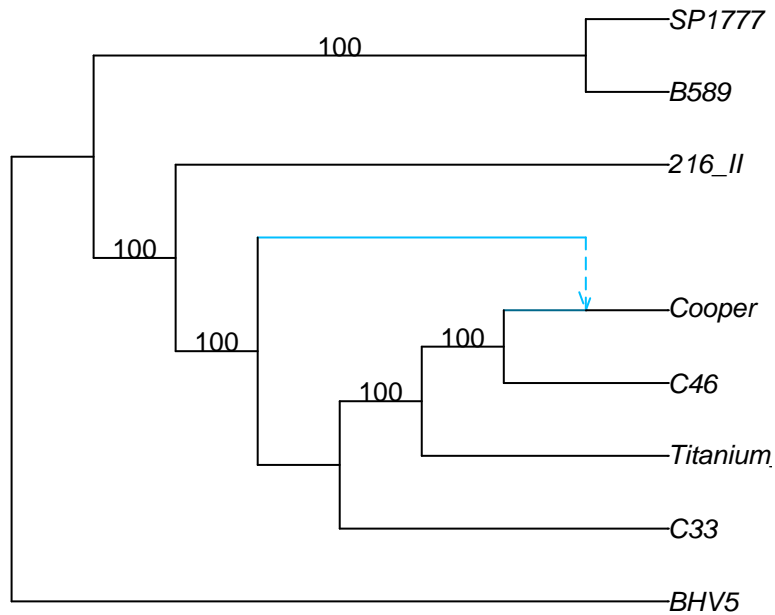
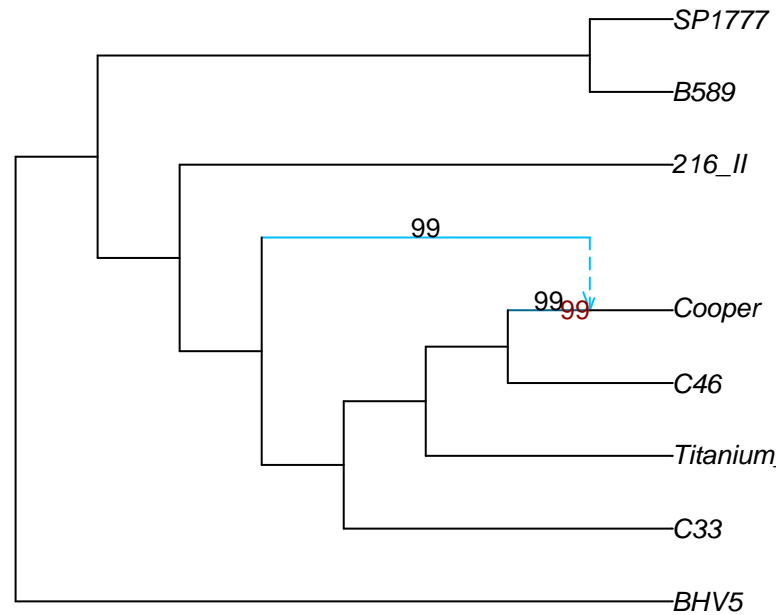


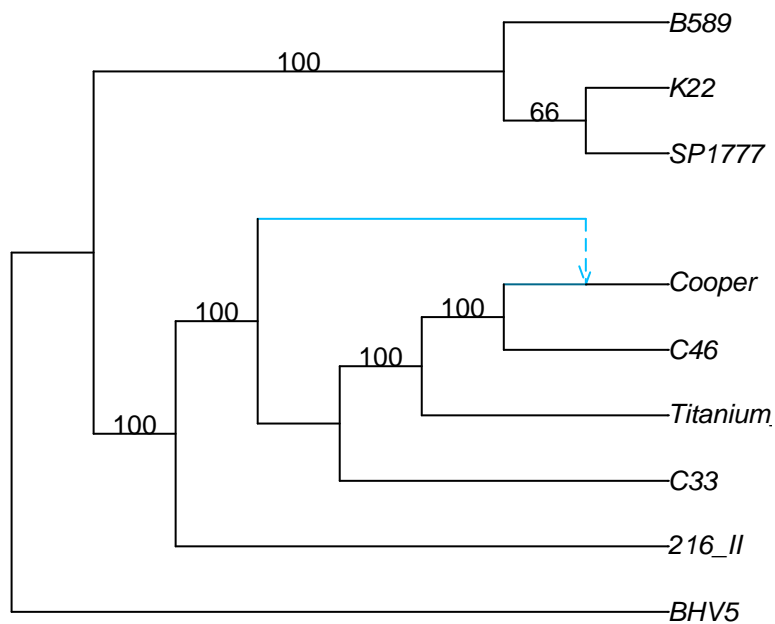
from SNPs, sets 1b (top) and 1c (bottom)



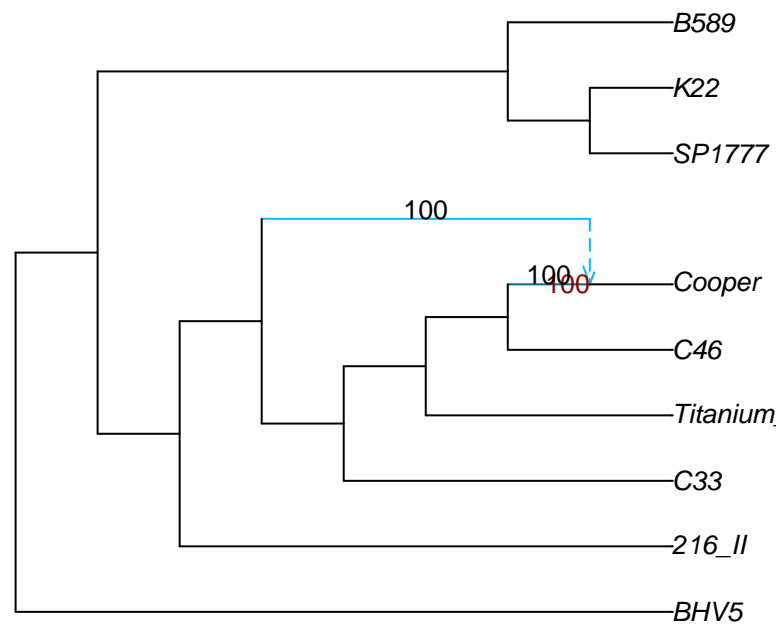
support for major tree edges



support for hybrid edges & full reticulation

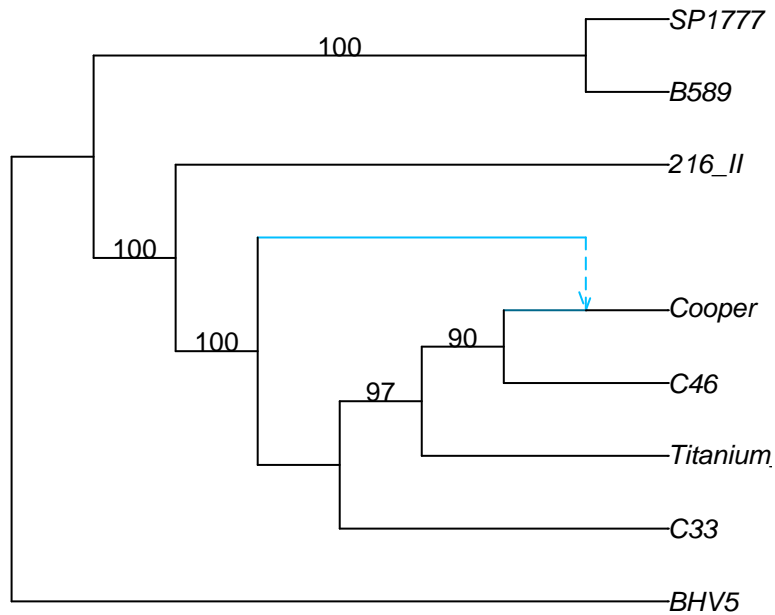


support for major tree edges

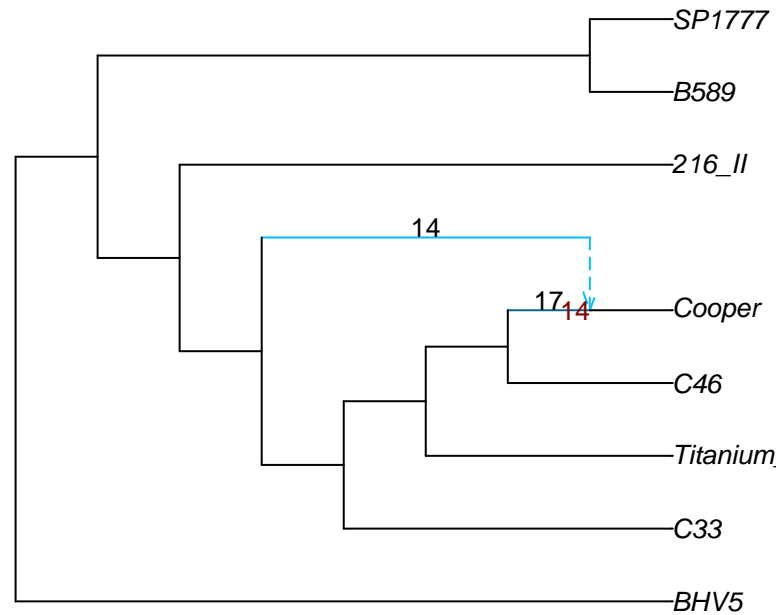


support for hybrid edges & full reticulation

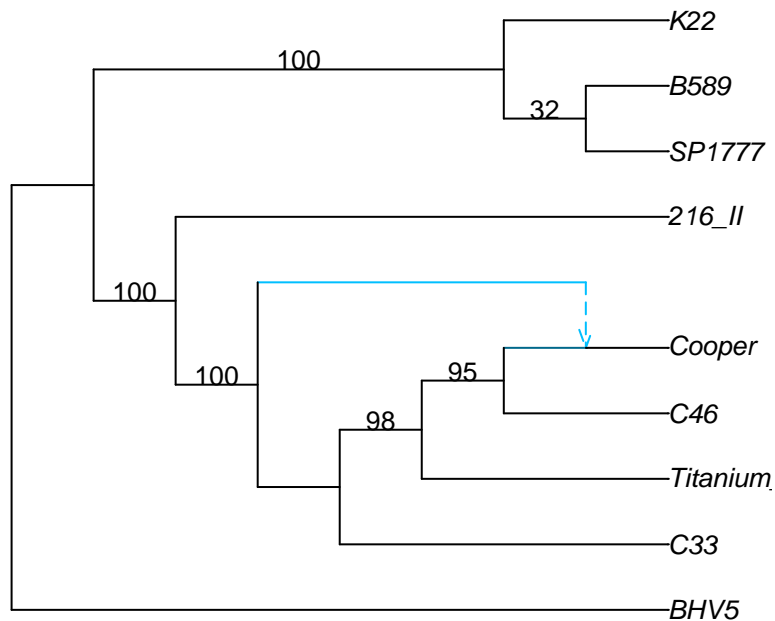
from genetrees, sets 1b (top) and 1c (bottom)



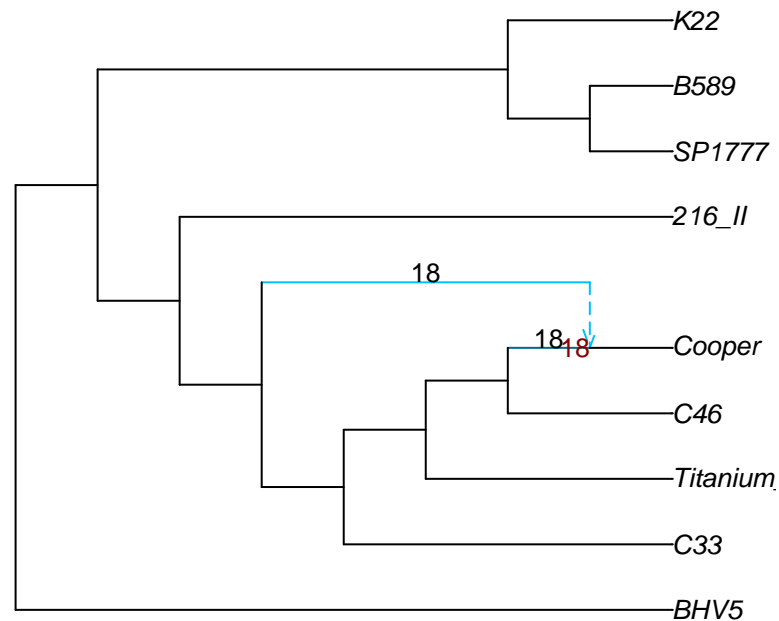
support for major tree edges



support for hybrid edges & full reticulation

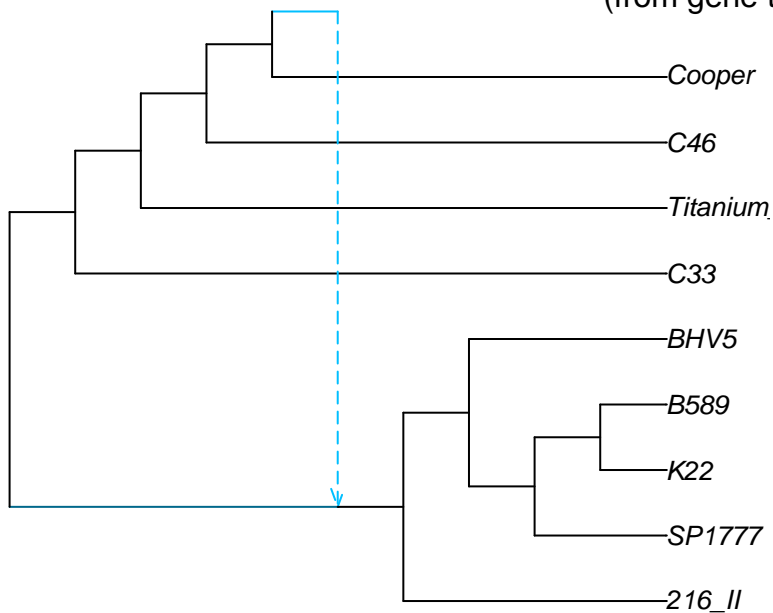


support for major tree edges

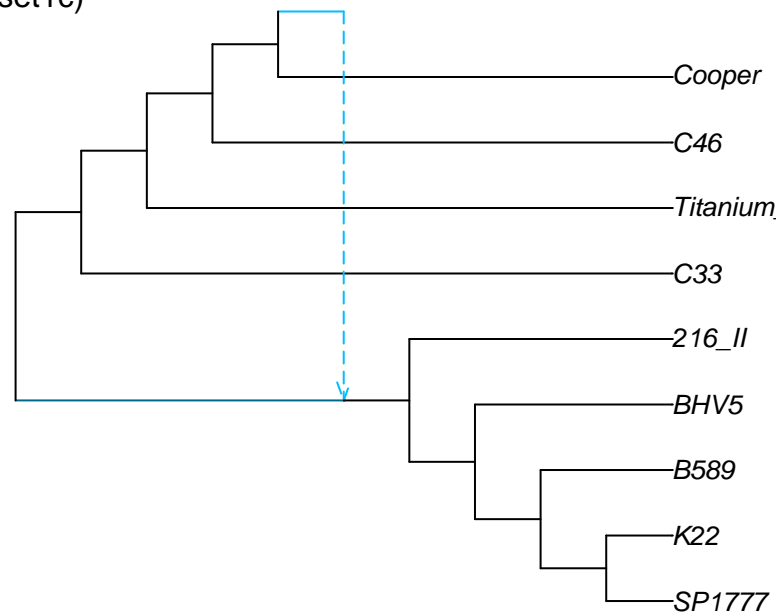


support for hybrid edges & full reticulation

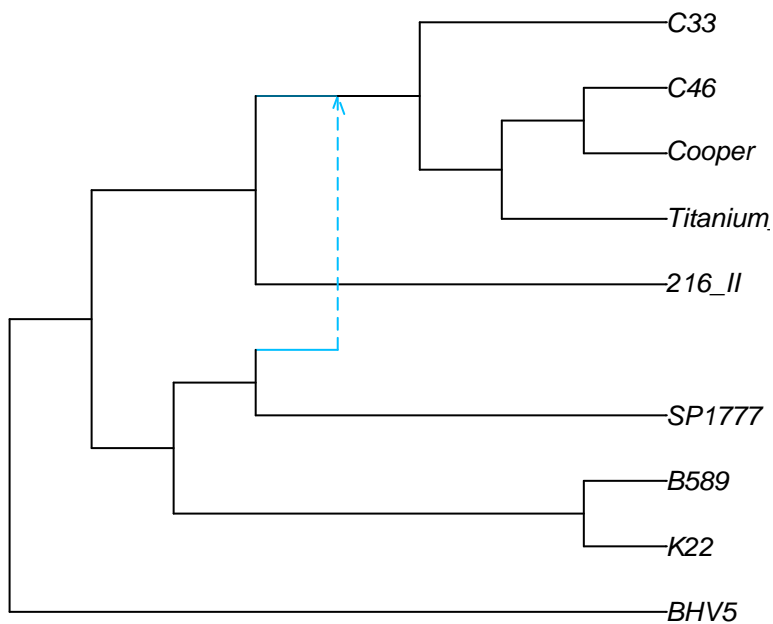
example bootstrap: often (~10%) the direction of gene flow is reversed  
(from gene trees, set1c)



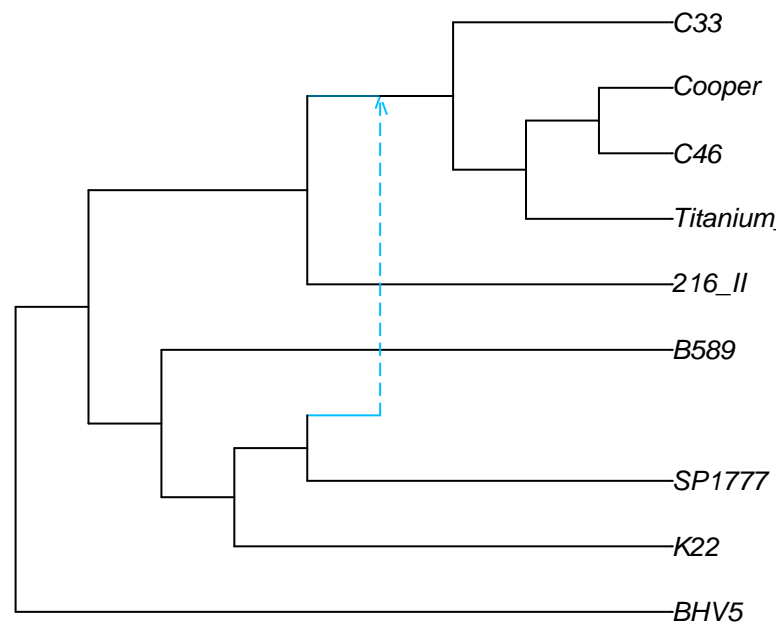
bootstrap replicate 3



bootstrap replicate 6



bootstrap replicate 29



bootstrap replicate 32