

Supplementary table1. Genomic regions analyzed.

Chr	Gene	Chr position		kb analyzed
		Start	End	
1	<i>EPHX1</i>	196,537,000	196,544,000	7
	<i>IL6R</i>	152,644,000	152,707,000	63
2	<i>SERPINE2</i>	216,693,000	216,710,000	17
	<i>SFTPB</i>	85,792,000	85,794,000	2
3	<i>PDZRN3</i>	73,515,000	73,755,000	240
4	<i>HHIP</i>	145,500,000	145,730,000	230
	<i>GYPA</i>	145,480,000	145,500,000	20
5	<i>ADAM19</i>	152,020,000	152,025,000	5
6	<i>TNF</i>	31,675,000	31,677,000	2
	<i>GPR126</i>	142,664,000	142,810,000	146
7	<i>SERPINE1</i>	75,024,000	75,027,000	3
	<i>IL6</i>	22,817,000	22,818,000	1
9	<i>TNFSF8</i>	116,747,000	116,761,000	14
10	<i>SFTPD</i>	81,687,000	81,700,000	13
	<i>SFTPD</i> cercano	75,484,000	75,633,000	149
13	<i>FOXO1</i>	40,020,000	40,140,000	120
16	<i>MMP15</i>	56,631,000	56,658,000	27
19	<i>TGFB1</i>	38,291,000	38,293,000	2
	<i>CYP2A6</i>	46,040,000	46,050,000	10
	<i>CYP2B6</i>	46,180,000	46,222,000	42
	<i>SLC7A9</i>	38,010,000	38,053,000	43
	<i>ZNF582</i>	61,585,000	61,597,000	12
20	<i>ADAM33</i>	3,596,000	3,610,000	14
21	<i>ADARB1</i>	45,434,000	45,480,000	46