



Figure 1: Differentiation, diversity and divergence for the S locus genomic region (chromosome 8) for *Antirrhinum majus pseudomajus* and *Antirrhinum majus striatum* using pooled whole genome data. For π_w (b): red is *Antirrhinum majus pseudomajus* and black is *Antirrhinum majus striatum*. Pink vertical lines indicate the positions of the 36 SLFs (F boxes) mapped to the reference genome. The two populations were separated by 20.9 km. The *A. m. pseudomajus* pool contained 50 plants, while the *A. m. striatum* 52 plants. Sliding window analysis based on a 10 kb window size with a 5 kb overlap. Gaps indicate areas of low mapping success.