

Figure S1

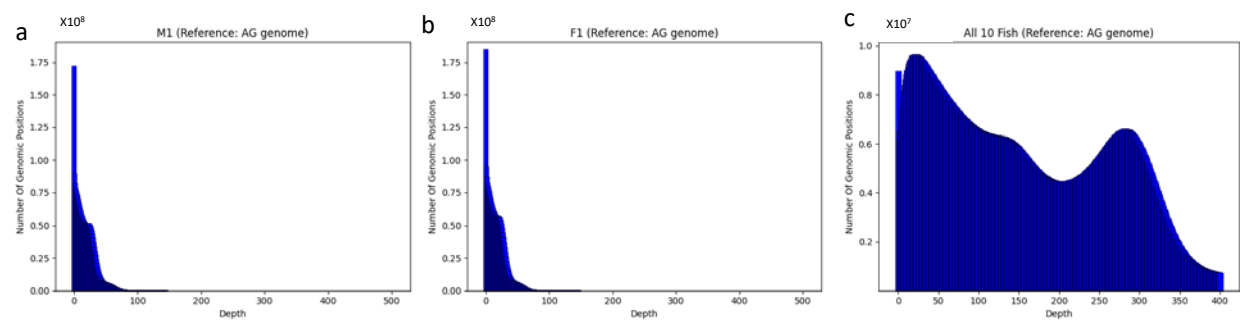


Figure S1: Distributions of sequencing depth over all genome positions when reads are aligned to the AG genome: (a) Male M1; (b) Female F1; (c) All samples combined.