CanReg analysis improvements

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1 Introduction

R is a powerfull open source freely available software package that could be coupled with CanReg to improve the analytical capabilites.

2 What data does registries store in CanReg5

Each record store (at least):

- Sex
- Incidence date
- Birth date (or age at the time of tumour)
- Coded address at the time of the tumour
- Topography, Morphology, Behaviour in ICD-O-3
- Most valid basis of diagnosis
- ICD10 and ICCC
- Date of last contact
- Vital status
- Source info:
 - Type of
 - Number of
 - (Dates)

Globally each registry also have population data sets (denominators).

3 What we do with that data now

- Incidence tables (Per 100.000 per cancer group, age group, ASR, CR etc.)
- Number of cases (per cancer group, age group)
- Population pyramids
- Frequencies by year

Otherwise data needs to be exported to be analysed in other software packages.

4 What we could do with that data

- Graphs:
 - Bar charts by cancer/sex
 - * incidence tables
 - * number of cases
 - Time trends
 - * ASRs (world) over time
 - * age spec rates over time
 - * age spec rates over cohort
- Quality indicators
 - Validity:
 - * DCO%
 - * PSU%
 - * MV%
 - * Compared with other reg (CI5 IX)
 - * DCO% over time
 - \cdot potentially with graphs
 - Completeness:
 - * Reference childhood incidence comparison
 - * Stability of rates over time by cancer/sex
 - · potentially with graphs
 - * Age specific rates by cancer/sex
 - * Sources
 - $\cdot\,$ number of sources per case
 - \cdot number of notifications per case
- Geographic stuff, maps?

5 What we could do if we linked it to other data

This is more for the future, but might be interesting...

- If linked to mortality data
 - M/I ratios as estimator of completeness

6 Technical aspects

Basically two main ways to do it.

6.1 Export the data from CanReg5 to files readable by R and then call R in batch mode

Prefered mehtod - more dynamic and loosely coupled. Easier to potentially reuse R code later.

6.2 Use libraries in Java (a Java to R bridge) to call R functions directly

Alternative method we might want to look into.