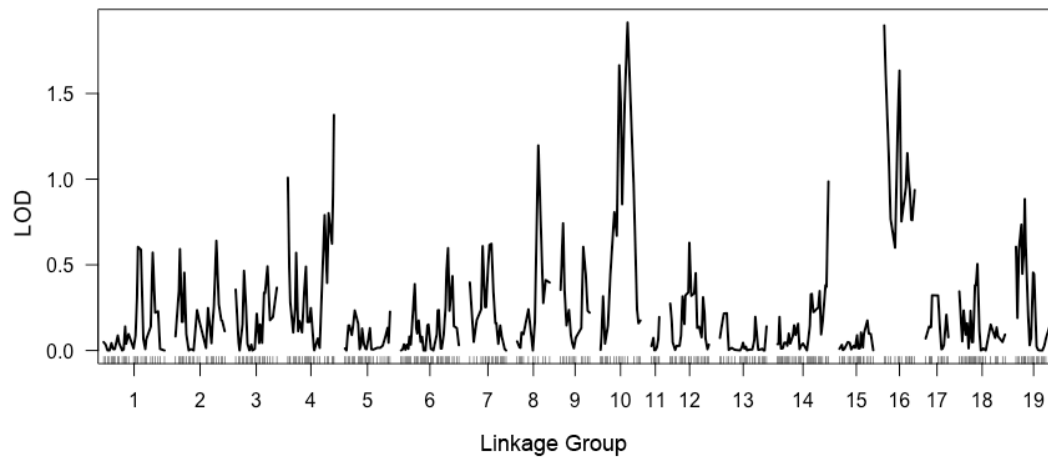




Additional file 3: Interval mapping of grape phylloxera resistance using the Riesling genetic map

Using the binary model, no LOD peaks above the threshold for phylloxera resistance were detected with the Riesling genetic map. The LOD threshold was 2.88, as determined by 1000 permutations. LOD score and linkage group number is shown on the y- and x-axis, respectively.