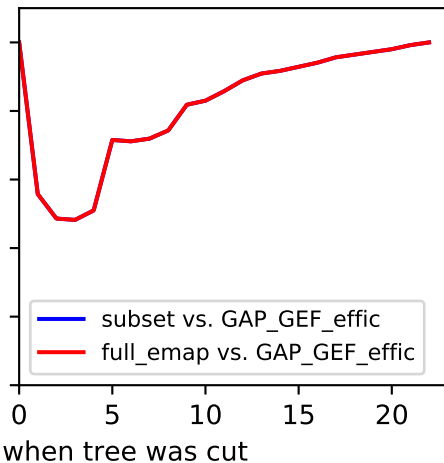
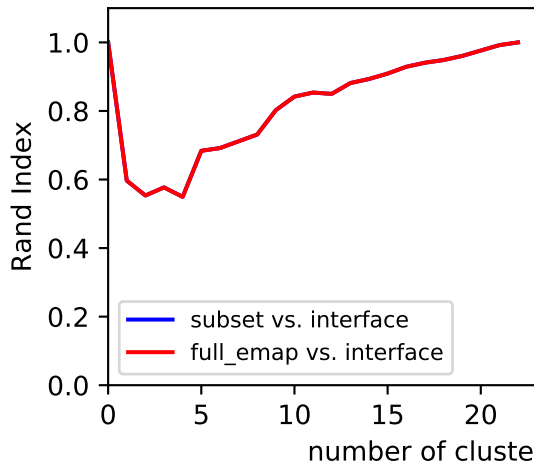


dataset = full_emap

Interface
 Δ to full_emap: 0.0

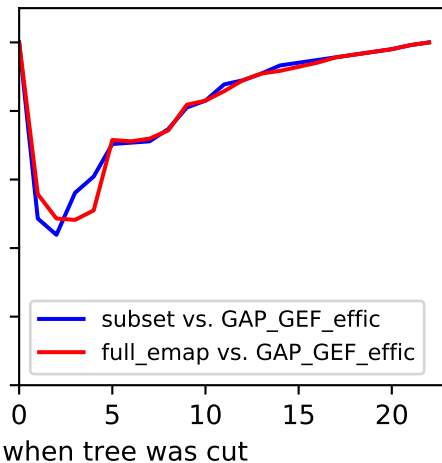
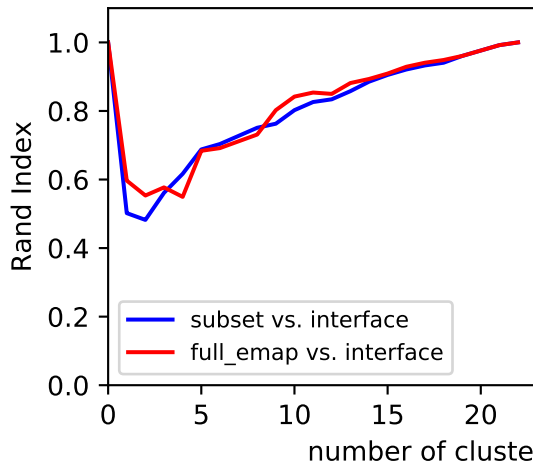
GAP/GEF effic
 Δ to full_emap: 0.0



dataset = SET3

Interface
 Δ to full_emap: -0.011

GAP/GEF effic
 Δ to full_emap: 0.004



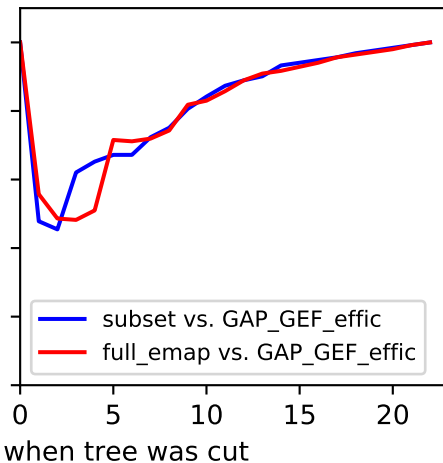
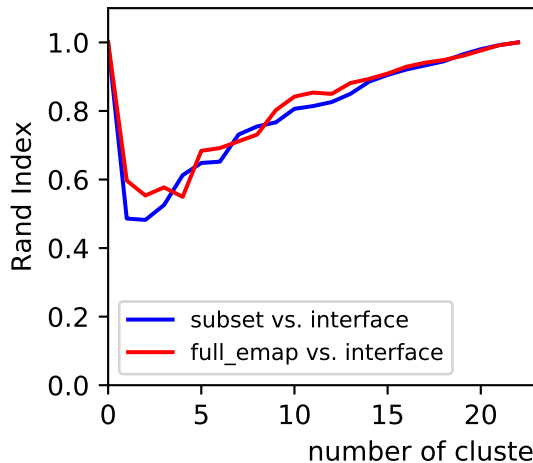
dataset = Arl GTPases

Interface

Δ to full_emap: -0.017

GAP/GEF effic

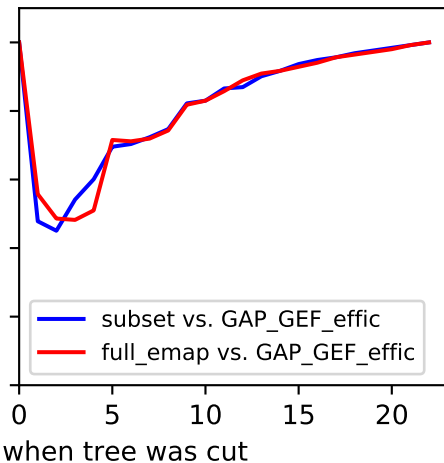
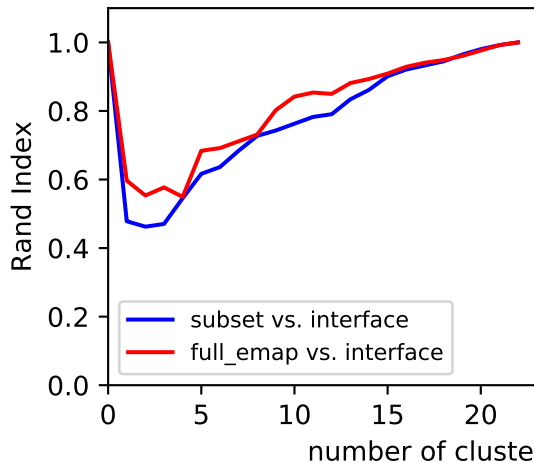
Δ to full_emap: 0.007



dataset = ADA

Interface
 Δ to full_emap: -0.037

GAP/GEF effic
 Δ to full_emap: 0.001



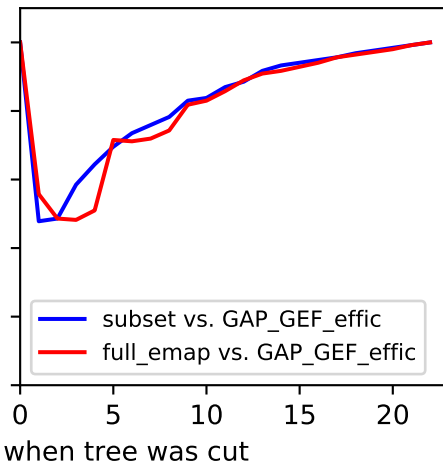
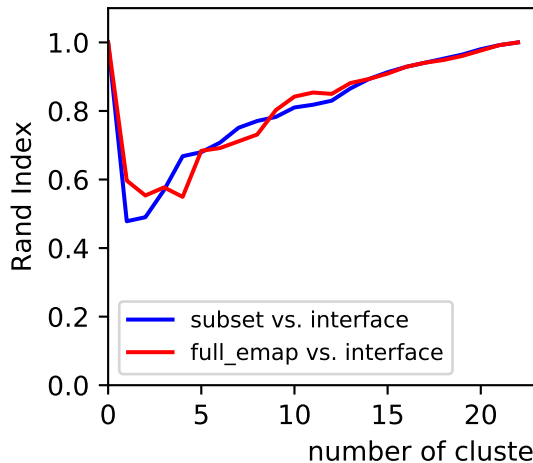
dataset = mitochondrial

Interface

Δ to full_emap: -0.004

GAP/GEF effic

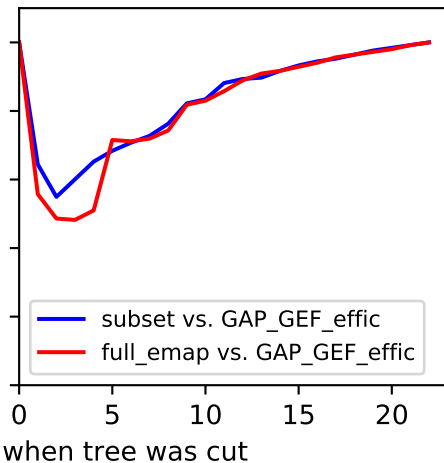
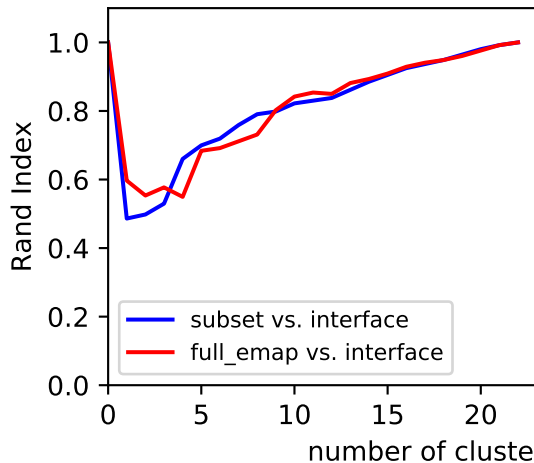
Δ to full_emap: 0.014



dataset = RIM101

Interface
 Δ to full_emap: -0.002

GAP/GEF effic
 Δ to full_emap: 0.019



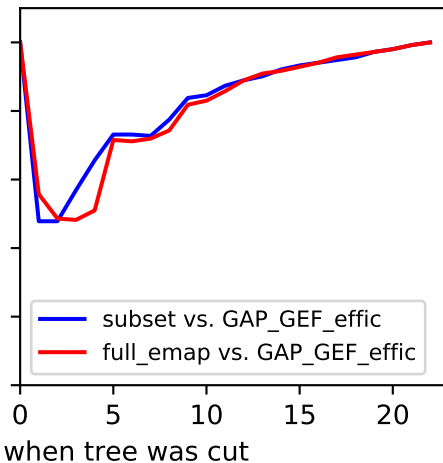
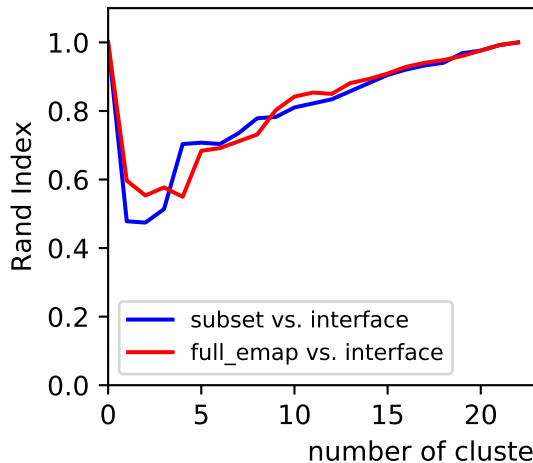
dataset = Elongator

Interface

Δ to full_emap: -0.007

GAP/GEF effic

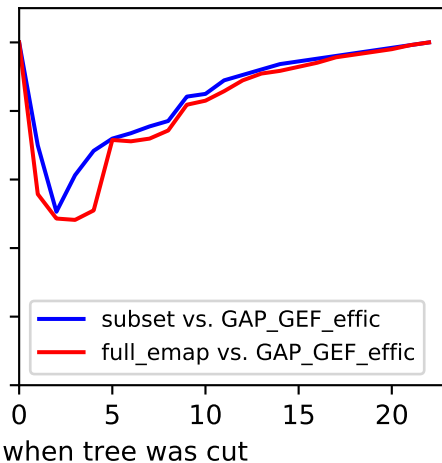
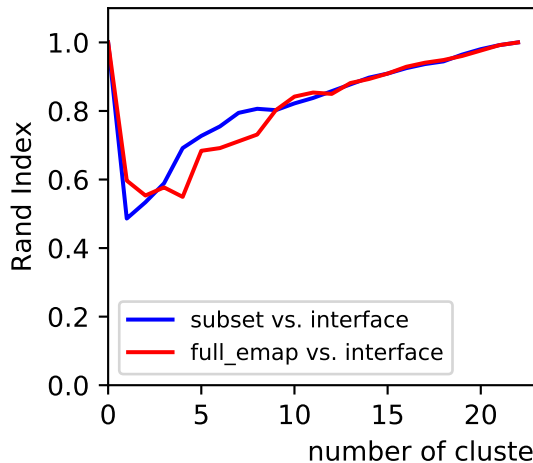
Δ to full_emap: 0.011



dataset = Rpd3L

Interface
 Δ to full_emap: 0.011

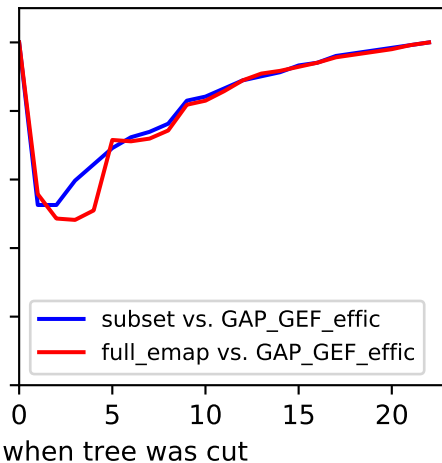
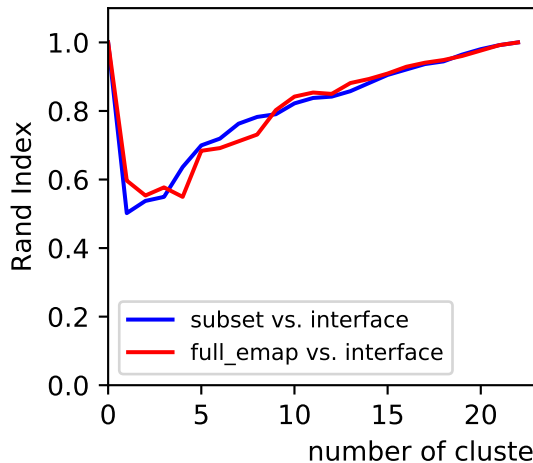
GAP/GEF effic
 Δ to full_emap: 0.031



dataset = HOG1

Interface
 Δ to full_emap: -0.0

GAP/GEF effic
 Δ to full_emap: 0.014



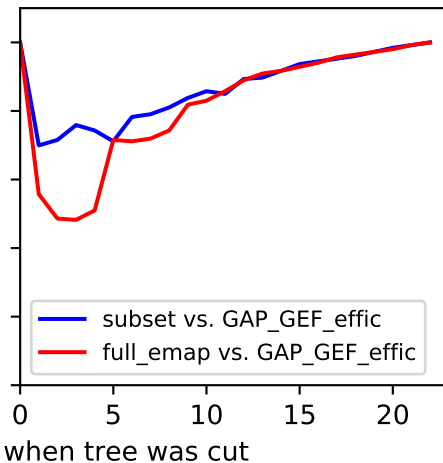
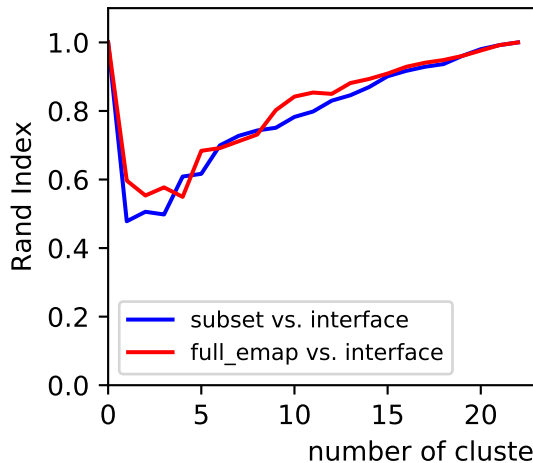
dataset = DNA Replication

Interface

Δ to full_emap: -0.022

GAP/GEF effic

Δ to full_emap: 0.049



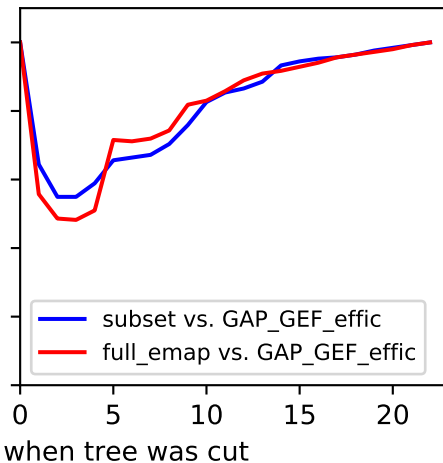
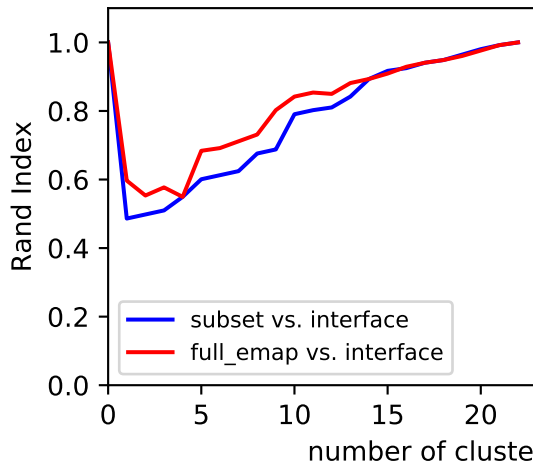
dataset = Peroxins

Interface

Δ to full_emap: -0.036

GAP/GEF effic

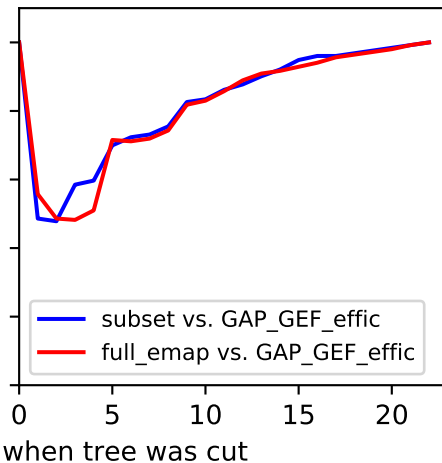
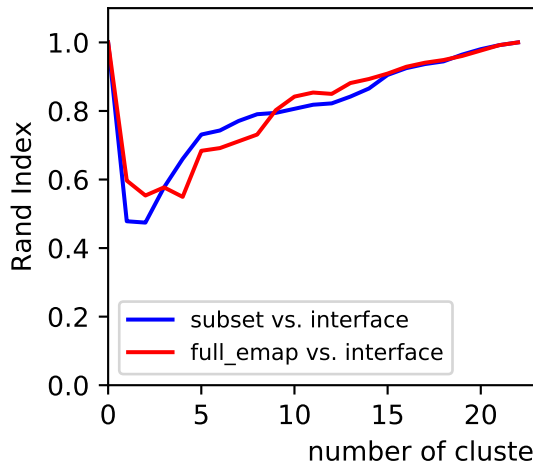
Δ to full_emap: 0.002



dataset = SWR1

Interface
 Δ to full_emap: -0.002

GAP/GEF effic
 Δ to full_emap: 0.008



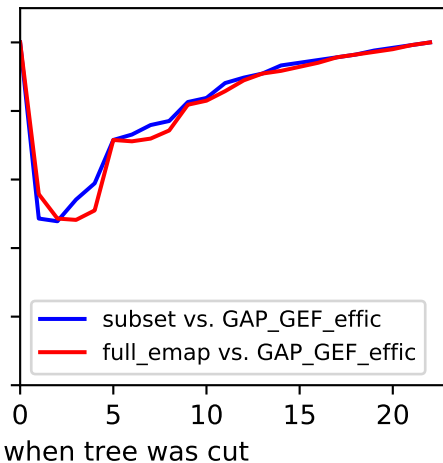
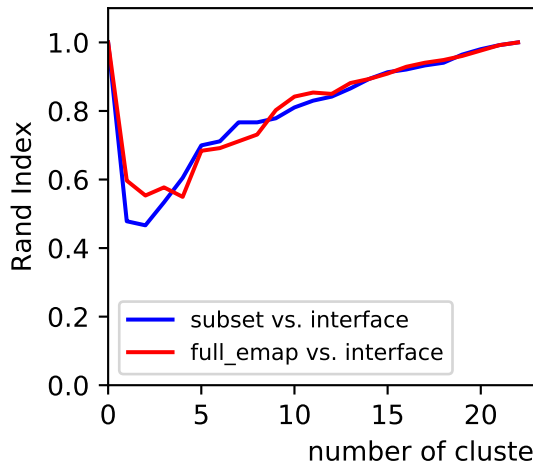
dataset = mRNA processing

Interface

Δ to full_emap: -0.008

GAP/GEF effic

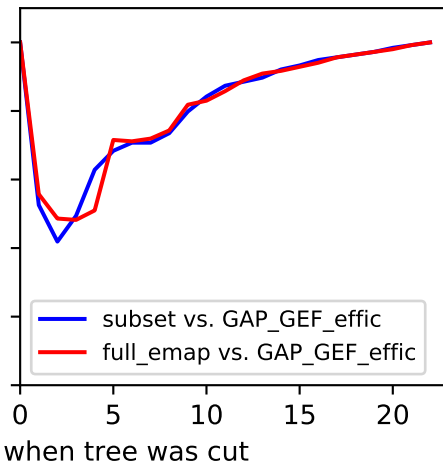
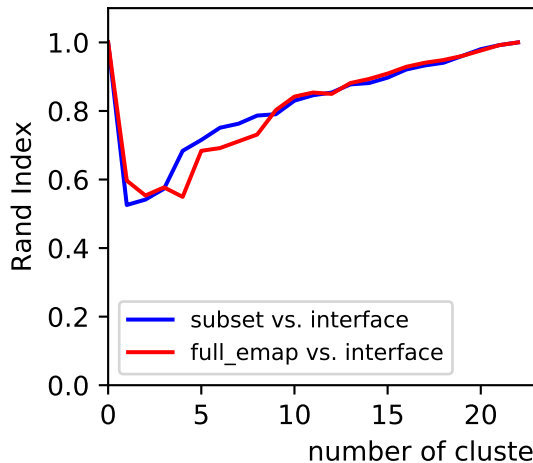
Δ to full_emap: 0.01



dataset = Dynactin

Interface
 Δ to full_emap: 0.007

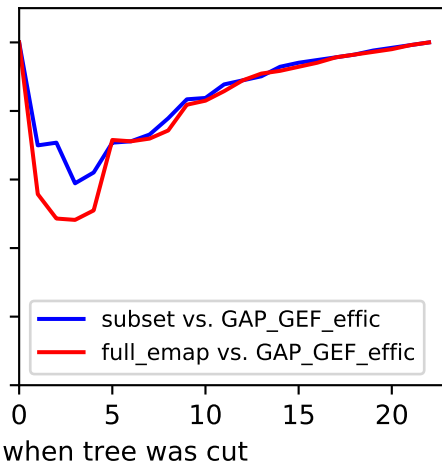
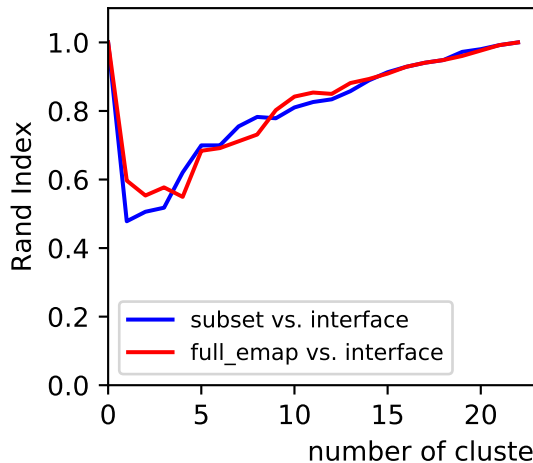
GAP/GEF effic
 Δ to full_emap: -0.001



dataset = ERMES

Interface
 Δ to full_emap: -0.006

GAP/GEF effic
 Δ to full_emap: 0.03



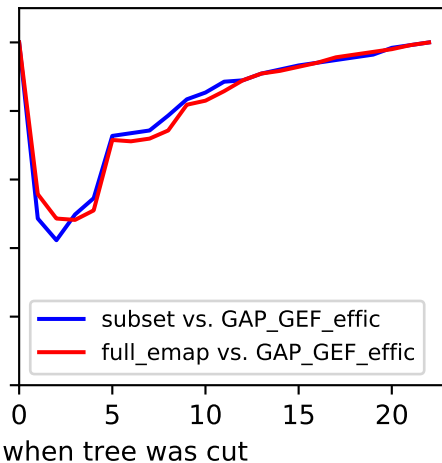
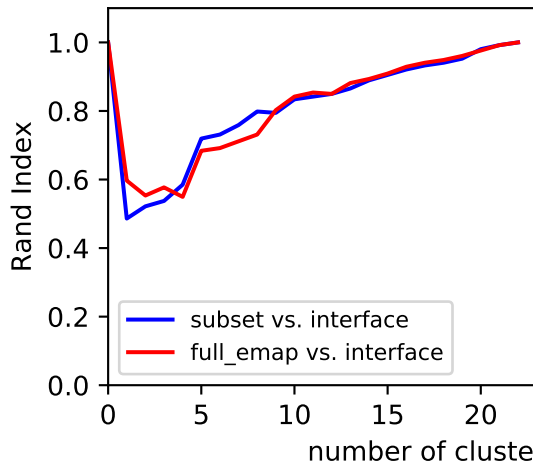
dataset = Cell Cycle Checkpoint

Interface

Δ to full_emap: -0.002

GAP/GEF effic

Δ to full_emap: 0.003



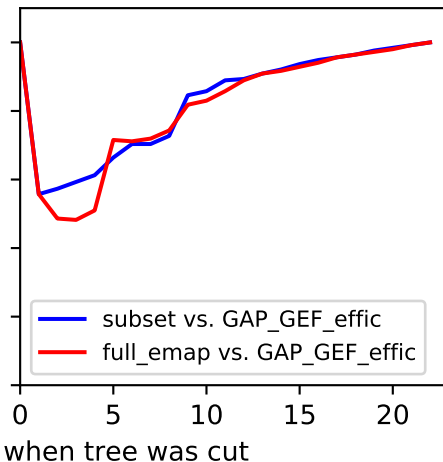
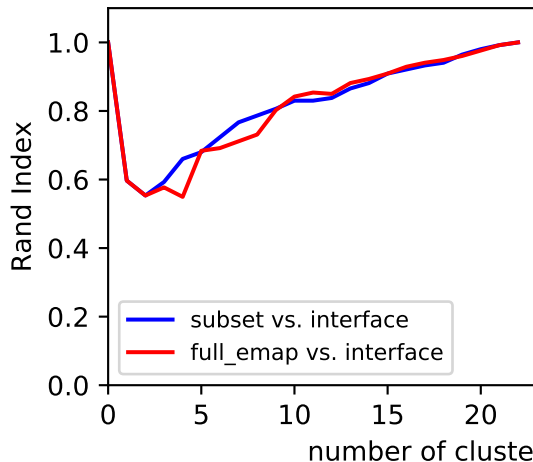
dataset = mRNA export

Interface

Δ to full_emap: 0.008

GAP/GEF effic

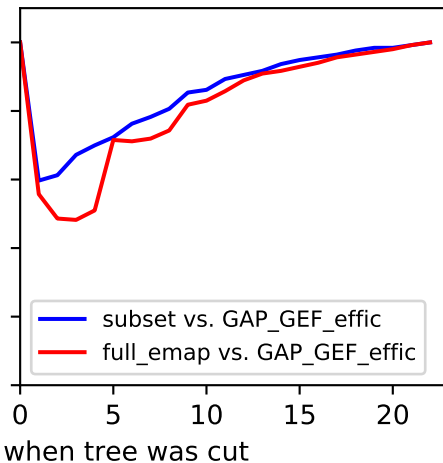
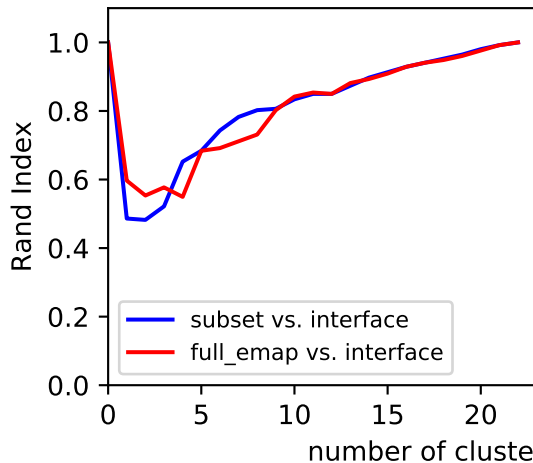
Δ to full_emap: 0.014



dataset = SAGA

Interface
 Δ to full_emap: 0.003

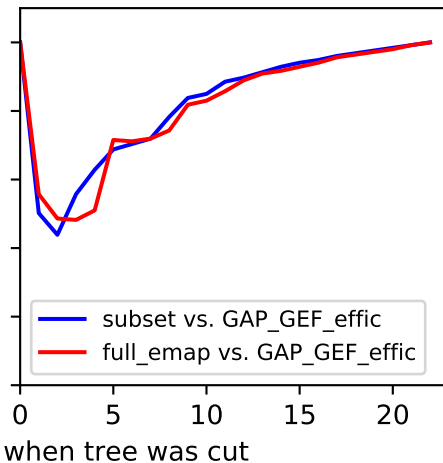
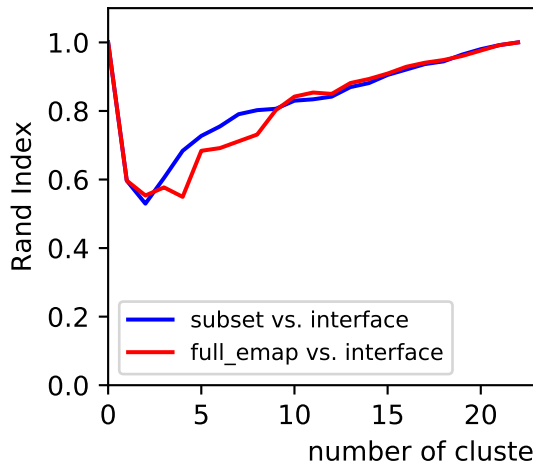
GAP/GEF effic
 Δ to full_emap: 0.041



dataset = Inner NPC

Interface
 Δ to full_emap: 0.014

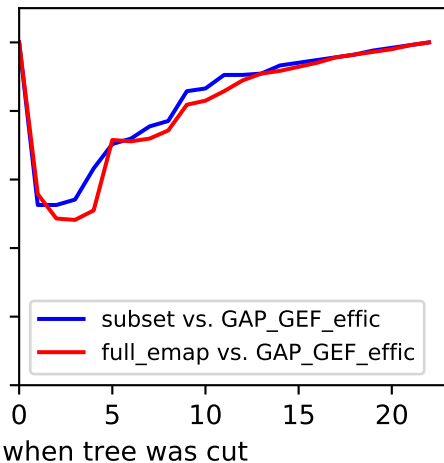
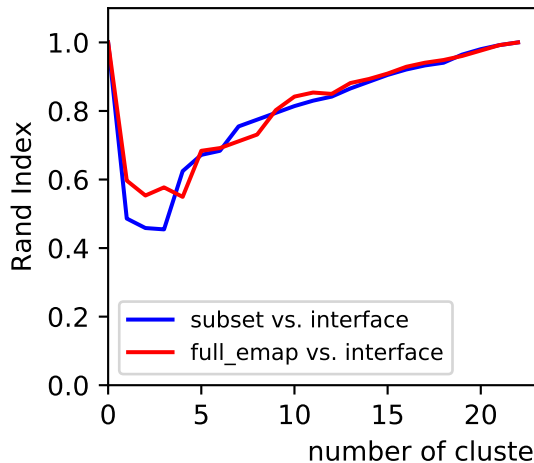
GAP/GEF effic
 Δ to full_emap: 0.01



dataset = SIT4

Interface
 Δ to full_emap: -0.013

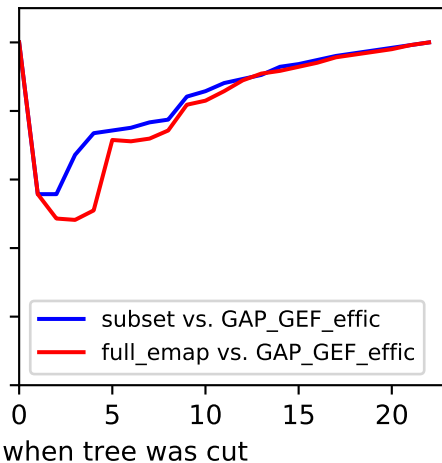
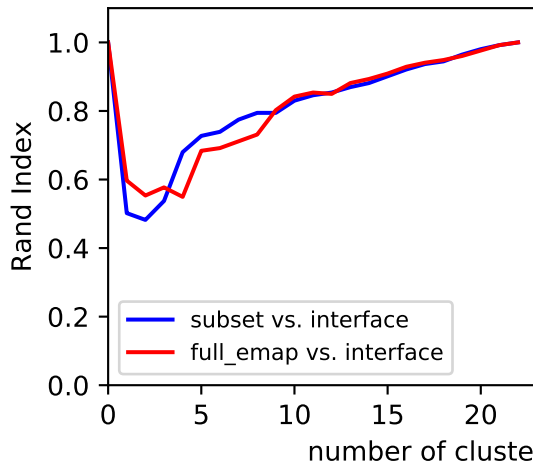
GAP/GEF effic
 Δ to full_emap: 0.019



dataset = APC

Interface
 Δ to full_emap: 0.003

GAP/GEF effic
 Δ to full_emap: 0.033



dataset = Urmylation

Interface
 Δ to full_emap: 0.02

GAP/GEF effic
 Δ to full_emap: 0.007

