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Homo uses four functions (i.e., xGaussian_Distribution_Tail, xChi_Square_Distribution_Tail, Sum_Poisson_Terms, and Sum_Over_Odd_Terms) from two source code files (i.e., gaussian_distribution_tail.c, and chi-square_distribution_tail.c) written by Dr Richard L. Horn (athretprogr@gmail.com).

Dr Horn granted CSIRO and USYD a license to use these source code files without restrictions.

The functions provided by Dr Horn replace some of the functions made available to the public from his Mathematics Source Library C & ASM (<http://www.mymathlib.com>)

Credits

This software was developed by Lars S Jermiin while employed at CSIRO and USYD. Currently, LSJ is at Research School of Biology, The Australian National University, Acton, ACT 2601, Australia (lars.jermiin@anu.edu.au).