## The License Agreement for PANDA and PANDA-view installation

This Agreement is a legal agreement on "protein quantitative analysis" software products (PANDA and PANDA-view, collectively called the "Software" below) between you (individual or a single entity) and National Center for Protein Sciences (Beijing)-Bioinformatics group. Once you install, copy or otherwise use this software product, you agree to be bound by the terms of the agreement. If you disagree with this agreement, you can not get the right to use this software product.

## 1 Software Product Protection Clause

- 1.1 This software product is developed by the National Center for Protein Sciences (Beijing)-Bioinformatics group. All title and intellectual property rights, which is generated by the software product (including, but not limited to the software products, including any relative images, data, texts and additional programs (dll, exe, etc.)), incidental help materials, and any copies of the Software Products is protected by the copyright laws in People's Republic of China and international copyright treaties and other intellectual property laws and treaties. Users only get the right to use this software product.
- 1.2 Users may not be detrimental to National Center for Protein Sciences (Beijing)-Bioinformatics group's rights and interests, and may not implement the following acts (including but not limited to):
  - delete the information of the copyright on the software and all the other copies
  - reverse engineering on the software, for example, disassemble or decompile the programs
  - modify or destroy the status quo of this software
- 1.3 If the user does not comply with any provision of this Agreement, National Center for Protein Sciences (Beijing)-Bioinformatics group has the right to immediately terminate this agreement, and the right to be held accountable through legal channels is reserved.
- 1.4 If you need to carry out commercial sales, reproduce and distribute, for example, preinstalled and bundled software, you must obtain National Center for Protein Sciences (Beijing)-Bioinformatics group's written authorization and permission.
- 1.5 Users who use this software should have the required operating conditions of the software product, where the user must ensure the right to use the Windows operating system and other software including Thermo MSFileReader. Within the maximum extent

permitted by applicable law, special, incidental, direct or indirect infringement or loss resulting from the use or inability to use this software product that occurs in any case are not liable for damages by National Center for Protein Sciences (Beijing)-Bioinformatics group. Even if the possibility of such damages are advised in advance.

1.6 The defined information include: text, software, sound, photograph, video, graphic, all the contents in the advertisement and we provide users with business information. All the elements are protected by copyright, trademark right, other intellectual property rights and ownership laws. Therefore, the user can only use the content under the authority by us and our advertisers, but not unauthorized copy, modify, compile the content, or create content-related derivative products.

## 2 User Notice

Reminding the user especially, the use of the software must be complied with the relevant state policies and laws, such as criminal law, national security law, privacy law, computer information system security protection regulations, intellectual property law, must protect the national interests and security. Users should take full responsibility for illegal use of the software.

- 2.1 National Center for Protein Sciences (Beijing)-Bioinformatics group appeal that all users follow our designated website or specify way to download and install the software products. Beware of non-designated site to download the software, so computer infection can damage a user's computer and retrieve user data privacy by malicious programs.
- 2.2 National Center for Protein Sciences (Beijing)-Bioinformatics group pay attention to user privacy protection, and ensure that the software product does not contain any function code ,which will track, monitor user's computer or operating behavior ,and we do not monitor user's online behavior, offline behavior or leak user's privacy.
- 3 National Center for Protein Sciences (Beijing)-Bioinformatics group draws particular attention to the user: in order to protect the autonomy of the business development and adjustment, National Center for Protein Sciences (Beijing)-Bioinformatics group has the right to modify or discontinue the service without noticing to the user, and National Center for Protein Sciences (Beijing)-Bioinformatics group exercise right to modify or discontinue the service without be responsible for users or any third party. Users must agree to the terms under the premise, then National Center for Protein Sciences (Beijing)-Bioinformatics group begins to provide services to users.
- 4 National Center for Protein Sciences (Beijing)-Bioinformatics group solemnly reminds users of paying attention to this Agreement, which exempts the responsibility of National Center for Protein Sciences (Beijing)-Bioinformatics

group and aggravates the obligation from the terms of users, please read carefully, consider the risks independently. Minors should read this agreement, accompanied by legal guardian. The right of final interpretation and modification of the provisions of the above content are owned by National Center for Protein Sciences (Beijing)-Bioinformatics group for all.

5 If you have any comment to National Center for Protein Sciences (Beijing)-Bioinformatics group and the software product, please contact National Center for Protein Sciences (Beijing)-Bioinformatics group (8610-61777053), Dr. Cheng Chang will be at your service (email: 1987ccpacer@gmail.com).

Address: NO. 38, Life Science Park Road, Changping District, Beijing, China. Code: 102206

National Center for Protein Sciences (Beijing)

Beijing Proteome Research Center

State Key Laboratory of Proteomics

**Bioinformatics Group** 

Dr. Cheng Chang