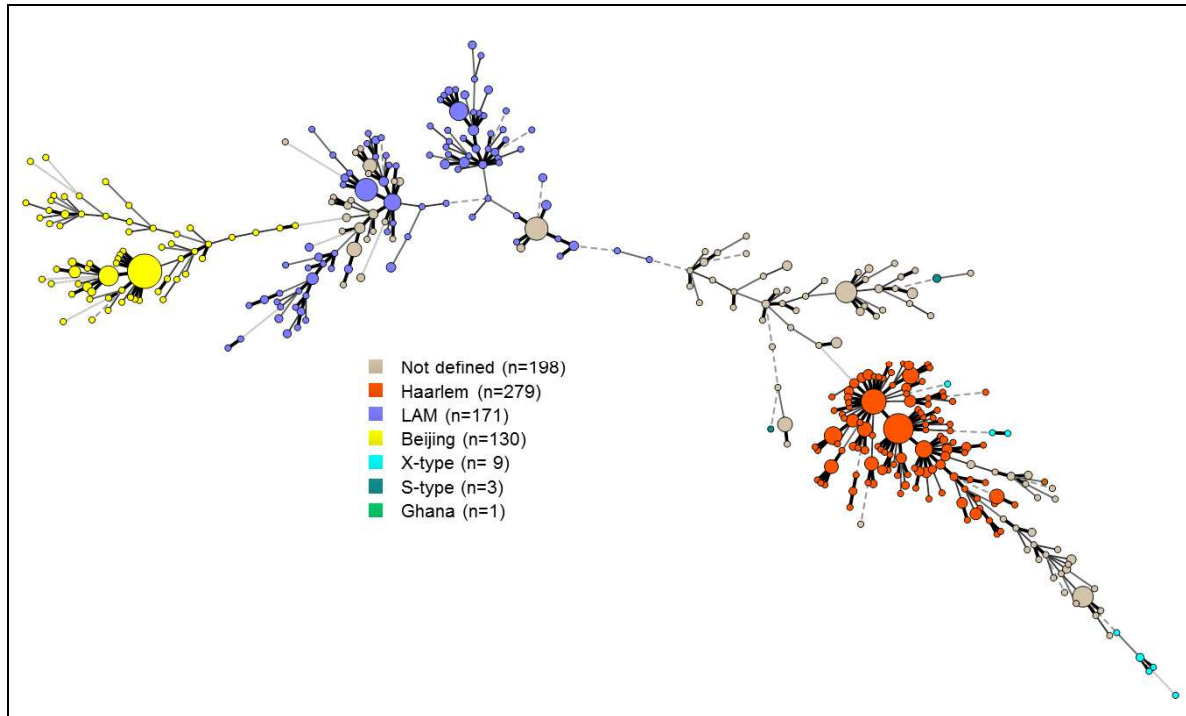


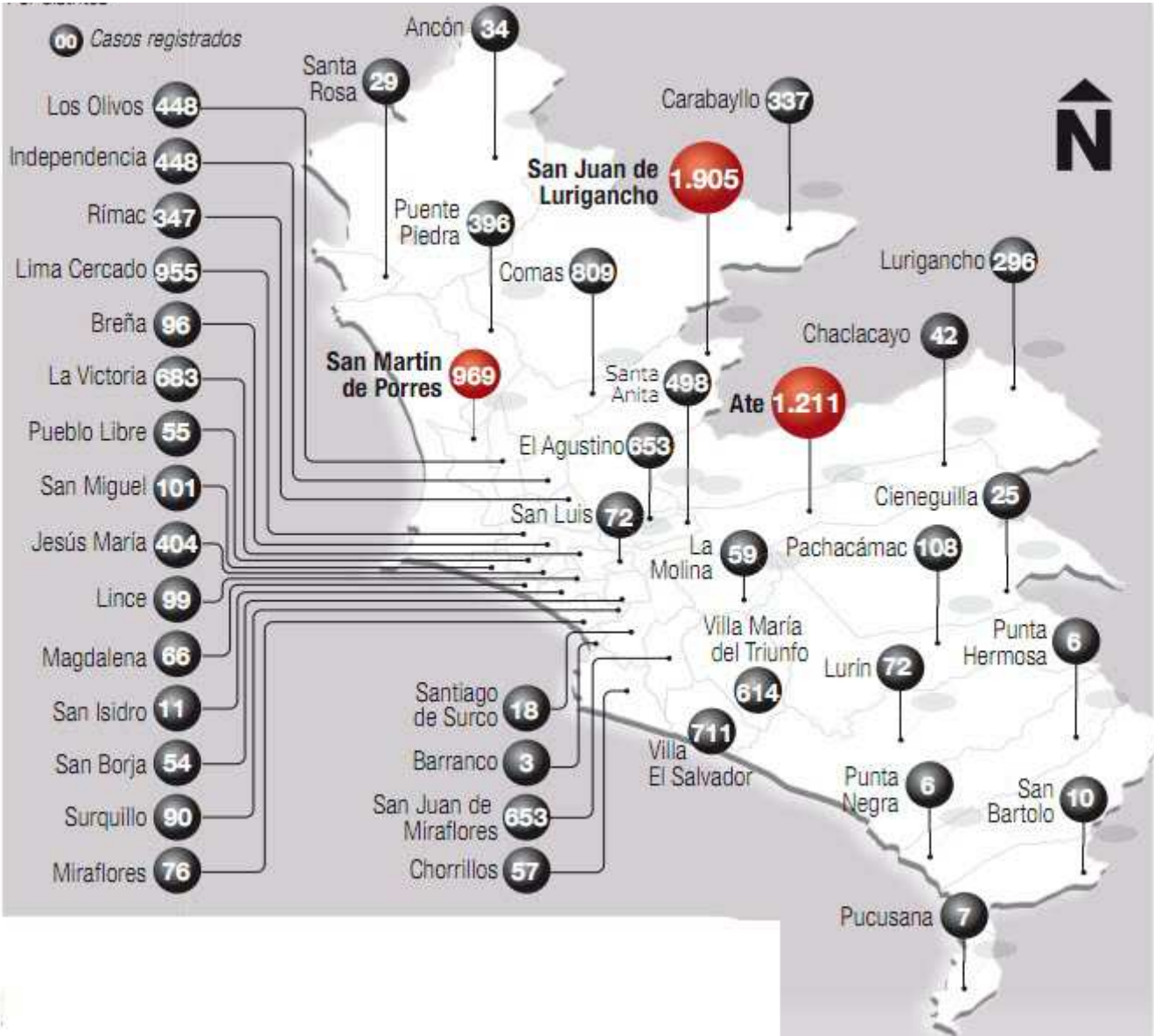
Supplementary Material 1 (S1):

Minimum Spanning tree based on the diversity of MIRU-15 data



The different complexes identified are coloured (maximum neighbour distance: four changes; minimum size: two MIRU-VNTR types) by the set of 15-loci among the 791 *M. tuberculosis* strains analyzed. The size of each circle is proportional to the number of MIRU-VNTR types belonging to a particular complex. LAM=*M. tuberculosis* Latin American Mediterranean.

Supplementary Material 3 (S3): Registered TB cases stratified by districts



Source: Ministry of Health. Peru. 2011.