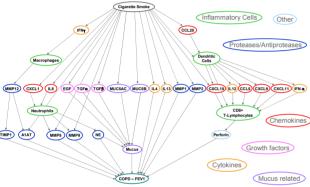
New Method to Capture Context-Dependent Quantitative Data to Build a Data Warehouse for Chronic Obstructive Pulmonary Disease

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Introduction

Chronic Obstructive Pulmonary Disease (COPD) is characterized by progressive airflow limitation associated with abnormal inflammatory response of the lung to noxious particles and gases. It is the fourth leading cause of death in the world. Comprehensive information related to COPD is available, but little is known about the disease mechanisms. In order to investigate disparate pieces of information, data must be collected, stored, and evaluated in a systematic way. Here, we present a new method to capture context-dependent quantitative data to build a data warehouse for COPD.

Current COPD Biological Network



Network composed of the various links used so far for the literature search.

Literature Search: Three Phases

Query phase: generation of a comprehensive list of articles from gueries including all synonyms for all specified links of the defined biological pathways (see above) to search in PubMed. Primary screening phase: at the abstract level to determine the relevance of the articles to the specified links. Secondary screening phase: at the full-text article level and based on inclusion/exclusion criteria to select the most relevant publications. (Screening phases performed by biologists.)

INCLUSION CRITERI A

- > English peer-reviewed articles relevant for the different biological links defined in the COPD Bayesian model (direct acyclic graph with probabilistic distribution)
- > Human, mouse, and rat studies

modifications with quantitative data

> COPD patients with or without co-morbidities > Articles dealing with cigarette smoke (CS) & CS constituents inducing mechanistic

EXCLUSION CRITERIA

- > Articles dealing ONLY with immortalized
- > Articles dealing ONLY with cell-free assays > Case reports, reviews, editorials, comments, and letters
- > Epidemiological data and meta-analysis.
- > Articles dealing ONLY with COPD patients with exacerbations AND/OR alpha-1 antitrypsin deficiency

Conclusion

We have set up a process to build a COPD data warehouse. The process captures quantitative data and associated context information, such as type of experiment (in vivo, in vitro), species used (human, mice, rat), characteristics of the study group (smoking status, demographic data, disease phenotype, etc.) and a scoring system (the Measurement Certainty Index) to evaluate the certainty of the data.

Data from 600 research articles have been recorded to date. This mine of information is easily retrieved and provides us with a tool to identify data gaps and putative biomarkers. Based on these data, an in silico COPD Bayesian network model for disease risk prediction is being built. This approach can be also adapted to other research areas

Literature Data Transfer Template General Information MCI score for the study as a whole Abstract Dialog Box to Generate Tables: Table name Type of experin Groups within the smoking/exposure status Variables within the groups Type of table to create OK Cancel Raw Data Tables Example of Raw Data Table statistics captured from the raw data Contains measurement certainty index that measures the weight of the evidence Sample size comments (if confusion) and and

Data Curation & General Process ICROSOF EXCEL PMI Curator Literature Automated data cleansing Search & ⇒ Manual data quality Data checks Literature data transfer templat Pipeline Pilot Step 3: Scripts Disease Model DWH Data Transfer Analysis Process

Reporting

Tool

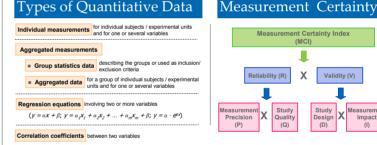
(ESQ)

■ Modelers

■ Biologists

X Measurement

Manual data accessment



COPD modeling

reporting, data mining,

COPD DWH in PMI Data Integration Platform

