

Table S1. Characteristics of the study population used in this study.

Variables	Discovery cohort		GWAS-NJMU ^a		GWAS-SLD ^b		GWAS-CADM ^c		GWAS-FLCCA ^d	
	Case (%)	Control (%)	Case (%)	Control (%)	Case (%)	Control (%)	Case (%)	Control (%)	Case (%)	Control (%)
Sample size	192	278	2,331	3,077	1,937	1,984	3,779	3,837	4,796	3,741
Sex										
Male	140 (72.9)	198 (71.2)	1,711 (73.4)	2,086 (67.8)	1,532 (79.1)	1,519 (76.6)	2,926 (77.4)	3,375 (88.0)	--	--
Female	52 (27.1)	80 (28.8)	620 (26.6)	991 (32.2)	405 (20.9)	465 (23.4)	853 (22.6)	462 (12.0)	4,796 (100.0)	3,741 (100.0)
Age (years)										
<60	102 (53.1)	144 (51.8)	1,111 (47.7)	1,429 (46.4)	423 (21.8)	502 (25.3)	1,402 (37.1)	1,315 (34.3)	2,408 (50.2)	2,023 (54.1)
≥60	90 (46.9)	134 (48.2)	1,220 (52.3)	1,648 (53.6)	1,514 (78.2)	1,482 (74.7)	2,377 (62.9)	2,522 (65.7)	2,388 (49.8)	1,718 (45.9)
Smoking status										
Never	63 (32.8)	146 (52.5)	838 (36.0)	1,755 (57.0)	138 (7.2)	636 (32.1)	n/a	n/a	4,796 (100.0)	3,741 (100.0)
Ever	129 (67.2)	132 (47.5)	1,493 (64.0)	1,322 (43.0)	1,787 (92.8)	1,343 (67.9)	n/a	n/a	--	--
Histology										
SCC	63 (32.8)	--	822 (35.3)	--	492 (25.4)	--	n/a	--	660 (13.8)	--
AD	129(67.2)	--	1304 (55.9)	--	795 (41.0)	--	n/a	--	3469 (72.3)	--
Others	0	--	205 (8.8)	--	650 (33.6)	--	n/a	--	667 (13.9)	--

^a GWAS-NJMU: Genome-wide Association Study at Nanjing Medical University;

^b SLD: Environment and Genetics in Lung Cancer Etiology in Italy;

^c CADM: Alpha-Tocopherol, Beta-Carotene Cancer Prevention Study in Finland, the Prostate, Lung, Colon, Ovary Screening Trial and the Cancer Prevention Study II Nutrition Cohort in the US.

^d FLCCA: Female Lung Cancer Consortium in Asia.

SCC: Squamous cell carcinoma; AD: Adenocarcinoma; Others: other subtypes or missing; n/a: not available.

Table S2. The association of significant SNPs identified by previous GWAS and fine-mapping studies with lung cancer risk in this study.

First author (Reference)	Previous Study				Targeted Sequence			
	SNP	Effect Allele	OR(95%CI) ^a	<i>P</i> ^a	EAF ^b		OR(95%CI) ^c	<i>P</i> ^c
					Case	Control		
Hu Z (21725308)	rs4488809	T	1.26 (1.21–1.32)	7.20×10^{-26}	0.55	0.47	1.31 (1.01-1.75)	0.044
Lan Q (23143601)	rs4488809	T	1.19 (1.12-1.26)	4.00×10^{-9}				
Miki (20871597)	rs10937405	C	1.31 (1.22–1.42)	7.26×10^{-12}	0.74	0.66	1.49 (1.11-2.04)	0.009
Shiraishi (22797724)	rs10937405	C	1.25 (1.19–1.32)	6.92×10^{-17}				
Wang Y (24880342)	rs13314271	T	1.13 (1.09-1.18)	7.00×10^{-10}	0.54	0.47	1.31 (1.00-1.72)	0.054

^a Retrieved from previous papers;

^b EAF: effect allele frequency;

^c Logistic regression with adjustment for age, gender and smoking status.

Table S3. Conditional analysis of GWAS SNVs by rs4488809 and rs10937405.

SNP	Ref/Eff Allele	MAF		Step 1		Step 2		Step 3		Step 4	
		case	control	OR (95%CI) ^a	<i>P</i> ^a	OR (95%CI) ^b	<i>P</i> ^b	OR (95%CI) ^c	<i>P</i> ^c	OR (95%CI) ^d	<i>P</i> ^d
rs4396880	G/A	0.26	0.35	0.64(0.47-0.87)	3.88×10 ⁻³	-	-	0.65(0.48-0.89)	6.10×10 ⁻³	-	-
rs3856776	C/T	0.13	0.08	1.84(1.19-2.85)	6.05×10 ⁻³	1.79(1.15-2.77)	9.52×10 ⁻³	-	-	-	-
rs4488809	T/C	0.45	0.53	0.76(0.57-0.99)	4.43×10 ⁻²	0.98(0.68-1.41)	0.912	0.8(0.61-1.06)	0.127	1.08(0.75-1.57)	0.678
rs10937405	C/T	0.26	0.34	0.67(0.49-0.9)	9.27×10 ⁻³	1.99(0.47-8.4)	0.348	0.68(0.5-0.93)	0.143	2.08(0.49-8.89)	0.325
rs13314271	C/T	0.54	0.47	1.31(1-1.72)	5.45×10 ⁻²	1.01(0.71-1.44)	0.963	1.23(0.93-1.63)	0.145	0.92(0.64-1.33)	0.659

^a adjusted for age, sex and smoking status.

^b adjusted for age, sex, smoking status and rs4396880.

^c adjusted for age, sex, smoking status and rs3856776.

^d adjusted for age, sex, smoking status, rs4396880 and rs3856776.