

Project Background: Rare Disease Algorithm Overview

(c) 2023 Genzyme Corporation. All rights reserved. Sanofi is a registered trademark of Sanofi or an affiliate.

For use by Sanofi Medical Affairs for scientific and medical discussions only. Do not photograph, copy or distribute.

()

MAT-US-2107797 v2.0 - P Expiration Date: 10/03/2025

Opportunity to further patient diagnosis using EHR data



For use by Sanofi Medical Affairs for scientific and medical discussions only. Do not photograph, copy or distribute.

3

Developing the Rare Disease Algorithm

A behind-the-scenes look



For use by Sanofi Medical Affairs for scientific and medical discussions only. Do not photograph, copy or distribute.

4

What's in the De-identified Database?



For use by Sanofi Medical Affairs for scientific and medical discussions only. Do not photograph, copy or distribute.

5

Algorithm (Model) Development – 3 Stages





How does a typical Gaucher patient present longitudinally within EHR Data?



NERVOUS SYSTEM

Ataxia Extrapyramidal disorder Bradykinesia Cranial nerve disorders Developmental retardation Dysphagia Gaze palsy Hearing impairment Hemiplegia/paresis Hydrocephalus Hypotonia Laryngeal spasms Mvoclonic seizure Nerve root compression Oculomotor apraxia Optic kinetic nystagmus Parkinsonism Strabismus Tonic-clonic seizure Tremor

DEVELOPMENT

Delayed puberty Delayed skeletal maturation Developmental regression Growth delay Short stature

BONES

Arthralgia Joint stiffness (e.g. arthrogryposis) Avascular necrosis Delayed skeletal maturation Erlenmeyer flask deformity Joint dislocation **Kyphosis** Osteoarthritis Osteopenia Osteolysis Osteoporosis Spinal deformation Bone pain Pathological fracture

HEART

Aortic and mitral valve calcification

HEPATIC

Cirrhosis Hepatic fibrosis Portal hypertension Hepatitis Elevated CRP Elevated ferritin

Renal

Hematuria Proteinuria

RESPIRATORY

Pulmonary fibrosis Pulmonary hypertension Respiratory failure Interstitial pulmonary abnormalities

BLOOD

Pancytopenia Thrombocytopenia Gingival bleeding

ORGANOMEGALY

Hepatomegaly Splenomegaly Ventriculomegaly

OPHTHALMOLOGIC

Corneal deposit & opacity Retinopathy (nondiabetic)

MALIGNANCY

Multiple myeloma Liver neoplasm Non-Hodgkin lymphoma Malignant melanoma Pancreatic cancer

PSYCHIATRIC

Dementia (non senile) Depression

PERINATAL

Hydrops fetalis Ichthyosis

Anemia Anemia

IMMUNOLOGY

Polyclonal gammopathy

Monoclonal gammopathy

GENERAL SIGNS

Abdominal pain Fatigue Fever

9

sanofi

The Two Models – How do they differ?



How does a typical Gaucher patient present longitudinally within EHR Data?

LITERATURE FEATURES	 Symptom from literature Enriched with SDS, labs, procedures
Data driven Features	 Features the model picked up as differentiators of Gaucher vs controls
DEMOGRAPHICS	• Region, race, gender, age
Healthcare Interactions	 Provider specialty Visits Encounters



Age Model

- Symptoms are derived with the age of first occurrence for each patient
- Favors features that have different ages of onset between the Gaucher and control patients
- Favors earlier onset
- Favors younger patients

Prevalence Model

- Symptoms are flagged with presence/absence
- Model favors features that have different prevalence between the Gaucher and control patients
- Favors accumulation of comorbidities
- Favors older patients

sanofi

Testing the Models

2 TEST

Based on features identified during the training, how well do they identify GD patients?



Random Gaucher patients

Patients N=1,000,000 (ratio 1:10,000)Gaucher patients N=100



Machine Learning models vs Each Other vs Rare Diagnostic Algorithm

* Choice of model(s) and threshold should be assessed for each database and context of the application

- There is flexibility in the application of the model that is <u>situation dependent</u>. We chose a threshold of **≥0.95 (95% probability)**.
- Consider what types of patients are *most likely* undiagnosed
- Testing capacities



Both models identify more known GD patients in the dataset, in addition to having a better known-GD-to-controls ratio, than the rule-based filter

**Adapted for RWD from published diagnostic algorithm/Cappellini Protocol (Mistry et al, 2011)

sanofi

Together or Separately?

* Choice of model(s) and threshold should be assessed for each database and context of the application

- There is flexibility in the application of the model that is <u>situation dependent</u>. We chose a threshold of ≥0.95 (95% probability).
- Consider what types of patients are most likely undiagnosed
- Testing capacities



- 2467 patients highly-ranked by either model
- 34 known GD patients (1.4%)





What's Next?

Implementation and Evaluation of a Rare Disease Algorithm to Identify Persons at Risk of Gaucher Disease Using Data from Electronic Health Records (EHRs) in the United States

sanofi

Clinical Trial Design





Study Objectives

Primary

econdary

Ň

Exploratory

Patient

Reach

- Estimate the positive predictive value (PPV), or diagnostic yield, of the Rare Disease Algorithm (RDA) for Gaucher Disease (GD) (i.e., the proportion of persons with GD out of all persons highly ranked by the RDA)
- Identify the magnitude of "unrecognized" GD (i.e., previously undiagnosed persons highly ranked by RDA who subsequently test positive for GD, out of all persons with GD)
- Describe characteristics (e.g., demographics, alternative and concomitant diagnoses, treatments, and patterns of provider visits) of all persons highly ranked by the RDA, by subgroups:
 - Persons previously diagnosed with GD
 - Persons previously undiagnosed with GD who subsequently test positive for GD
 - Persons previously undiagnosed with GD who subsequently test negative for GD
- Describe characteristics (e.g., demographics, alternative and concomitant diagnoses, treatments, and patterns of provider visits) of all persons highly ranked by the RDA, by subgroups:) of persons highly ranked by the RDA who are negative in confirmatory testing for GD but positive for ASMD
- Determine the diagnostic yield for ASMD among patients highly ranked by the RDA
- Explore the relationship of ASMD biomarker lysosphingomyelin in the diagnostic process for ASMD
- Describe the time and resources on a per-role basis (e.g., physician, study coordinator, bioinformatics) needed for the following:
 - Identifying potential predictors (demographics, diagnoses, treatments, lab values and visits) of those persons highly ranked by the RDA not previously diagnosed with GD who are found through confirmatory testing to be GD-positive (if sample size allows)
 - Describe IDN characteristics including demographics, academic vs. non-academic, EHR type, physician specialties, etc.

sanofi

Bioinformatics

Big Data

Clinical

Champion

Next Steps: Deploying RDA in a Compliant Manner

- Share algorithm with US Health Systems
- Access via secured URL or API
- Hands off approach
- Provide patient readout that fosters transparency with implementing the algorithm
- Gather feedback on implementation and patients identified with Gaucher
- Information **will not be shared** with Commercial colleagues



Questions





Clinicaltrial.gov

Patrick Pavlick C: 908.842.4390 E: patrick.pavlick@sanofi.com



Model Development

