Figure S4  Dotplot comparisons between M. truncatula and G. max in regions surrounding M. truncatula LPs. Panels show syntenic regions of G. max chromosomes 8 a) and 15 b) compared against a 2 Mb region in A17 chromosome 4, corresponding to 0.5% of the total M. truncatula genome. Syntenic regions of G. max chromosomes 8 c) and 18 d) are compared against a 3-4 Mb region of A17 chromosome 7. Red dots indicate regions of sequence similarity with a forward orientation, while blue dots indicate reverse matches. The LP region in M. truncatula is shown with green rectangles.