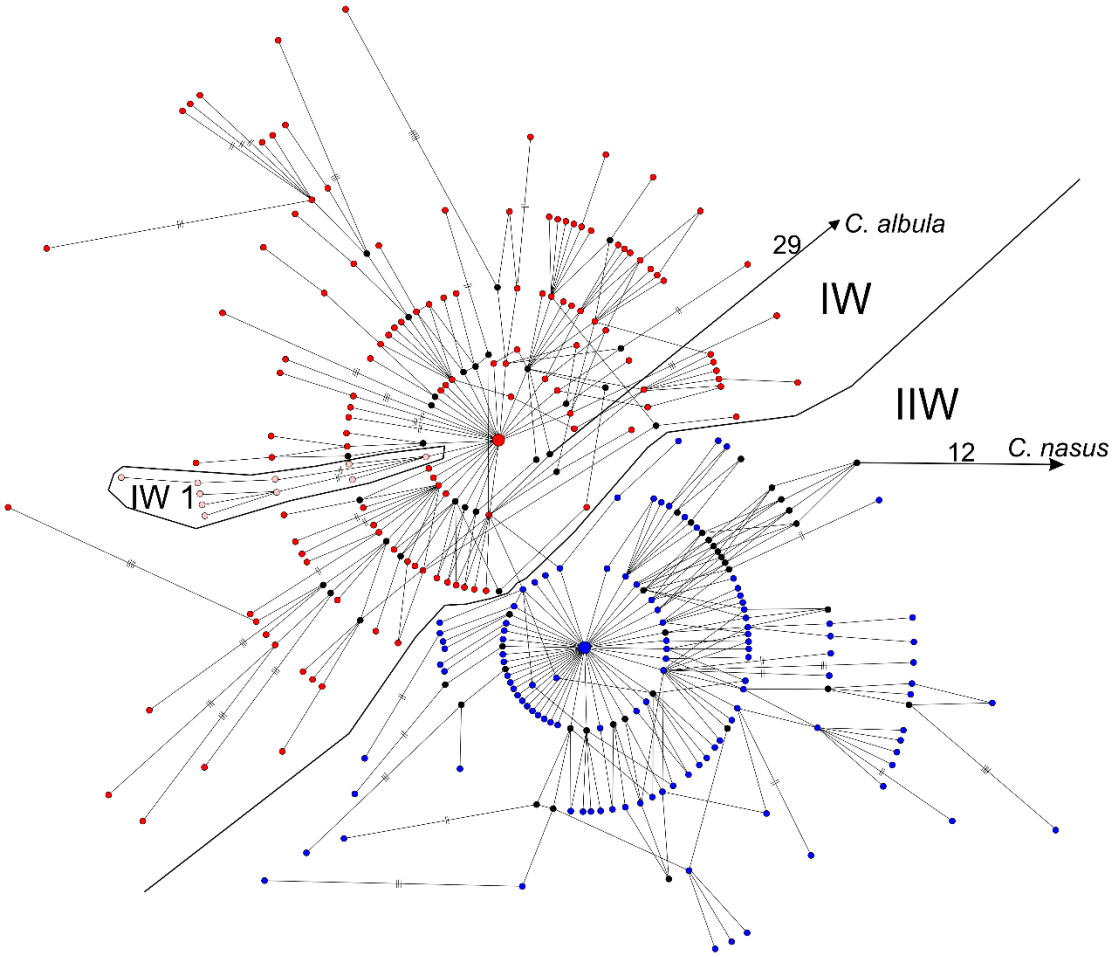

Figure S3. Median-joining haplotype network of the whitefish based on the *nd1* gene sequences. If the number of nucleotide substitutions between the haplotypes is more than 1, it is indicated next to the segment connecting the haplotypes (by short lines or numbers). In some cases, the proportions in the distances between haplotypes could not be preserved due to the two-dimensionality of the figure. The diameter of the circle does not correspond to the frequency of the haplotype; black circles are median vectors denoting not revealed or extinctic haplotypes. IW, IW1, and IIW–discussed haplogroups. The directions of the network to the other coregonid species, *C. albula* and *C. nasus*, are shown.