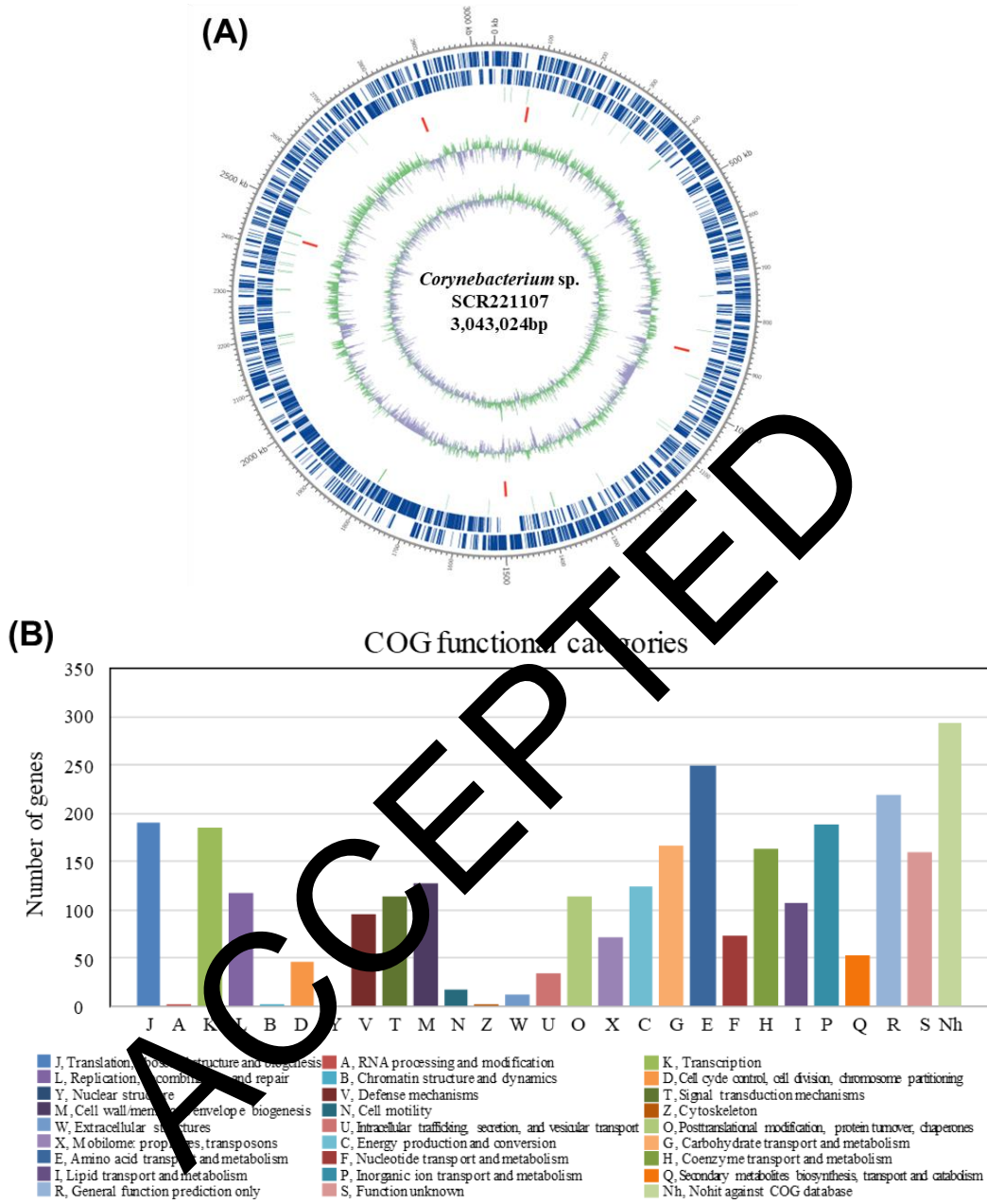


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129 **Figure 1.** Genome map of *Corynebacterium* sp. SCR221107 (A) and the functional categorization of
130 predicted coding sequences (B).

131 Marked characteristics are shown from the outside to the center: coding sequence (CDS) on the forward
132 strand, CDS on the reverse strand, tRNA, rRNA, guanine + cytosine (GC) content, and GC skew.