CPGview is developed based on open-source software packages, including gggenes, ggplot2, circlize, etc., the package of R language and the main biology package biopython in Python language. These R language packages are freely available on the Comprehensive R Archive Network (CRAN) under the GPL-2 license at https://cran.r-project.org/; the R platform and all R dependencies are similarly available from CRAN. Python biopython package and corresponding documentation are licensed under the PSF License Agreement(<https://docs.python.org/3/license.html#psf-license>).

The Singularity container used in this procedure and documentation are subject to the BSD license at https://docs.sylabs.io/guides/3.6/user-guide/license.html.