**Supplementary file 1** Summary of identified WRKY33 binding sites within the Arabidopsis genome 14 h post *B. cinerea* 2100 infection.



Note:

1) A peak region in a replicate is counted as reproducible, if it overlaps with a peak region in the other replicate by at least 50% of the length of the smaller peak region.

2) A peak region is counted as high confidence region (within a sample), when it is identified against all (two/three) negative controls used for peak calling.

3) A peak region in the pooled sample is counted as consistent, if it overlaps with a reproducible peak region in both of the original replicates by at least 50% of the length of the smaller region.

- For both replicates, peak calling was performed against two different negative controls: a mock-treated ChIP control (mock\_IP) and a Bc-treated input control. Bc\_input samples were identically processed as the IP samples except that no HA-antibody was added in the ChIP assay.

- For the WT-type negative controls (WT\_IP\_Bc\_vs\_mock), no peaks were observed, therefore only one replicate (replicate 2) was performed and analyzed here.