

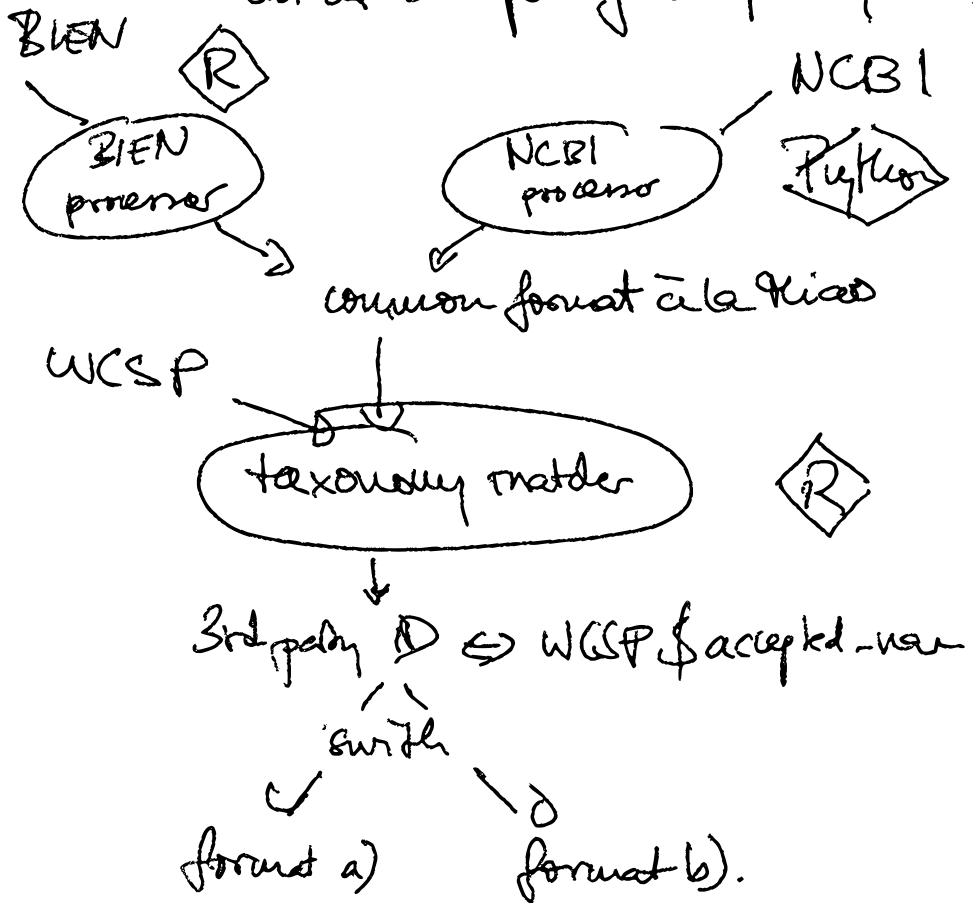
MVP:

TAXONOMY MATCHER

- input: 3rd party taxonomy data frame
+ WCSF

output: a) same data frame annotated with
WCSF ID.

b) WCSF data frame annotated with
list of corresponding 3rd party IDs.



Rough project backlog

- NCBI proems:
 - fix authority issue
 - add fields.
 - parse things into columns.
- B1BN proems
 - B1BN download
 - write it into ind. parry things into columns.
- Develop taxonomy matcher:
 - develop pseudocode.
 - implement in R.

next meetings: 5/2 + 12/2

MVP by 19/2

