**Supplementary Table 1 -** Demographic and clinical characteristics for veterans in the MVP PAD genome-wide association study

	<b>White</b>		<b>Black</b>		<b>Hispanic</b>	
	PAD	PAD	PAD	PAD	PAD	PAD
	Cases	Controls	Cases	Controls	Cases	Controls
Veterans, N	24,009	150,983	5,373	42,485	1,925	18,285
Age ± SD, years	$74.4 \pm$	$66.9 \pm$	69.6 ±	$60.4 \pm$	71.6 ±	59.0 ±
	9.4	13.2	9.4	11.6	9.6	14.8
Male, n (%)	23,416	138,753	5,166	36,188	1,885	16,400
	(97.5%)	(91.9%)	(96.1%)	(85.2%)	(97.9%)	(89.7%)
Statin Therapy	12,351	61,575	3,401	15,787	1,340	7,460
Prescription, n (%)	(51.4%)	(40.8%)	(63.3%)	(37.2%)	(69.6%)	(40.8%)
Diabetes, n (%)	13,314	45,307	3,568	16,242	1,431	11,943
	(55.4%)	(30.0%)	(66.4%)	(38.2%)	(74.3%)	(65.3%)
Mean Max LDL-C	$142.0 \pm$	$139.6 \pm$	149.3 ±	141.9 ±	145.5 ±	141.5 ±
$\pm$ SD, mg/dL	38.9	37.8	43.7	40.0	40.4	38.1
Former Smoker, n	14,056	80,809	2,696	17,729	1,144	8,660
(%)	(58.5%)	(53.5%)	(50.1%)	(41.7%)	(59.4%)	(47.4%)
Current Smoker, n	6,643	25,040	1,954	10,988	413	2,788
(%)	(27.7%)	(16.6%)	(36.4%)	(25.8%)	(21.4%)	(15.2%)
Variants Passing Quality Control	20,323,458		32,412,077		31,270,737	

Abbreviations: MVP, Million Veteran Program; PAD, Peripheral Artery Disease; SD, Standard Deviation; LDL-C, Low-Density Lipoprotein Cholesterol; mg/dL, milligrams per deciliter; Max, Maximum

**Supplementary Table 3 -** Odds ratios and two-sided logistic regression P values for three lead autosomal, genome-wide associated PAD variants previously reported in the literature and the lead variant and logistic regression odds ratios and two-sided P values in MVP (N = 31,307 PAD cases and 211,753 controls).

Published Lead Variant	Chr	Locus	Published OR	Published P Value	MVP Lead Variant*	MVP OR	MVP P Value	Ref
rs9584669	13	IPO5/ RAP2A	1.72	6.8x10 <sup>-14</sup>	rs9584742	1.09	0.003	4
rs6842241	4	<i>EDNRA</i>	1.18	5.3x10 <sup>-9</sup>	rs6841581	1.07	1.9x10 <sup>-9</sup>	4
rs10757269	9	9p21	NA**	$2.7x10^{-9}$	rs1537372	1.11	$2.0 \times 10^{-28}$	3

<sup>\*</sup> Defined as the lead variant in MVP within a 500kB window around previously published lead variant

Abbreviations: Chr, chromosome; OR, Odds Ratio; Ref, Reference; GWAS, Genomewide Association Study; MVP, Million Veteran Program; PAD, Peripheral Artery Disease

<sup>\*\*</sup> Previously reported based on a GWAS of ankle-brachial index (ABI) as a continuous trait

**Supplementary Table 13 -** Overlap of CAD and PAD diagnoses in the 174,992 white veterans in the MVP PAD GWAS analysis

	CAD Case	CAD Control	<u>Total</u>
PAD Case	15,342	8,667	24,009
PAD Control	35,073	115,910	150,983
<u>Total</u>	50,415	124,577	174,992

Abbreviations: CAD, Coronary Artery Disease, PAD, Peripheral Artery Disease, GWAS, Genome-wide Association Study; MVP, Million Veteran Program

**Supplementary Table 15 -** Overlap of LAS and PAD diagnoses in the 174,992 white veterans in the MVP PAD GWAS analysis

	LAS Case	LAS Control	<u>Total</u>
PAD Case	519	23,490	24,009
PAD Control	186	150,797	150,983
<u>Total</u>	705	174,287	174,992

Abbreviations: LAS, Large Artery Stroke, PAD, Peripheral Artery Disease, GWAS, Genome-wide Association Study; MVP, Million Veteran Program

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