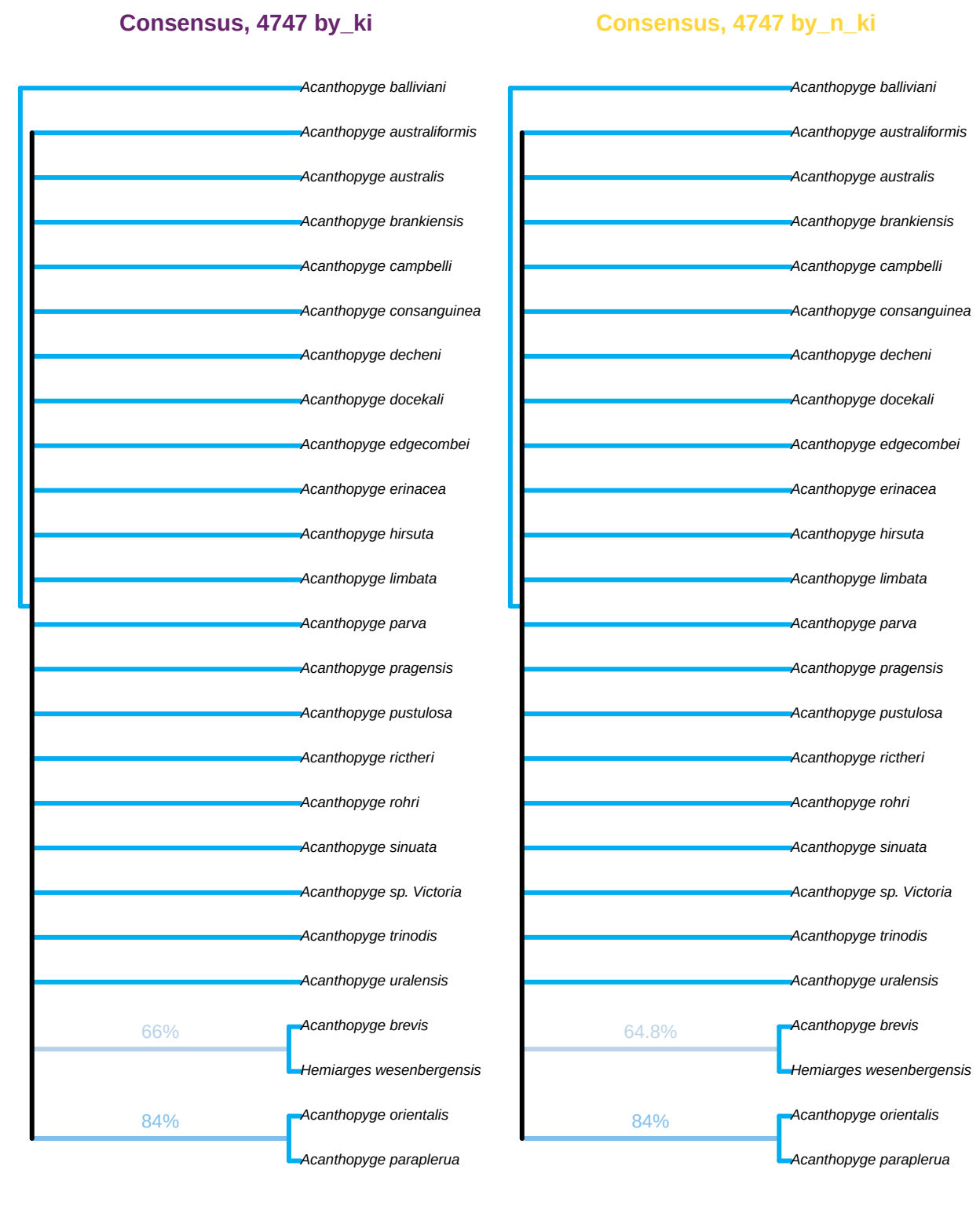
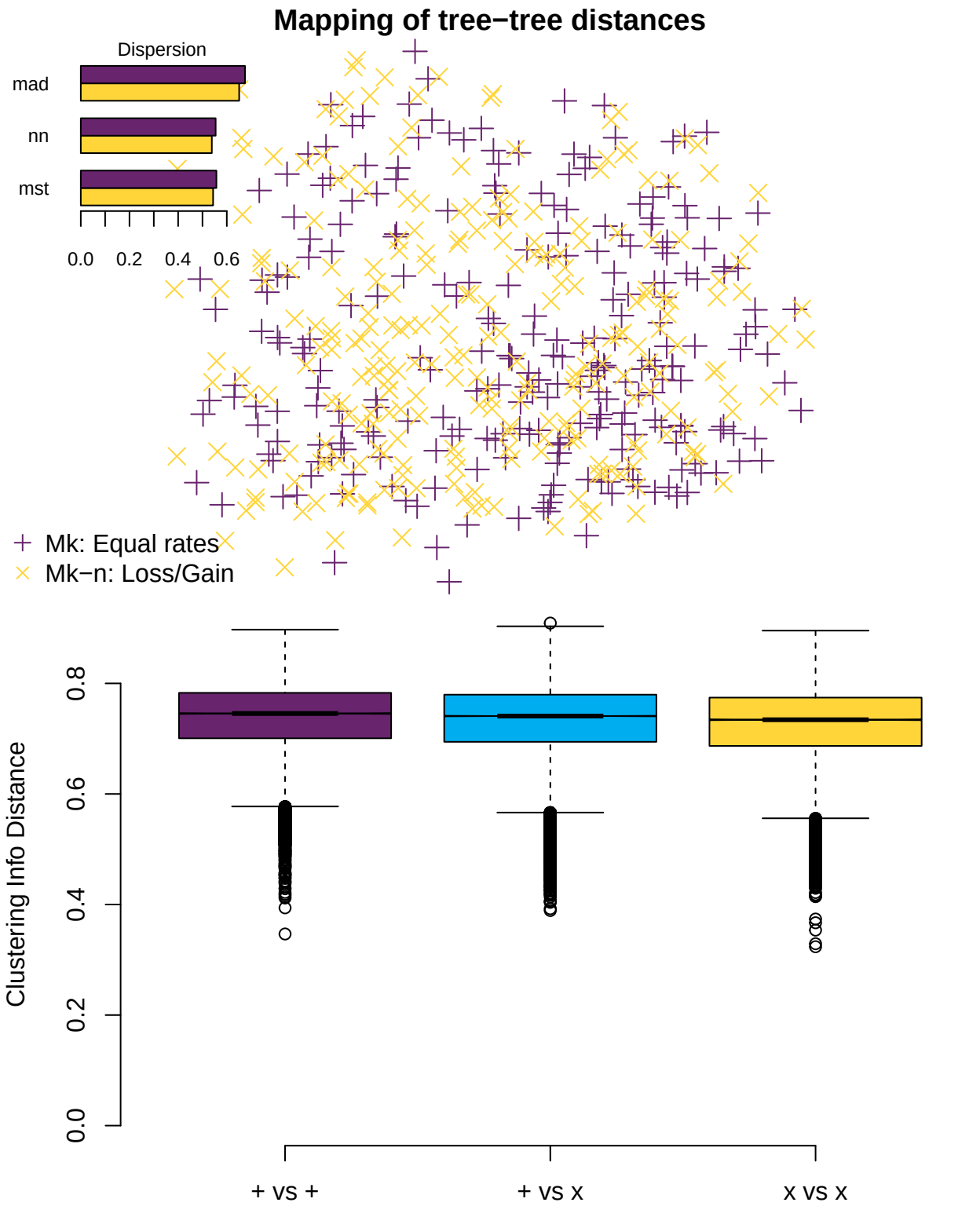

Consensus, 4747 by_ki
Acanthopyge balliviani
Acanthopyge australiformis
Acanthopyge australis
Acanthopyge brankiensis
Acanthopyge campbelli
Acanthopyge consanguinea
Acanthopyge decheni
Acanthopyge docekali
Acanthopyge edgecombei
Acanthopyge erinacea
Acanthopyge hirsuta
Acanthopyge limbata
Acanthopyge parva
Acanthopyge pragensis
Acanthopyge pustulosa
Acanthopyge rictheri
Acanthopyge rohri
Acanthopyge sinuata
Acanthopyge sp. Victoria
Acanthopyge trinodis
Acanthopyge uralensis
66%
Acanthopyge brevis
Hemiarges wesenbergensis
84%
Acanthopyge orientalis
Acanthopyge paraplerua
Consensus, 4747 by_n_ki
Acanthopyge balliviani
Acanthopyge australiformis
Acanthopyge australis
Acanthopyge brankiensis
Acanthopyge campbelli
Acanthopyge consanguinea
Acanthopyge decheni
Acanthopyge docekali
Acanthopyge edgecombei
Acanthopyge erinacea
Acanthopyge hirsuta
Acanthopyge limbata
Acanthopyge parva
Acanthopyge pragensis
Acanthopyge pustulosa
Acanthopyge rictheri
Acanthopyge rohri
Acanthopyge sinuata
Acanthopyge sp. Victoria
Acanthopyge trinodis
Acanthopyge uralensis
64.8%
Acanthopyge brevis
Hemiarges wesenbergensis
84%
Acanthopyge orientalis
Acanthopyge paraplerua


Mapping of tree−tree distances
Dispersion
mad
nn
mst
0.0
0.2
0.4
0.6
Mk: Equal rates
Mk−n: Loss/Gain
Clustering Info Distance
0.0
0.2
0.4
0.6
0.8
+ vs +
+ vs x
x vs x

Silhouette plot
n = 512
2 clusters C_j
j : n_j | $ave_{i \in Cj}$ s_i
1 : 256 | −0.007
2 : 256 | 0.008
0.0
0.2
0.4
0.6
0.8
1.0
Silhouette width s_i
Average silhouette width : 0

	Burnin	PSRF	ESS	FrechESS	mdPsESS	steppingStn	pathSmpl
Mk: Equal rates	0.05	1.003	306.16	629.929	370.816	−273.198	−273.11
Mk−n: Loss/Gain	0.1	1.00349	296.45	697.142	523.575	−273.521	−273.29