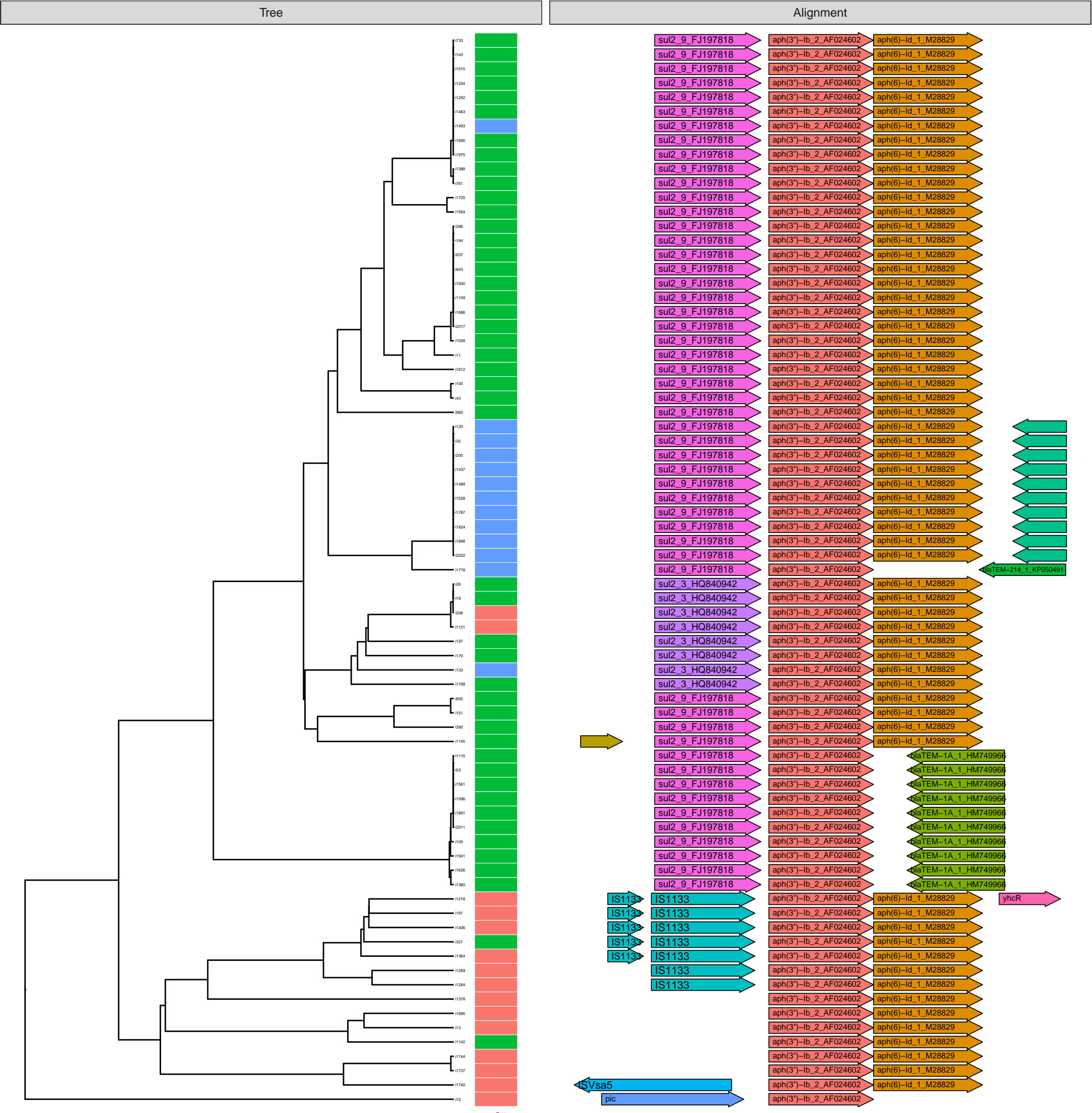


Cluster 23 – aph – distance tree based on only flanking region sequences



Target – Reference DB % identity

- aph(3'')-lb_2_AF024602_100_%
- aph(3'')-lb_5_AF321551_100_%
- aph(3'')-lb_5_AF321551_99.88_%

Gene

- aph(3'')-lb
- aph(6'')-ld_1
- blaTEM-185_
- blaTEM-1A_1
- blaTEM-214_
- blaTEM-95_1
- IS1133
- ISVsa5
- pic
- sul2_3_HQ84
- sul2_9_FJ19
- yhcR