Patient: PD9772

PD9772 : evolutionary trajectories
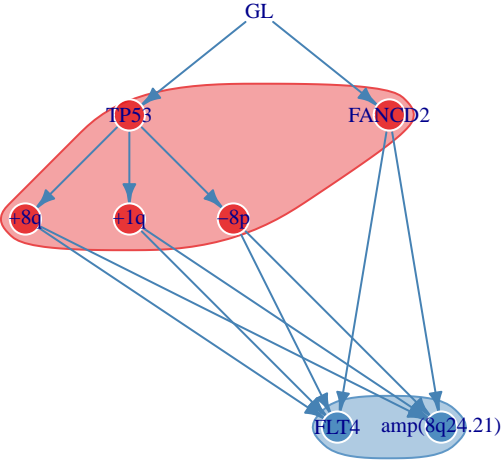
PD9772 : information transfer

Score
- *f* = 2e−01
- *g* = 1

Violations
- None

GL

1

2

GL

TP53   FANCD2

+8q   +1q   −8p

FLT4   amp(8q24.21)

GL

27   2

TP53 [27, 3]   FANCD2 [2, 1]

16   8   8

+8q [24, 9] +1q [18, 4] −8p [13, 4]   1   1

1   7   4   3   1   3

FLT4 [1amp(8q24.21) [3, 5]
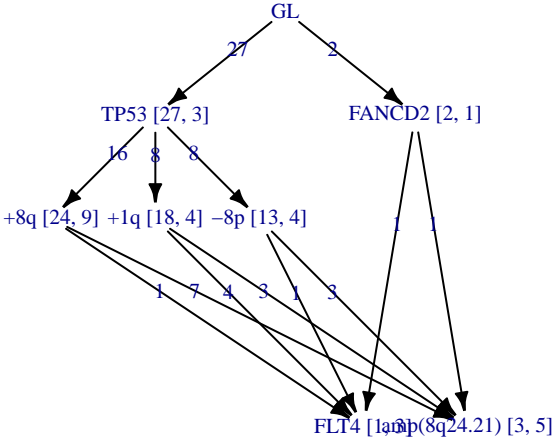
Ranked 1/1   Exploded trajectories   [n m]: n clonal, m subclonal (driver)   ... : k patients