

**Additional file 2 Nucleotide diversity ( $\pi$ ) values for subpopulations within Colombia**

<b>n</b>	<b>Location</b>	<b>Ss</b>	<b>S</b>	<b>Ps</b>	<b>H</b>	<b>k</b>	<b><math>\pi</math></b>
14	North-west	9	1	8	6	3.6	0.0032 (0.0005)
3	Midwest	7	7	0	2	4.7	0,0042 (0.0019)
4	South-west	1	1	0	2	0.5	0.0004 (0.0002)
9	South-east	9	1	8	4	3.3	0.0029 (0.0011)

n: number of isolates, Ss: number of segregant sites, S: number of singleton sites, Ps: number of parsimony-informative sites, H: number of haplotypes, *k*: average number of nucleotide differences by sequence pairs,  $\pi$ : nucleotide diversity per site.