

Figure S2 Detailed representation of the four causal models fitted to individual data for the *Viola cazortensis* plants considered in this study, linking multilocus genotypes based on herbivory-related AFLP markers (PC_{afp1} - PC_{afp3} scores), multilocus epigenotypes (PC_{msap1} - PC_{msap3} scores), and magnitude of browsing damage ($R_{Herbivory}$, residuals after controlling for the average effect of substrate). Variable line widths of arrows reflect differences in absolute values of path coefficients, as shown by the scale at bottom.

