

ANKRO11 - TBC1XR1

- human

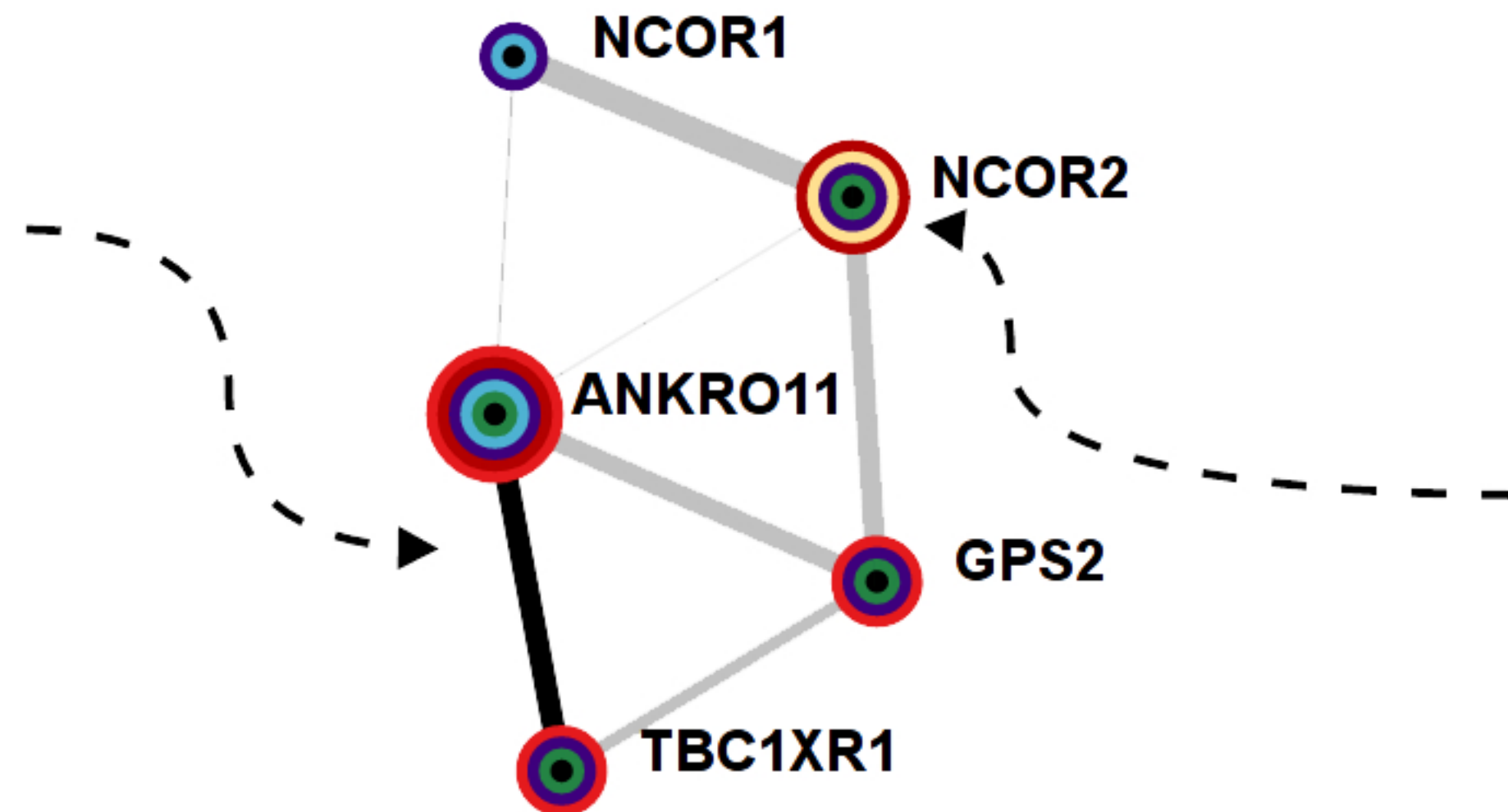
- rat

- mouse

- zebrafish

- fly

- yeast



b

NCOR2

- | | |
|-----------------------------|----|
| ● nonsense | 20 |
| ● splice site | 18 |
| ● frame shift | 14 |
| ● non conservative missense | 7 |
| ● conservative missense | 3 |
| ● synonymous | 2 |

investigating sequence variation and interaction networks (or pathways). Here the

thickness/colour density of graph edges reflects the level of phylogenetics support for them, as show in (a). Nodes are surrounded by coloured circles representing the types of variations (b) observed for that gene. For the set of patients/strains being examined, (c) shows how variations are distributed across individuals, or instance showing that, in particular cancer types, mutations in members of a complex tend to be mutually exclusive (Sander et al. 2013).

Variations × Individuals C

[illegible]