

grab_published_metadata.sh file attached.

Attached is the script (with some comments!) and an example output set.

When the tool is run (be sure to change the variable 'basedir' on line 16 to reflect your fork of the yang github repo).

When the tool is run, the output is placed into a directory, one html file for each published IEEE YANG module under 'published' - if the latest revision, as grabbed from the yangcatalog does not match what is the github/fork repo, a bold line is added towards the top of the html output file indicating the apparent discrepancy.