

Obtain Node Information from NCBI Entrez

The image shows the NCBI Entrez Gene web interface. On the left, the 'Entrez, The Life Sciences Search Engine' logo is visible. Below it, the 'Data Source' is set to 'NCBI Entrez Gene'. The 'Key Attribute' section shows 'Attribute: ID' and 'Data Type: Entrez Gene ID'. The 'Available Annotation Category' section has checkboxes for 'Summary', 'Publications', 'Phenotypes', 'Pathways', and 'General Protein Information', all of which are checked. A white arrow points from the 'Pathways' checkbox to a network diagram on the right. The network diagram, titled 'PPAR-Gamma from NCBI', shows a complex web of nodes (represented by blue circles with numbers) connected by lines. A yellow circle highlights a specific node. Below the network diagram is a 'Data Panel' with a table showing gene information. The table has columns for 'ID', 'GO Term: Biological Process', and 'GO Term: Cellular Component'. The first row shows '5469' and '(androgen receptor signaling pathway, fat cell d...'. A 'Node Attribute' dropdown menu is open, showing a list of GO terms related to the selected node. The 'Network Attribute Browser' is also visible, showing a list of attributes for the network.

NCBI Entrez Gene

Entrez, The Life Sciences Search Engine

Data Source: NCBI Entrez Gene

Key Attribute: Attribute: ID, Data Type: Entrez Gene ID

Available Annotation Category: ☒ Summary, ☒ Publications, ☒ Phenotypes, ☒ Pathways, ☒ General Protein Information

Reset, Cancel

PPAR-Gamma from NCBI

Node Attribute: ☐ androgen receptor signaling pathway, ☐ fat cell differentiation, ☐ positive regulation of transcription from RNA polymerase II promoter, ☐ regulation of transcription, DNA-dependent, ☐ transcription, ☐ transcription initiation from RNA polymerase II promoter

Network Attribute Browser