# SuperQuant node installation

## **IMPORTANT NOTE**

The node was built for Thermo Proteome Discoverer 2.0.x and can't work in earlier versions of the software.

# **Installation process**

- 1. Close Proteome Discoverer if it is open.
- 2. Copy the node dll file (**SuperQuantNode.dll**) and exclusion list file (**cf.exclude**) to the folder with Proteome Discoverer. The default location is:
  - C:  $\Program Files\Thermo\Proteome Discoverer 2.0\System\Release$

The path on your system might be slightly different. You might need to use administrator account to copy the files.

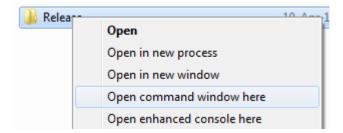
3. Use either of two options (step 4) or (step 5).

#### 4. Step 4

a. Open the command window and move to the Proteome Discoverer folder.



In Windows Explorer you can use Shift + right click on the folder and select **Open command** window here. The default location is mentioned at step 1.



b. Run the following command

Thermo. Magellan. Server -install

c. You should see the console window with the list of all nodes found in the system. You should see **Complementary Finder** and **MSn Deconvolution** nodes in the list.

```
- - X
C:\windows\system32\cmd.exe
  NFO - ApplicationExtensionManager: Found 68 extensions
NFO - ProcessingNodeManager: Searching for Nodes in directory: C:\Program File
\Thermo\Proteome Discoverer 2.0\System\Release
NFO - ProcessingNodeManager: Found 53 processing nodes
NFO - ProcessingNodeManager: Node: Complementary Finder registered
NFO - ProcessingNodeManager: Node: MSn Deconvolution registered
                                                                                          Spectrum File Injection (non mode) register
                     rocessingnouerianager.
                                                                           noue -
  NFO
                   ProcessingNodeManager: Node: Spectrum File Injection (Spectrum Mode) reg
  stered
                                                                            Node: Ga MM Finder registered
                   ProcessingNodeManager:
                                                                                         Spectrum Scalpel registered
MS Amanda registered
ptmRS registered
Annotation DB Updater registered
To ProteinCenter registered
Controlled Protein List Retrieval registere
                   ProcessingNodeManager:
ProcessingNodeManager:
ProcessingNodeManager:
ProcessingNodeManager:
ProcessingNodeManager:
ProcessingNodeManager:
                                                                           Node:
Node:
Node:
   NF0
   NFO
NFO
                                                                           Node:
                                                                            Node:
                                                                           Node:
                   ProcessingNodeManager:
                                                                                           ProteinCenter Annotation registered
                   ProcessingNodeManager:
                                                                                          Spectrum Library Remover registered
FASTA File Creator registered
FASTA File Remover registered
Spectrum Library Creator registered
Spectrum Confidence Filter registered
                                                                           Node:
Node:
   NF0
                                                                            Node:
                                                                           Node:
                                                                           Node:
Node:
Node:
                                                                                           FASTA Index Creator registered
                                                                                           Sequest HT registered
Fixed Value PSM Validator registered
                                                                           Node:
Node:
Node:
                                                                                          Mascot registered
Target Decoy PSM Validator registered
Percolator registered
                   ProcessingNodeManager:
                                                                           Node: MSPepSearch registered
```

#### NOTE:

In some cases the execution of the command results in the following warning and the installation fails

```
WARN [1] Services. ProcessingNodeManager - Could not load this assembly "SuperQuantNode.dll" since the following exception is raised

System. IO. FileLoadException: Could not load file or assembly 'file:///C:\Program

Files\Thermo\Proteome Discoverer 2.0\System\Release\SuperQuantNode.dll' or one of its dependencies. Operation is not supported. (Exception from HRESULT: 0x80131515)

File name: 'file:///C:\Program Files\Thermo\Proteome Discoverer

2.0\System\Release\SuperQuantNode.dll' ---> System.NotSupportedException: An attempt was made to load an assembly from a network location which would have caused the assembly to be sandboxed in previous versions of the .NET Framework. This release of the .NET Framework does not enable CAS policy by default, so this load may be dangerous. If this load is not intended to sandbox the assembly, please enable the loadFromRemoteSources switch. See http://go.microsoft.com/fwlink/?LinkId=155569 for more information.
```

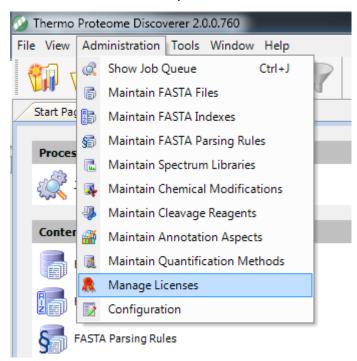
The reason is the policy of .NET environment that prevents libraries downloaded from web sources to be executed. To override this policy you should add the following line to Thermo. Magel lan. Server. exe. config file that could be found in the Proteome Discoverer folder. Since this is XML file you should put it in <configuration> <runtime> section of it.

```
<loadFromRemoteSources enabled="true"/>
```

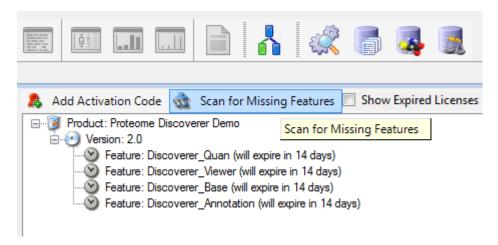
The final result should similar to this.

## 5. Step 5

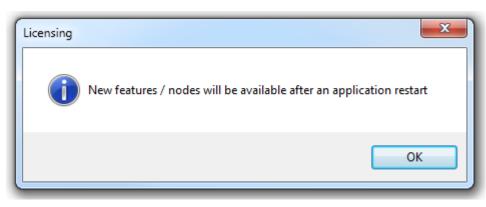
- a. Run Proteome Discoverer.
- b. In the main menu open **Administration** and then find **Manage Licenses**.



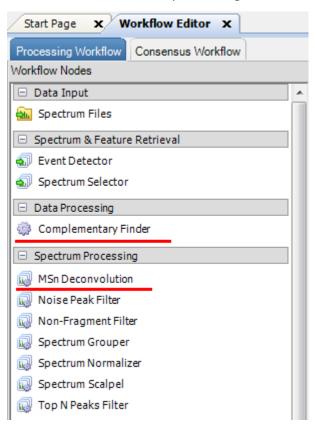
c. Click **Scan for Missing Features** on the top of the window.



d. After a while you will see the following notification



- 6. Restart Proteome Discoverer
- 7. You should find two new processing nodes: **Complementary Finder** and **MSn Deconvolution**



8. If you used Step 5 and can't see the nodes in the list afterwards, consider reading NOTE from the step 4.