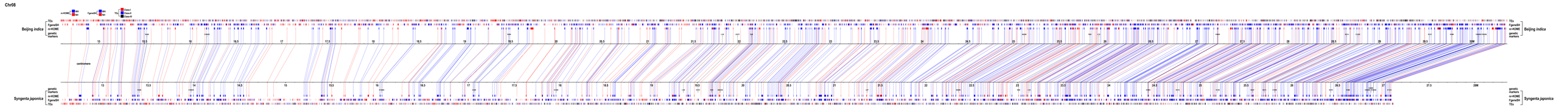
Chr08
m-KOME
Bin
Gap
FgenesH
Bin
Gap
TEs
Class-I
Class-II
Class-III
Beijing indica
TEs
FgenesH
m-KOME
genetic markers
15
15.5
16
16.5
17
17.5
18
18.5
19
19.5
20
20.5
21
21.5
22
22.5
23
23.5
24
24.5
25
25.5
26
26.5
27
27.5
28
28.5
29
29.5
30M
centromere
12
13.5
14
14.5
15
15.5
16
16.5
17
17.5
18
18.5
19
19.5
20
20.5
21
21.5
22
22.5
23
23.5
24
24.5
25
25.5
26
26.5
27
27.5
28M
Syngenta japonica
genetic markers
m-KOME
FgenesH
TEs