# Storage of data on the Bioimg server

## a. Exporting the data:
**1.** Create in "D:\UserData\" a folder with your exact User name (in small letters!)
**2.** Export your data from BD FACSDiva in FCS file format or in Experiment format (depending on requirements) into the folder created.
**3.** Wait for the data to be backed up by the Bioimg server (up to two hours)

## B.1. Accessing the data: (PC)
**1**. Right click on "My Computer"
**2.** Choose "Map Network Drive"
**3.** Choose "Drive: Z:"
**4.** Folder:  \\bioimg\User name
**5.** Click on "Finish"
**6.** Wait a few seconds until "**Connect to bioimg**" window appears
**7**. Write your **User Name**
**8**. Write the Password **User Name1**
**9.** The folder of your user name on bioimg data will open.
Copy the folder required by dragging it.

## B.2. Accessing the data: (MAC)
1.  Go to "Go>Connect to server"
2.  Write: "smb://bioimg"
**3.**  Wait a few seconds until **Connect to bioimg** window appears
**4**.  Write your **User Name**
**5**.  Write the Password **User Name1**

## C. Disconnect from bioimg
**1.** Open "My Computer"
**2.** Right click on your User Name on 'BioImaging Data (bioimg)' (Z:) and disconnect
**3.** On the MAC put the connected server in the trash

## D. Accessing and managing the data on the BioImaging Storage Server
**1**. Open the chrome internet browser and go to "bioimg.weizmann.ac.il"
**2.** Logon with your User name and your regular Weizmann password
**3.** Use "annotate Raw Data" to add experiment to the database by assigning it to a project.
**4.** Use "Frequently asked Questions" for help