

**Figure 4-figure supplemental 1.** Sharing pattern percentages by pattern type

Sharing depth	Sharing pattern	Percentage of shared variants							
		Chromosome							genome
		1	2	3	4	5	6	7	
shallow	within N.Am	36.4	36.8	37.8	32.9	37.4	36.7	37.2	36.8
	within Asia	10.7	10.6	11.0	12.7	10.6	10.7	10.1	10.8
	within Europe	5.5	5.1	4.4	5.2	4.7	5.2	4.6	4.9
deep	N.Am and Asia	7.4	7.1	7.1	7.6	7.2	6.8	7.0	7.1
	Europe and N.Am	3.3	3.6	3.7	3.7	3.3	3.9	3.3	3.5
	Europe and Asia	14.7	14.2	14.7	13.2	15.1	14.5	15.5	14.7
very deep	among all	22.0	22.6	21.4	24.6	21.7	22.3	22.2	22.2