SNPs to discriminate different classes of hybrid between wild Atlantic salmon and aquaculture escapees: Supplementary Figures.

Figure S1: Relationship between true B allele frequency obtained through individual genotyping, and B allele frequency estimated by allelotyping, for the four New Teno Mainstem pools. Top: raw estimated frequencies before PPC correction, 199,297 SNPs; middle: estimated frequencies following PPC correction, 199,297 SNPs; bottom: 200 SNPs used for hybrid discrimination.

Figure S1

