

Disease mapping RIF: a tool for epidemiologists

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- Development: Small Area Health Statistics Unit (SAHSU)
- Free and integrated in ESRI[®] ArcGIS
- Aim: automated generation and representation of classical epidemiological and public health indicators based on routinely collected health and population data
- Altered and adapted for:
 - European countries: EUROHEIS (European Health and Environment Information System for Exposure and Disease Mapping and Risk Assessment)
 - US Centers for Disease Control and prevention(CDC) : EPHT (Environmental Public Health Tracking) Network

- Two types of analysis
 - Risk analysis around putative hazardous sources
 - Disease mapping

Disease mapping via the RIF

4 characteristics to be defined:

- Geographical level and reference area
 - Ward level, South East England
- Health outcomes
- Population
- Gender, age groups, adjustment covariates

Results presented as maps and/or tables

- Mortality/incidence rates
- SMR (Standardized Mortality Ratio)
- smoothed SMR (empirical Bayes estimation)



Excel table: same results + 95% CI + expected numbers

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35	01.014.01	17900.2	01.014.017900	01.014 01	52	34.5484077	1.50513449	1.124103905	1.97378127	1.456185938	66.09731435	48.86443587	87.29281234	30.7513202	1.69098431	1.26290513	2.21749829	1.60433159	64.7437771	47.9428628	85.40798277
36	01.014.01	18000.1	01.014.018000	01.014 01	18	22.3829553	0.80418335	0.476612666	1.270967109	0.864759211	35.38470645	20.09411681	57.17358814	22.8694526	0.78707612	0.46647378	1.24393008	0.8477235	35.5570111	20.192353	57.45143934
37	01.014.01	18000.2	01.014.018000	01.014 01	24	18.0919222	1.32655887	0.849937326	1.973809059	1.282535486	50.63483432	31.38267185	76.77919504	18.624387	1.28863302	0.82563791	1.91737854	1.24801023	50.2549273	31.1392932	76.21388314
38	01.014.01	18000.3	01.014.018000	01.014 01	14	25.2761773	0.55388122	0.302814778	0.929333568	0.651076831	24.90471527	13.20440575	42.40171138	29.2985096	0.47784001	0.26124196	0.80174727	0.57502227	22.3402134	11.9865855	37.82334122
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Disease mapping issues

- Small areas and/or rare diseases
 - Low counts of observed cases/ small populations
 - Numerical instability of SMRs
 - Extreme risks associated to low populated areas
 - Small SE in highly populated areas: SE(log(SMR)) α E^{-1/2}
 - Adjacent areas might show opposite risks
- Mapping the SMRs not allow to detect any spatial pattern

Smoothing the SMRs via Bayesian hierarchical models

• model BYM (Besag et al, 1991)

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O_i \sim Poisson (E_i \rho_i)
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 $Log(\rho_i) = \alpha + U_i + V_i$

- O_i, E_i: observed and expected number of cases in area i
- ρ_i: unknown relative risk in area i
- U: Spatially structured random effects \rightarrow local smoothing
- V: Unstructured random effects \rightarrow global smoothing
- RIF runs externally the BYM model using WinBUGS or INLA

..\..\RIF\example_DiseaseMapping_RIF\RIF v3.13 link to INLA.wmv

RIF issues

- Limitations
 - Single model
 - Poisson model with BYM distribution for residual effects
 - Prior distribution on precision parameters: Gamma(0.5, 0.0005)
 - Fixed number of iterations for WinBUGS
 - No convergence diagnostics tool
- Advantages
 - Interface easy to use
 - Inpout files saved -> re-run modified models
 - Support (Linda Beale)

Improvements in process

Protocol developed for an atlas (South East England), in collaboration with Nicky Best

With WinBUGS and INLA

- **1**. Parameters of interest
- 2. Sensitivity analyses
- **3.** Convergence Monitoring for WinBUGS / accuracy of the approximation for INLA
- 4. Comparison of the results WinBUGS / INLA (?) if the user wants to use both softwares

1. Parameters of interest

- Actual parameters
 - Smoothed SMR: overall risk + specific area risk (exp(BYM))
 - Smoothed RR: specific area risk (exp(BYM))
 - Posterior probabilities
- Added parameters: quantitative summaries of the spatial variability
 - **R.90.10**: ratio between the 90th and 10th percentile of the posterior distribution of the smoothed SMR/RRs

> Indicator of the variability across the study region

 Spatial fraction: relative contribution of spatial vs unstructured heterogeneity to the overall variability

2. Sensitivity analyses (default)

Priors for the precisions (N=4)

- Gamma distributions
- Truncated Normal distribution



Structure on the residuals (N=3)

- Heterogeneity only
- CAR model only
- BYM (heterogeneity + CAR)

Comparison of 12 models: -DIC -R.90.10 -posterior variance summaries

3. checking CV/accuracy of the approximation

Visual checks

- posterior densities of:
 - RR for 5 areas based on the expected numbers

(2.5, 25, 50, 75 and 97.5 percentile of the distribution of the expected numbers)

- variance parameters
- Autocorrelation (MCMC) of variance parameters

	WinBUGS	INLA				
Pop-up windows	Gelman and Rubin diagnostics R<1.05	Number of replicates >2 expected number of parameters/size of data				
with warnings	Monte Carlo error <5% of sd	Symmetric Kullback-Leibler divergence (SKLD) between the Gaussian and the Laplace approximation				

Suggestions and comments are welcome!