Figure S3. Phylogenetic tree of some human small GTPases. All bootstrap values are included. For information on how the tree has been generated check **Materials and Methods** in

I

the main article. All sequence accessions are listed in **Table S2**. The representation has been generated with Dendroscope [1].

References

1. Huson DH, Richter DC, Rausch C, Dezulian T, Franz M et al. (2007) Dendroscope: An interactive viewer for large phylogenetic trees. BMC Bioinformatics 8: 460.