

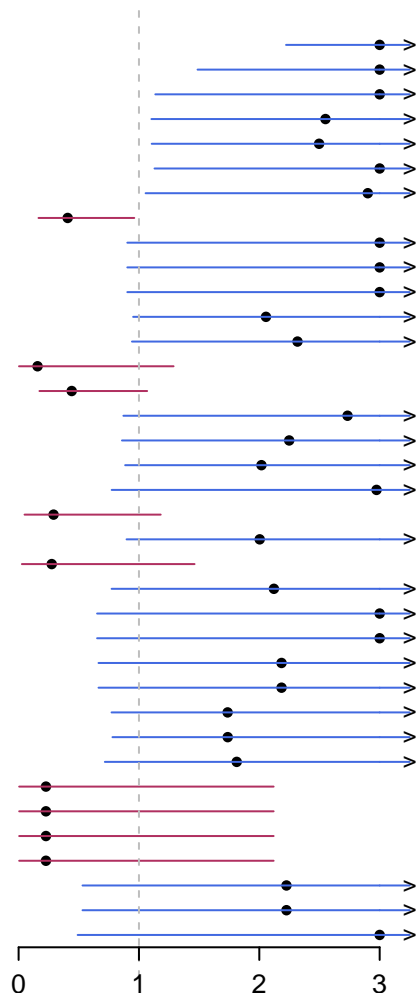
Relapse (n = 68) v/s LymphNode (n = 57)

Relapse

LymphNode

OR

P-value



*EIF4E*

0

8

Inf

\*\*

*DICER1*

21

34

3.275

\*\*

*FH*

0

5

Inf

\*

*NOTCH3*

15

24

2.55

\*

*APC*

17

26

2.497

\*

*MTAP*

1

7

9.239

\*

*ATRX*

8

16

2.901

\*

*PIK3CA*

27

12

0.408

\*

*KDM5C*

4

10

3.371

\*

*MET*

4

10

3.371

\*

*PIK3CG*

4

10

3.371

\*

*KMT2C*

26

32

2.055

NS

*SEMA3E*

12

19

2.317

NS

*DOT1L*

7

1

0.157

NS

*MAP3K1*

24

11

0.441

NS

*INPP4B*

6

12

2.733

NS

*TET2*

10

16

2.248

NS

*PKHD1*

17

23

2.018

NS

*BRIP1*

4

9

2.974

NS

*GZMH*

11

3

0.29

NS

*PALB2*

19

25

2.003

NS

*ERBB2*

8

2

0.275

NS

*NCOR1*

9

14

2.121

NS

*MYCN*

2

6

3.842

NS

*RAD17*

2

6

3.842

NS

*ATR*

6

10

2.185

NS

*GID4*

6

10

2.185

NS

*CIC*

19

23

1.737

NS

*AMER1*

20

24

1.738

NS

*MAGI2*

12

16

1.812

NS

*ARID2*

5

1

0.227

NS

*EML4*

5

1

0.227

NS

*FGFR4*

5

1

0.227

NS

*MAP3K13*

5

1

0.227

NS

*PHLPP2*

4

7

2.226

NS

*PMS2*

4

7

2.226

NS

*FGF12*

2

5

3.145

NS

Odds ratio with 95% CI

(1 = no effect, < 1 Relapse has more mutants)