Evolutionary History and Ongoing Transmission of Phylogenetic Sublineages of *Mycobacterium tuberculosis* Beijing Genotype in China

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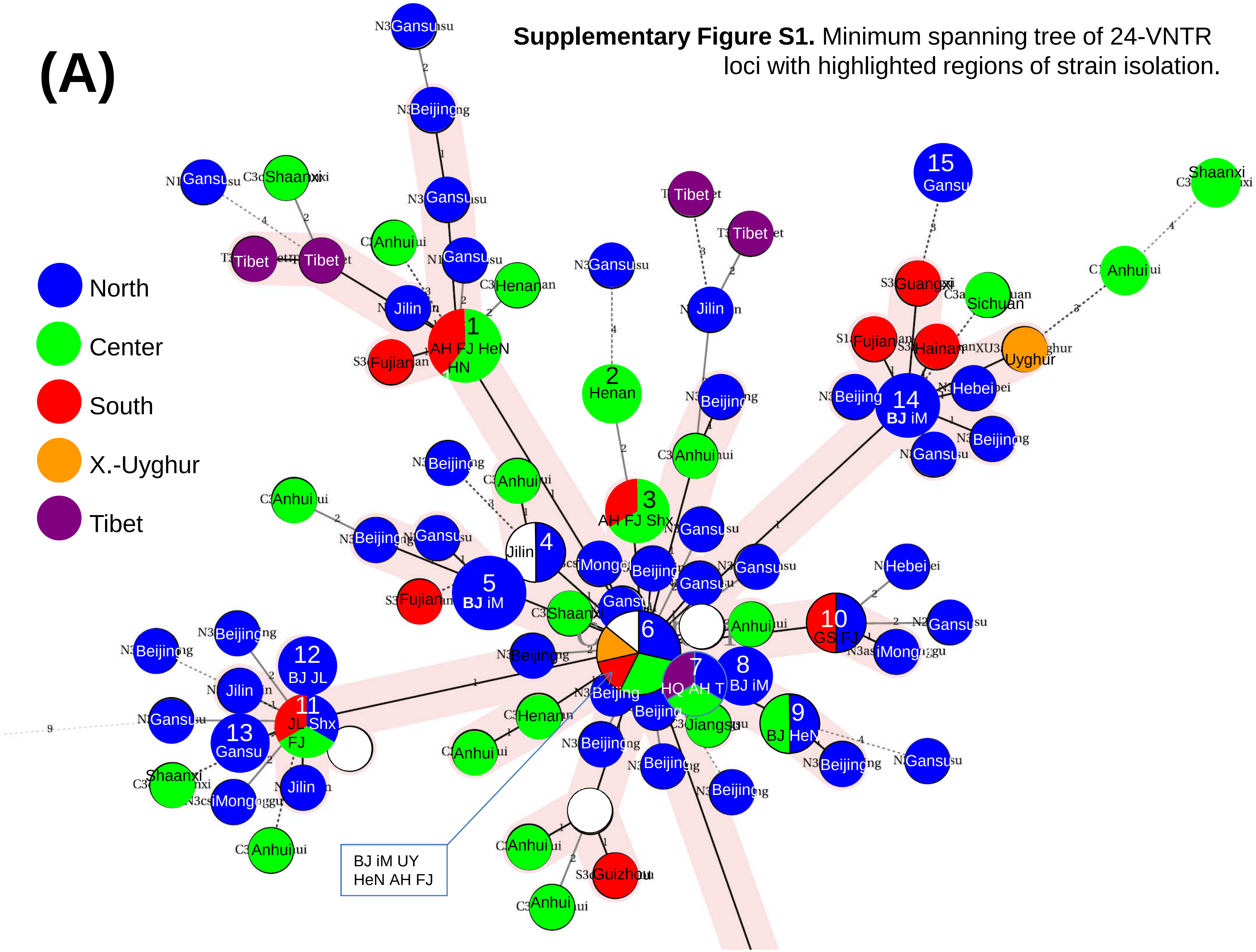
Supplementary Table S2. Genetic diversity of local *M. tuberculosis* Beijing genotype populations.

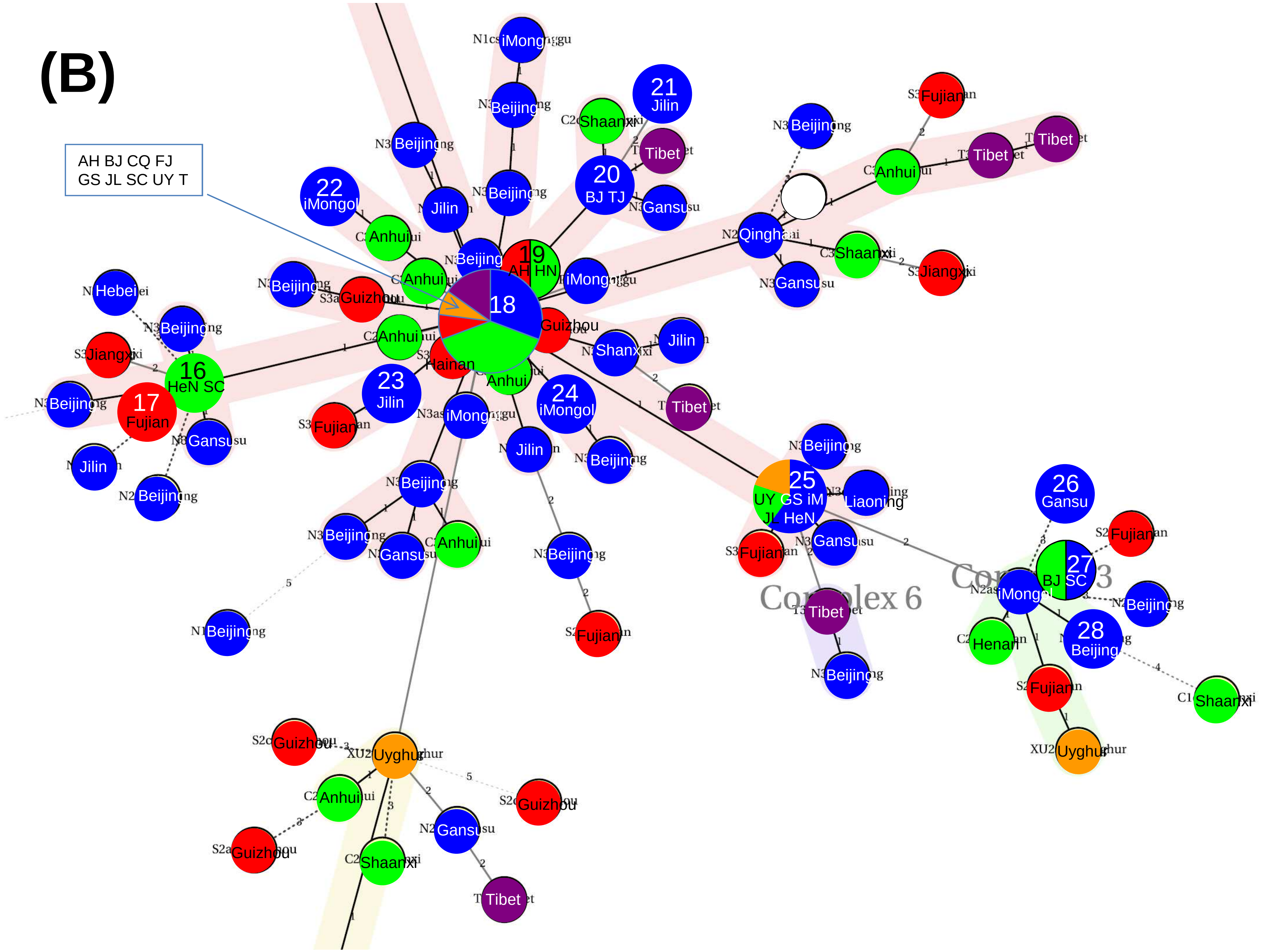
Province, region	H, mean diversity of 24 loci	N, number of SDLV branches /other branches	Cumulative index of diversity, H/N	Distance from Guizhou*, km
Tibet	0.2	19/10=1.9	0.105	1405
Gansu	0.28	13/19=0.68	0.412	986
Manchuria	0.19	25/8=3.13	0.061	2649
Guizhou	0.27	1/11=0.09	3	0
Fujian	0.20	10/8=1.25	0.160	1364
Anhui	0.17	16/8=2	0.085	1189 1260

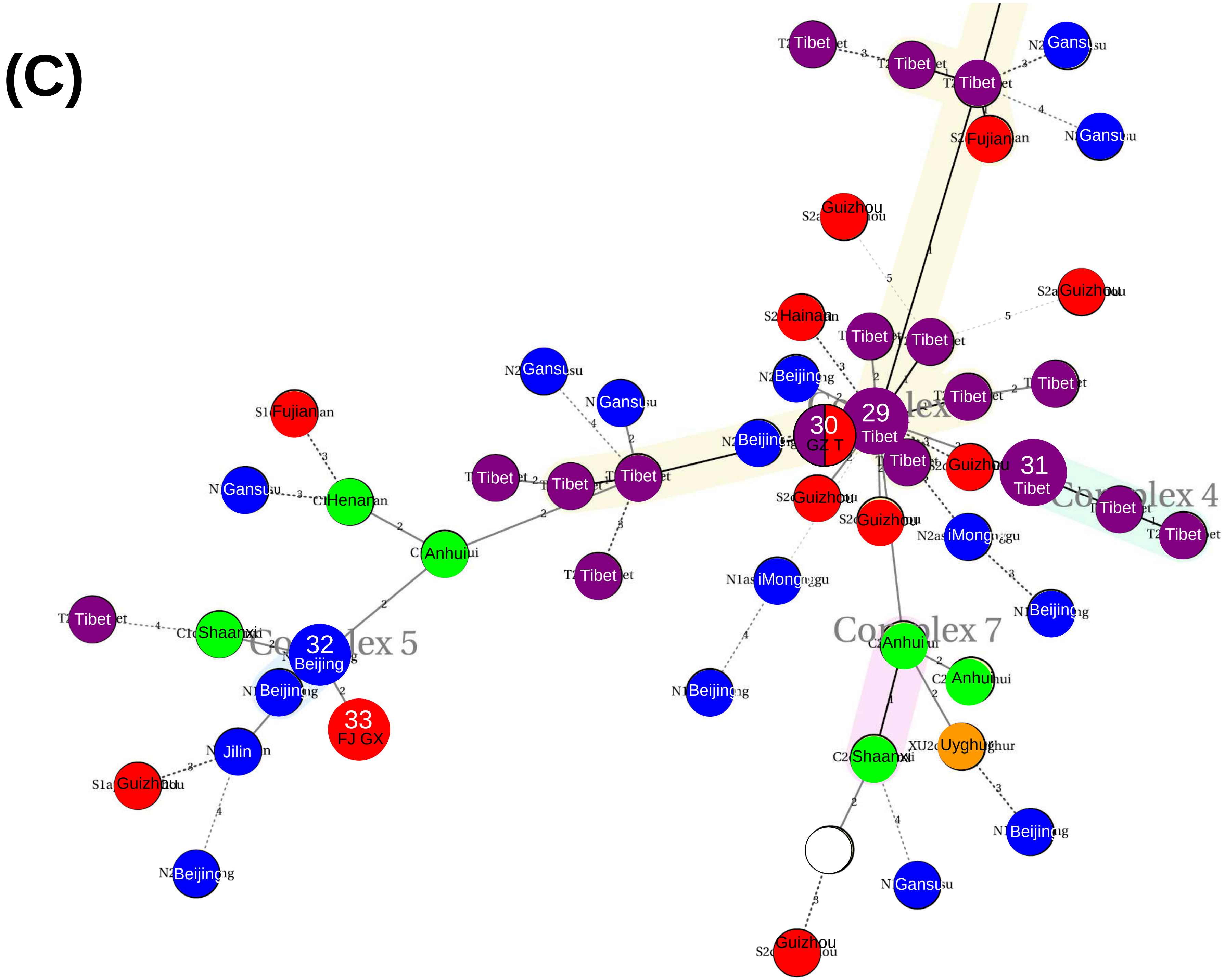
*Distance between Bijie (the largest city in Guizhou) and the largest city of the respective province/region, calculated using: http://www.distancefromto.net/



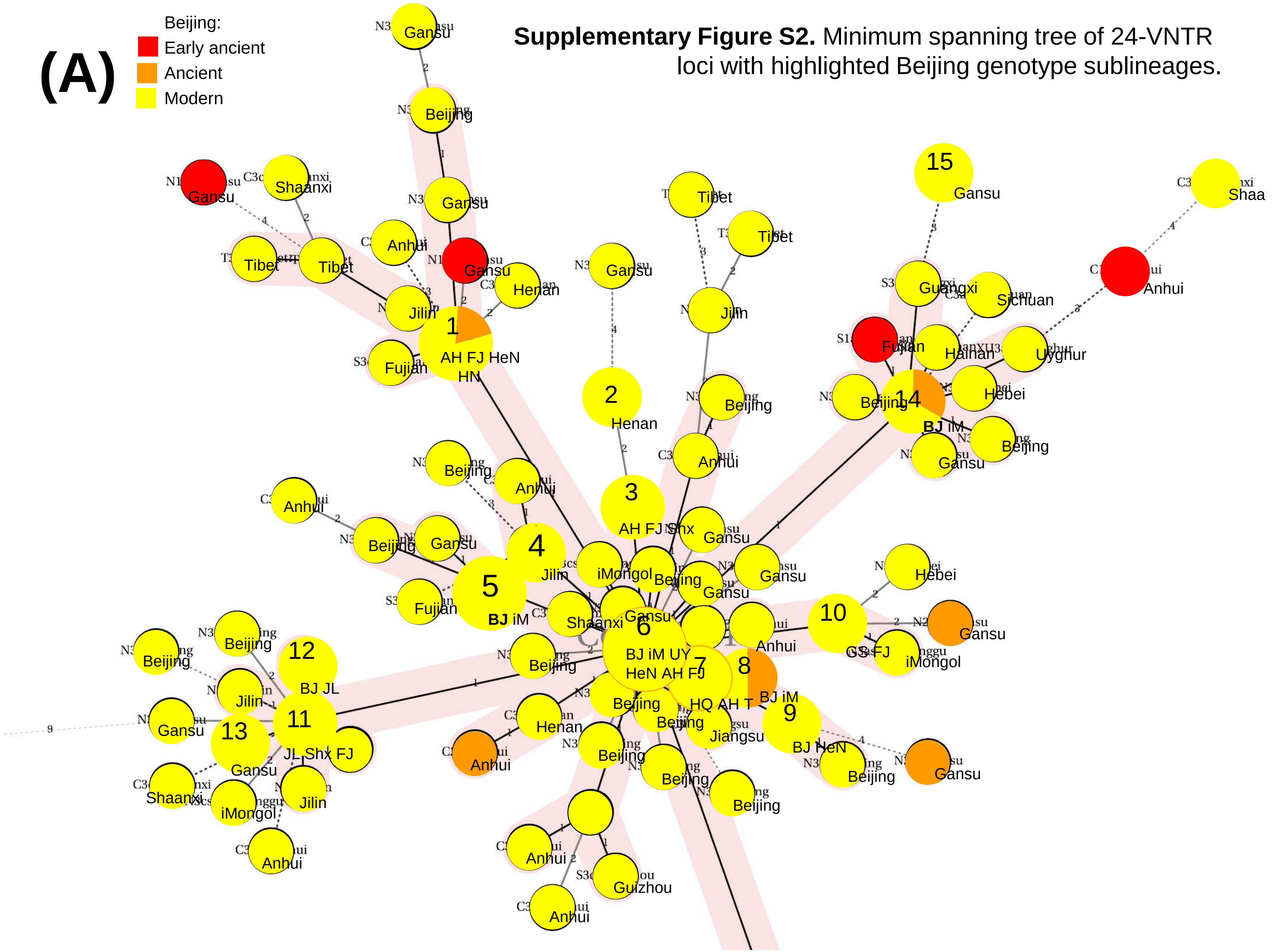
BJ iM UY
HeN AH FJ

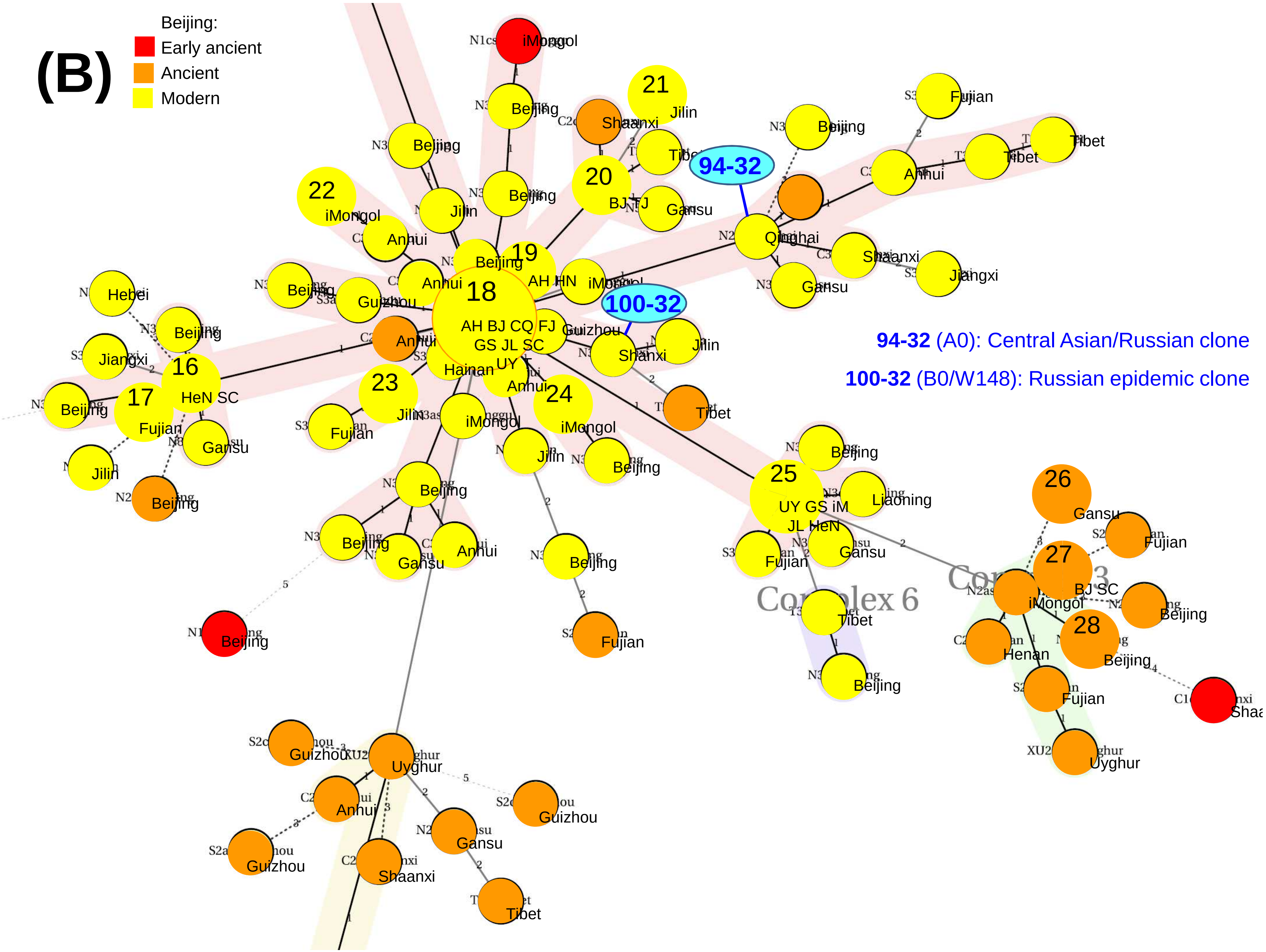
Supplementary Figure S1. Minimum spanning tree of 24-VNTR loci with highlighted regions of strain isolation.


(B)
AH BJ CQ FJ
GS JL SC UY T
18
19
AH HN
20
BJ TJ
21
Jilin
22
iMongol
23
Jilin
24
iMongol
25
UY GS iM
JL HeN
16
HeN SC
17
Fujian
26
Gansu
27
BJ SC
28
Beijing
Complex 6
Complex 3
Beijing
Jilin
Anhui
Guizhou
Hainan
Shaanxi
Tibet
Gansu
Qinghai
Jiangxi
Fujian
Hebei
Shanxi
Liaoning
Henan
iMongol
Uyghur


(C)
Complex 4
Complex 5
Complex 7
29
Tibet
30
GZ T
31
Tibet
32
Beijing
33
FJ GX
Tibet
Gansu
Fujian
Guizhou
Hainan
Beijing
Henan
Anhui
Shaanxi
Jilin
iMong
Uyghur
1
2
3
4
5

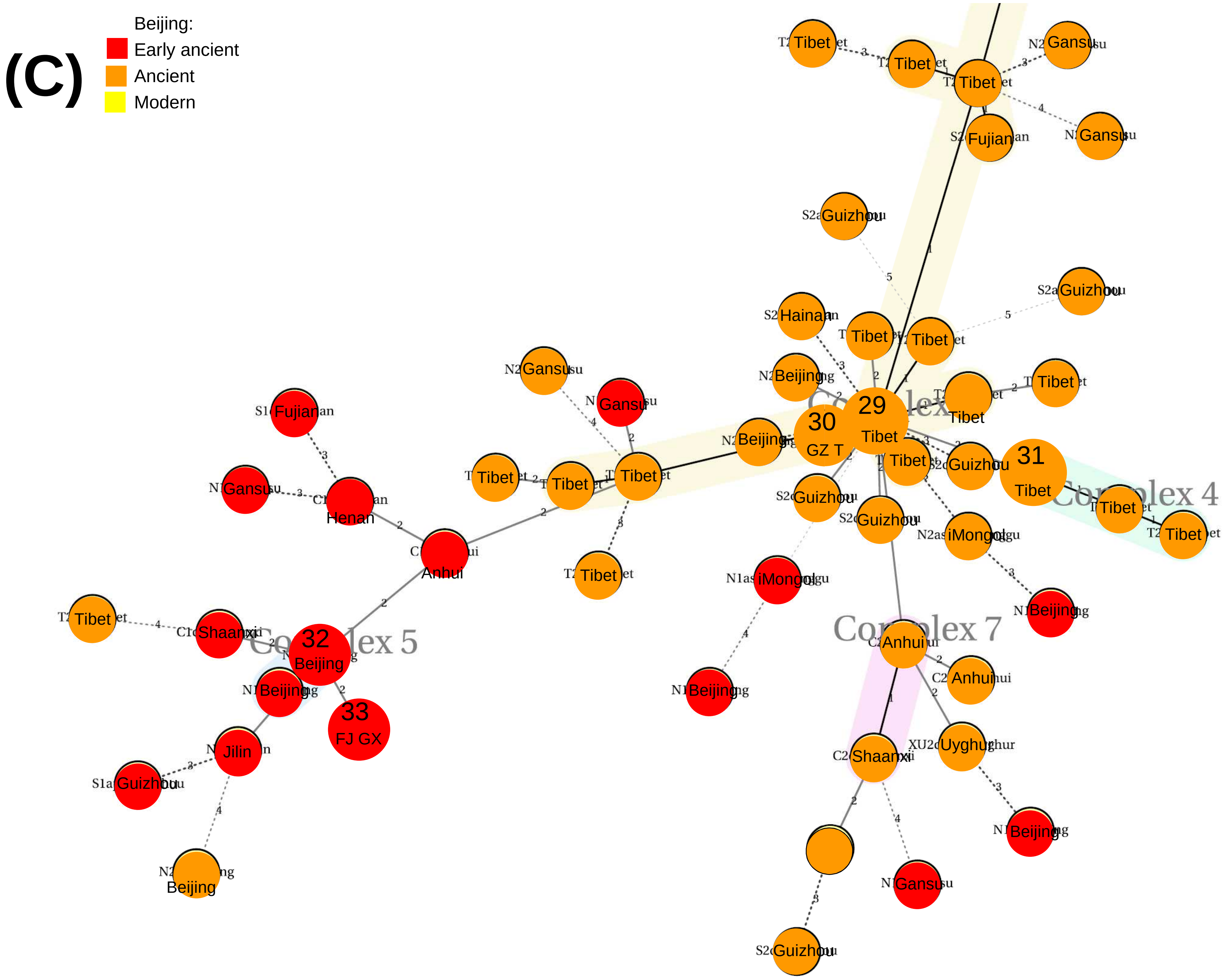
Supplementary Figure S2. Minimum spanning tree of 24-VNTR loci with highlighted Beijing genotype sublineages.



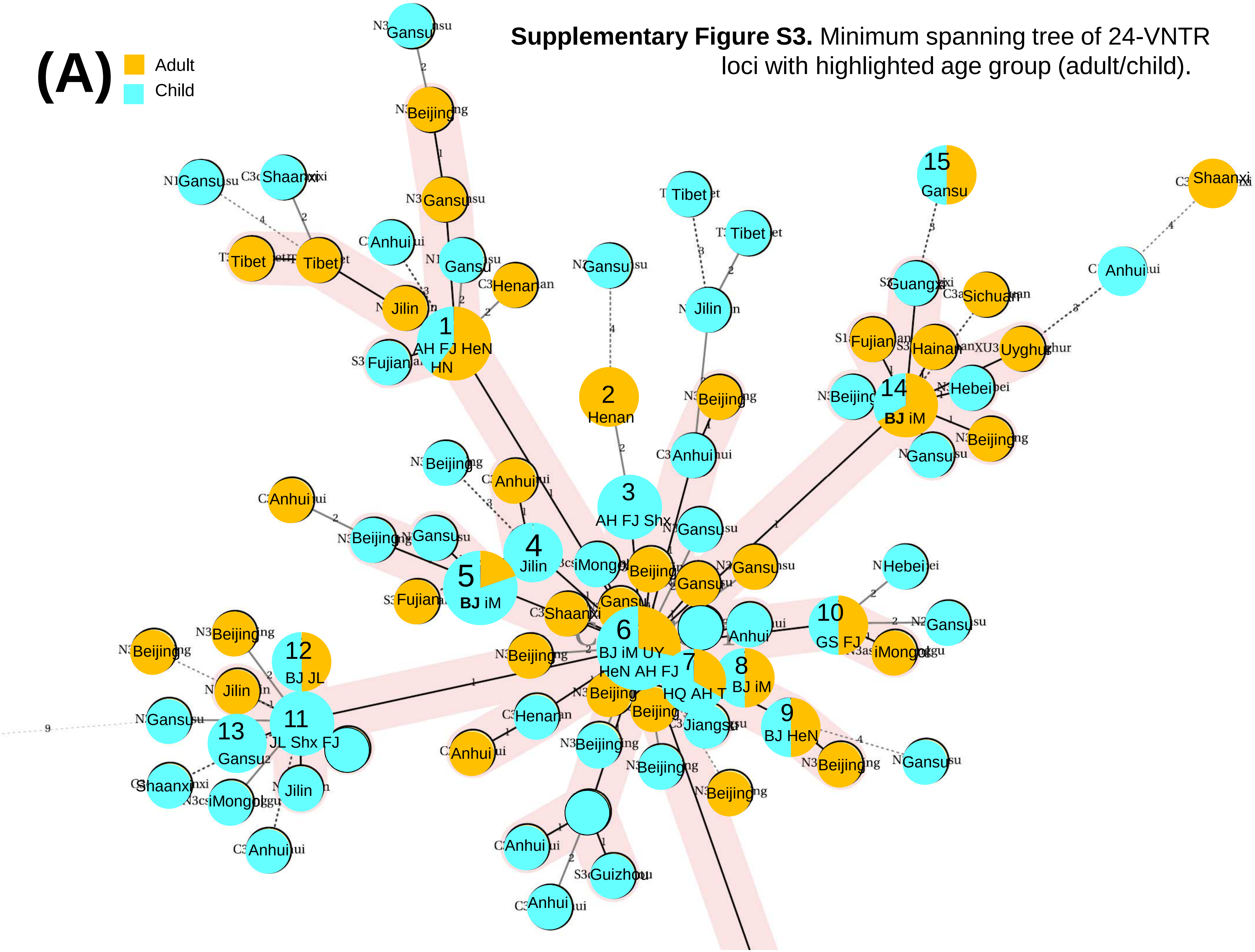


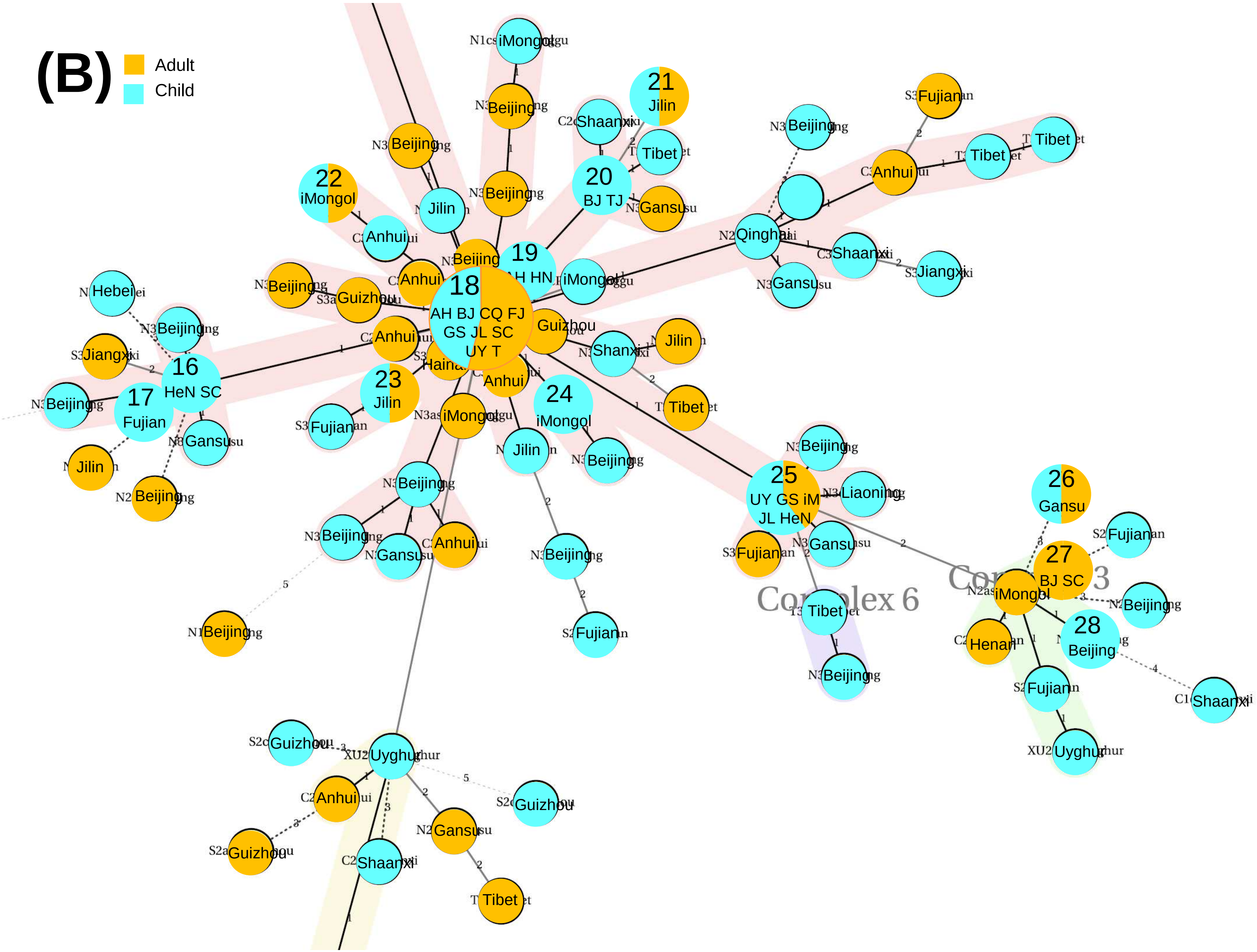
94-32 (A0): Central Asian/Russian clone

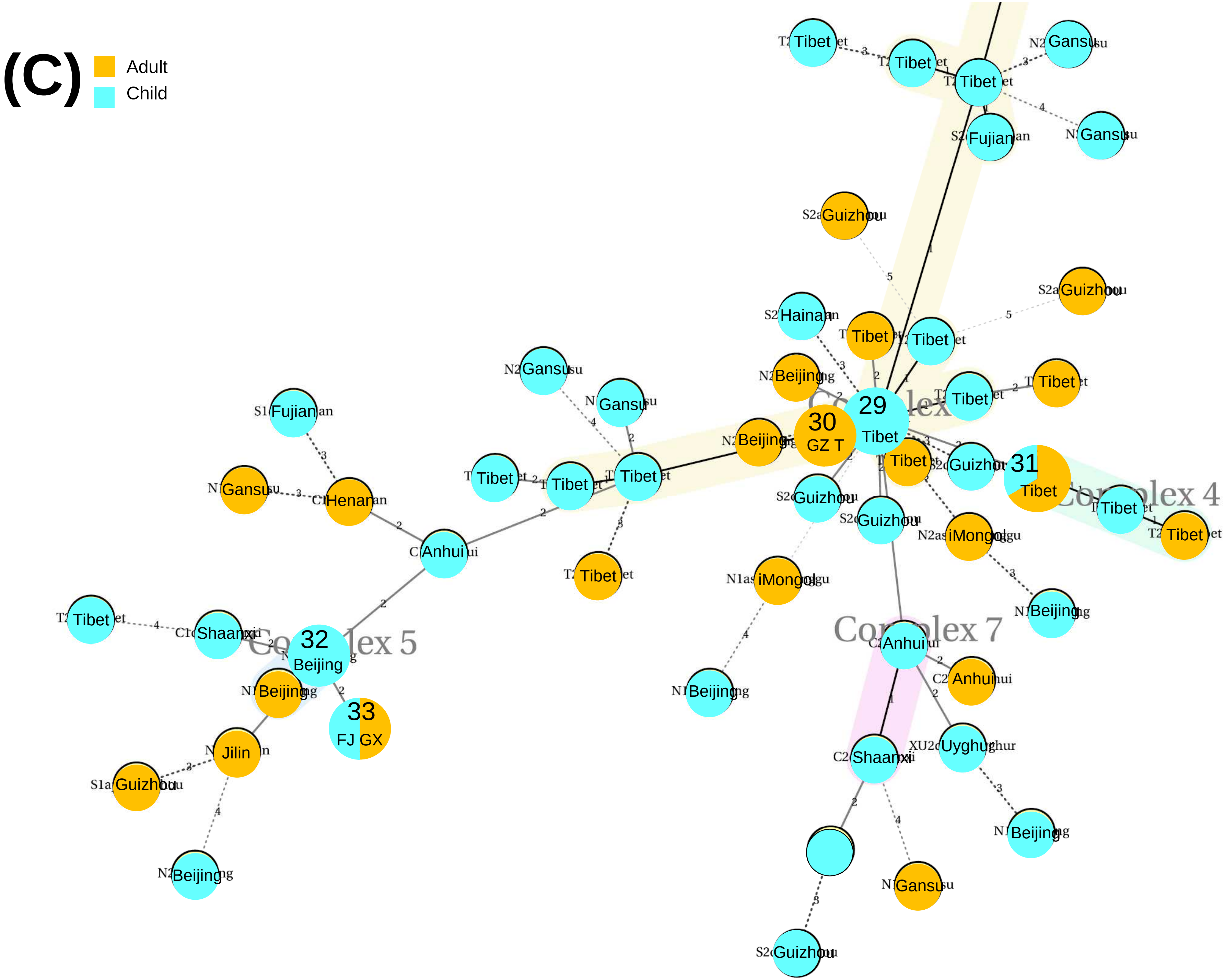
100-32 (B0/W148): Russian epidemic clone


(C)
Beijing:
🟥 Early ancient
🟧 Ancient
🟨 Modern
Tibet
Gansu
Fujian
Guizhou
Hainan
Beijing
29
Tibet
30
GZ T
31
Tibet
Complex 4
iMongol
Complex 7
Anhui
Shaanxi
Uyghur
Henan
Complex 5
32
Beijing
33
FJ GX
Jilin
1
2
3
4
5

Supplementary Figure S3. Minimum spanning tree of 24-VNTR loci with highlighted age group (adult/child).

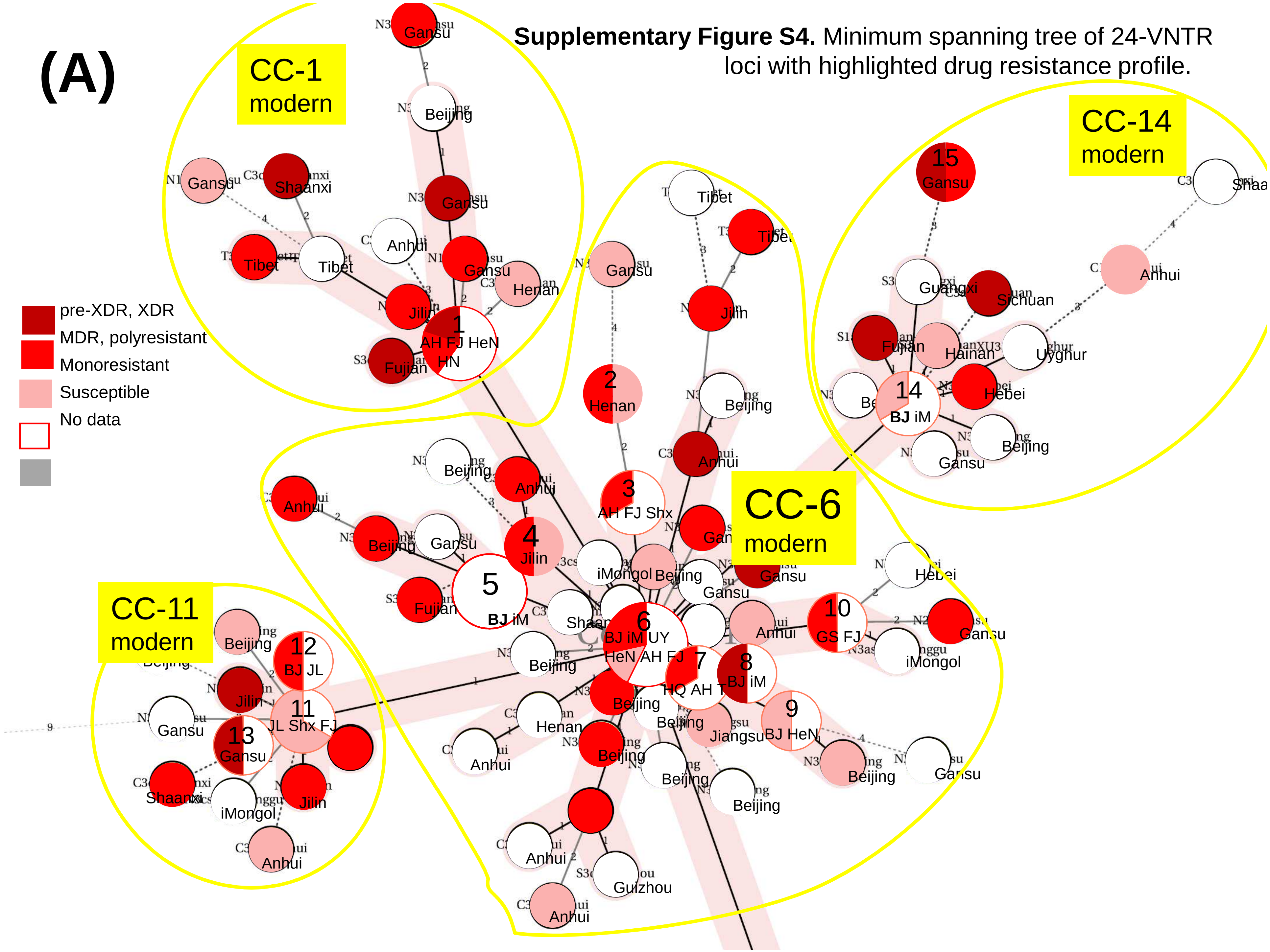


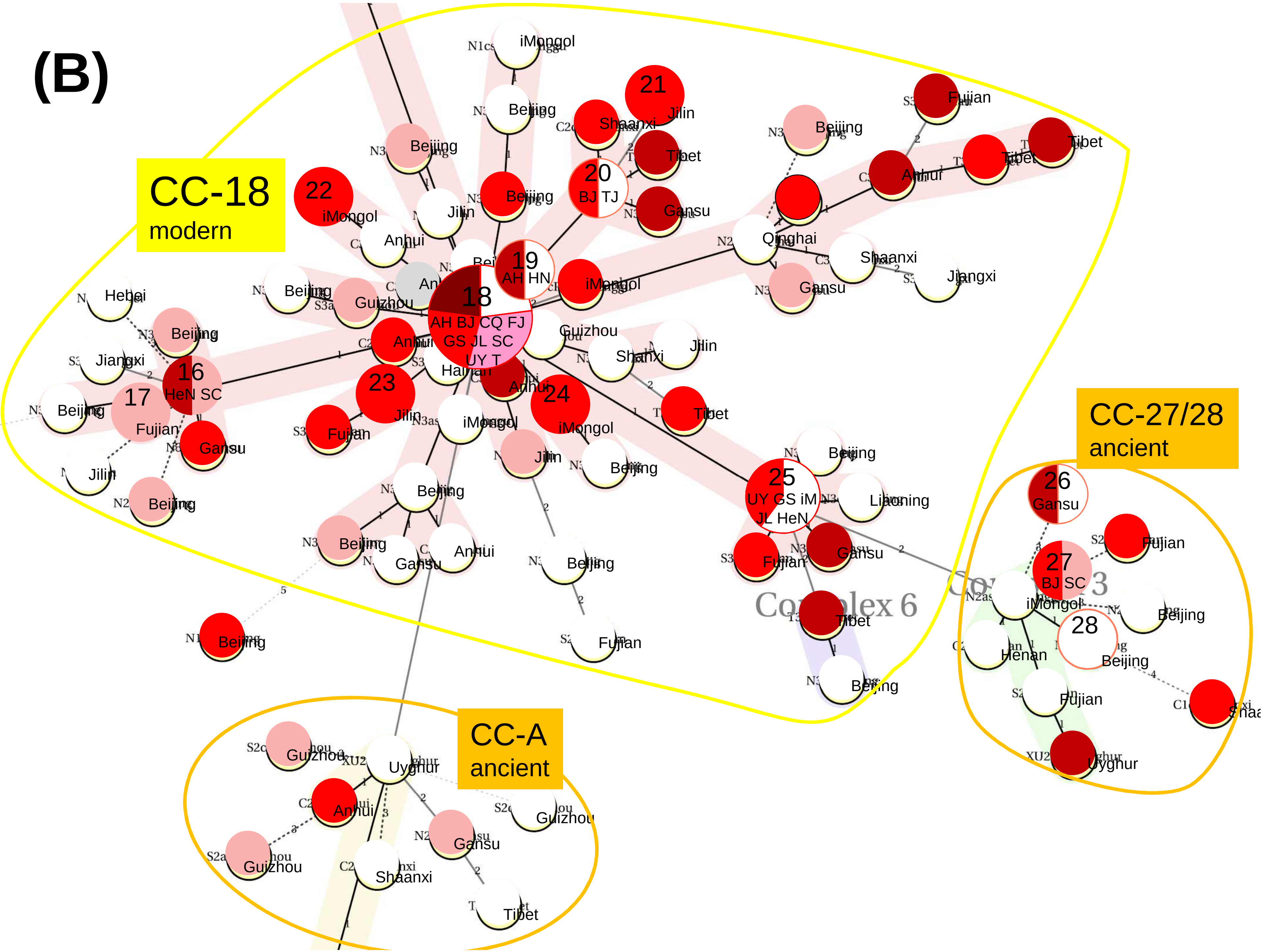

(B)
Adult
Child
18
AH BJ CQ FJ
GS JL SC
UY T
16
HeN SC
17
Fujian
19
AH HN
20
BJ TJ
21
Jilin
22
iMongol
23
Jilin
24
iMongol
25
UY GS iM
JL HeN
26
Gansu
27
BJ SC
28
Beijing
Complex 6
Complex 3
Beijing
iMongol
Jilin
Anhui
Guizhou
Tibet
Gansu
Shaanxi
Qinghai
Jiangxi
Fujian
Hebei
Hainan
Shanxi
Liaoning
Henan
Uyghur
1
2
3
4
5


(C)
Adult
Child
29
Tibet
30
GZ T
31
Tibet
32
Beijing
33
FJ GX
Complex 4
Complex 5
Complex 7
Tibet
Gansu
Fujian
Guizhou
Hainan
Beijing
iMongol
Anhui
Henan
Shaanxi
Jilin
Uyghur

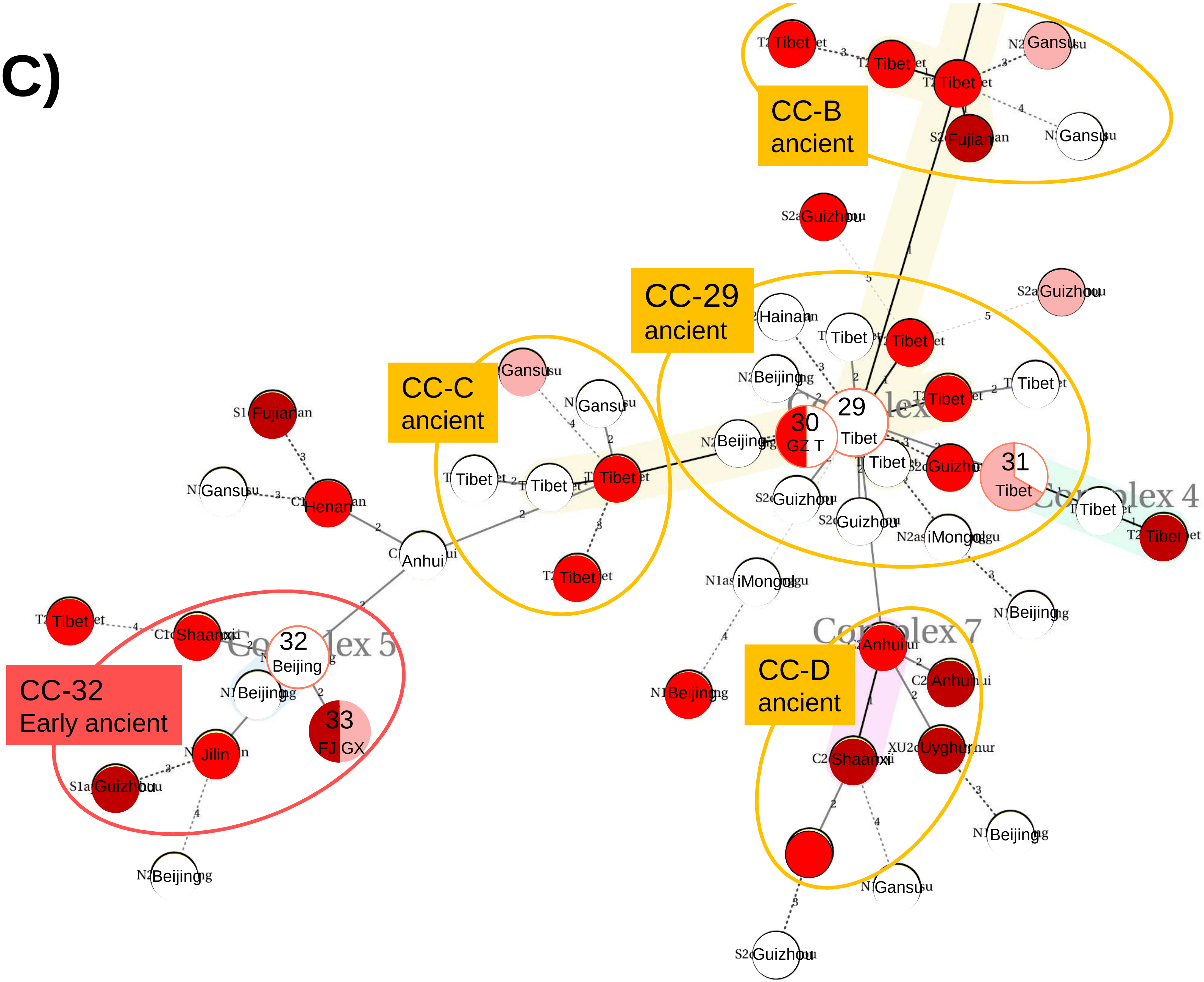
Supplementary Figure S4. Minimum spanning tree of 24-VNTR loci with highlighted drug resistance profile.

(A)

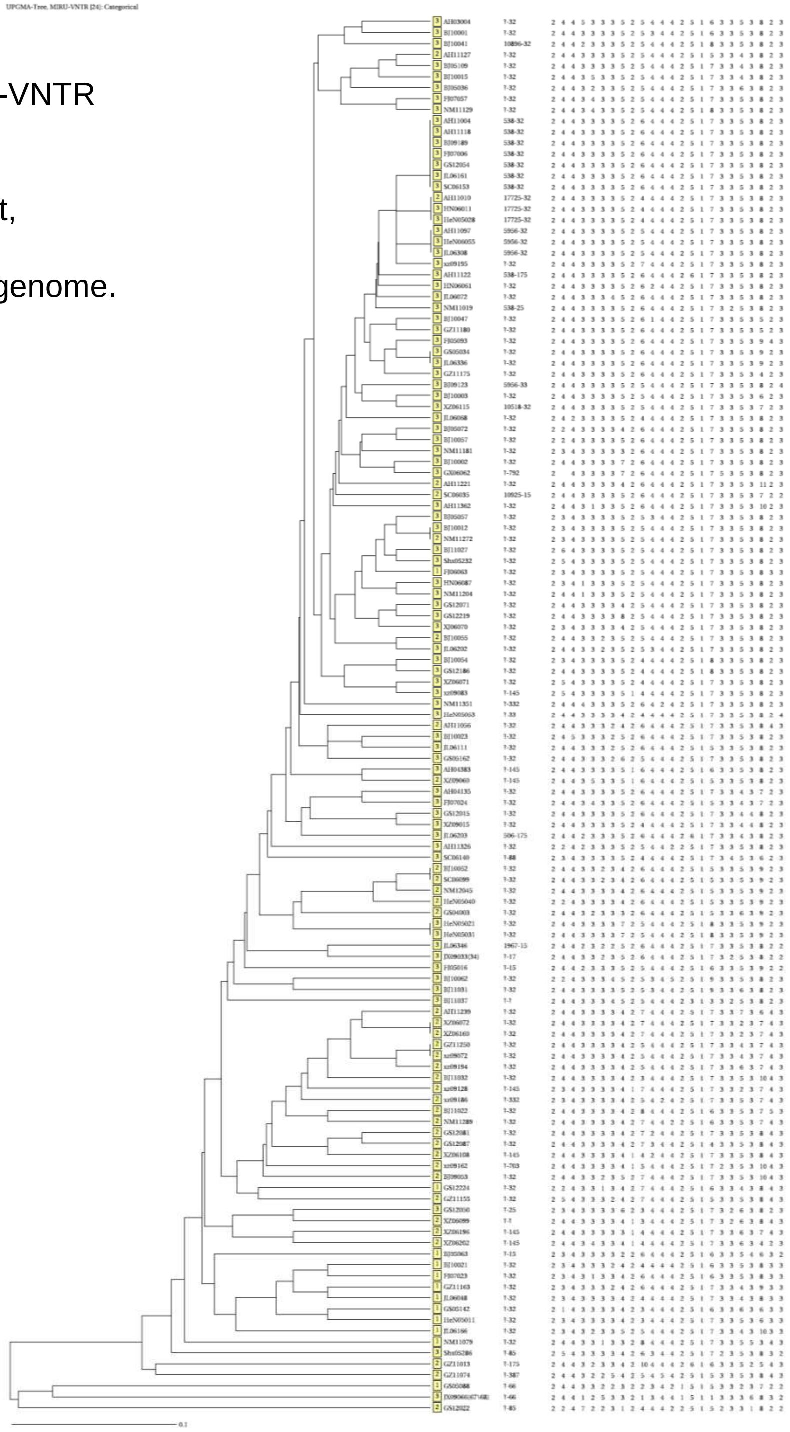



(B)
CC-18
modern
CC-27/28
ancient
CC-A
ancient
Complex 6
Complex 3
18
AH BJ CQ FJ
GS JL SC
UY T
19
AH HN
20
BJ TJ
21
22
23
24
25
UY GS iM
JL HeN
16
HeN SC
17
26
27
BJ SC
28
iMongol
Beijing
Jilin
Anhui
Guizhou
Hebei
Jiangxi
Fujian
Gansu
Hainan
Shaanxi
Tibet
Qinghai
Liaoning
Henan
Uyghur

(C)


CC-B
ancient
CC-29
ancient
CC-C
ancient
CC-32
Early ancient
CC-D
ancient
29
Tibet
30
GZ T
31
Tibet
32
Beijing
33
FJ GX
Complex 4
Complex 5
Complex 7
Tibet
Gansu
Fujian
Guizhou
Hainan
Beijing
iMongol
Henan
Anhui
Shaanxi
Jilin
Uyghur

Supplementary Figure S5. UPGMA dendrogram of 24-MIRU-VNTR profiles of M. tuberculosis Beijing genotype isolates from adult TB patients in China.
Number code designates Beijing sublineages: 1 - early ancient, 2 - ancient, 3 - modern. Mlva15-9 code was assigned by MIRU-VNTRplus.org. 24-loci order according to their order in genome.



Supplementary Figure S6. UPGMA dendrogram of 24-MIRU-VNTR profiles of M. tuberculosis Beijing genotype isolates from children TB patients in China.
Number code designates Beijing sublineages: 1 - early ancient, 2 - ancient, 3 - modern. Mlva15-9 code was assigned by MIRU-VNTRplus.org. 24-loci order according to their order in genome.