Extending Bioconductor to exposome data analysis

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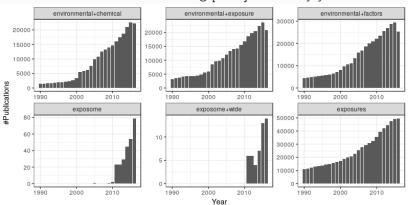
July 26th, 2017

Barcelona Institute for Global Health

Environmental Health

The exposome encompasses the totality of human environmental [...] exposures from conception onwards, complementing the genome. Dr. Christopher Wild, "Complementing the Genome with an Exposome"

Publications containing [term] in title by year



Environmental Health

The European Commission awarded two large-grants to pursue exposome-related research (2012):

- The **HELIX project**, lead by *Barcelona Institute for Global Health*, will attempt to develop an early life exposome, noting that the first exposures occur during development.
- The EXPOSOMICS, lead by Imperial College London, will use smartphones that utilize GPS and environmental sensors to assess exposures in adulthood.

Late:

- Health and Environment-Wide Associations based on Large Scale population Surveys (HEALS)
- HERCULES that aims to provide expertise to develop and refine new tools and technologies to assess the exposome

The HELIX Project



Cohort	# Sample
BIB	14 000
EDEN	2 000
INMA	2 500
KANC	4 000
MoBa	8 000
RHEA	1 500

From the total $(+30\ 000)$, 1 200 will get omic data

rexposome project

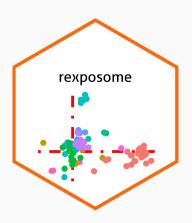
rexposome project1

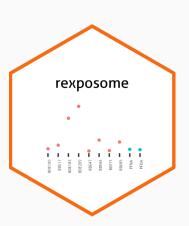
Aims to incorporate the exposome into Bioconductor ecosystem.

- rexposome (under revision): package for exposome data management, exposome characterization and testing exposome - health outcome association.
- omicRexposome (under revision): package for testing exposome - omic association (exposome and exposome cluster) and integration.

¹Hernandez-Ferrer C, et al.; Comprehensive analysis of the exposome, exposome-health associations and omics intermediates; [submitted]

rexposome project





Contact Information

Any questions?

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```
1  data = rnorm(100)
2  attempts = 0
3    pvalue = 0
4  while( pvalue > 0.05 ) {
5     attempts = attempts + 1
6     cases = sample(data,50)
7     controls = data[ (data %in% cases]
8    pvalue = t.test(cases,controls)5.value }
9    cat("Congratulations With p = "r.onund(pvalue,3),"you achieved scientific success in",attempts,"attempts\n")
```

How to achieve scientific success in 10 lines of R code!

Leon Eyrich Jessen (@jessenleon) - October 23th, 2016