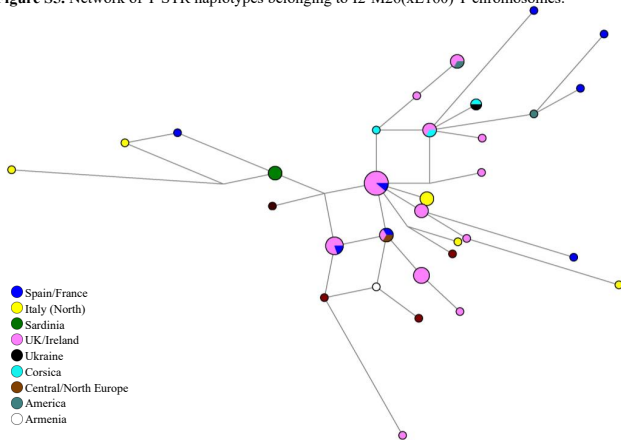


Figure S5. Network of Y-STR haplotypes belonging to I2-M26(xL160) Y chromosomes.



Network analysis was performed on Sardinian samples in comparison with all the I2-M26(xL160) samples available in the literature (Table S8). Circles and colored sectors are proportional to the number of subjects, with the smallest circle equal to one Y chromosome; connecting lines are proportional to the number of mutations.