

Instructions for using the GenomicLand software

- **Step 1:** Download the R software on the link <https://cran.r-project.org/bin/windows/base/old/3.5.1/> (**version 3.5.1**).

- **Step 2:** Install the R software as instructed in the topic **Installation and other instructions**.

- **Step 3:** Set environment variables
 - i. In Search, search for and then select: System (Control Panel);
 - ii. Click the **Advanced system settings** link;
 - iii. Click **Environment Variables**;
 - iv. Under **System variables**, select **Path** and click on **edit**;
 - v. Click **New**, and **add or concatenate** (**use ;**) the folder address for R to there (for example, C:\Program Files\R\R-3.5.0\bin\x64);
 - vi. Click **OK**. Close all remaining windows by clicking **OK**.

- **Step 4:** Fill the form (https://docs.google.com/forms/d/e/1FAIpQLSfEvdJgrY-IFY8mG-bgv3_nccr11P2NRafI60Thga382E2yWA/viewform?usp=sf_link).

- **Step 5:** You will receive in the email that you informed the link to download.

- **Step 6:** Download the GenomicLand software.

- **Step 7:** Unzip the file. Open the extracted folder, find the file GenomicLand.exe, and click Sending a shortcut to the Desktop.

After completing these steps, the GenomicLand software will be ready for use.