



Supplementary Figure S7 Representative genome-wide QTL evidence output of FlexQTL™ for resistance to *Phytophthora cactorum*. The X-axis represents cumulative genetic distance, and the Y-axis represents traces of QTL models (A) and posterior intensity (B). The number in the yellow is the Bayes Factor, a measure of decisive evidence (BF >10), for the QTL for a single run. These figures are representative of FlexQTL™ outputs obtained from four independent runs