

Documentation

Functions

There is only one function that drives this website: `runQuery()`. `runQuery()` is a function that is ran onclick of the button "Medication query" this function reads the dropdown selectors, forms a sparql query out of them dispatches the queries and forms the incoming JSON file to match the expected data structure of D3's `forcegraph()` function.

```
/*runQuery() is a function that is ran onclick of the button "Medication query"
 * this function reads the dropdown selectors, forms a sparql query out of them dispatches the queries and forms
 * the incoming JSON file to match the expected data structure of D3's forcegraph() function.*/
function runQuery() {
  class SPARQLQueryDispatcher {
    constructor(endpoint) {
      this.endpoint = endpoint;
    }

    query(sparqlQuery) {
      const fullUrl = this.endpoint + '?query=' + encodeURIComponent(sparqlQuery);
      const headers = {'Accept': 'application/sparql-results+json'};

      return fetch(fullUrl, {headers}).then(body => body.json());
    }
  }

  //get the selected dropdown items
  var d1 = document.getElementById("ddDisease1");
  var disease1 = d1.value;
  var d2 = document.getElementById("ddDisease2")
  var disease2 = d2.value;

  //log the diseases ids to the console
  console.log(disease1)
  console.log(disease2)

  //set the endpointURL to wikidata sparql
  const endpointUrl = 'https://query.wikidata.org/sparql';

  //SPARQL Query for getting every drug that treats disease 1
  const sparqlQuery1 = `SELECT DISTINCT ?drug ?drugLabel ?disease ?diseaseLabel
    WHERE
    {
      wd:${disease1} wdt:P2176 ?drug.

      BIND(wd:${disease1} as ?disease)
      SERVICE wikibase:label { bd:serviceParam wikibase:language "[AUTO_LANGUAGE],en" }
    }`;

  //Dispatch query for disease 1
  const queryDispatcher1 = new SPARQLQueryDispatcher(endpointUrl);

  //add to the previous query every drug that treats disease 2
  queryDispatcher1.query(sparqlQuery1).then(
    function (results1){
      //SPARQL Query for getting every drug of for treating disease 2
      const sparqlQuery2 = `SELECT DISTINCT ?drug ?drugLabel ?disease ?diseaseLabel
        WHERE
        {
          wd:${disease2} wdt:P2176 ?drug.

          BIND(wd:${disease2} as ?disease)
          SERVICE wikibase:label { bd:serviceParam wikibase:language "[AUTO_LANGUAGE],en" }
        }`;

      //Dispatch query for disease 2
      const queryDispatcher2 = new SPARQLQueryDispatcher(endpointUrl);

      /* The next part of the code;
      * In a loop, sets;
      * every disease to group 1,
```

```

* every drug for disease 1 to group 3,
* every drug for disease 2 to group 4
* and drugs that treat both as group 5
* every group will later be assigned to a different color in the graph.
* The nodes are placed in the networkData dictionary
*
* After the nodes are set, the edges between those nodes are extracted from the json file and inserted
* into the networkData dictionary, along with the nodes.
*/

queryDispatcher2.query(sparqlQuery2).then(
  function (results2){
    var networkData = {}
    networkData["nodes"] = [];
    networkData["links"] = [];
    var drugSet = new Set();

    //set both drug nodes to group 1
    var disease1Node = {"id":results1.results.bindings[0].diseaseLabel.value, "group":1}
    var disease2Node = {"id":results2.results.bindings[0].diseaseLabel.value, "group":1}
    networkData["nodes"].push(disease1Node,disease2Node)

    // for each drug in the results of query 1
    for(var i =0; i < results1.results.bindings.length;i++){
      if(!drugSet.has(results1.results.bindings[i].drugLabel.value)){

        //set node to group 3 if treats disease 1
        var drugNode = {"id":results1.results.bindings[i].drugLabel.value, "group":3}
        networkData["nodes"].push(drugNode)
        drugSet.add(results1.results.bindings[i].drugLabel.value)

      }
      //add link of the drug to the dictionary
      var diseasedrugLink = {"source":results1.results.bindings[0].diseaseLabel.value,
        "target":results1.results.bindings[i].drugLabel.value, "value":1 }
      networkData["links"].push(diseasedrugLink)
    }

    // for each drug in the results of query 2
    for(var j =0; j < results2.results.bindings.length;j++){
      if(!drugSet.has(results2.results.bindings[j].drugLabel.value)) {

        //set node to group 4 if treats disease 2
        var drug2Node = {"id": results2.results.bindings[j].drugLabel.value, "group": 4}
        networkData["nodes"].push(drug2Node)
        drugSet.add(results2.results.bindings[j].drugLabel.value)
      }else{
        for(var k =0; k < networkData["nodes"].length; k++){
          if(networkData["nodes"][k].id === results2.results.bindings[j].drugLabel.value){
            networkData["nodes"][k].group = 5; //set node to group 5 if treats both
          }
        }
      }
      //add link of the drug to the dictionary
      var disease2drugLink = {"source":results2.results.bindings[0].diseaseLabel.value,
        "target":results2.results.bindings[j].drugLabel.value, "value":1 }
      networkData["links"].push(disease2drugLink)
    }

    // form a ForceGraph with the data in the dictionary
    const chart = ForceGraph(networkData, {
      nodeId: d => d.id,
      nodeGroup: d => d.group,
      nodeTitle: d => `${d.id}\n${d.group}`,
      linkStrokeWidth: l => Math.sqrt(l.value),
      width: 600,
      height: 600,
    });

    // send the ForceGraph after removing the previous chart, if any
    const div = document.getElementById('chart');
    div.remove();
    var elemDiv = document.createElement('div');
    elemDiv.id = "chart"
    document.body.appendChild(elemDiv);
    elemDiv.appendChild(chart);
  }
);

```

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}    })
```