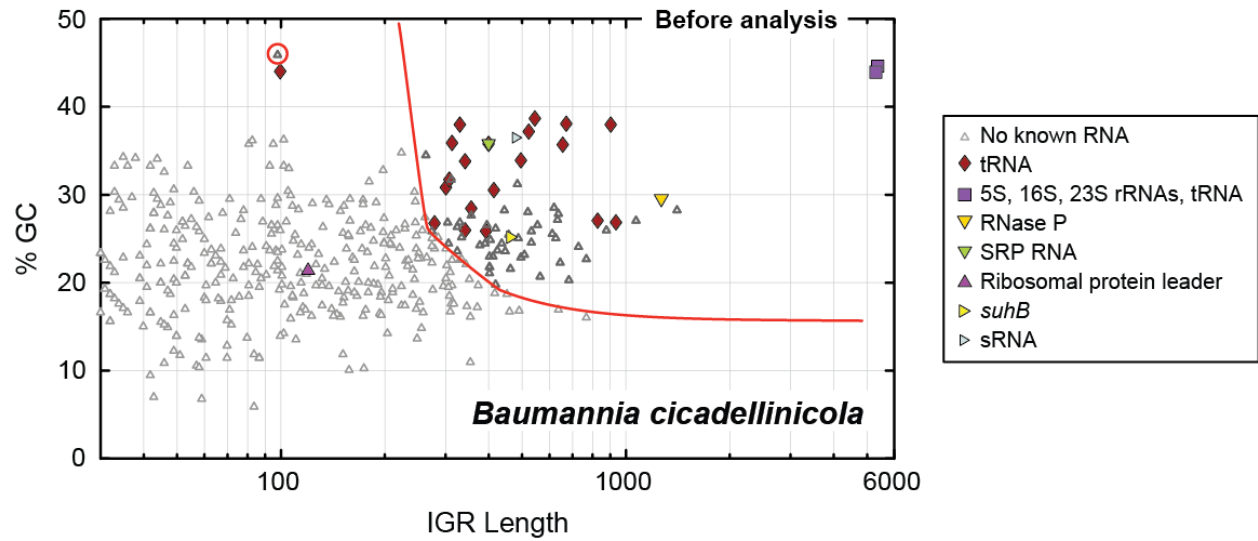




**Figure S5 | Plots of the IGRs from the *B. cicadellinicola* genome sorted based on IGR length and GC content.** Details are as described in the legend to **Fig. 2a**. In addition to IGRs selected with the main boundary (red line), the IGR within the red circle was also analyzed.