

# API Documentation

## Service API Documentation

*Developers can reuse the function of our software quickly by use Service API.*

### 1 Get minimal gene set

URL	/api/cegRecord/fetchCeg.php
GET	id
Return	Result in json style. Status 0 means execute successfully, and 1 means error. Data is the minimal gene set filtered.
Example	http://yourHost.com/api/cegRecord/fetchCeg.php?id=01020304

### 2 Compare two minimal gene set

URL	/api/cegRecord/compare.php
GET	id
Return	Result in json style. Status 0 means execute successfully, and 1 means error. Data is the minimal gene set. Statics is the statistical value.
Example	http://yourHost.com/api/cegRecord/compare.php?id=01020304+05060708

### 3 Download minimal gene set with CSV format

URL	api/download/cegCSV.php
GET	id
Return	File with CSV format.
Example	http://yourHost.com/api/download/cegCSV.php?id=01020304

### 4 Download minimal gene set with XML format

URL	api/download/cegXML.php
GET	id
Return	File with XML format.
Example	http://yourHost.com/api/download/cegXML.php?id=01020304

## 5 Get module with whole minimal gene set

<b>URL</b>	<b>api/kegg/entireModule.php</b>
<b>GET</b>	<b>id</b>
<b>Return</b>	Result in json style. Status 0 means execute successfully, and 1 means error. Data is the URL of module results.
<b>Example</b>	<a href="http://yourHost.com/api/kegg/entireModule.php?id=01020304">http://yourHost.com/api/kegg/entireModule.php?id=01020304</a>

## 6 Get pathway with whole minimal gene set

<b>URL</b>	<b>api/kegg/entirePathway.php</b>
<b>GET</b>	<b>id</b>
<b>Return</b>	Result in json style. Status 0 means execute successfully, and 1 means error. Data is the URL of pathway results.
<b>Example</b>	<a href="http://yourHost.com/api/kegg/entireModule.php?id=01020304">http://yourHost.com/api/kegg/entireModule.php?id=01020304</a>

## 7 Get module with gene set that only about metabolism

<b>URL</b>	<b>api/kegg/metabolicModule.php</b>
<b>GET</b>	<b>id</b>
<b>Return</b>	Result in json style. Status 0 means execute successfully, and 1 means error. Data is the URL of module results.
<b>Example</b>	<a href="http://yourHost.com/api/kegg/metabolicModule.php?id=01020304">http://yourHost.com/api/kegg/metabolicModule.php?id=01020304</a>

## 8 Get pathway with gene set that only about metabolism

<b>URL</b>	<b>/api/kegg/metabolicPathway.php</b>
<b>GET</b>	<b>id</b>
<b>Return</b>	Result in json style. Status 0 means execute successfully, and 1 means error. Data is the URL of pathway results.
<b>Example</b>	<a href="http://yourHost.com/api/kegg/metabolicPathway.php?id=01020304">http://yourHost.com/api/kegg/metabolicPathway.php?id=01020304</a>

## Service Page Documentation

*Service Page can be visited directly to view the corresponding result.*

### 1 Page of Module with whole minimal gene set

URL	<b>/app/page/mapEntireModule.php</b>
GET	<b>id</b>
Return	Module list
Example	<b><a href="http://yourHost.com/app/page/mapEntireModule.php?id=01020304">http://yourHost.com/app/page/mapEntireModule.php?id=01020304</a></b>

### 2 Page of pathway with whole minimal gene set

URL	<b>/app/page/mapEntirePathway.php</b>
GET	<b>id</b>
Return	Pathway list
Example	<b><a href="http://yourHost.com/app/page/mapEntireModule.php?id=01020304">http://yourHost.com/app/page/mapEntireModule.php?id=01020304</a></b>

### 3 Page of module with gene set that only about metabolism

URL	<b>/app/page/mapMetaModule.php</b>
GET	<b>id</b>
Return	Module list
Example	<b><a href="http://yourHost.com/app/page/mapMetaModule.php?id=01020304">http://yourHost.com/app/page/mapMetaModule.php?id=01020304</a></b>

### 4 Page of pathway with gene set that only about metabolism

URL	<b>/app/page/mapMetaPathway.php</b>
GET	<b>id</b>
Return	Module list
Example	<b><a href="http://yourHost.com/app/page/mapMetaPathway.php?id=01020304">http://yourHost.com/app/page/mapMetaPathway.php?id=01020304</a></b>