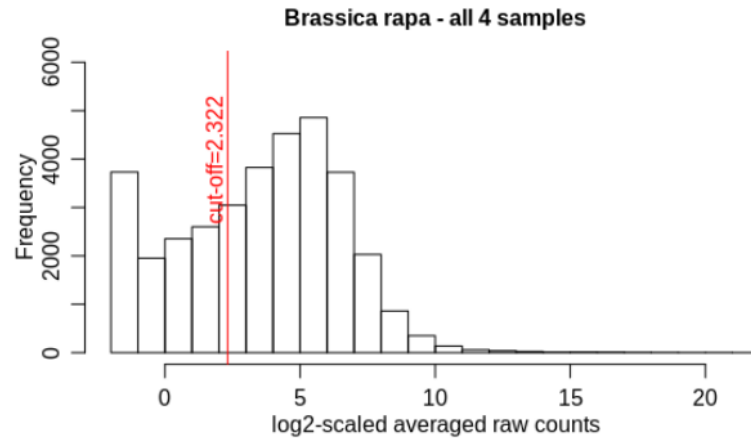


Figure S1. Intermediates from the evaluation of sequencing data related to *X. campestris* pv. *campestris* (Xcc) genome. **(A)** Occurrence of 310 contigs that possess mapped reads in the negative Xcc sample. **(B)** Comparison of four normalisation methods and rlog normalized counts of reads mapped within individual Xcc variants **C** normalized counts of mapped reads for individual Xcc variants as generated by individual normalization methods.

A.



B.

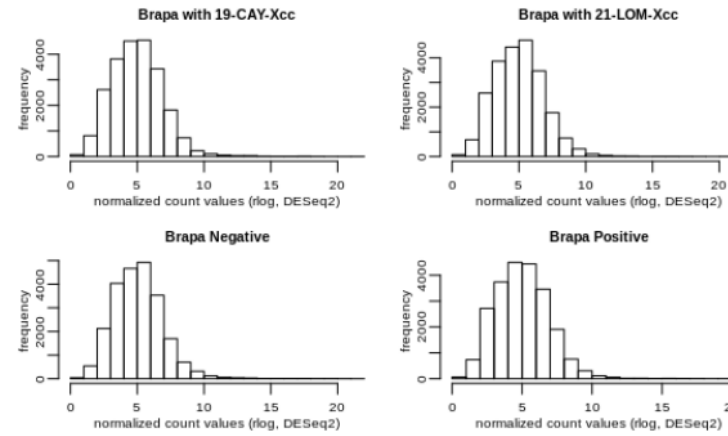


Figure S2. Intermediates from the evaluation of sequencing data related to *B. rapa* genome. **(A)** log2-scaled mean values of mapped reads per contig. **(B)** distributions of rlog normalised values computed for individual variants.



Figure S3. Divergent virulence of some epigenetically treated strains of *X. campestris* pv. *campestris* registered 15 days after inoculation. Positive control represent plants inoculated by untreated Xcc strain, remaining quintuplets represents plants inoculated by Xcc strains treated by CAY and LOM chemicals.