



#### reaction blank node

- ("index" : int/string)



### reaction edge

- id : int/string)
- source : string target : string quantity : int



#### metabolit node

- ("index" : int/string)
  name : string
  quantity\_compartment A : uint
  quantity\_compartment B : uint
  quantity\_compartment X : uint



## gen node

 $\hbox{- index}: int$ 

- name : string

# gen edge

- id : int/string)

- source : string target : string and : Array<gen node index>



reaction graph



subsystem graph



compartment graph





