p7 "we used a used Genome-wide," do you need two used?

p12 "three of the hotspots have an overrepresentation of photosynthesis-related functions."  Downregulation of photosynthesis genes is a hallmark of plant immune process (e.g. Bilgin et al., 2010: <https://onlinelibrary.wiley.com/doi/full/10.1111/j.1365-3040.2010.02167.x>

Jiang et al., 2017:

<https://link.springer.com/article/10.1007%2Fs11103-017-0617-5>

).  You may want to add a line.

p16 "the identify of" right wording?

p18 in Experimental design, please give more information on the inoculation experiment such as the length of incubation after inoculation and disease phenotypes at the time of harvest for RNAseq.

Fig. 6 Gene linked to eQTL.  I wonder if you can add published QTL regions such as lesion size or virulence on the map.

p12 "no specific molecular insights arose largely because the majority of genes had no annotation".  Are genes without annotation enriched?  In general, filamentous fungus-specific genes involved in niche adaptation or development, those not found in yeasts, are largely unannotated.  You may want to check if genes with signal peptides for secretion are enriched.  Many effectors and plant cell wall degrading enzymes are secreted.

<http://www.cbs.dtu.dk/services/SignalP/>

Genes for secreting proteins and those expressed in planta are often enriched in subtelomeric regions and/or gene sparse high-speed regions (I think Cuomo et al., Science 2007; Raffaele et al., Science 2010), which are discussed later.

P15 "it would predict clustering of the great majority of eGWA hits to a few locations"  Isn't it also possible that not eQTL but their downstream genes are in the high-speed genomic regions?

Table 1 There is not much overlap between Botrytis and Arabidopsis transcripts.  For instance, the first one, Bcin01g01610 does not associate with Bot but lots of Arabidopsis mRNA.  Can you explain how the SNP influence those host genes without changing its own transcriptome?

Please let me know if you have any questions.