

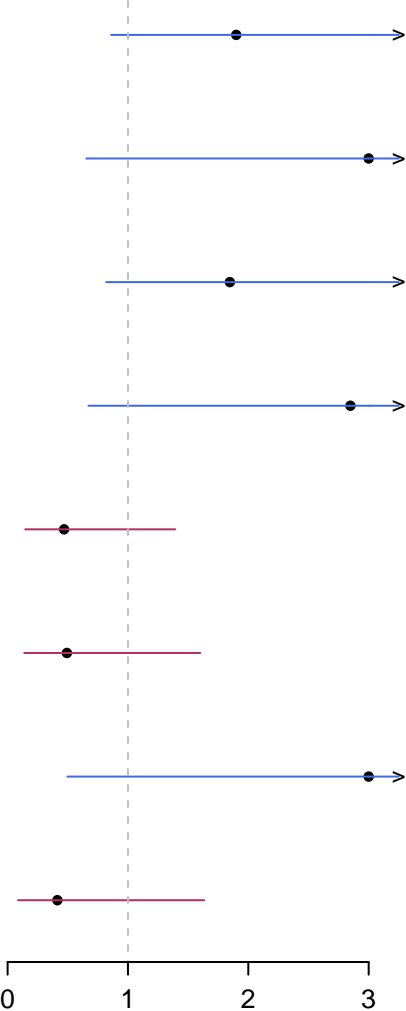
Relapse (n = 68) v/s Primary (n = 77)

Relapse

Primary

OR

P-value



KMT2C

15

27

1.9

NS

RAD17

1

6

5.607

NS

PKHD1

14

25

1.847

NS

AMER1

3

9

2.848

NS

CIC

12

7

0.469

NS

POLE

10

6

0.493

NS

SEMA3A

1

5

4.612

NS

ERBB2

8

4

0.413

NS

Odds ratio with 95% CI
(1 = no effect, < 1 Relapse has more mutants)