中国科学院计算技术研究所
INSTITUTE OF COMPUTING TECHNOLOGY , CHINESE ACADEMY OF SCIENCES

pFind Studio

pFind 3 User Guide

Version 3.1.2

pFind Group

2018.4

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- Before software installation
- Software registration
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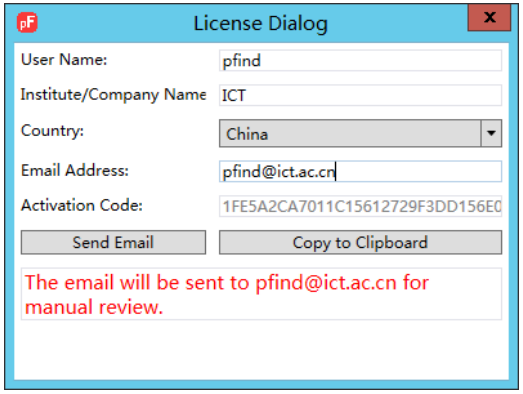
Before software installation

- **Please install 64-bit MSFileReader first**

- **Windows 7 and above**
  - **64 bit version**
  - **.NET Framework 4.5 environment**

Software registration

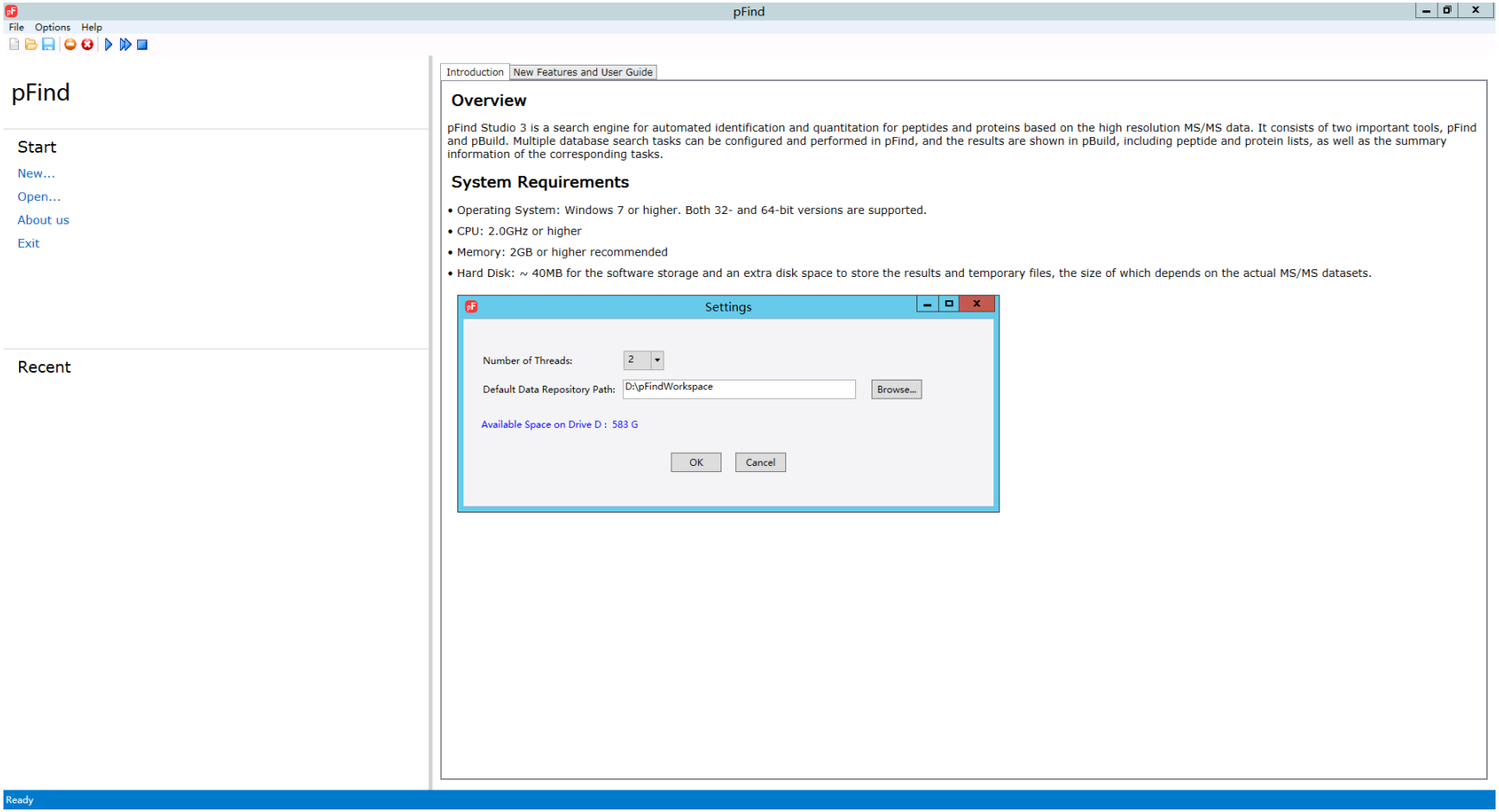
- Click pFind.exe and fill in personal information in the following information panel.



- Send the information to pfind@ict.ac.cn
- Apply for pFind.license.
- Put the pFind.license into pFind installation directory under the \pFindStudio\bin folder.

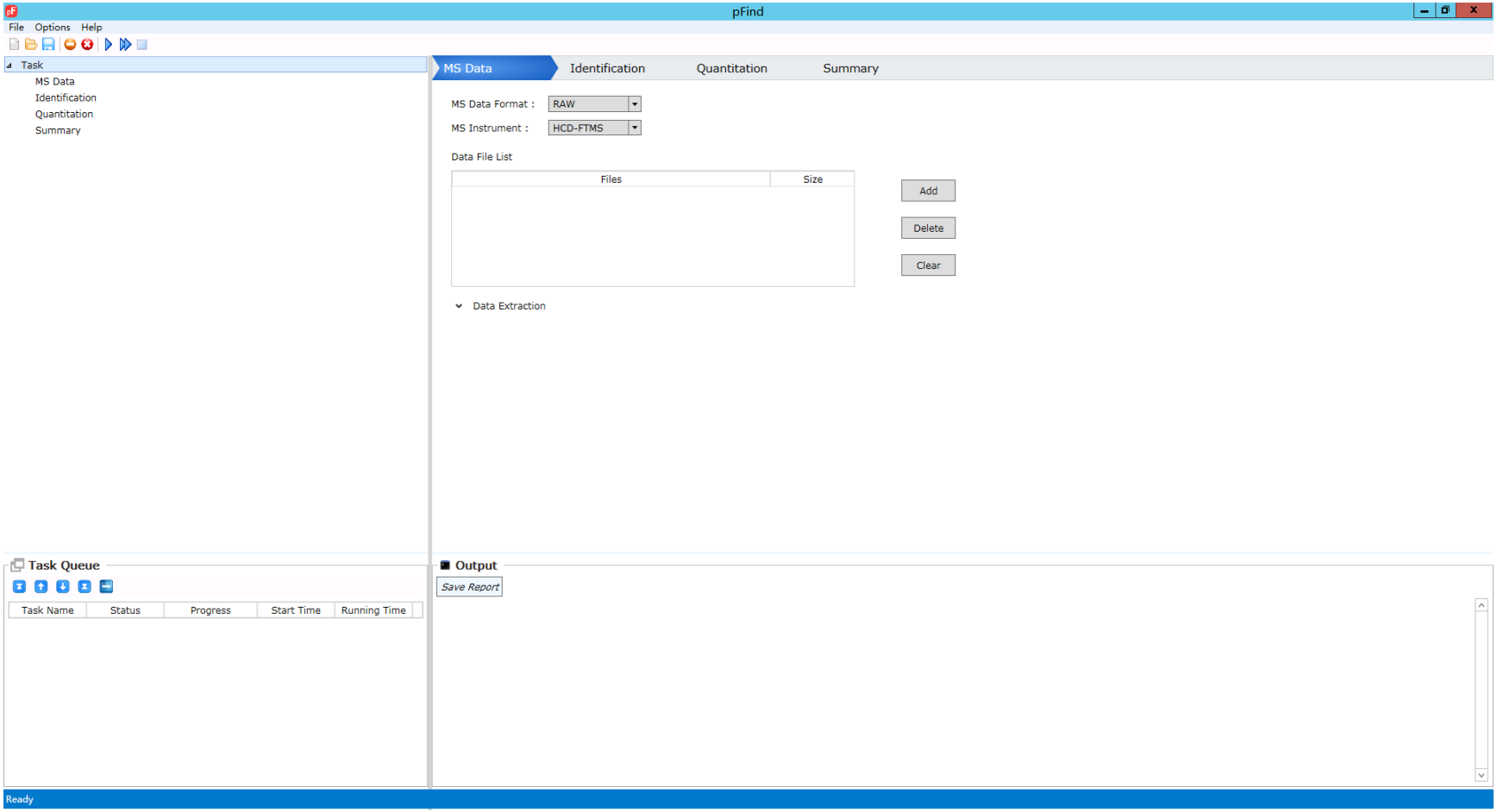
Main interface of pFind 3

- **Select the number of threads and search task storage path according to your computer situation.**



Create a new search task

❖ **Click File→New**



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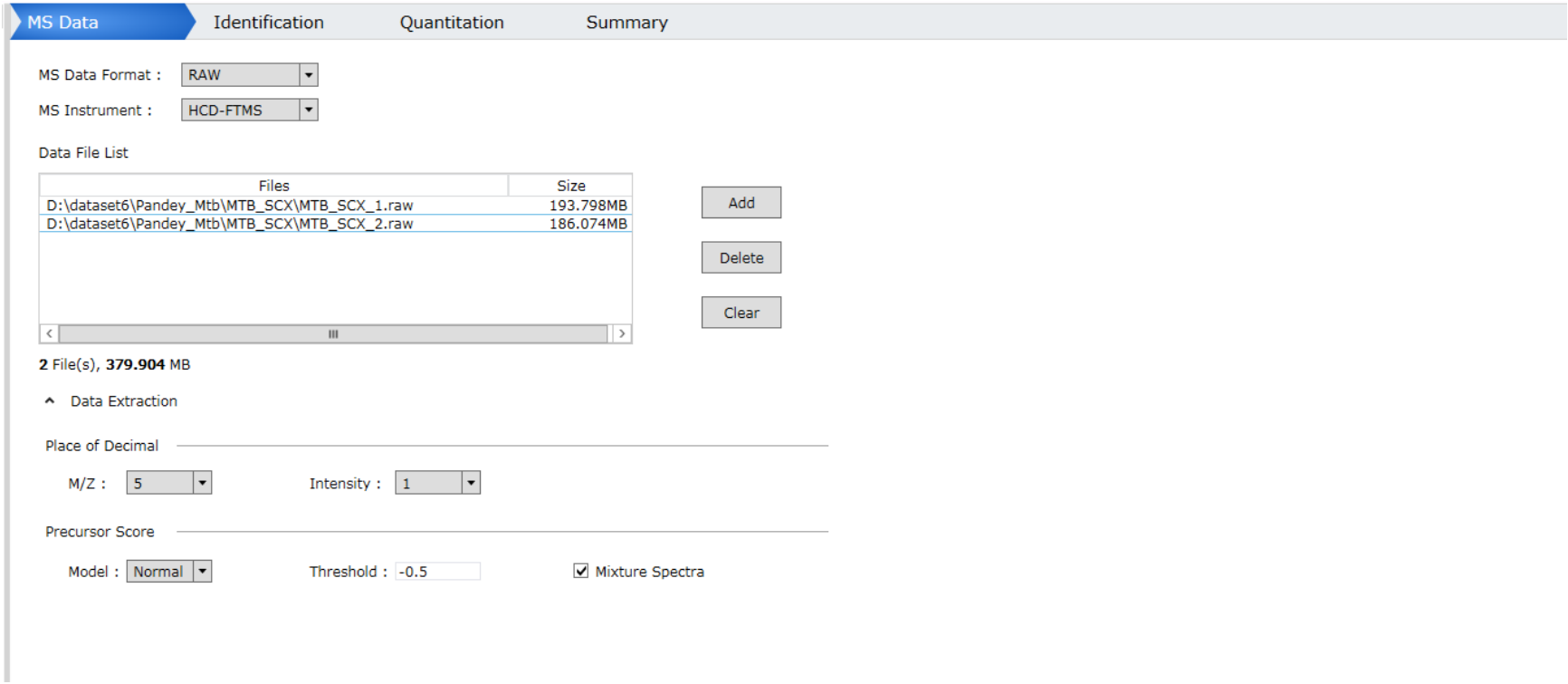
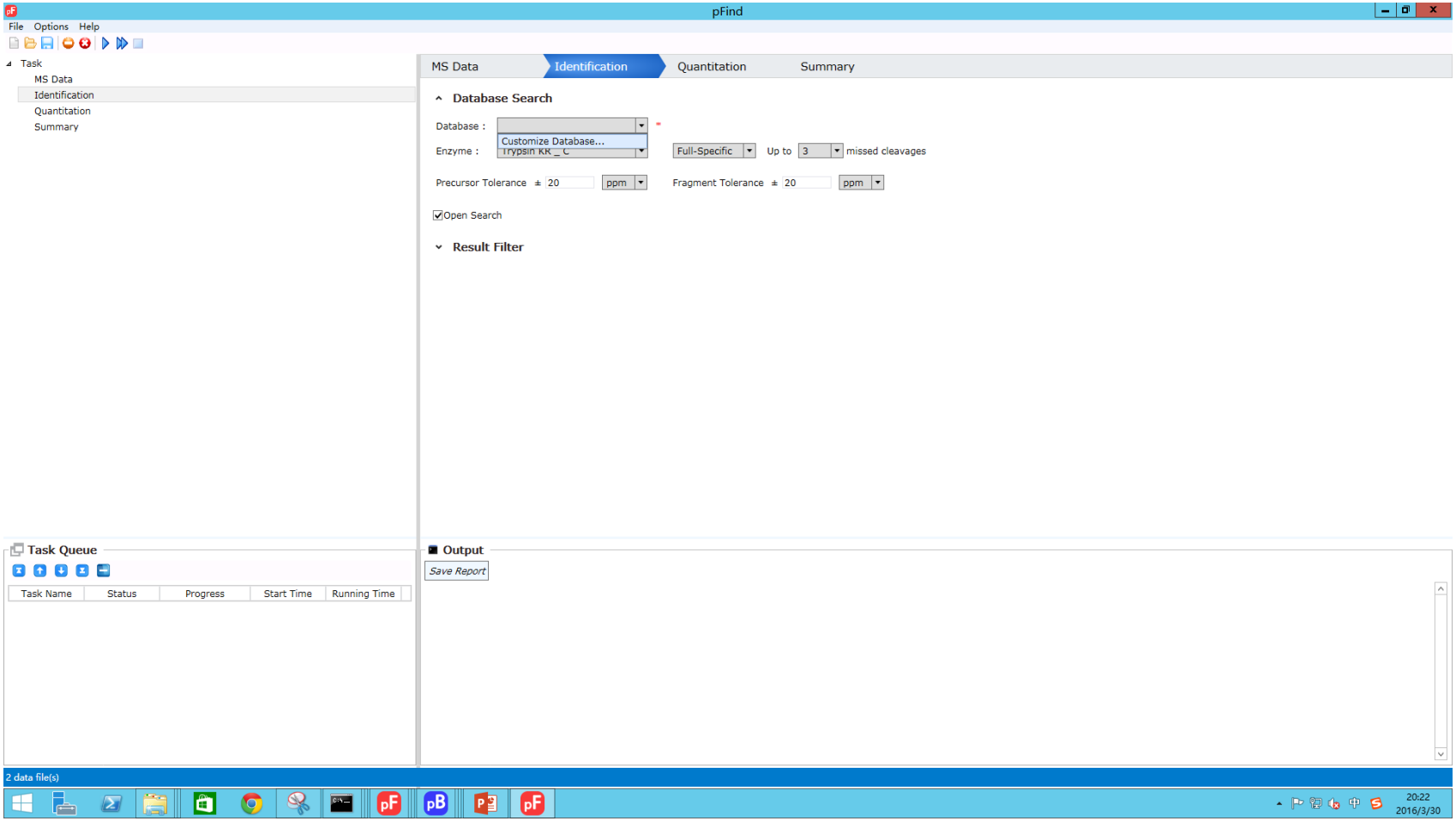
1. Data import and preprocess.

- **You can choose MS data format and other data preprocessing type in MS Data panel.**



2. Set search parameters

❖ A) Select and import database.



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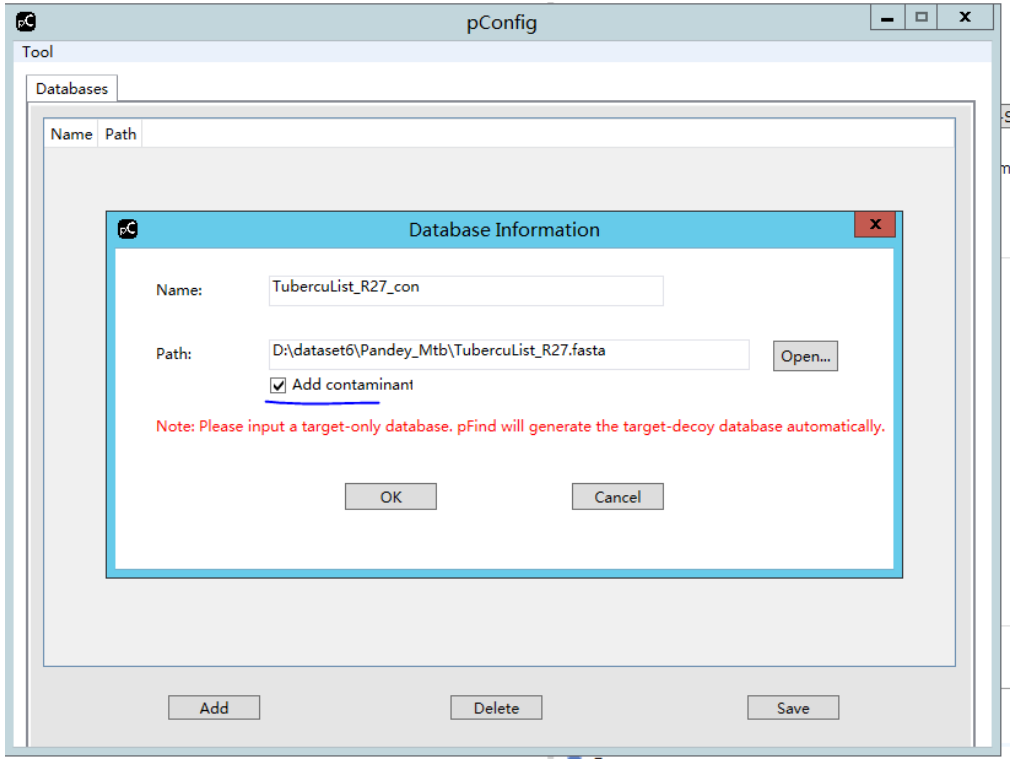
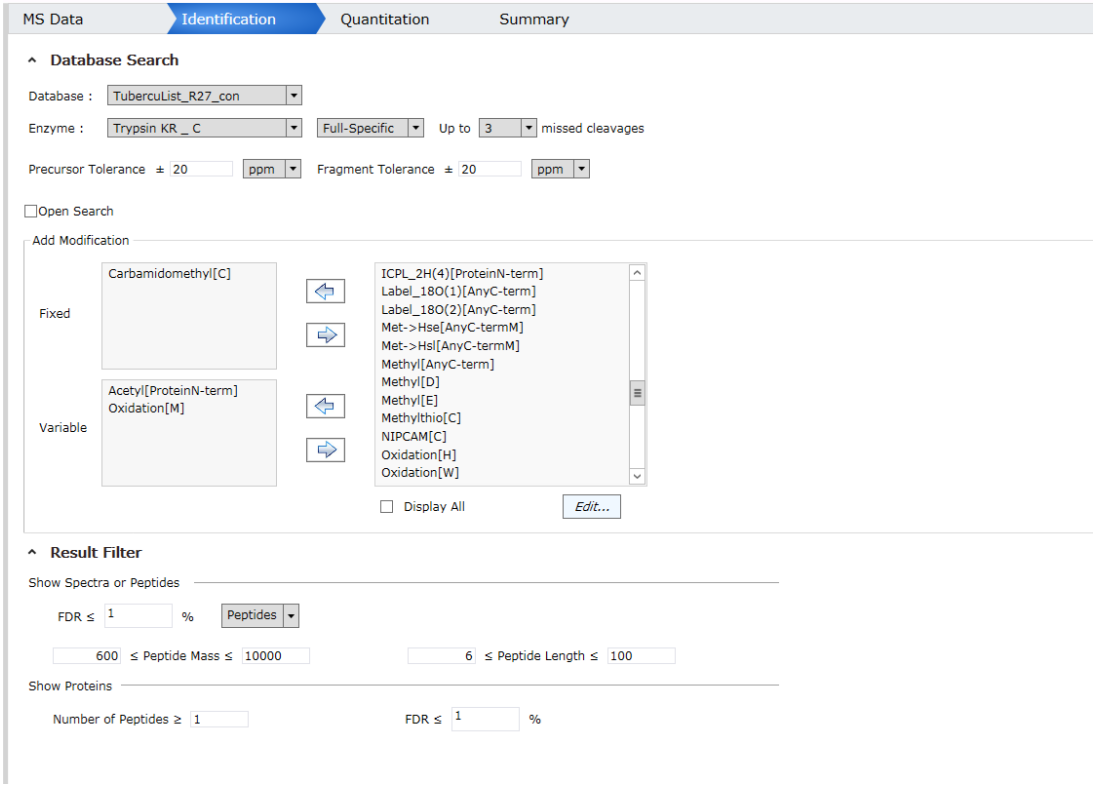
2. Set search parameters

- **A) Select and import database.**
  - **Add contaminated proteins to the database if it doesn't contain them.**



2. Set search parameters

❖ **B) Select the appropriate error range and set the suitable modification and filtering parameters.**



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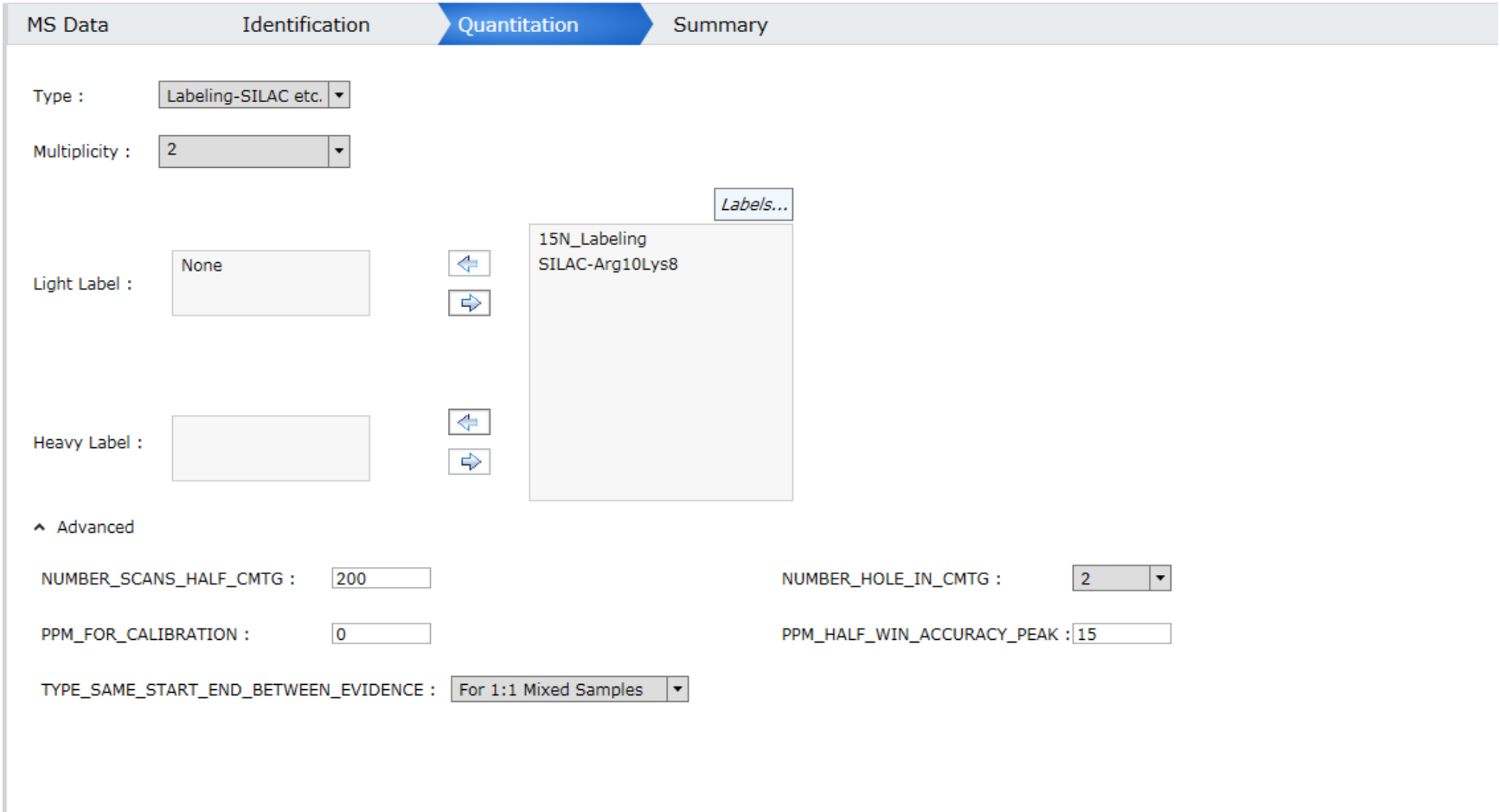
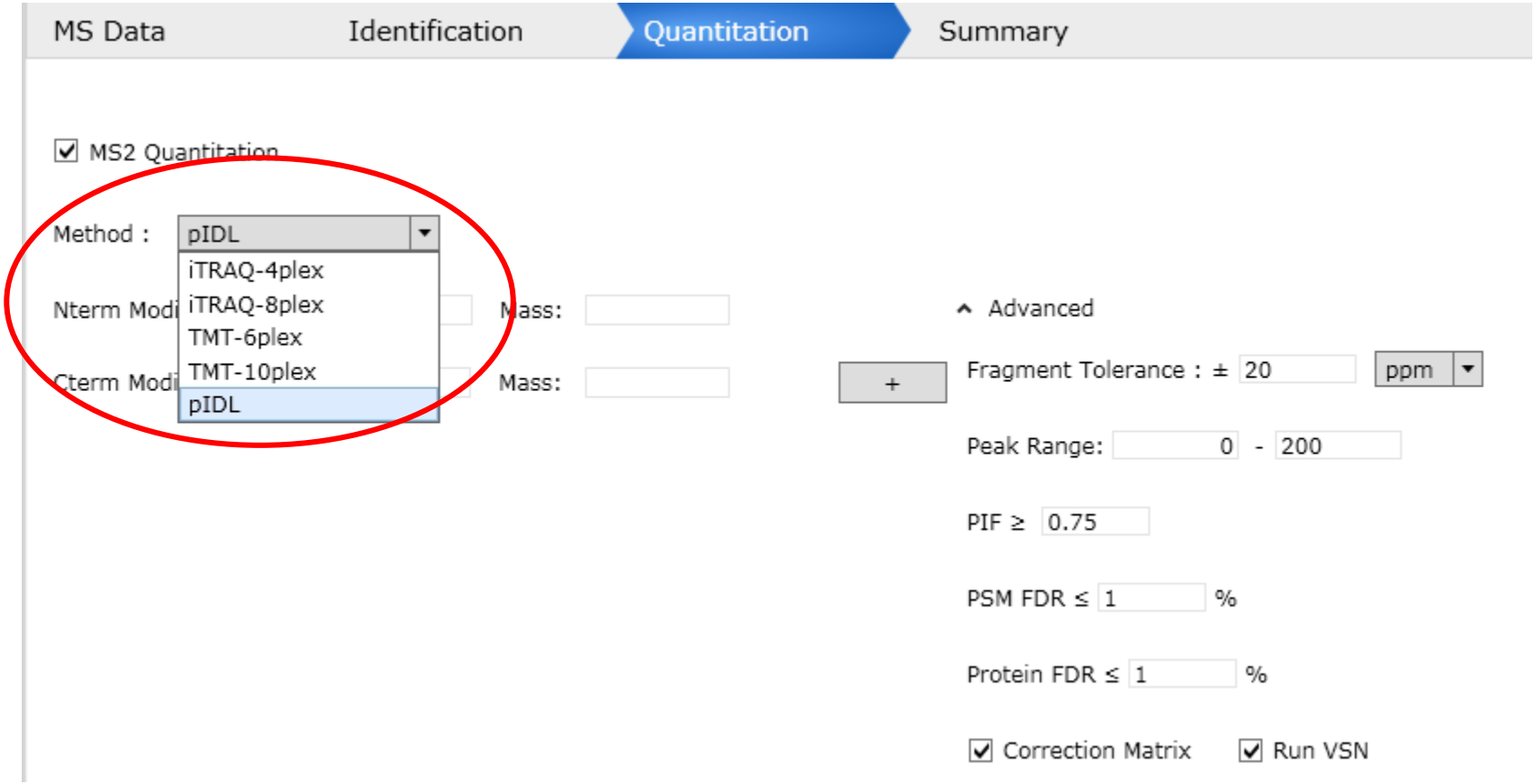
3. Set quantification parameters – MS1

- **SILAC and 15N labeling are supported if necessary**



3. Set quantification parameters - MS2



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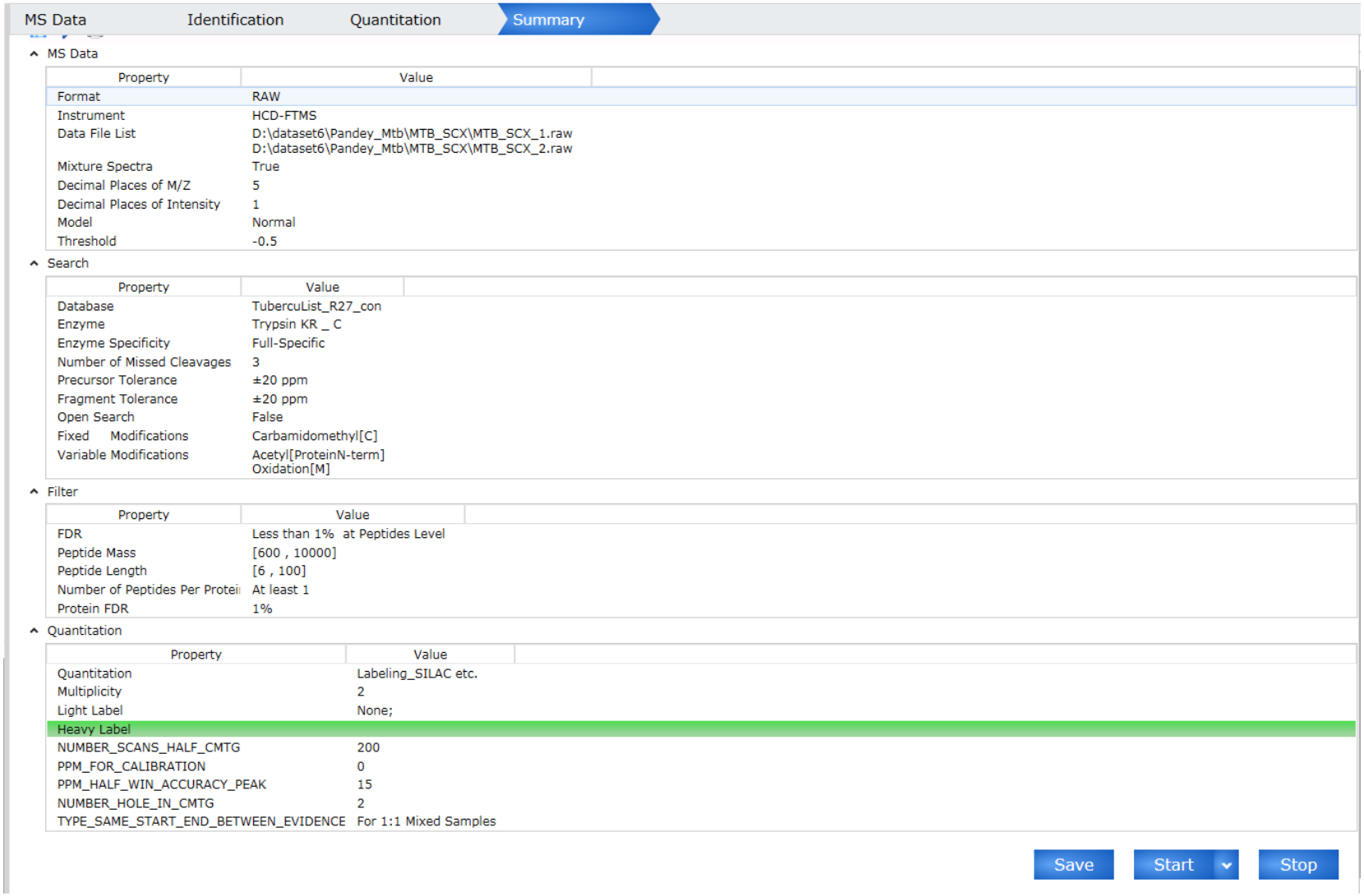
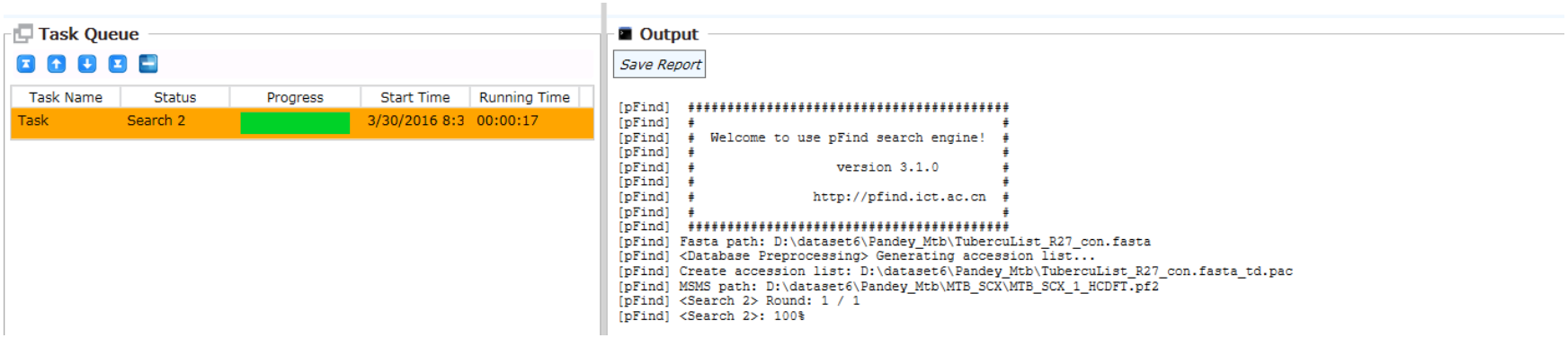
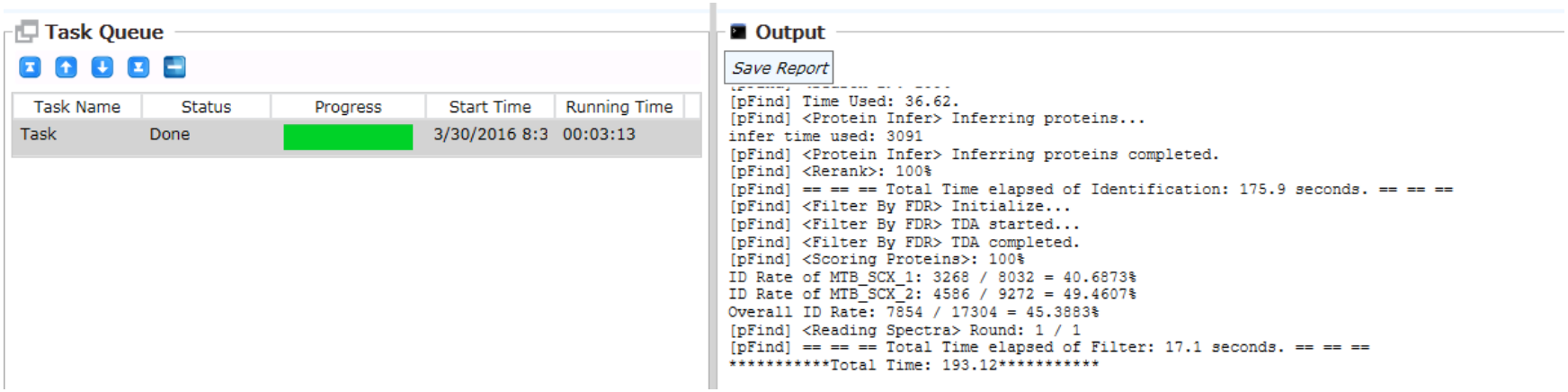
4. Check parameters and run tasks



Searching…


Task Queue
Task Name
Status
Progress
Start Time
Running Time
Task
Search 2
3/30/2016 8:3
00:00:17
Output
Save Report
[pFind] ##
[pFind] #   #
[pFind] # Welcome to use pFind search engine! #
[pFind] #   #
[pFind] #   version 3.1.0   #
[pFind] #   #
[pFind] #   http://pfind.ict.ac.cn #
[pFind] #   #
[pFind] ##
[pFind] Fasta path: D:\dataset6\Pandey_Mtb\TubercuList_R27_con.fasta
[pFind] <Database Preprocessing> Generating accession list...
[pFind] Create accession list: D:\dataset6\Pandey_Mtb\TubercuList_R27_con.fasta_td.pac
[pFind] MSMS path: D:\dataset6\Pandey_Mtb\MTB_SCX\MTB_SCX_1_HCDFT.pf2
[pFind] <Search 2> Round: 1 / 1
[pFind] <Search 2>: 100%

Searching completed

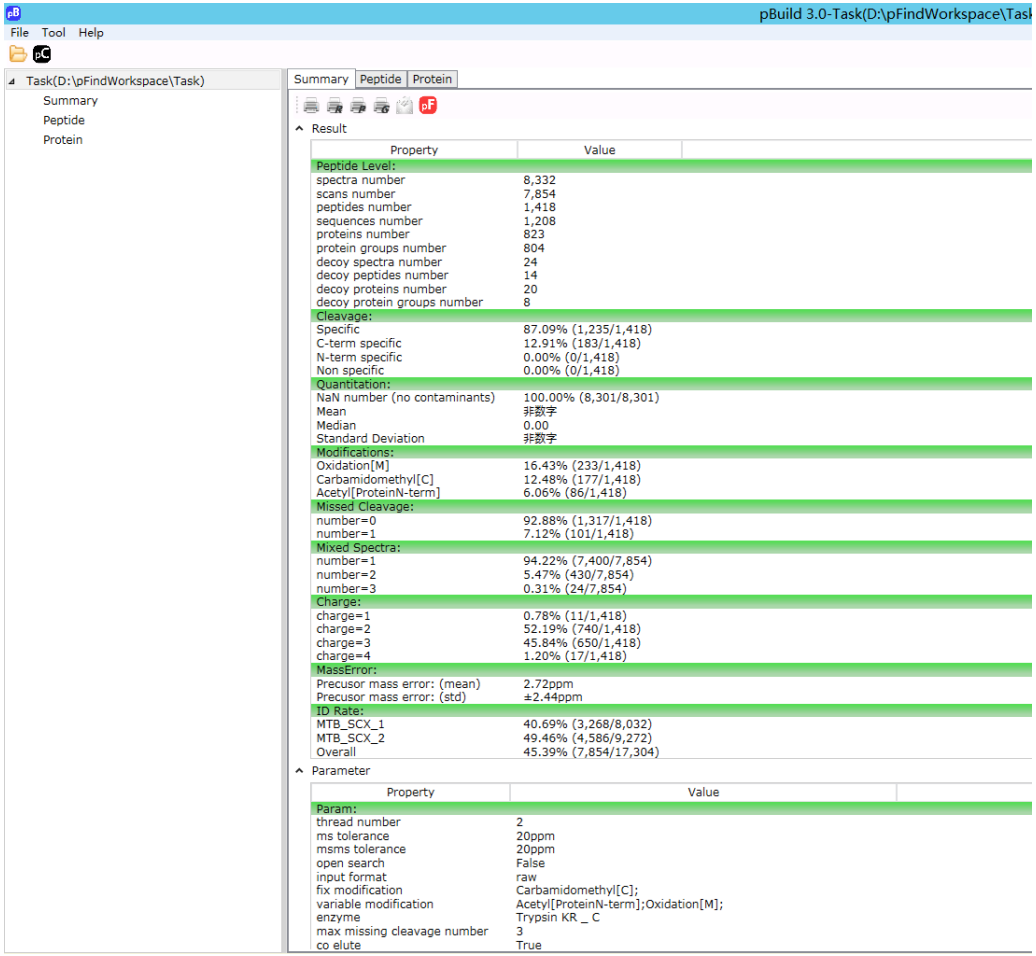


Contents of search results files

- **pFind.spectra**
  - Search results of each PSM. Such as protein names, modifications, E-value and so on.
- **pFind.protein**
  - Protein group information and identified PSMs.
- **pFind.summary**
  - Statistical information of modifications, identification rate, missed cleavage and so on.

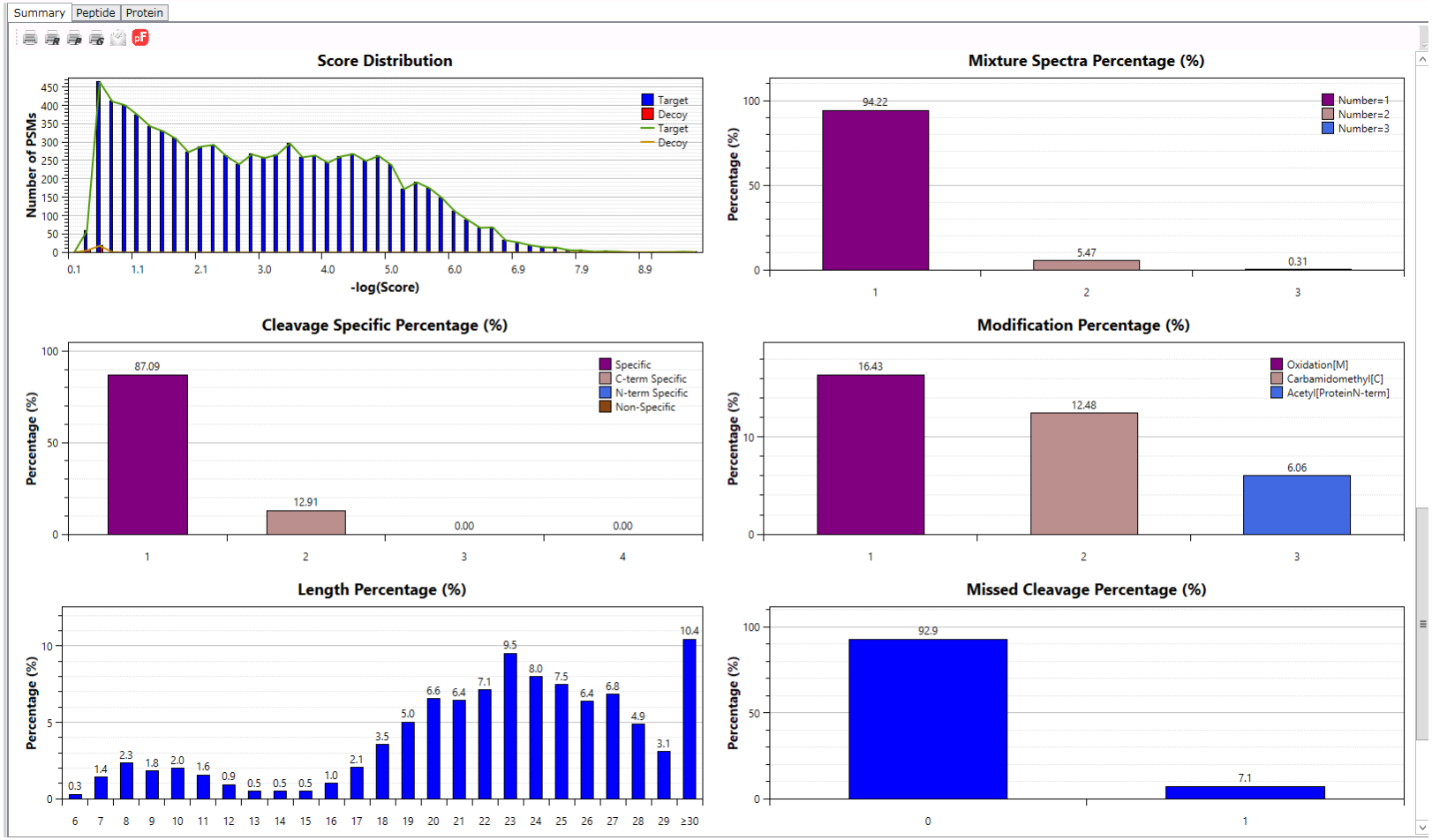
View search results

❖ pBuild: Result statistics



View search results

❖ pBuild: The summary panel



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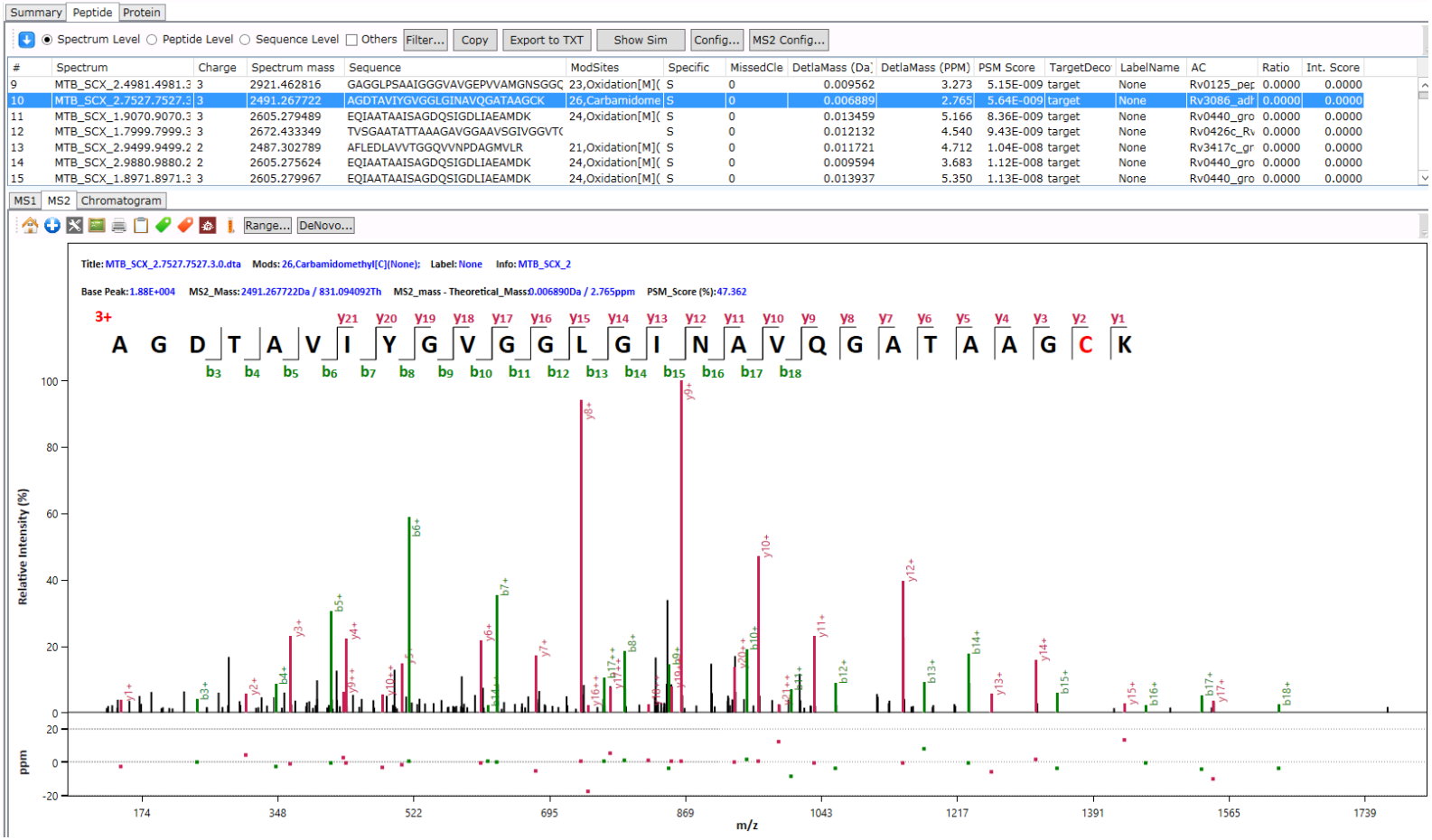
View search results

❖ pBuild: Peptide spectrum matching



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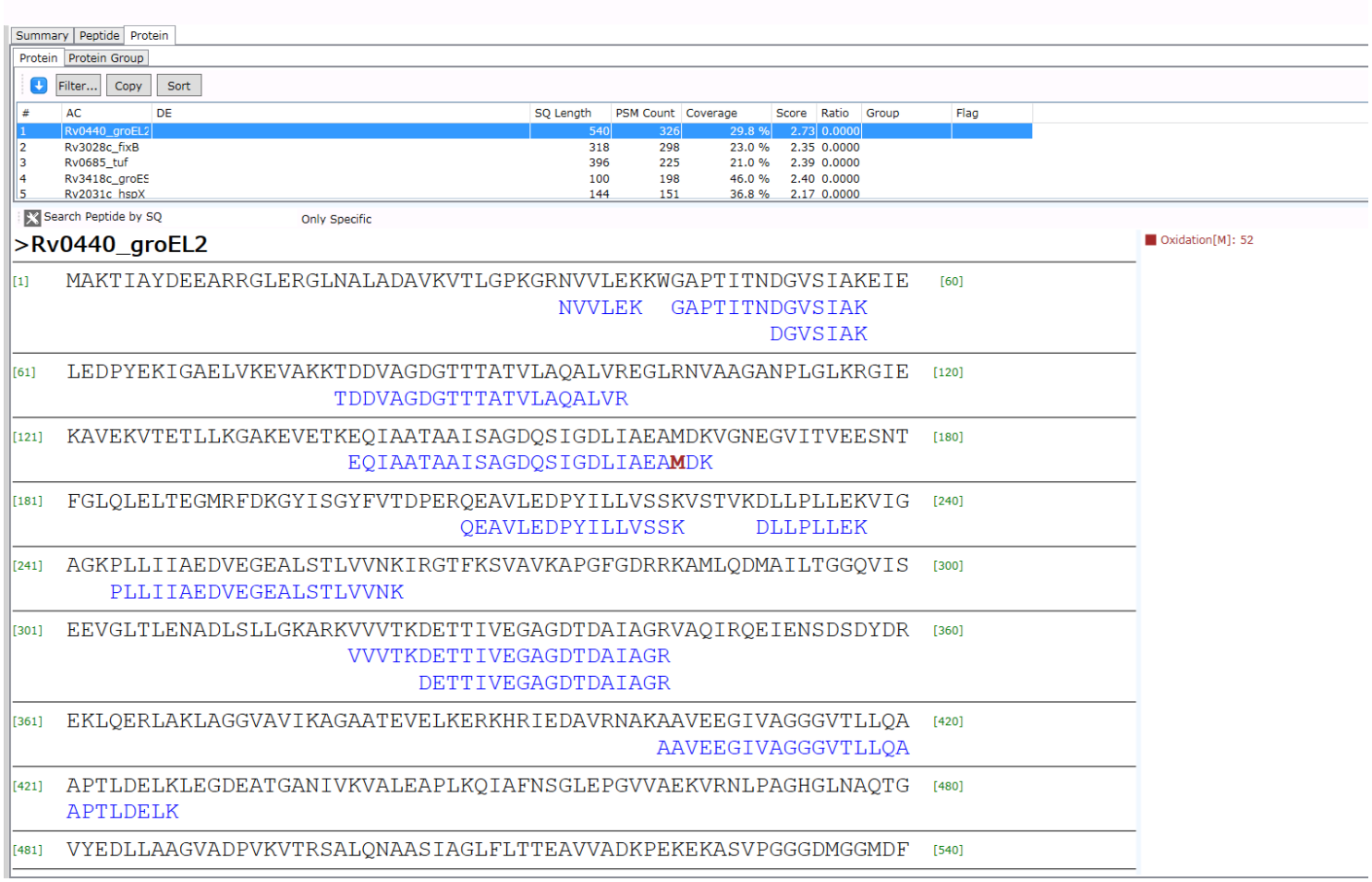
View search results

❖ pBuild: Protein coverage



- **Thank you for using pFind 3!**

- **If you have any questions, please contact pfind@ict.ac.cn.**