Possible Measures:

* Maxflow -> number of possible paths
* Distance to first common parent
* Depth difference
* Similarities in nomenclature
* Number of subsumers
* Number of different axes/concepts traversed

Open points:

* **SNOMED data (.csv with information about graph, https://github.com/antsh3k/Clinical-Coding-Pilot)**
* **Input graph into Python (hint: IDs)**