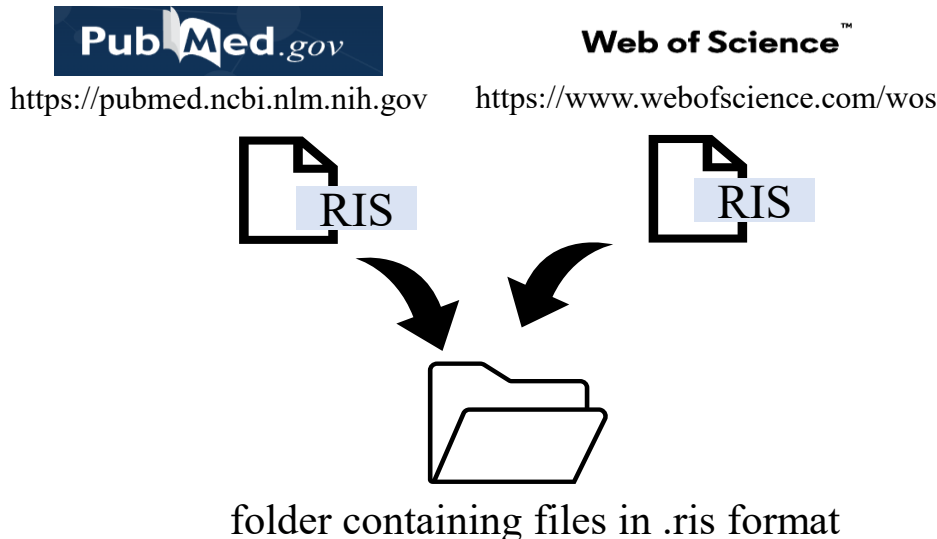


# 1. Download your bibliographic references from different databases

## 2. Convert files to .ris format

→ use <https://www.bibtex.com/c/bibtex-to-ris-converter/>

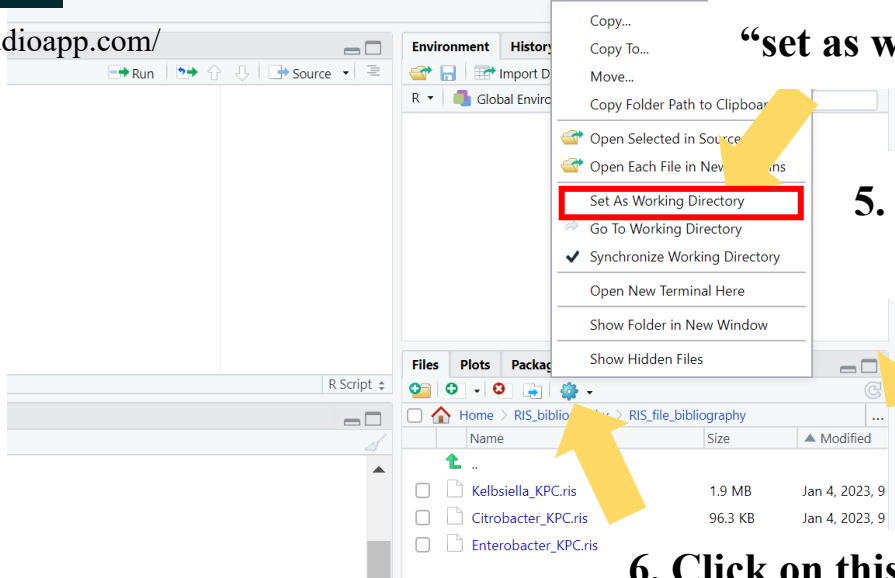
## 3. Place ris format files in the same file



## 4. Start Rstudio



<https://rstudioapp.com/>



## 8. Copy and paste this code available at this link:

[Prisma\\_Method/Script\\_Prisma.R at main · agodmer/Prisma\\_Method \(github.com\)](https://github.com/agodmer/Prisma_Method/blob/main/Script_Prisma.R)



## 9. Run the code

## 10. See number of references per file and duplicates eliminated according the doi

```
[1] "Number of references per file:"
[1] "1 Citrobacter_KPC.ris"
[1] 35
[1] "Number of references per file:"
[1] "2 Enterobacter_KPC.ris"
[1] 68
[1] "Number of references per file:"
[1] "3 Kelbsiella_KPC.ris"
[1] 732
[1] "Number of unique references:"
[1] "Number of duplicates:"
[1] 78
```

## 11. Select your abstract according to revtools package

