

BC|GENOME Administration for both user account and data management tasks and logging of system events

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This document describes the tools for user and dataset management and for viewing BC|GENOME system logs.

The first chapter contains general information for all users about accessing BC|GENOME main page and about changing user's password.

From the second chapter onwards tools for database administrators are described with the information about user profiles, and the changes logged in BC|GENOME.

User login

To log in to BC | GENOME:

When you receive your ID and password you can log in to BC | GENOME.

- 1. Go to https://server/bcos/ (replace server with the name or IP address of your own server). The account information is provided by your local database owner (usually the project PI).
- 2. Enter your user ID.
- 3. Enter your password.

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Figure 1. BC|GENOME login page.

NOTE: You can enter an incorrect user name and password <10 times before access to BC | GENOME is locked.

Website certificate

To login, users go to the BCOS main page using the following address: https://server/bcos/.

When accessing the system for the first time the browser programs may show a warning message and ask whether to trust the provider of the certificate or not. This so called website certificate, displayed on the screen before logging into the database, is to notify the user of a guarantee of the secure data transfer. The certificate can either be provided by a third-party company, who sells the certificates, or the web certificate can be provided privately. As BC products are not public services, the website certificate is by default provided by Biocomputing Platforms Ltd. With both public and private certificates, the method of encryption (SSL, Secure Sockets Layer (SSL) technology) is exactly the same.

If the certificate is private, as with BC products, the browser program will show a warning message and ask whether to trust the provider of the certificate or not. To be able to proceed the certificate needs to be accepted for the BCOS URL (see Figure for details). If you want purchase the public certificate please contact support@bcplatfors.com.

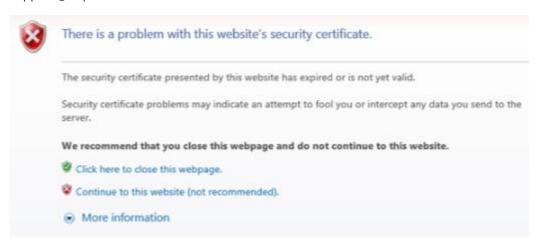


Figure 2. When you see the browser warning similar to this one about the untrusted certificate when accessing BC|GENOME select Continue to this website and accept the access to the BC|GENOME login page.

Changing of BC | GENOME user password

For safety reasons it is recommended to change the BCOS password regularly.

If your institution provides a centralized authentication system, and it has been integrated with BC|GENOME, follow the instructions provided by your local IT team.

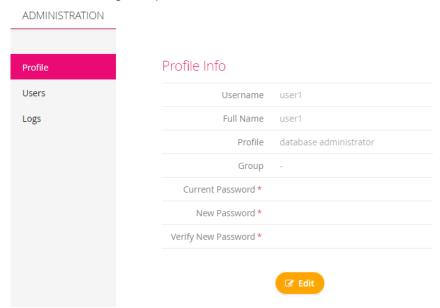
To change your password:

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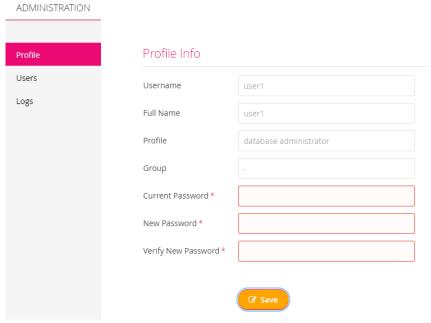
1. Go to **Administration** in the main page of BC | GENOME.



2. Press Edit to change the password



3. Type your current password



4. Type the new password and verify the change by retyping the password.

A password must be 7 (min) - 32 (max) characters long, and contain a combination of capital (A-Z) and lower case letters (a-z), and numbers (0-9).



NOTE: A good password is not easy to guess and should not be written down.

- 5. Press Save to store the new password
- 6. Press **OK**

When the new password matches the required criteria, you get a message to say your password is changed.

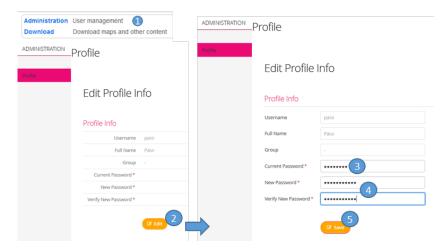


Figure 3. Changing user's password.

Database administration tools

In BC | GENOME Administration you can find a list of tools for user and data management tasks (see Figure 4):

- **Profile** for changing a user's password (not applied for the DBowner account)
- Users tool available for both DBowner and database administrator to generate new user accounts
- Groups tool for DBowner to manage user groups in BC | GENOME (DB2 role)
- Datasets list tool for DBowner to manage ownership of datasets
- Logs tool for DBowner (extendable for database administrator) to search for log information in BC|GENOME

The following chapters describe the tools in Administration.

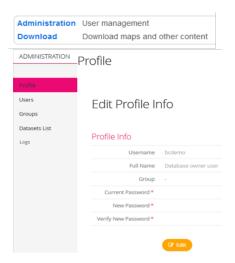




Figure 4. Administration tools of DBowner in BC|GENOME.

Creating a new user account with a profile

The DBowner and database administrator users can generate new user accounts for the BC|GENOME user interface. Technically, BC|GENOME user is a local linux user account. If you have a centralized user authentication system in use (for example, lightweight directory access protocol (LDAP), active directory (AD)), contact support@bcplatforms.com for more details about the configuration.

When a new user account is created in BC|GENOME, one of five user profiles must be specified (see Table 1). These profiles are specified in BC|GENOME to provide a set of tools to match a user's work profile and privileges. Table 1 describes user profiles and privileges.

Table 2 describes the data management and analysis tools for different user profiles.. In addition to profiles, the data permission tool can be used to manage user access to stored data in datasets in view, add and/or edit access level.

Table 1. User profiles in BC|GENOME

| Profile | Privileges | Notes |
|------------------------------|--|---|
| DBowner (set by BC support) | Tools for user, group and dataset permission management | Account credentials are provided by BC support. Not for data management and analysis work |
| Database administrator | User creation and tools for data management and analysis work | Cannot be used to change other user password |
| Researcher | Tools for data management and analysis tools | Profile for research work |
| Researcher with restrictions | Tools for data management and analysis but no tools for exporting data from the database | Analysis results can be exported from the result archive |
| Data entry | Tools for managing data entries in clinical work | BC CLIN user profile |
| Data viewer | Only for accessing and exporting data | Data sharing profile for collaboration purposes |



Table 2. Matrix for profile-specific user and data management permissions and tools

| Profile / Action | DBowner | Database administrator | Researcher | Researcher with restriction | Data entry | Data viewer |
|-----------------------------|---------|---------------------------|------------|-----------------------------------|------------|-------------|
| User managem | ent | | | | | |
| User Creation | Yes | Yes | No | No | No | No |
| User group management | Yes | No | No | No | No | No |
| Change any password | Yes | No | No | No | No | No |
| Change own password | No | Yes | Yes | Yes | Yes | Yes |
| Logs | Yes | No * | No | No | No | No |
| Data managem | ent | | | | | |
| Data structures | No | Yes | Yes | Yes | No | No |
| Folder management | Yes | Yes | Yes | Yes | No | No |
| Dataset creation | No | Yes | Yes | Yes | No | No |
| Dataset permission | Yes | Yes | Yes | Yes | No | No |
| Dataset removal | Yes | Yes | Yes | Yes | No | No |
| Data upload / edit | | | | | | |
| Single entry | No | Yes | Yes | Yes | Yes | No |
| File(s) upload | No | Yes | Yes | Yes | Yes | No |
| Delete entry / - ies* | No | Yes | Yes | Yes | No | No |
| Dataset export | No | Yes | Yes | No | No | Yes |
| Data queries | No | Yes | Yes | Yes | Yes | No |
| Download services | Yes | Yes | Yes | Yes | Yes | Yes |
| Data analysis | | | | | | |
| Statistical analyses | No | Yes | Yes | Yes | No | No |



| Storing | Yes | Yes | Yes | Yes | Yes | No |
|-----------------|-----|-----|-----|-----|-----|-----|
| scripts | No | Yes | Yes | Yes | No | No |
| Running scripts | INO | 163 | 163 | 163 | INO | NO |
| Running | No | Yes | Yes | Yes | Yes | Yes |
| reports | | | | | | |
| Result archive | Yes | Yes | Yes | Yes | Yes | Yes |
| sharing | | | | | | |

^{*} Contact BC support to change the default setting.

Changes in user accounts

When a BC|GENOME user account is created in BC|GENOME, BC support makes all user account modifications, such as:

- Change of user ID
- Change of user description
- Change of user profile
- Removal of user account from graphical user interface

Therefore, BC|GENOME is able to securely maintain the integrity of the system logs that are collected and stored.

Administration tools for user management

The DBowner and database administrator can add users, edit existing user profiles, manage groups and dataset lists using the Administration tools.

A regular user can only edit their own user profile(s) and logs.

From the main page, go to the **Administration** page to access the available tools.

Manage users

Select **Users** in the Administration page.

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Figure 5. Tools in Administration / Users

To change a user's password

- 1. Select the user you want to edit from the list of users.
- 2. Select **Edit**.
- 3. Type a new user password.
- 4. Type it again to verify the password.



Change user's password

User ID bcedu Full Name BC edu User's Profile researcher User's password * Verify user's password *



To add a new user:

Both DBowner and database administrator can add all new users and all users except administrators,

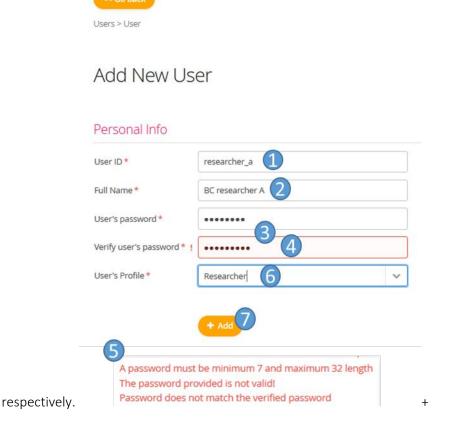


Figure 6. Specifying the user account information in Administration.

- 1. Select Add.
- 2. Type a User ID.

The user ID can contain 2 – 16 alphanumeric characters and underscores (_).

- 3. Type a full name for the user (max. 65 characters).
- 4. Type a user password.

A password must be 7 (min) - 32 (max) characters long, and contain a combination of capital (A-Z) and lower case letters (a-z), and numbers (0-9).

NOTE: A good password is not easy to guess and should not be written down.

- 5. Retype the password to verify it.
 - If there is no match between the passwords a red exclamation mark (!) sign is shown.
 - If the password violates password specifications, a pop-up help is shown.
- 6. Select a user profile from the drop-down menu (see Table 1 for a list of user profiles).
- 7. Press **Save** to store the new password.

Manage groups

In BC|GENOME, dataset user permissions in groups are managed by DBowner. Instead of granting dataset permissions individually, read or read/write permissions can be granted to a group of users. Technically, a group



is a type of role (R) in IBM's DB2 database

(https://www.ibm.com/support/knowledgecenter/en/SSEPGG_10.5.0/com.ibm.db2.luw.sql.rtn.doc/doc/r00522_86.html). One user can belong to several groups.

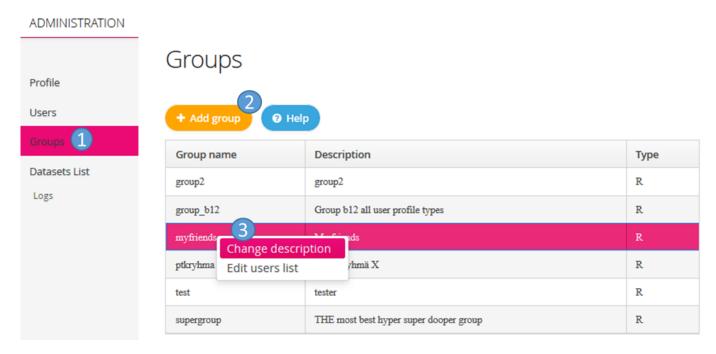


Figure 7. Group management tools.

To add a group:

- 1. Select **Groups**.
- 2. Select Add group.



Figure 8. Add a new user group

3. Type a name for the group.

A group name can contain 1-128 alphanumeric characters and underscores (_), and the name cannot start with a number.

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- 4. Type a description for the group.
- 5. Select **Add** to create a new group.

When a group has been specified (type of role, R) click the right mouse button to open the action menu for:

- Change description Edit the group description
- Edit users list To add or remove users from groups

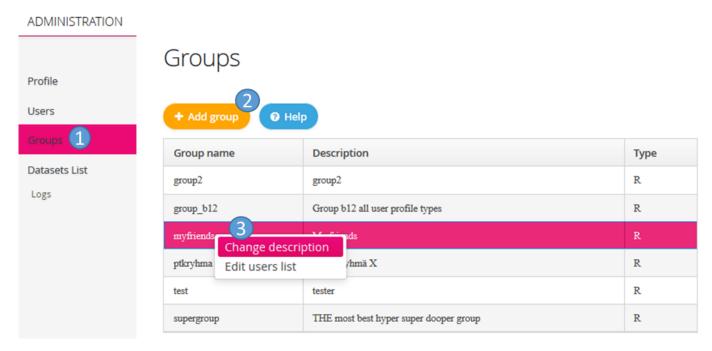


Figure 9. Edit a group.

To manage users in groups:

- 1. Select a group to manage.
- 2. Select one or more (CTRL+click) users to add to a group.

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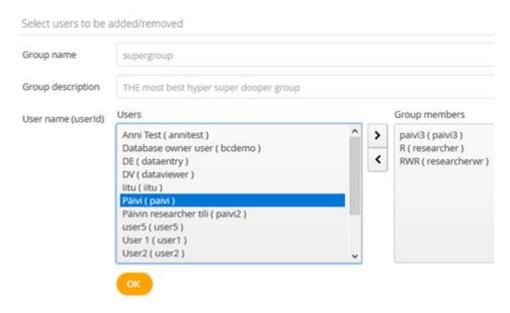
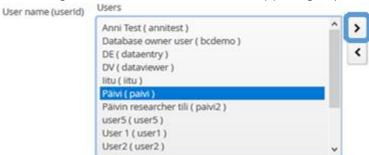


Figure 10. Administration / Groups: Edit user lists

3. Use the right arrow to add the selected user(s) to a group.



4. Use the left arrow to remove users from a group. You can select one or more subjects to be removed from a group.



5. Press **OK** to store changes.

If you want to remove unnecessary groups from graphical user interface, contact support@bcplatforms.com

Manage datasets

DBowner can grant ownership of a dataset to another user in BC|GENOME. The datasets list tool shows all datasets in the database.



To change ownership of a dataset:

- 1. Select Datasets List.
 - A list of datasets in BC | GENOME is shown.
- 2. Select a dataset whose ownership you want to change.
- 3. Right-click the dataset.
- 4. Select Change ownership.

Menu to specify up to 600 lines per page

Tools for browsing rows shown in the browser page

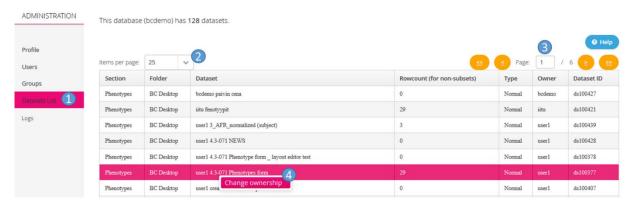


Figure 11. Administration / Datasets List.

Browser for BC | GENOME log information

The Logs tool is an administrative tool for viewing changes that are made in the BC|GENOME database.

DBowner can view system logs using Administration / Logs tool by events (Table 3). BC support can also configure the database administrator to access the Logs tool. Logs shown in BC|GENOME are collected from the BC|GENOME application server system log files in:

- /var/log/bcos/messages
- /var/log/bcos/gened.log
- /var/log/httpd/ssl_access_log
- /var/log/httpd/ssl error log
- /var/log/bcos/vaadin-servlet.log
- /var/log/bcos/vaadin-events.log

Figure 12 shows the features of Administration / Logs:

- 1. System logs for viewing change log information.
- 2. Refresh tool for updating log information.
- 3. Clear filters in the log table.
- 4. Click to view the menu to specify the columns displayed in the Logs table:
 - Event
 - Timestamp

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- Grantor
- Grantee
- Database
- Message
- Job id
- Client IP
- Type
- 5. Filter the Logs view by events, (see Table 3).
- 6. Press the column title to sort information in either ascending or descending order, or type a field value to search for.
- 7. Press the **Event** field to view the summary collected from column fields.

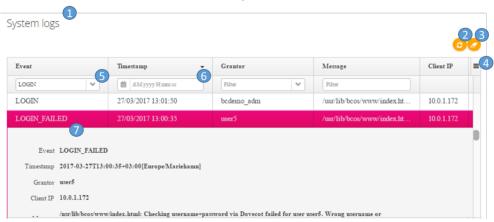


Figure 12. Administration / Logs for viewing system logs.

Table 3 shows the events listed when you select the **Event** drop-down menu in the **Logs** page.

Table 3. System events generated in BC|GENOME Administrations / Logs.

| E | Author to DOLOFNOME |
|----------------------------|---|
| Event | Action in BC GENOME |
| Password_change | Administration: Profile / Edit; Users / |
| | Edit |
| User Create | Administration: Users / add |
| User_access | Datasets / permissions change (access |
| _ | / revoke) |
| Create_user_group | Administration: Groups /new |
| Remove_user_from_group | Administration: Groups / edit |
| Dataset_create | Datasets / new |
| Dataset_moved | Datasets / move to trash, or Folders / |
| _ | move datasets |
| Dataset_interactive_export | Datasets / export to the next page |
| Backup_sys_tables | DB2 backup script for system tables |
| | started / completed |
| Empy_Trash | Indication of a job completed when |
| | using the Tools / empty trash tool |
| Fileserver_upload | Tools / file transfer |
| Dataset_privilege | When creating a new dataset |

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| Logout | User logout |
|-----------------------------------|---|
| Login | User login |
| Login_failed | Login with a wrong user ID / password |
| Drop_table | Tools / empty trash for a dataset |
| Drop_view | Tools / empty trash for a subset |
| Create_view | Dataset / subset |
| Create_subset | Dataset / subset |
| Job_started | Job submitted in the analysis GUI: Data input / large files; analysis jobs |
| Job_failed | Tools / cancel jobs when the job of software is in run |
| Job_success | Completed analysis job(s) in Tools / result archive |
| Data_viewed | Subjects: Multi-subject browser or Phenotypes: Datasets / new datagrid |
| Result_folder_command | Download of a file in Tools / result archive; or delete result folder content; or user shared result folder content |
| Result_folder_upload_to_a_dataset | Data input / one file, or edit row, or add row, or delete row |
| User_logout | Event generated by the logout tool |
| Open_dataset | Datasets / open |
| Snapshot_create | Datasets / snapshot |
| Add_user_to_group | Administration: Groups / add groups |