1.66x10 <sup>-07</sup>
Age.at.recruitment.0.0_scaled	-0.8	0.06	-12.8	<2x10 <sup>-16</sup>
PC1_scaled	0.19	0.08	2.52	0.01
PC2_scaled	-0.07	0.07	-0.95	0.34
PC3_scaled	0.11	0.07	1.74	0.08
PC4_scaled	4.1x10 <sup>-03</sup>	0.06	0.07	0.95
PC5_scaled	-0.03	0.07	-0.5	0.62
PC6_scaled	-0.13	0.07	-1.92	0.06
PC7_scaled	-0.07	0.07	-1.1	0.27
PC8_scaled	-0.05	0.06	-0.87	0.38
PC9_scaled	0.12	0.08	1.53	0.13
PC10_scaled	-0.22	0.07	-3.15	1.61x10 <sup>-03</sup>
Centre11002	0.69	0.52	1.31	0.19
Centre11003	0.11	0.52	0.21	0.83
Centre11004	-0.27	0.51	-0.53	0.59
Centre11005	1.15	0.52	2.2	0.03
Centre11006	0.03	0.5	0.07	0.95
Centre11007	1.21	0.47	2.6	0.01
Centre11008	-0.14	0.46	-0.3	0.76
Centre11009	-0.56	0.47	-1.2	0.23
Centre11010	0.17	0.44	0.39	0.7
Centre11011	0.2	0.45	0.44	0.66
Centre11012	1.54	0.57	2.69	0.01

Centre11013	0.36	0.46	0.78	0.43
Centre11014	-0.22	0.46	-0.49	0.63
Centre11016	-0.02	0.45	-0.05	0.96
Centre11017	-0.09	0.48	-0.18	0.86
Centre11018	1.07	0.48	2.22	0.03
Centre11020	0.64	0.48	1.33	0.18
Centre11021	0.48	0.47	1.01	0.31
Centre11022	-0.62	0.95	-0.65	0.52
Centre11023	-3.52	1.51	-2.33	0.02
PRS_std:adopted	-0.33	<mark>0.09</mark>	<mark>-3.65</mark>	2.66x10 <sup>-04</sup>
PRS_std:Sex.0.0	-0.06	0.09	-0.66	0.51
PRS_std:Age.at.recruitment.0.0_scaled	-0.03	0.04	-0.64	0.52
PRS_std:PC1_scaled	-0.04	0.05	-0.93	0.35
PRS_std:PC2_scaled	-0.03	0.05	-0.69	0.49
PRS_std:PC3_scaled	-0.1	0.04	-2.21	0.03
PRS_std:PC4_scaled	-0.05	0.04	-1.08	0.28
PRS_std:PC5_scaled	-0.01	0.05	-0.25	0.80
PRS_std:PC6_scaled	0.03	0.05	0.72	0.47
PRS_std:PC7_scaled	0.02	0.05	0.38	0.70
PRS_std:PC8_scaled	0.02	0.04	0.37	0.71
PRS_std:PC9_scaled	1.7x10 <sup>-04</sup>	0.05	4x10 <sup>-03</sup>	1.00
PRS_std:PC10_scaled	3.6x10 <sup>-03</sup>	0.05	0.08	0.94
PRS_std:Centre11002	0.37	0.36	1.03	0.30
PRS_std:Centre11003	0.46	0.36	1.28	0.20
PRS_std:Centre11004	-0.27	0.37	-0.74	0.46
PRS_std:Centre11005	0.76	0.37	2.09	0.04
PRS_std:Centre11006	0.25	0.35	0.73	0.47
PRS_std:Centre11007	0.25	0.33	0.75	0.45
PRS_std:Centre11008	0.01	0.33	0.04	0.97

PRS_std:Centre11009	0.14	0.33	0.43	0.67
PRS_std:Centre11010	-0.14	0.31	-0.45	0.65
PRS_std:Centre11011	0.08	0.31	0.26	0.8
PRS_std:Centre11012	0.45	0.39	1.16	0.25
PRS_std:Centre11013	0.03	0.32	0.11	0.92
PRS_std:Centre11014	0.19	0.33	0.59	0.56
PRS_std:Centre11016	0.26	0.32	0.8	0.43
PRS_std:Centre11017	0.11	0.35	0.33	0.74
PRS_std:Centre11018	0.35	0.34	1.03	0.31
PRS_std:Centre11020	0.39	0.34	1.16	0.24
PRS_std:Centre11021	0.21	0.33	0.63	0.53
PRS_std:Centre11022	-0.21	0.62	-0.34	0.73
PRS_std:Centre11023	0.14	1.03	0.14	0.89
adopted:Sex.0.0	-0.27	0.17	-1.58	0.12
adopted:Age.at.recruitment.0.0_scaled	-0.23	0.09	-2.70	0.01
adopted:PC1_scaled	-0.03	0.09	-0.37	0.71
adopted:PC2_scaled	0.04	0.10	0.46	0.64
adopted:PC3_scaled	0.03	0.09	0.36	0.72
adopted:PC4_scaled	-0.07	0.09	-0.85	0.39
adopted:PC5_scaled	0.11	0.09	1.16	0.25
adopted:PC6_scaled	0.05	0.10	0.49	0.62
adopted:PC7_scaled	-0.02	0.10	-0.25	0.80
adopted:PC8_scaled	0.12	0.09	1.40	0.16
adopted:PC9_scaled	-0.15	0.09	-1.60	0.11
adopted:PC10_scaled	0.12	0.09	1.28	0.20
adopted:Centre11002	-0.44	0.72	-0.61	0.54
adopted:Centre11003	-0.61	0.70	-0.87	0.39
adopted:Centre11004	0.10	0.71	0.13	0.89
adopted:Centre11005	-1.12	0.73	-1.52	0.13

adopted:Centre11006	0.07	0.68	0.10	0.92
adopted:Centre11007	-1.42	0.64	-2.23	0.03
adopted:Centre11008	-0.03	0.63	-0.06	0.96
adopted:Centre11009	0.27	0.63	0.43	0.67
adopted:Centre11010	-0.68	0.59	-1.14	0.26
adopted:Centre11011	-0.42	0.60	-0.70	0.48
adopted:Centre11012	-0.73	0.77	-0.95	0.34
adopted:Centre11013	-0.88	0.62	-1.43	0.15
adopted:Centre11014	0.07	0.64	0.12	0.91
adopted:Centre11016	-0.29	0.62	-0.46	0.64
adopted:Centre11017	0.02	0.67	0.03	0.97
adopted:Centre11018	-0.23	0.66	-0.35	0.73
adopted:Centre11020	-0.59	0.66	-0.91	0.36
adopted:Centre11021	-0.53	0.65	-0.82	0.41
adopted:Centre11022	0.96	1.29	0.74	0.46
adopted:Centre11023	1.04	2.15	0.48	0.63

Table S4: Comp	varison of adopted	l and non-adopte	d individuals f	or mean year.	s of education p	er decile of	polygenic
score for educat	tion.						

score for eaucation.								
	Ado	pted		Non-ad	Non-adopted			
Decile	Mean education years	SE	N	Mean education years	SE	N	zdiff	zdiffp
1	-0.26	0.04	723	-0.48	0.04	559	-3.97	7.07x10 <sup>-05</sup>
2	-0.18	0.04	710	-0.27	0.04	571	-1.74	0.081
3	-0.20	0.04	672	-0.27	0.04	609	-1.21	0.225
4	-0.09	0.04	640	-0.11	0.04	641	-0.50	0.619
5	-0.02	0.04	638	-0.12	0.04	643	-1.95	0.051
6	0.02	0.04	623	0.002	0.04	658	-0.37	0.708
7	0.12	0.04	625	0.09	0.04	656	-0.65	0.515
8	0.15	0.04	583	0.21	0.04	698	1.11	0.269
9	0.27	0.04	562	0.26	0.04	719	-0.05	0.962
10	0.36	0.04	535	0.46	0.03	746	1.93	0.053

Figure S1: genetic correlations between traits in the LD Hub database and years of education (plus standard errors) for adopted (red) and for non-adopted (blue) individuals. The figure only shows results for traits that showed significant genetic correlations with education in the sample of adoptees at p < 0.05.



Due to the relatively small sample of adoptees, only 4 traits were significantly genetically correlated with education years in the adopted sample after correcting for multiple testing by using a stringent p-value of 0.000202 (0.05/247): years of schooling (2016), college completion, intelligence, and years of schooling (proxy). All traits in this figure also showed significant genetic correlations with education years in the non-adopted sample at p<0.000202, except height of females age 10 and males at age 12 (p=0.0004), Alzheimer's disease (p=0.0002) and difference in height between adolescence and adulthood (p=0.7265). None of the genetic correlations with education were significantly different between adoptees and non-adopted individuals at p<0.000202.

Table S5: Genetic correlations with binary adopted/not adopted variable, from LD Hub, ranked by p-value. The table only shows correlations that remained significant after a Bonferroni correction for multiple testing (p < 0.0002 i.e. 0.05/247).

Trait	Genetic correlation	se	р
Years of schooling 2016	-0.5216	0.0615	2.28x10 <sup>-17</sup>
Age of first birth	-0.6589	0.0836	3.23x10 <sup>-15</sup>
College completion	-0.5584	0.0882	2.41x10 <sup>-10</sup>
Years of schooling (proxy cognitive performance)	-0.522	0.0851	8.38x10 <sup>-10</sup>
Years of schooling 2013	-0.4739	0.0823	8.39x10 <sup>-09</sup>
Depressive symptoms	0.5176	0.1026	4.60x10 <sup>-07</sup>
Intelligence	-0.306	0.0679	6.66x10 <sup>-06</sup>
Obesity class 1	0.2796	0.0652	1.77x10 <sup>-05</sup>
Waist-to-hip ratio	0.2507	0.0626	6.23x10 <sup>-05</sup>
Obesity class 2	0.3368	0.0851	7.58x10 <sup>-05</sup>
Extreme bmi	0.2943	0.0748	8.32x10 <sup>-05</sup>
Mothers age at death	-0.6567	0.1679	9.14x10 <sup>-05</sup>
Body mass index	0.2486	0.0648	0.0001
Insomnia	0.4258	0.1098	0.0001

Note: These significant negative genetic correlations with educational attainment, age of motherhood, and poor physical health agree with evidence that adoptees are more likely than individuals from comparable origins to be born to young mothers, to have had a low birthweight, and to have received suboptimal obstetric care (Maughan et al. 1998), although the two groups were highly similar apart from these factors. The fact that adoptee status had a low heritability (6%; liability scale) means that interpretation of these genetic correlations should be cautious.

Figure S2: Comparing the variance explained in years of education by polygenic scores (PRS) for years of education, per year-of-birth grouping, for adoptees and non-adopted individuals. 95% CIs were obtained by bootstrapping with 1000 replications.



Any systematic differences between adoptive and biological families are likely to have changed over time. The variance explained by polygenic scores is more variable in the group of individuals reared by their biological relatives, but generally increases over time. The plot shows that prediction is stronger for adoptees in the youngest age group (although this increase was not significant). Due to increasing use of contraceptives across the 20th century, adoption became less related to young motherhood and more related to removal of children from high risk environments. Adoptions occurring later in the twentieth century were more likely to involve older children rather than babies (Maughan et al. 1998; Keating 2008). We speculate that these changes could mean that younger adoptees in our sample are less generalisable, more likely to have come from high risk environments, and more likely to have spent more time with their biological relatives, experiencing passive gene-environment correlation. However, other than the non-significant spike in the youngest age group of adoptees, polygenic prediction remains stable across cohorts (at ~4%). It could be argued that the variance explained by polygenic scores in adoptees provide a rough benchmark of 'direct' genetic influence remaining throughout periods of great social change.

Table S6: sample sizes by year-of-birth grouping.

	Adopted	Non-adopted group d (N=6500)
>=1936 <=1940	341	448
>=1941 <=1945	1687	1413
>=1946 <=1950	1312	1503
>=1951 <=1955	839	1124
>=1956 <=1960	781	867
>=1961 <=1965	869	796
>=1966 <=1970	482	349

The post-war spike in the number of adoptions (and number of births generally) is consistent with the 'baby scoop' and the 'baby boom'.

## References

Keating, J. 2008. A Child for Keeps: The History of Adoption in England, 1918-45. Springer.

Maughan, B., Collishaw, S. and Pickles, A. 1998. School achievement and adult qualifications among adoptees: a longitudinal study. *Journal of Child Psychology and Psychiatry, and Allied Disciplines* 39(5), pp. 669–685.