

Figure 4. Proportion of chromosome length and female relative depth compared with SNP-loading range. A) Proportion of genome against 95% range of SNP-loading values for PC1. Dotted black line is Z-chromosomal, dashed grey line is autosomal and full black line is all scaffolds pooled. B) Relative scaffold depth the female sequenced to a high depth against 95% SNP-loading range. Open grey circles are autosomal scaffolds and full black dots are Z-chromosomal. The vertical line in both plots represents 3 SDs above the mean. For legibility, panel B was limited on the y axis to 0-1.5. Two scaffolds had relative depth greater than 1.5 (both >15), an autosomal scaffold around 0.025 SNP loading range, and a Z-chromosomal scaffold around 0.15.