

S1 Table. Cohort characteristics

Sr. No.	Study Cohorts	N	Women [%]	MTHFR Genotype		GRS2 [Mean (S.D.)]	Age [Mean (S.D.)]	Gran [Mean (S.D.)]			Lym [Mean (S.D.)]	Mono [Mean (S.D.)]	Study Design	Pathophysiology			
				%CT	%TT												
CHARGE consortium																	
1	RS-III-I	729	54.0	41.6 (303)	8.6 (63)	1.09 (0.19)	59.86 (8.16)	4.21 (1.51)			2.42 (0.67)	0.42 (0.42)	Prospective, population-based	Mainly healthy			
2	LBC1921	436	60.3	45.0 (196)	12.4 (24)	1.13 (0.19)	79.10 (0.60)	4.6 (1.34)	0.17 (0.17)	0.02 (0.04)	1.71 (1.12)		0.54 (0.18)	Prospective, population-based	Mainly healthy		
3	LBC1936	905	49.4	43.2 (391)	11.4 (103)	1.17 (0.18)	69.60 (0.80)	4.5 (1.54)	0.17 (0.13)	0.04 (0.04)	1.90 (1.43)		0.53 (0.19)	Prospective, population-based	Mainly healthy		
4	KORA	1,799	51.1	45 (779)	10.6 (183)	1.11 (0.18)	60.98 (8.89)	0.33 (0.09)			0.10 ^W (0.07)	0.16 ^X (0.06)	0.02 ^Y (0.02)	0.05 ^Z (0.03)	0.12 (0.03)	Prospective, population-based	Mainly healthy
5	FHS	2,408	54.3	58.9 (1206)	18.2 (306)	1.12 (0.17)	66.4 (8.93)	0.49 (0.09)			0.10 ^W (0.06)	0.19 ^X (0.06)	0.02 ^Y (0.02)	0.04 ^Z (0.03)	0.13 (0.03)	Prospective, population-based	Mainly healthy
BIOS consortium																	
6	RS-II-3/RS-III-2	722	57.8	43.2 (312)	12.3 (89)	1.15 (0.19)	67.60 (5.98)	48.32 ^N (7.04)	2.74 ^E (0.73)	0.68 ^B (0.18)	39.66 (9.71)		7.19 (1.30)	Prospective, population-based	Mainly healthy		
7	LLS	739	51.7	48.0 (355)	11.8 (87)	1.14 (0.18)	58.98 (6.70)	58.14 ^N (7.93)	2.63 ^E (1.17)	0.64 ^B (0.25)	29.52 (6.18)		5.68 (1.16)	Prospective, family-based	Mainly healthy		
8	LLD	744	57.8	44.2 (329)	9.9 (74)	1.13 (0.17)	45.56 (13.32)	54.15 ^N (7.22)	2.93 ^E (1.49)	0.54 ^B (0.25)	33.47 (7.10)		8.14 (1.53)	Prospective, population-based	-		
9	NTR	691	68.5	41.8 (289)	10.3 (71)	1.12 (0.18)	33.95 (12.11)	52.59 ^N (7.66)	2.81 ^E (1.21)	0.47 ^B (0.41)	35.11 (7.81)		7.72 (2.08)	-	-		
10	CODAM	161	45.3	43.5 (70)	12.4 (20)	1.14 (0.17)	65.40 (6.77)	40.67 ^N (6.97)	2.80 ^E (0.63)	0.75 ^B (0.20)	51.11 (14.06)		7.95 (1.63)	Prospective, Observational	Mildly increased DM2 /CVD risk factors		
Other cohorts																	
11	MARTHA	349	78.5	48.1 (168)	11.7 (41)	1.14 (0.18)	43.77 (14.09)	4.58 (0.38)			1.87 (0.62)		0.32 (0.13)	Retrospective	Patients with Venous Thromboembolism		
12	F5L	211	52.1	48.8 (103)	20.9 (44)	1.15 (0.18)	39.7 (16.80)	0.56 (0.13)			0.04 ^W (0.04)	0.22 ^X (0.08)	0.10 ^Y (0.05)	0.10 ^Z (0.03)	0.08 (0.03)	Pedigree-based	Mainly healthy, probands with VTE

^N Neutrophils, ^E Eosinophils, ^B Basophils (Imputed)

^W CD8T, ^X CD4T, ^Y NK, ^Z Bcell (Imputed)