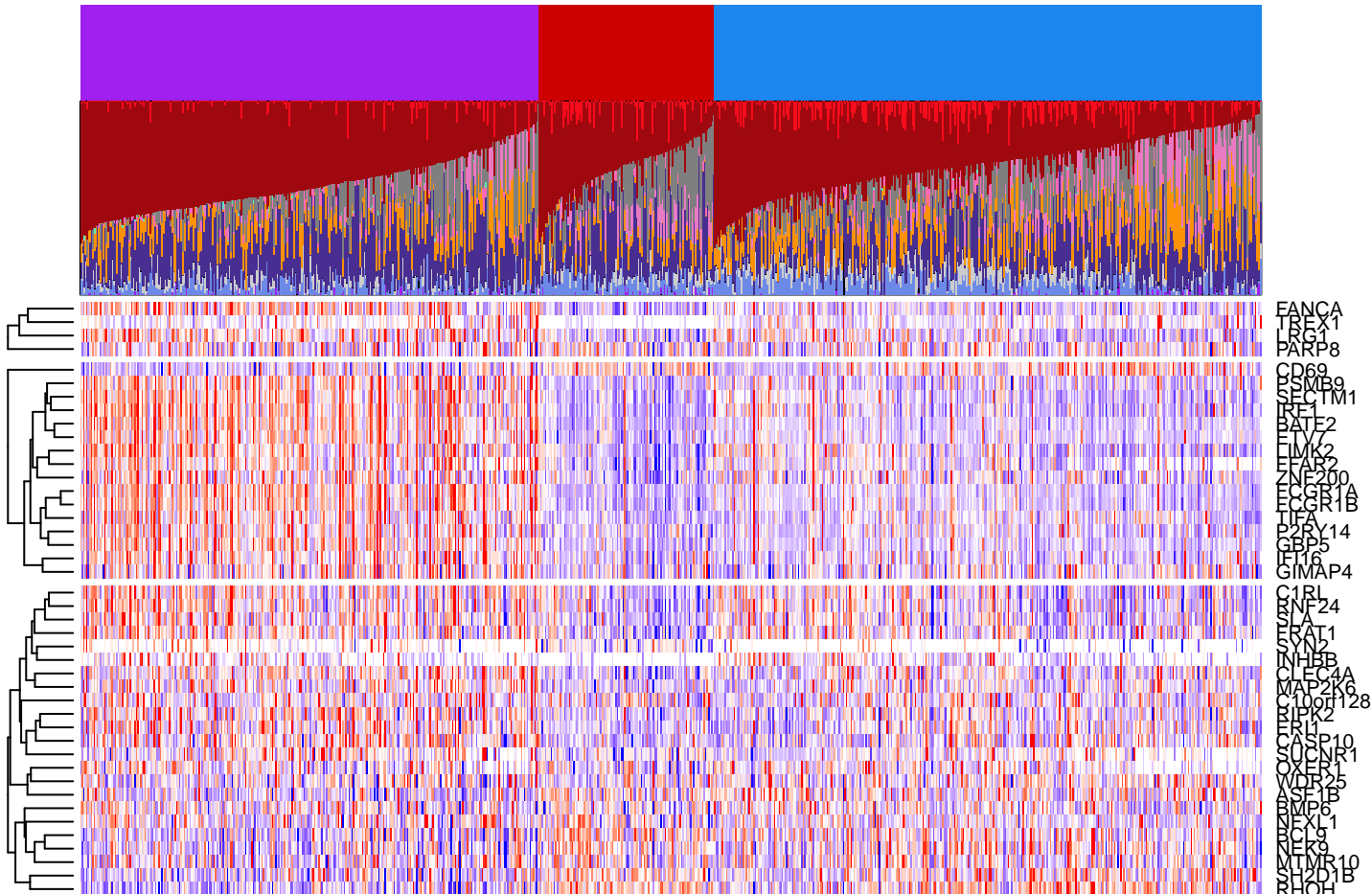


A phylogenetic tree showing the evolutionary relationships between various species. The tree is rooted on the left and branches out to the right. The species names are listed at the tips of the branches. A scale bar at the bottom indicates the genetic distance in substitutions per site, with a value of 0.1.



FANCA  
 BEX1  
 PARP8  
 CD68  
 PSMTB9  
 SECTM1  
 IRE1  
 BATE2  
 LMK2  
 ELK2  
 ZNF200  
 ECGR1B  
 TIFA  
 P2RY14  
 GBP5  
 IFT16  
 GIMAP4  
 C1RL  
 NF24  
 SMA  
 RAT1  
 SNO2  
 INHBB  
 CLEC4A  
 MAP2K6  
 C10orf128  
 RER1  
 PKP2  
 CASP10  
 SUCNR1  
 OXER1  
 VDRP25  
 BMP1B  
 BMP61  
 CXCL5  
 RIKK8  
 NTKR10  
 ZNF201B  
 SHCH

