## SynGO

**RRID:** SCR_017330  
**Type:** Tool

### Proper Citation

SynGO (RRID:SCR_017330)

### Resource Information

<table>
<thead>
<tr>
<th>URL</th>
<th><a href="https://syngoportal.org/">https://syngoportal.org/</a></th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Proper Citation:</strong></td>
<td>SynGO (RRID:SCR_017330)</td>
</tr>
<tr>
<td><strong>Description:</strong></td>
<td>Evidence based, expert curated knowledge base for synapse. Universal reference for synapse research and online analysis platform for interpretation of omics data. Interactive knowledge base that accumulates available research about synapse biology using Gene Ontology annotations to novel ontology terms.</td>
</tr>
<tr>
<td><strong>Synonyms:</strong></td>
<td>Synaptic Gene Ontologies</td>
</tr>
<tr>
<td><strong>Resource Type:</strong></td>
<td>analysis service resource, service resource, ontology, data or information resource, production service resource, data analysis service, controlled vocabulary</td>
</tr>
<tr>
<td><strong>Defining Citation:</strong></td>
<td>PMID:31171447</td>
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<tr>
<td><strong>Keywords:</strong></td>
<td>Synapse, evidence, curated, base, reference, analysis, omics, data, ontology, gene, annotation</td>
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<tr>
<td><strong>Funding Agency:</strong></td>
<td>Stanley Center for Psychiatric Research at The Broad Institute of MIT and Harvard, European Union, CERCA Program/Generalitat de Catalunya, NINDS, German Federal Ministry of Education and Research</td>
</tr>
<tr>
<td><strong>Availability:</strong></td>
<td>Free, Freely available</td>
</tr>
<tr>
<td><strong>Resource Name:</strong></td>
<td>SynGO</td>
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<tr>
<td><strong>Resource ID:</strong></td>
<td>SCR_017330</td>
</tr>
</tbody>
</table>

### Ratings and Alerts
No rating or validation information has been found for SynGO.

No alerts have been found for SynGO.

**Data and Source Information**

*Source:* SciCrunch Registry

**Usage and Citation Metrics**

We found 35 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [RRID](https://rrid.org).


Akingbuwa WA, et al. (2022) Ultra-rare and common genetic variant analysis converge to implicate negative selection and neuronal processes in the aetiology of schizophrenia. Molecular psychiatry, 27(9), 3699.


