Patient: CRUK0003

Score
- $f = 8e{-}01$
- $g = 0.8$

Violations
- 8 R6

GL

1

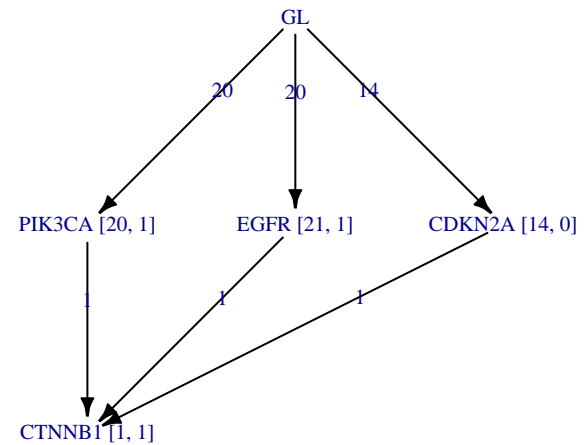6   4

8   2

5   3
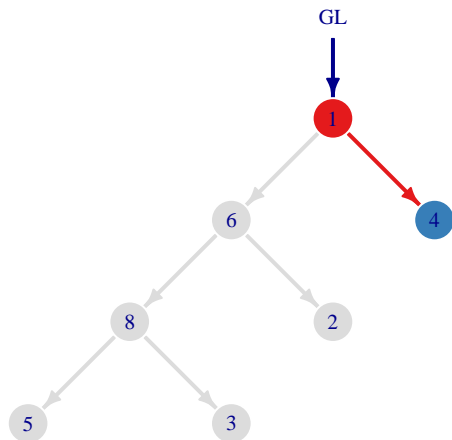
Ranked 1/1

CRUK0003 : evolutionary trajectories

GL

PIK3CA   EGFR   CDKN2A

6   CTNNB1

8   2

5   3
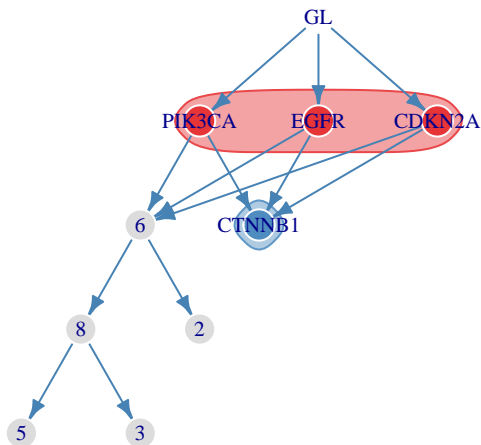
Exploded trajectories

CRUK0003 : information transfer

GL

20   20   14

PIK3CA [20, 1]   EGFR [21, 1]   CDKN2A [14, 0]

1   1   1

CTNNB1 [1, 1]

[n m]: n clonal, m subclonal (driver)   ... : k patients