

S3 Table. Details and notes on the sources for the sample size, the number of genome-wide significant hits, and the PGS R^2 , for large-scale GWAS efforts to date for height, BMI, *EduYears*, and self-rated health, reported in the same order as in Table 2.

Study	N	C^*	Number of hits						PGS R^2		
			Section	Page	Note	Section	Page	Note	Section	Page	Note
Wood <i>et al.</i> (2014) [1]			Abstract	1173		Results	1173		Abstract	1173	
Allen <i>et al.</i> (2010) [2]			Abstract	832	SI	2 Stages 1–2			Abstract	832	1175 Approximation least stringent p -value threshold
Weedon <i>et al.</i> (2008) [3]			Abstract	575	Results	576			Abstract	577	Figure 1a
Locke <i>et al.</i> (2015) [4]			Abstract	197		Results	197		Abstract	197	833 Approximation least stringent p -value threshold
Speliotes <i>et al.</i> (2010) [5]			Abstract	937		Results	937		Abstract	937	580 PGS based on 20 SNPs
Willer <i>et al.</i> (2008) [6]			Results	26	Stage 1	Results	26	Stage 1	Results	26	Figure 3c
Okbay <i>et al.</i> (2016) [7]			SI	16		SI	12,16		SI	16	Approximation
Okbay <i>et al.</i> (2016) [7]			Abstract	539		SI	12		Abstract	539	n.a.
Rietveld <i>et al.</i> (2013) [8]			Abstract	1467		Results	1467		Results	1467	SI
Harris <i>et al.</i> (2016) [9]			Abstract	1		Abstract	1		Abstract	1	Results
											1469 Approximation average

* C denotes the number of studies in the meta-analysis; C is slightly subjective (e.g., RS I, II, and III can be considered as one study or as three).

References

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