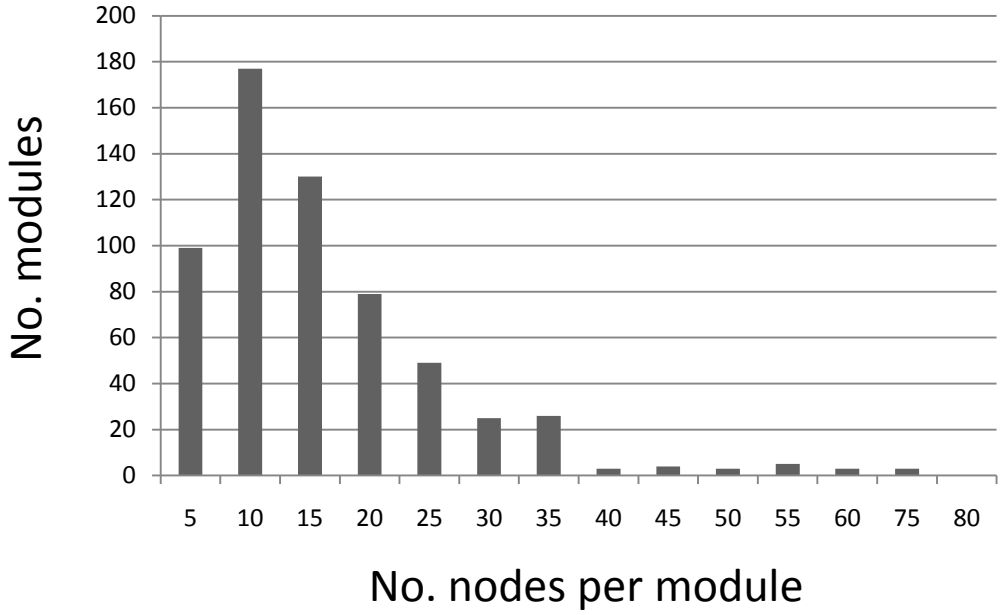

Figure S1. Distribution of number of probes per module.

Supplemental Figure S1

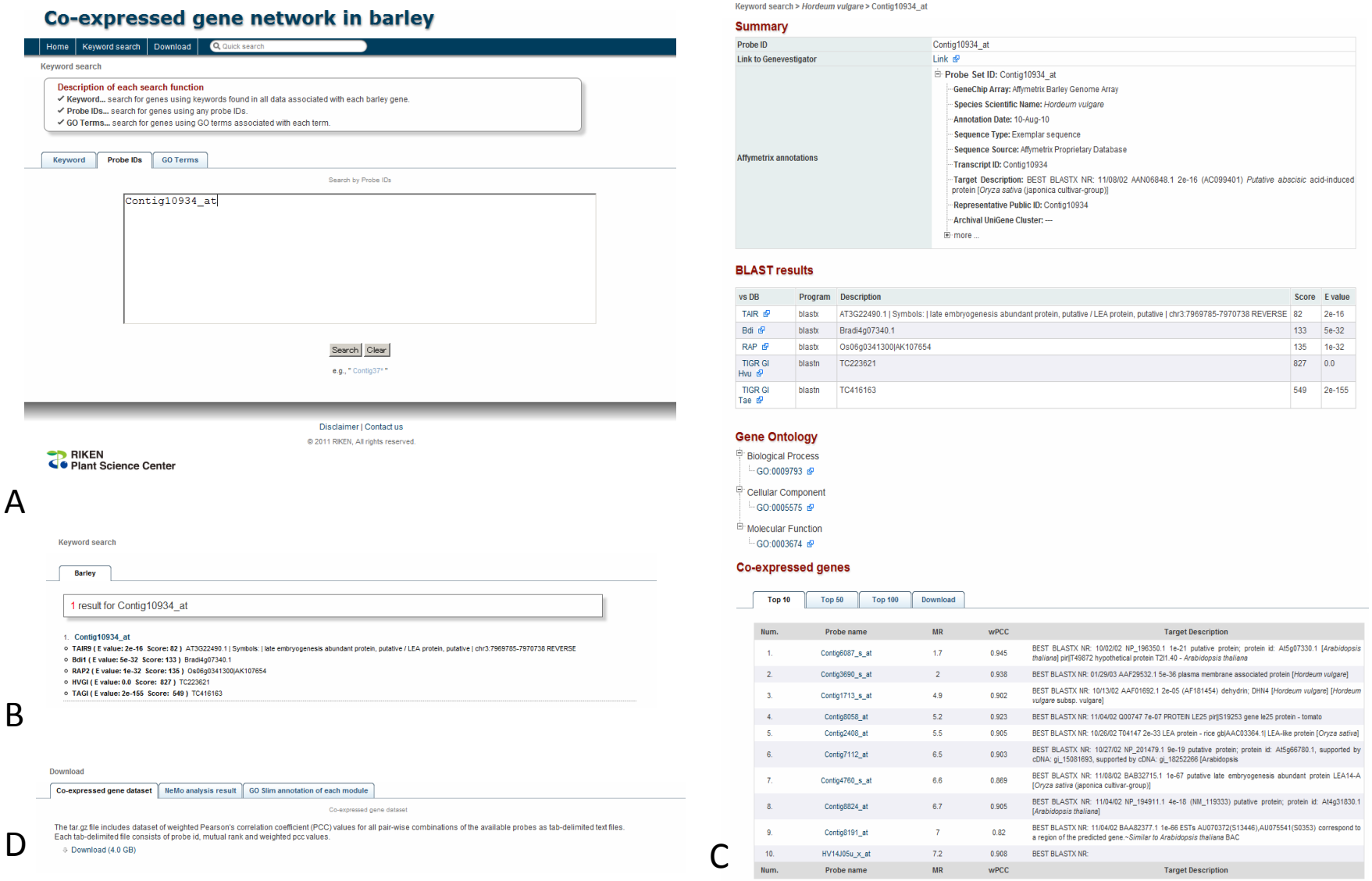

Figure S2. The interface of the web site used to provide datasets of the co-expression network of barley genes (http://coexpression.psc.riken.jp/barley/). (A) The interface provides search queries for the keywords, probe identifiers, and GO terms associated with the probes. (B) The search results are listed with annotations based on similarity searches. Users can navigate to the detailed annotation pages to browse the related annotations. (C) The detailed annotation pages provide summarized basic information on each of the probes, hyperlinks to Genevestigator, GO annotation, and ranked probes that are co-expressed. (D) The web site also provides a user interface for downloading archived datasets of the co-expressed genes in barley.