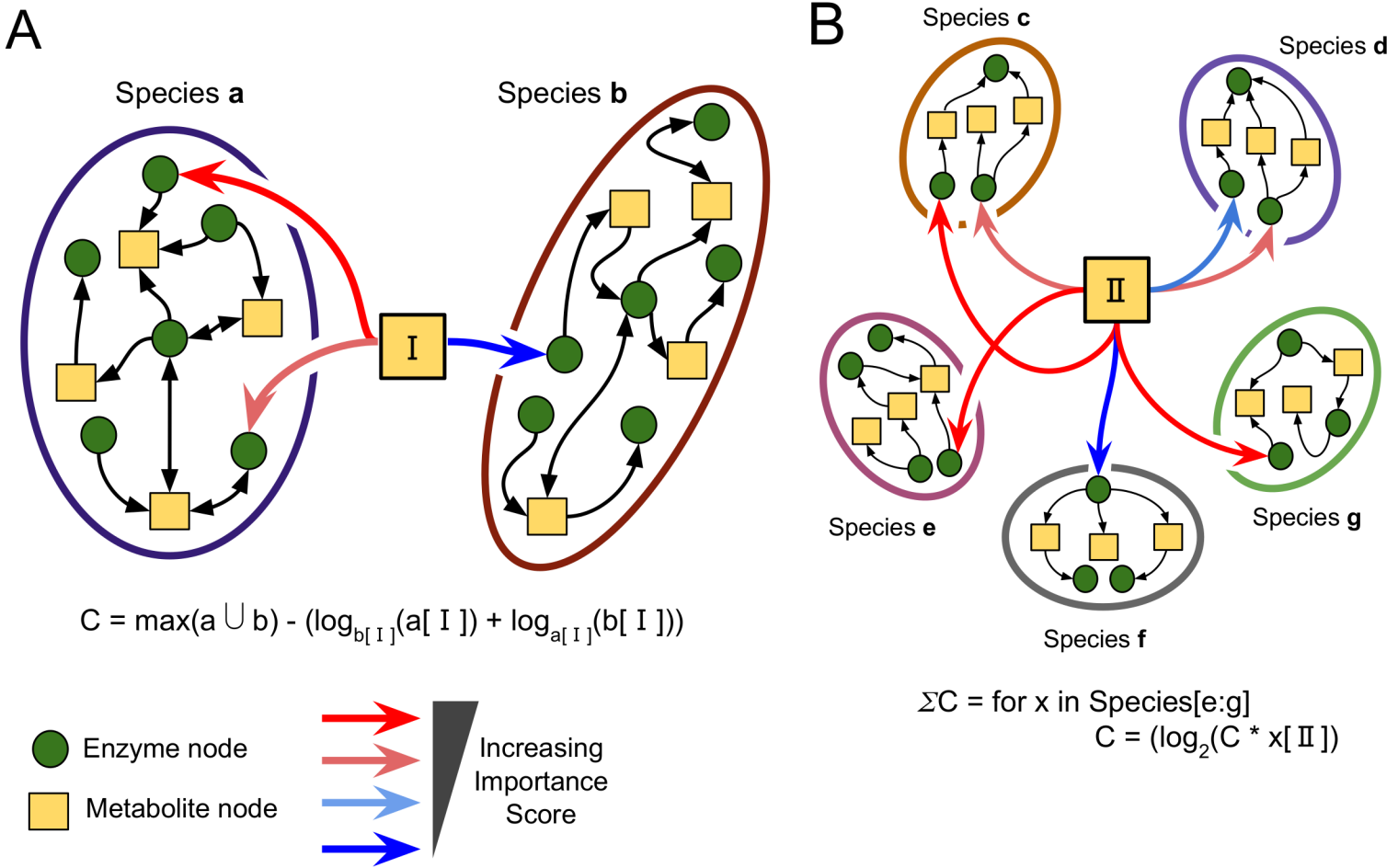
A
Species a
Species b
I
C = max(a ∪ b) - (log_b[I](a[I]) + log_a[I](b[I]))
Enzyme node
Metabolite node
Increasing
Importance
Score
B
Species c
Species d
II
Species e
Species g
Species f
ΣC = for x in Species[e:g]
C = (log_2(C * x[II])