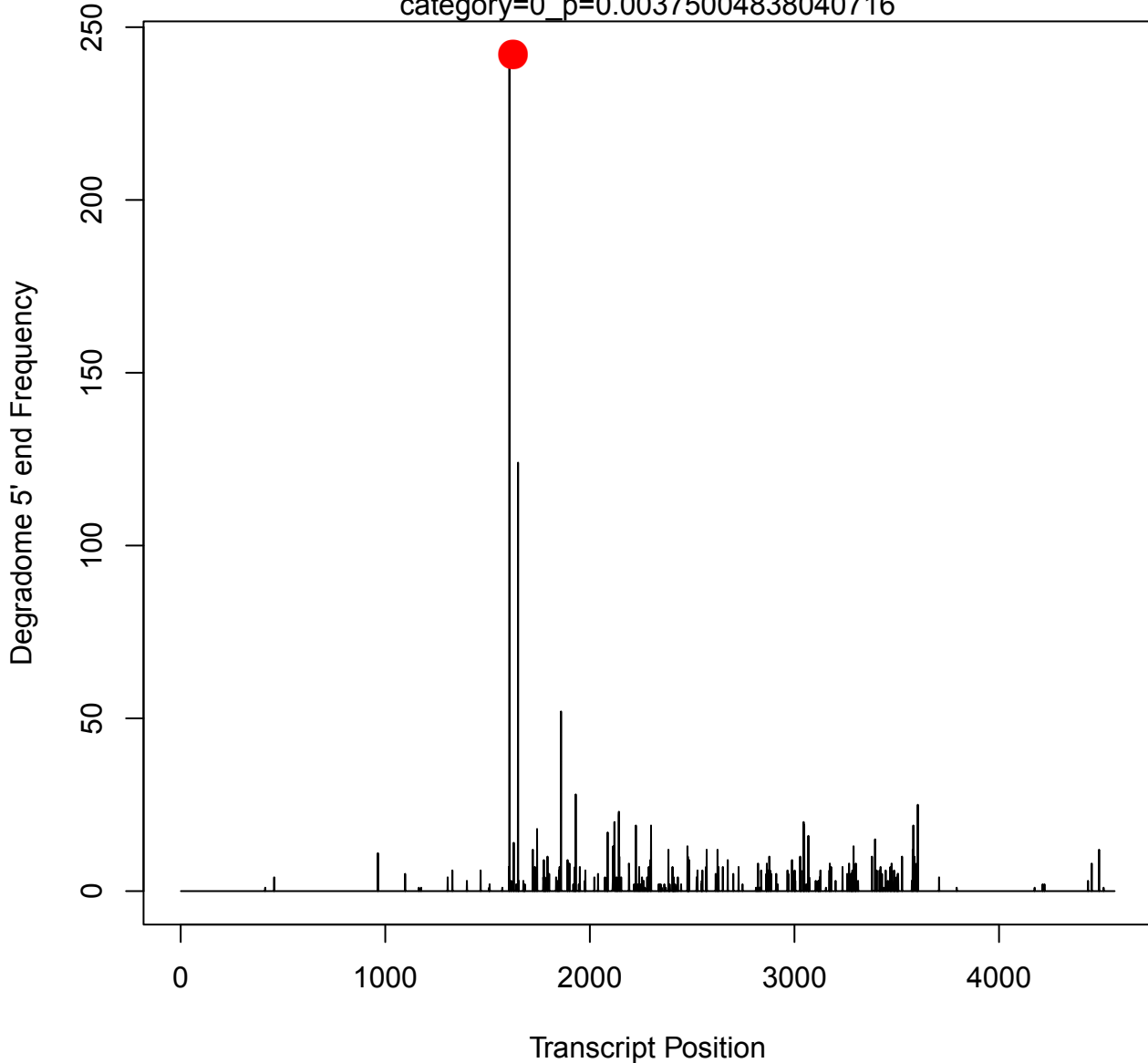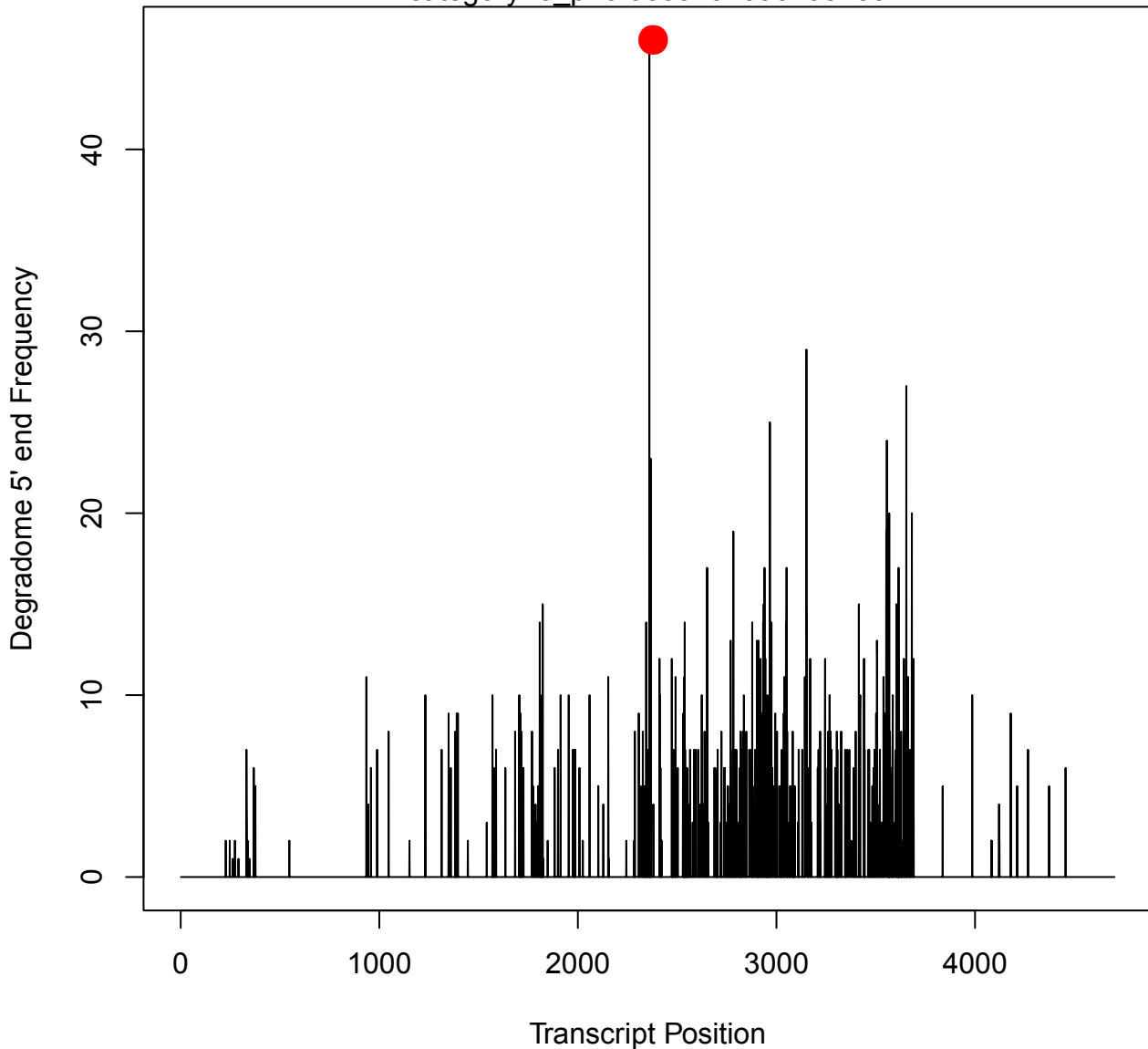**T=TC.CA05g06820 (CNL)_Q=can−miR482c_S=1607**

category=0_p=0.00375004838040716
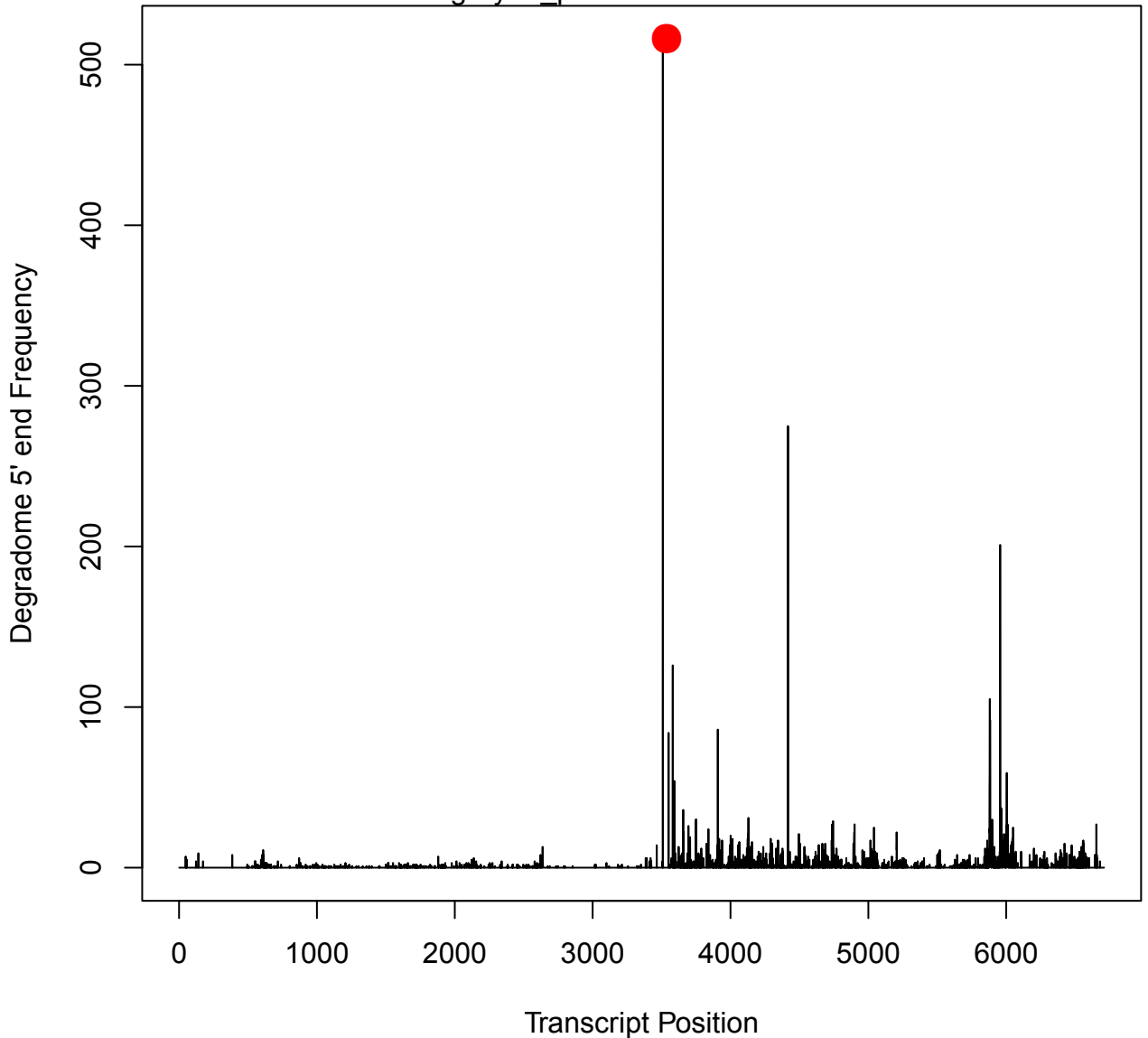
T=TC.CA11g01410 (CNL)_Q=can−miR6027-3p_S=2360

category=0_p=0.0683132896105783
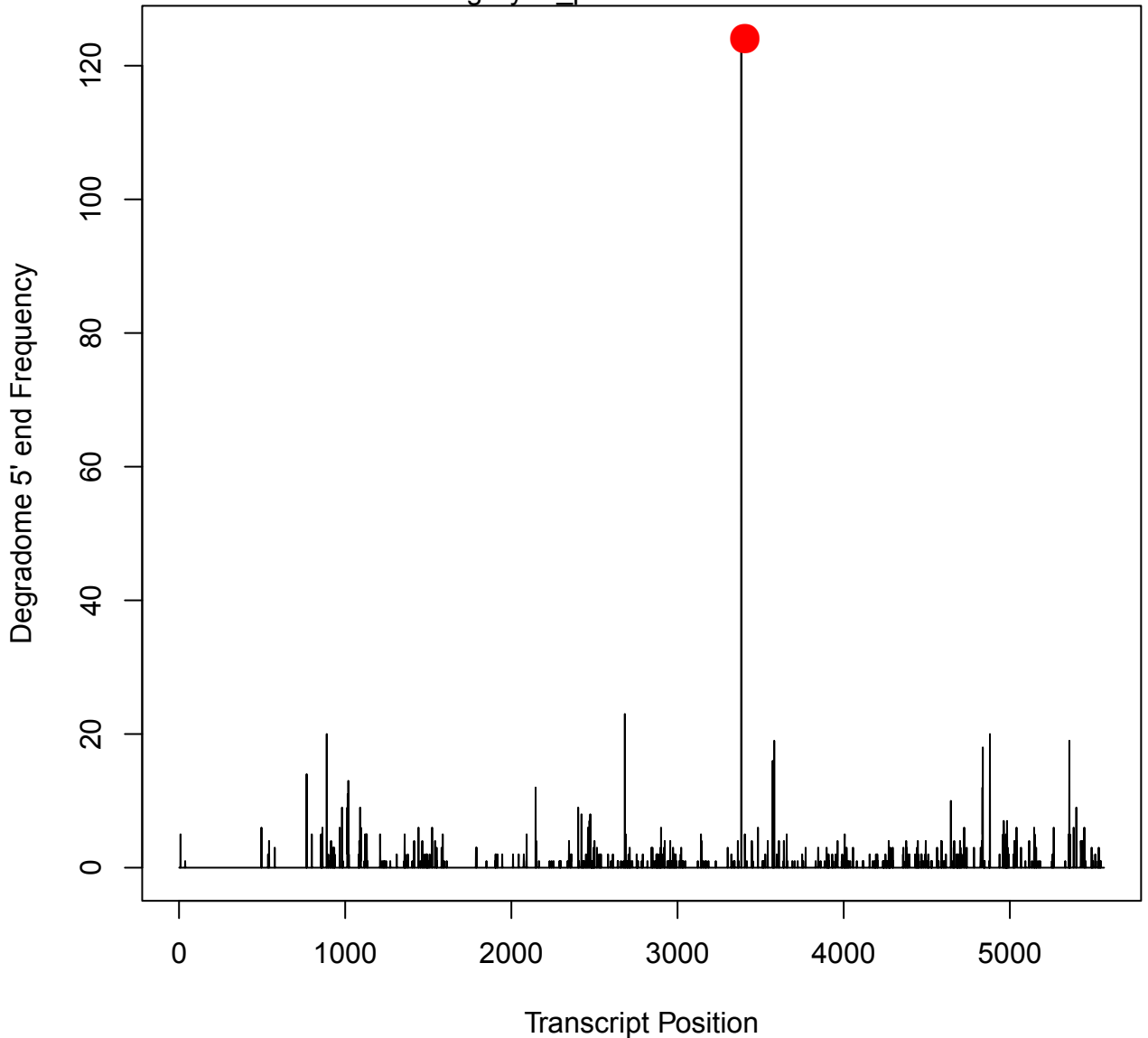
T=TC.CA08g01140 (RLP)_Q=can−miR−n026_S=3509

category=0_p=0.00437368779436897

T=TC.CA06g02150 (CNL)_Q=can−miR−n033a_S=3384

category=0_p=0.001876785351832

T=TC.CA03g01330 (CNL)_Q=can−miR6024-3p_S=2621
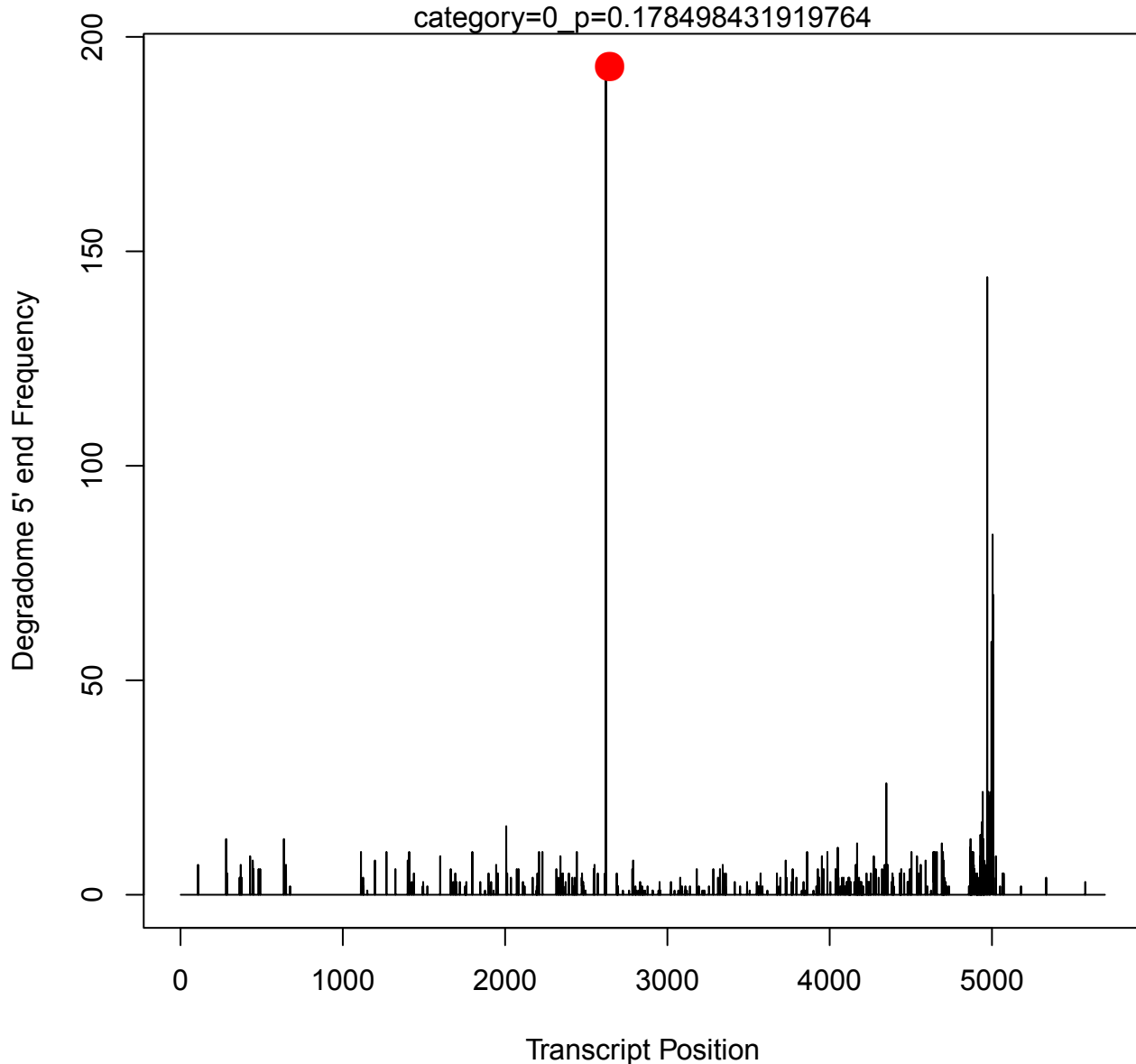
category=0_p=0.178498431919764

Figure S3. Representative examples of miRNAs cleaving disease-resistance genes

Plot showes the result of degradome analysis. X-axis and y-axis indicate the 5' end position of the mapped degradome reads and abundance of mapped degradome reads, respectively. Red dot indicates the cleavage site in the transcript by miRNA.