

# Moduł 12

## Przypadek użycia 1

### COVID Data hub



iBigWorld:  
Innovations for Big Data in a Real World

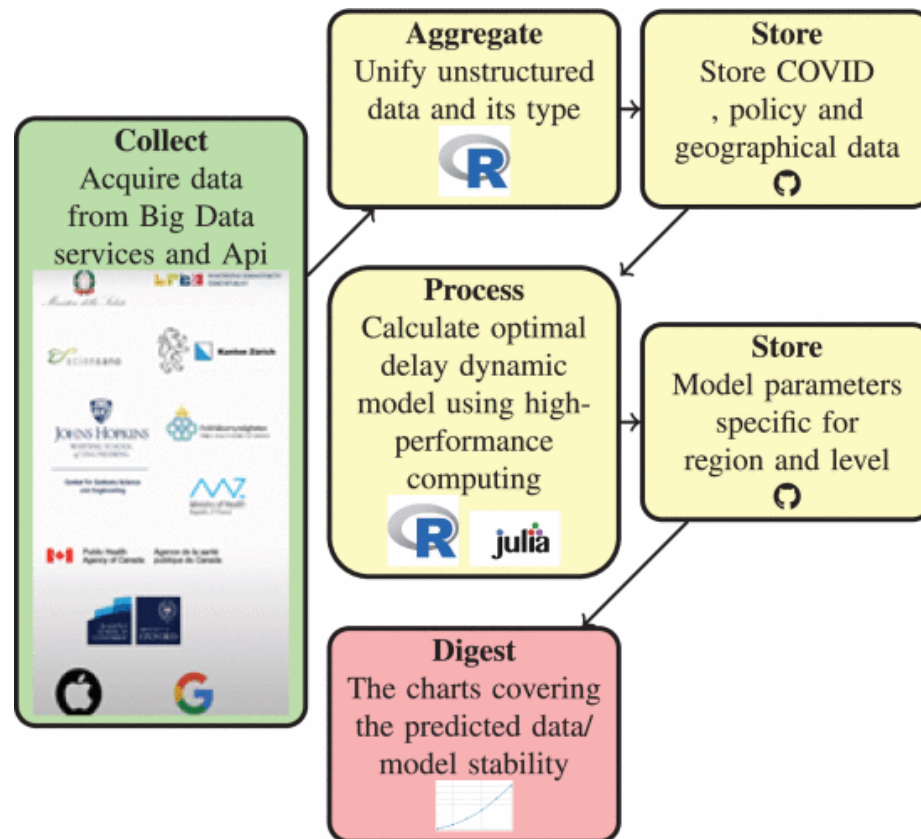
Zespół UBB

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# Potok Big Data

- COVID-19 Big Data Hub umożliwia nam przewidywanie rozwoju pandemii z uwzględnieniem wielu szczepów wirusa i opóźnień zakaźności
- Modele dynamiczne z dwoma szczepami z rozproszonymi opóźnieniami zostały dopasowane do szeregów czasowych pobranych z centrum danych COVID
- Wykorzystano dane na poziomie krajowym, regionalnym i powiatowym, które są płynnie zintegrowane z otwartymi danymi Banku Światowego, raportami mobilności Google, raportami mobilności Apple
- Identyfikacja parametrów została zrealizowana za pomocą algorytmu COBYLA
- Symulacje zostały zaimplementowane za pomocą modułu Julia.

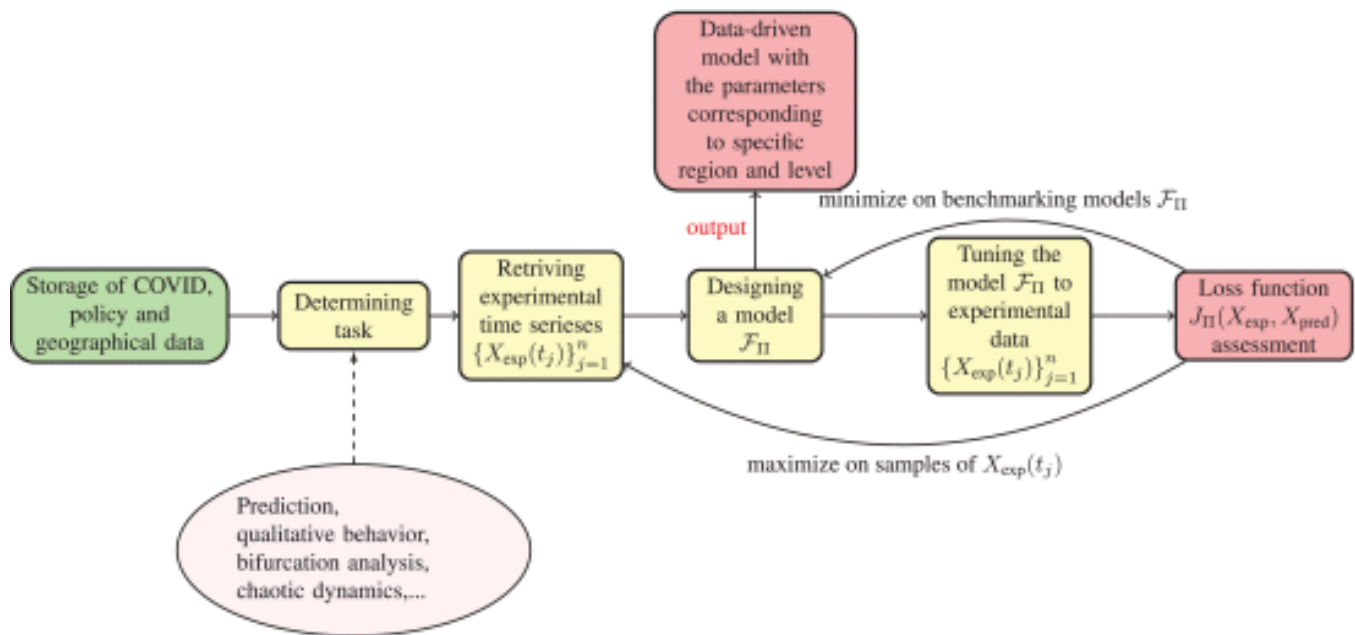


# Pobranie danych

- `library("COVID19")`
- `###COUNTRY#####`
- `x <- covid19(country = countryname, level = 1)`
- `###REGION#####`
- `x <- covid19(country = countryname, level = levelname)`
- `x <- x[which(x$administrative_area_level_2=="Texas"),]`
- `###COUNTY#####`
- `x <- covid19(country = countryname, level = levelname)`
- `x <- x[which(x$administrative_area_level_2=="Rheinland-Pfalz" & x$administrative_area_level_3=="SK Speyer"),]`

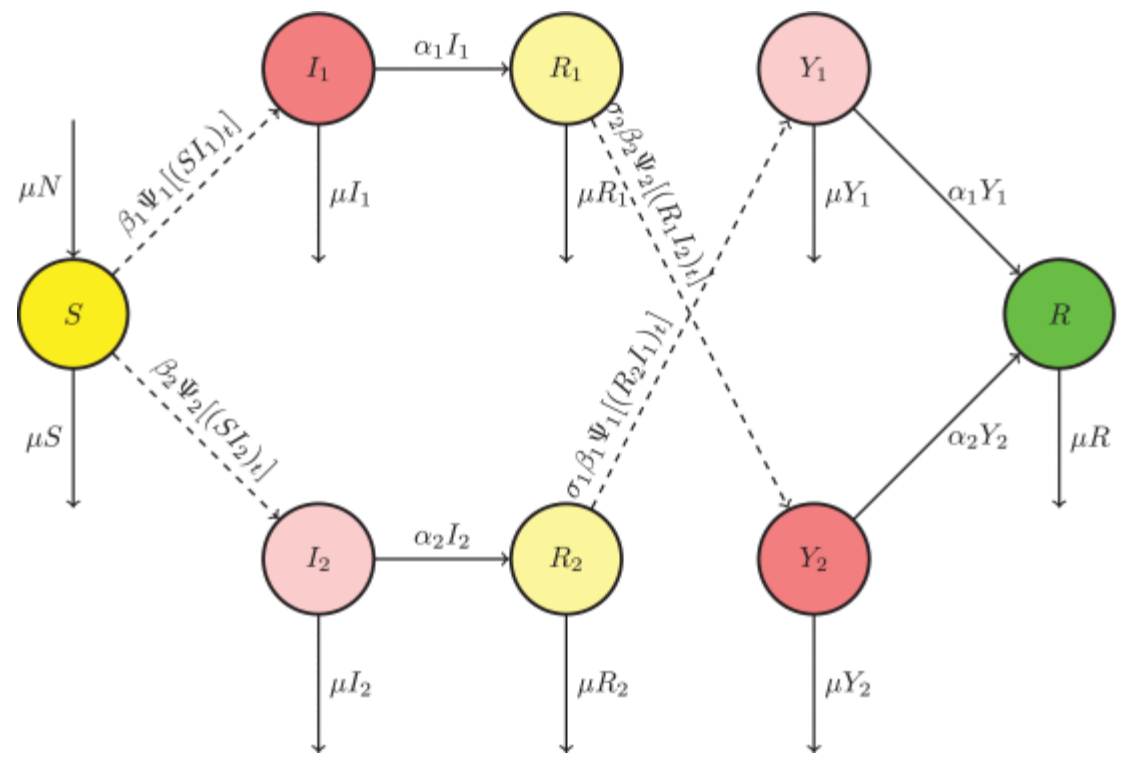
# Potok Big Data

- Potok rozwiązania problemu SciML



# Model

- Strojienie modelu



# Potok Big Data

- Strojenie modelu

**Input data:**  $X_{exp}, \Pi_{lower}, \Pi_{upper}, \Pi_{init}$

**Result:**  $\Pi_{opt}$

form the initial simplex  $C_{init}$  with the vertices

$\Pi_0^{init}, \Pi_1^{init}, \dots, \Pi_{27}^{init};$

**repeat**

for the current simplex  $C$  calculate the values

$\hat{\Phi}_C(\Pi_i), i = \overline{0, 27};$

search the vertex  $\Pi_l$  determined by the equation

$\hat{\Phi}_C(\Pi_l) = \min \{ \hat{\Phi}_C(\Pi_i), i = \overline{0, 27} \};$

calculate new vertex as

$\Pi_{new} := -\theta \Pi_l + (1 + \theta) \frac{1}{27} \sum_{i=\overline{0, 27}, i \neq l} \Pi_i$ , where reflection coefficient  $\theta \in (0, 1)$  being chosen as small as possible in order  $\hat{\Phi}_C(\Pi_l)$  not were the least calculated function value so far;

form modified simplex  $C_{new}$  replacing vertex  $\Pi_l$  with  $\Pi_{new}$ ;

search  $\Pi_{opt}$  as a solution of the problem of linear optimization

$$\left. \begin{array}{l} \text{minimize } \hat{\Phi}_{C_{new}}(\Pi), \\ \text{subject to } \Pi \in C_{new} \end{array} \right\} \quad (16)$$

**until** stop condition;

**return**  $\Pi_{opt}$ ;



# Potok Big Data

- Strojenie modelu

**Input data:**  $X_{exp}, \Pi_{lower}, \Pi_{upper}, \Pi_{init}$

**Result:**  $\Pi_{opt}$

form the initial simplex  $C_{init}$  with the vertices

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**until** stop condition;

**return**  $\Pi_{opt}$ ;



# Wysoko wydajne operacje w Julia

- `f <- JuliaCall::julia_eval("function f(du, u, h, p, t)`
- `mu=p[1]`
- `beta_1=p[2]`
- `alpha_1=p[3]`
- `a_1=p[4]`
- `m_1=p[5]`
- `tau_m_1=p[6]`
- `beta_2=p[7]`
- `alpha_2=p[8]`
- `a_2=p[9]`
- `m_2=p[10]`
- `tau_m_2=p[11]`
- `sigma_1=p[12]`
- `sigma_2=p[13]`
- `n=100`
- `c=0.01`
- `rmse_1=sqrt((m_1+1)/a_1^2)`
- `tau_M_1_tilde=sqrt(rmse_1^2 / c)`
- `tau_M_1=tau_m_1+(m_1+1)/a_1+tau_M_1_tilde`
- `rmse_2=sqrt((m_2+1)/a_2^2)`
- `tau_M_2_tilde=sqrt(rmse_2^2 / c)`
- `tau_M_2=tau_m_2+(m_2+1)/a_2+tau_M_2_tilde`
- `lags_1=range(1/n, length=n, stop=tau_M_1*(1-1/n))`
- `lags_2=range(1/n, length=n, stop=tau_M_2*(1-1/n))`



# Wysoko wydajne operacje w Julia (kontynuacja)



- `N=p[14]`
- `S_0=p[15]`
- `L_1_0=p[16]`
- `R_1_0=p[17]`
- `L_2_0=p[18]`
- `R_2_0=p[19]`
- `Y_1_0=p[20]`
- `Y_2_0=p[21]`
- `beta_1_tilde=p[22]`
- `omega_beta_1=p[23]`
- `beta_2_tilde=p[24]`
- `omega_beta_2=p[25]`
- `alpha_1_tilde=p[26]`
- `omega_alpha_1=p[27]`
- `alpha_2_tilde=p[28]`
- `omega_alpha_2=p[29]`
- `du[1]= mu*(N-u[1]) - beta(beta_1,beta_1_tilde,omega_beta_1,t)*u[1]*u[2] - beta(beta_2,beta_2_tilde,omega_beta_2,t)*u[1]*u[4]`
- `du[2]=beta(beta_1,beta_1_tilde,omega_beta_1,t)*tau_M_1*(1/n)*sum([psi(a_1,m_1,tau_m_1,tau)*h(p,t-tau)[1]*h(p,t-tau)[2] for tau in lags_1)`  
`- mu*u[2] - alpha(alpha_1,alpha_1_tilde,omega_alpha_1,t)*u[2]`
- `du[3]=alpha(alpha_1,alpha_1_tilde,omega_alpha_1,t)*u[2] - (mu+sigma_2*beta(beta_2,beta_2_tilde,omega_beta_2,t)*u[4])*u[3]`
- `du[4]=beta(beta_2,beta_2_tilde,omega_beta_2,t)*tau_M_2*(1/n)*sum([psi(a_2,m_2,tau_m_2,tau)*h(p,t-tau)[1]*h(p,t-tau)[4] for tau in lags_2)`  
`- mu*u[4] - alpha(alpha_2,alpha_2_tilde,omega_alpha_2,t)*u[4]`
- `du[5]=alpha(alpha_2,alpha_2_tilde,omega_alpha_2,t)*u[4] - (mu+sigma_1*beta(beta_1,beta_1_tilde,omega_beta_1,t)*u[2])*u[5]`
- `du[6]=sigma_1*beta(beta_1,beta_1_tilde,omega_beta_1,t)*tau_M_1*(1/n)*sum([psi(a_1,m_1,tau_m_1,tau)*h(p,t-tau)[5]*h(p,t-tau)[2] for tau in lags_1) - mu*u[6] - alpha(alpha_1,alpha_1_tilde,omega_alpha_1,t)*u[6]`
- `du[7]=sigma_2*beta(beta_2,beta_2_tilde,omega_beta_2,t)*tau_M_2*(1/n)*sum([psi(a_2,m_2,tau_m_2,tau)*h(p,t-tau)[3]*h(p,t-tau)[4] for tau in lags_2) - mu*u[7] - alpha(alpha_2,alpha_2_tilde,omega_alpha_2,t)*u[7]`
- `end"`



# Wysoko wydajne operacje w Julia (kontynuacja)



```
• h <- JuliaCall::julia_eval("function h(p, t)
•   [p[15],p[16],p[17],p[18],p[19],p[20],p[21]]
•   end")
• psi <- JuliaCall::julia_eval("function psi(a, m, tau_m, tau)
•   if tau <= tau_m
•     res = 0
•   else
•     res = (a^(m+1)/gamma(m+1))*(tau-tau_m)^m * exp(-a*(tau-tau_m))
•   end
•   res
• end")
• beta <- JuliaCall::julia_eval("function beta(beta_i,beta_i_tilde,omega_beta_i,t)
•   beta_i + beta_i_tilde*sin(2*pi*t/omega_beta_i)
• end")
• alpha <- JuliaCall::julia_eval("function alpha(alpha_i,alpha_i_tilde,omega_alpha_i,t)
•   alpha_i + alpha_i_tilde*sin(2*pi*t/omega_alpha_i)
• end")
```



# Wizualizacja

