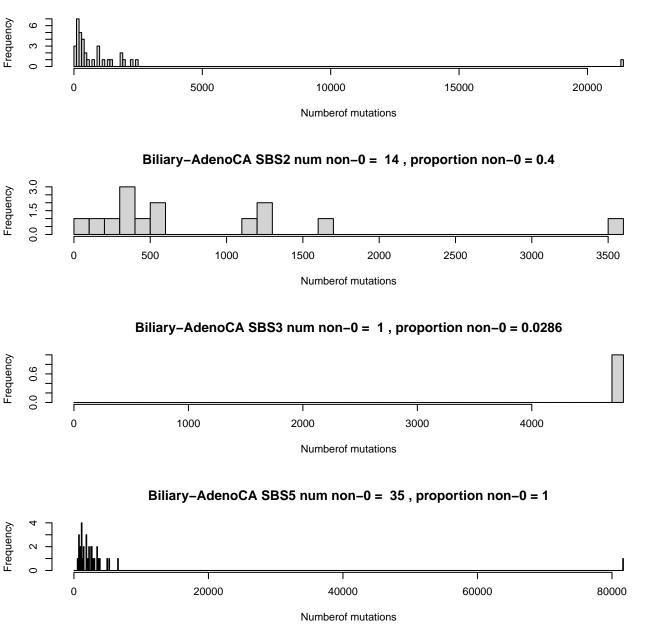
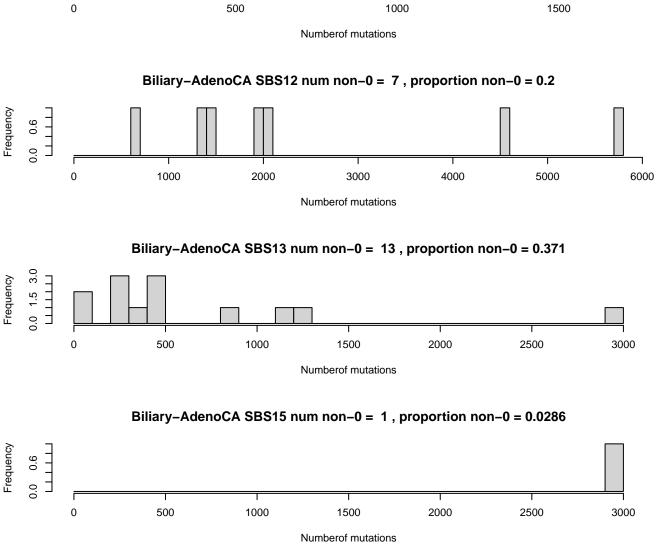
Biliary-AdenoCA SBS1 num non-0 = 35, proportion non-0 = 1



Biliary-AdenoCA SBS9 num non-0 = 1, proportion non-0 = 0.0286

0.0 0.6



Biliary-AdenoCA SBS17a num non-0 = 2 , proportion non-0 = 0.0571

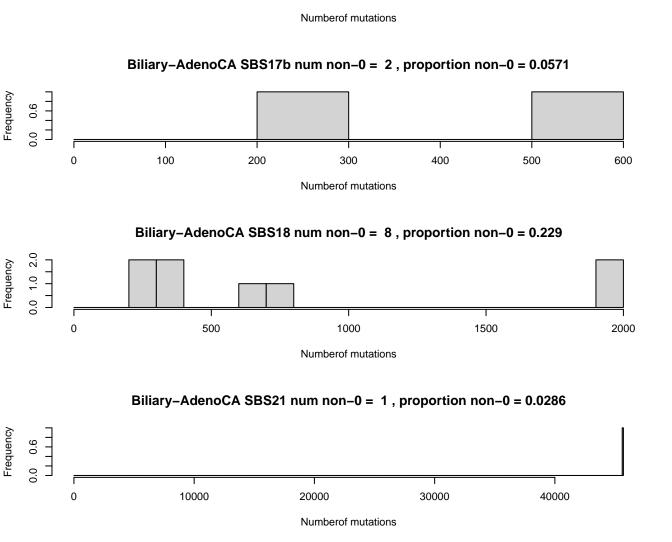
100

150

200

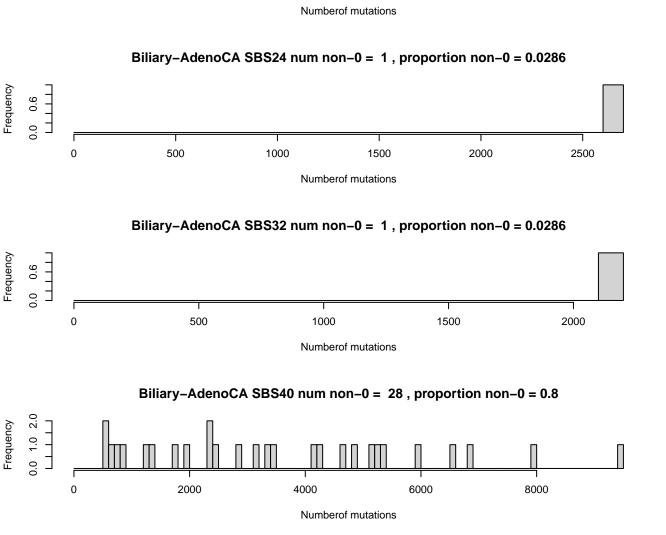
50

0.0 0.6

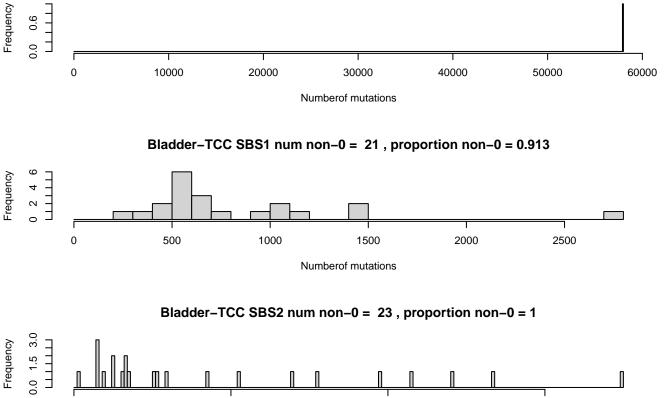


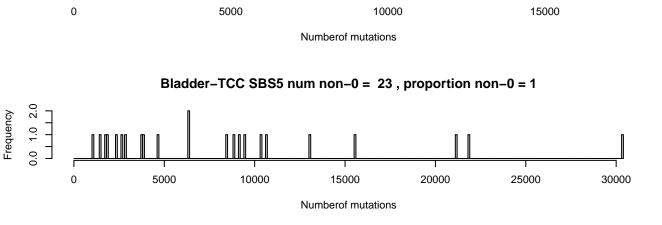
Biliary-AdenoCA SBS22 num non-0 = 1 , proportion non-0 = 0.0286

0.0 0.6

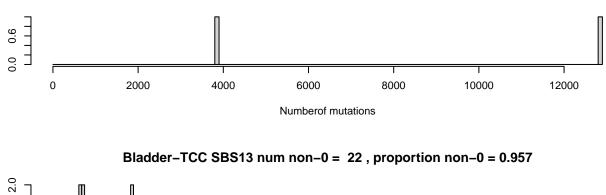


Biliary-AdenoCA SBS44 num non-0 = 1 , proportion non-0 = 0.0286





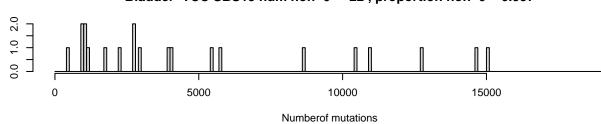
Bladder-TCC SBS8 num non-0 = 2, proportion non-0 = 0.087

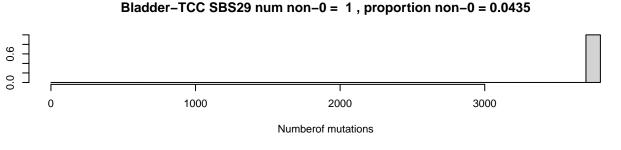


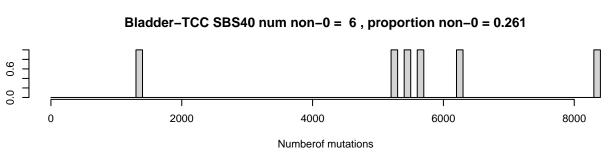
Frequency

Frequency

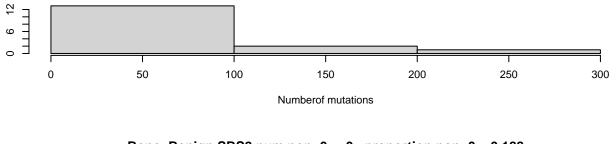
Frequency







Bone-Benign SBS1 num non-0 = 16, proportion non-0 = 1



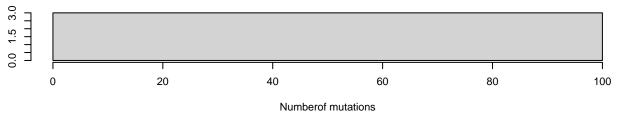
Frequency

Frequency

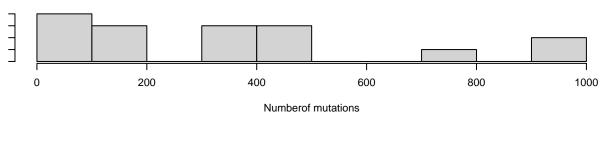
Frequency

Frequency

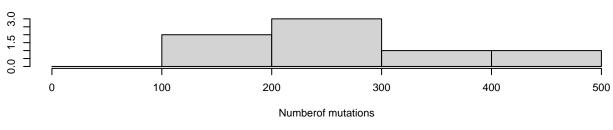
Bone–Benign SBS2 num non–0 = 3, proportion non–0 = 0.188



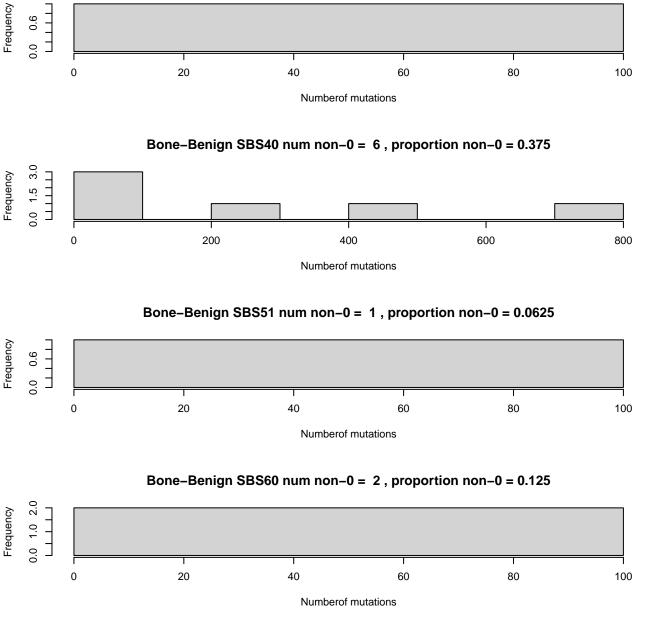
Bone-Benign SBS5 num non-0 = 16, proportion non-0 = 1



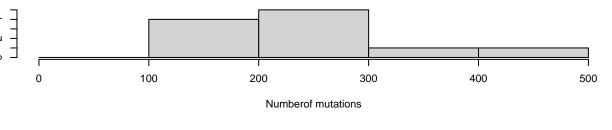
Bone-Benign SBS8 num non-0 = 7, proportion non-0 = 0.438



Bone-Benign SBS13 num non-0 = 1, proportion non-0 = 0.0625



Bone-Epith SBS1 num non-0 = 11, proportion non-0 = 1



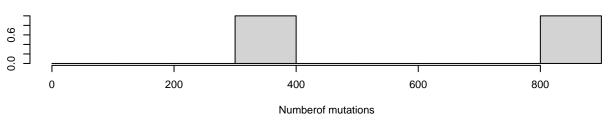
Frequency

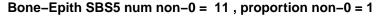
Frequency

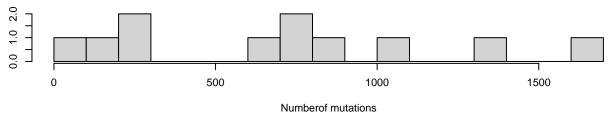
Frequency

Frequency

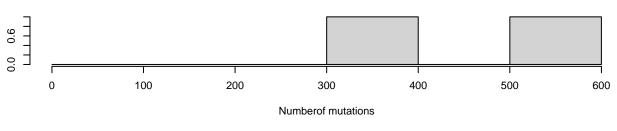
Bone–Epith SBS2 num non–0 = 2, proportion non–0 = 0.182



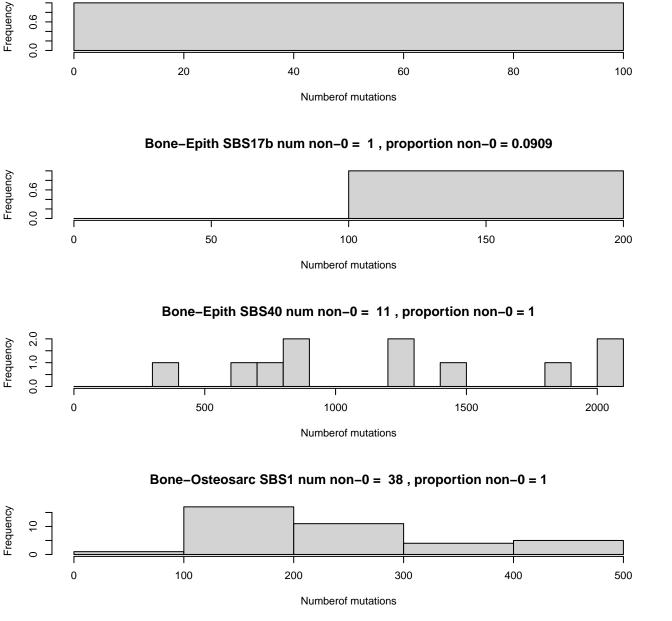




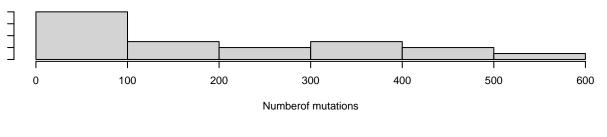
Bone-Epith SBS13 num non-0 = 2, proportion non-0 = 0.182



Bone-Epith SBS17a num non-0 = 1, proportion non-0 = 0.0909



Bone-Osteosarc SBS2 num non-0 = 19, proportion non-0 = 0.5



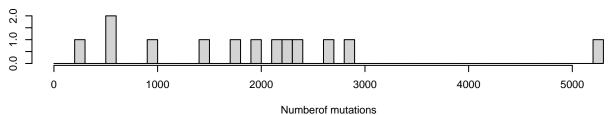
Frequency

Frequency

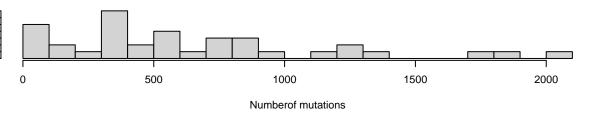
Frequency

Frequency

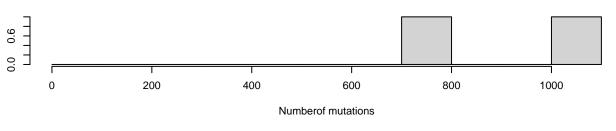
Bone–Osteosarc SBS3 num non–0 = 13, proportion non–0 = 0.342



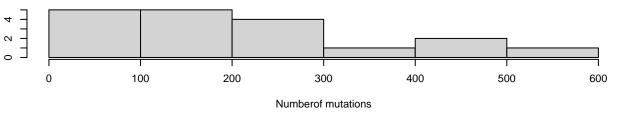
Bone–Osteosarc SBS5 num non–0 = 36, proportion non–0 = 0.947



Bone–Osteosarc SBS8 num non–0 = 2, proportion non–0 = 0.0526



Bone-Osteosarc SBS13 num non-0 = 18, proportion non-0 = 0.474



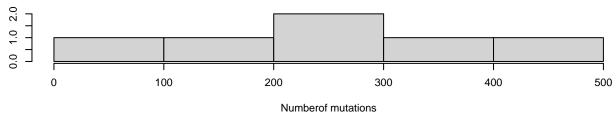
Frequency

Frequency

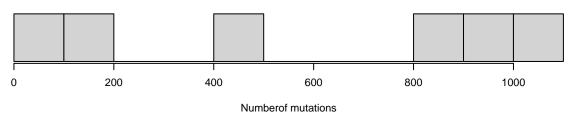
Frequency

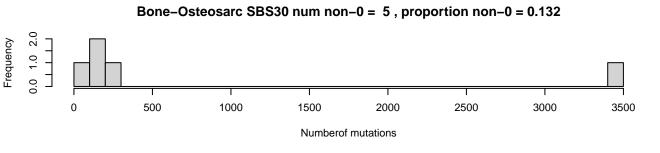
9.0 0.0

Bone–Osteosarc SBS17a num non–0 = 6, proportion non–0 = 0.158

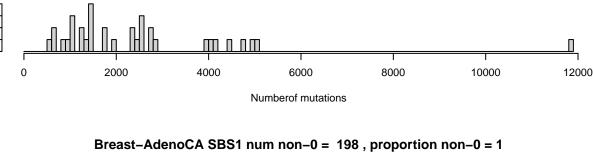


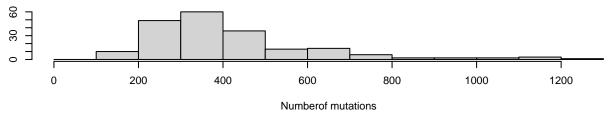
Bone–Osteosarc SBS17b num non–0 = 6, proportion non–0 = 0.158

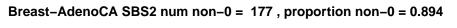


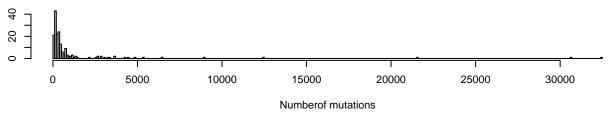


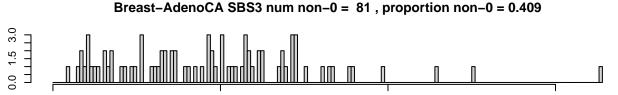
Bone-Osteosarc SBS40 num non-0 = 35, proportion non-0 = 0.921











10000

15000

Numberof mutations

5000

Frequency

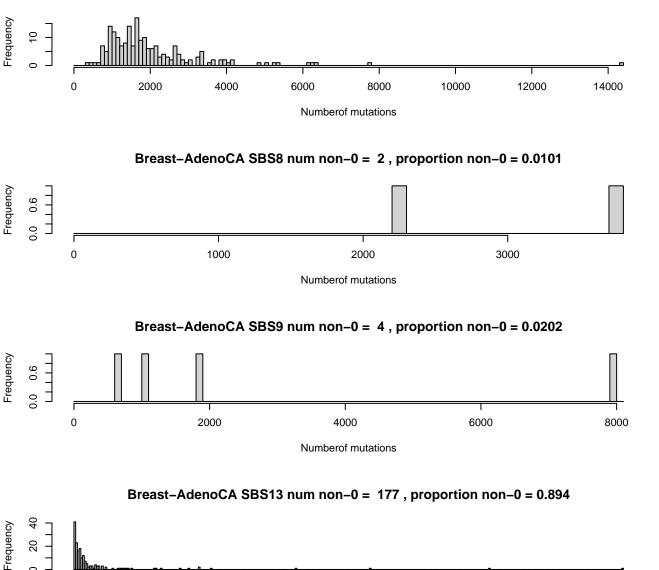
Frequency

-requency

Frequency

0

Breast-AdenoCA SBS5 num non-0 = 198, proportion non-0 = 1



Number of mutations

Breast-AdenoCA SBS17a num non-0 = 4 , proportion non-0 = 0.0202

Frequency

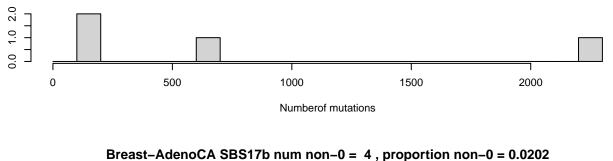
Frequency

Frequency

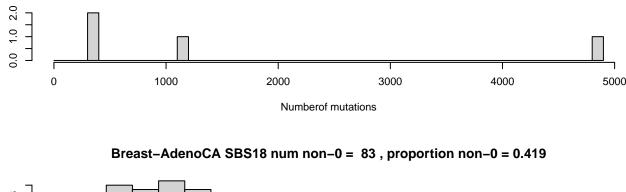
Frequency

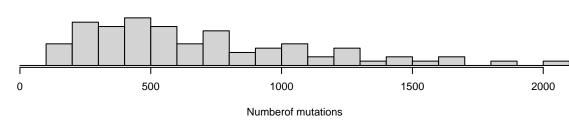
0.0 0.6

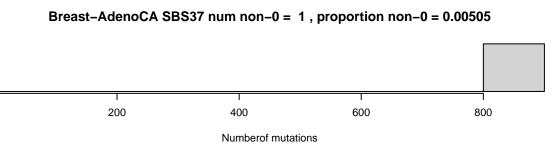
0



Breast-AdenoCA SBS17b num non-0 = 4, proportion non-0 = 0.0202







Breast-AdenoCA SBS40 num non-0 = 1, proportion non-0 = 0.00505

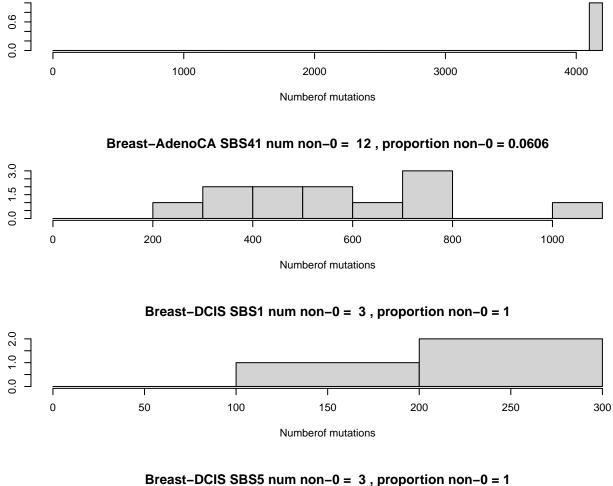
Frequency

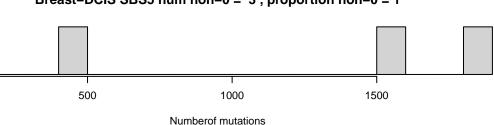
Frequency

Frequency

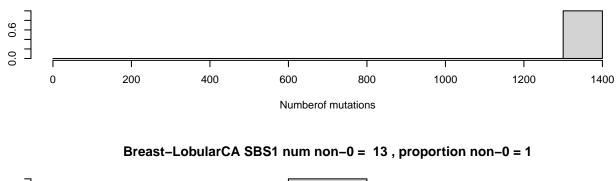
9.0 0.0

0





Breast-DCIS SBS40 num non-0 = 1, proportion non-0 = 0.333



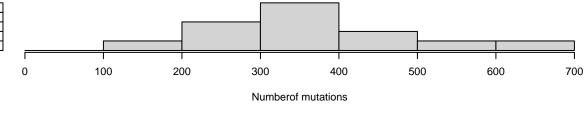
Frequency

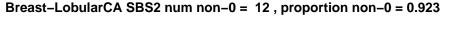
Frequency

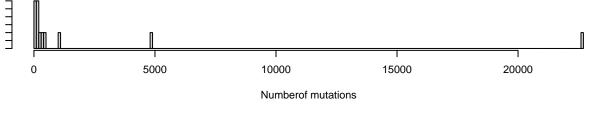
0.0 1.5 3.0 Frequency

Frequency

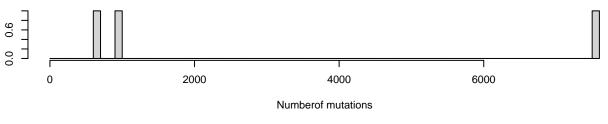




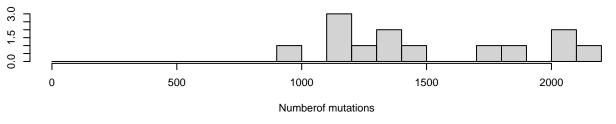




Breast-LobularCA SBS3 num non-0 = 3, proportion non-0 = 0.231



Breast-LobularCA SBS5 num non-0 = 13 , proportion non-0 = 1



Frequency

Frequency

0.0 1.0 2.0

0

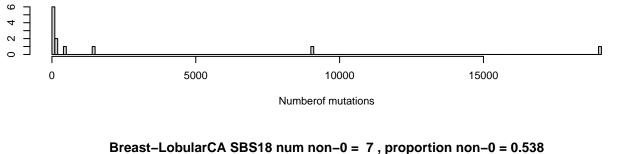
0

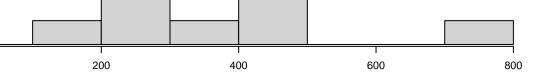
Frequency

Frequency

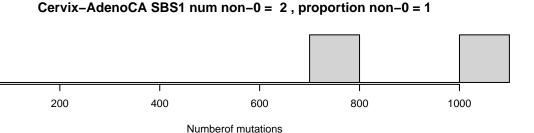
0.0 0.6

Breast-LobularCA SBS13 num non-0 = 12 , proportion non-0 = 0.923

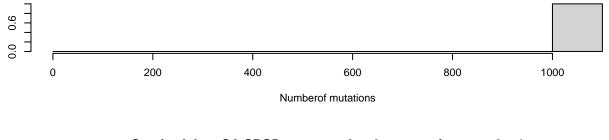




Number of mutations



Cervix-AdenoCA SBS2 num non-0 = 1 , proportion non-0 = 0.5



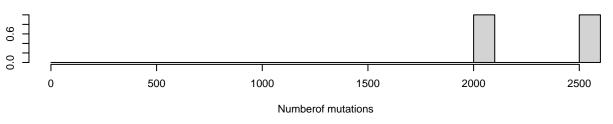
Frequency

Frequency

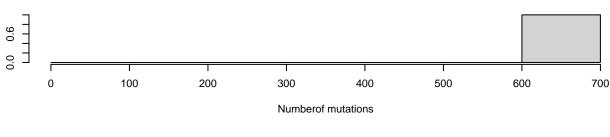
Frequency

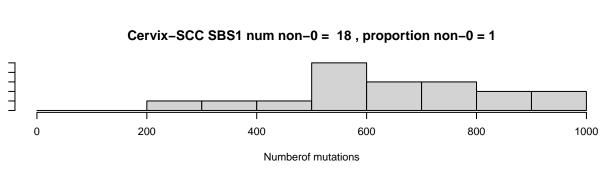
Frequency

Cervix-AdenoCA SBS5 num non-0 = 2, proportion non-0 = 1

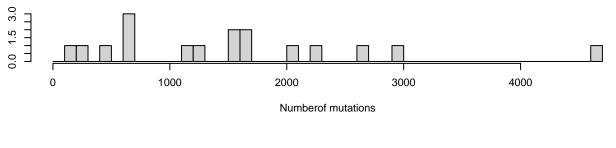








Cervix-SCC SBS2 num non-0 = 17, proportion non-0 = 0.944



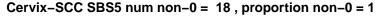
Frequency

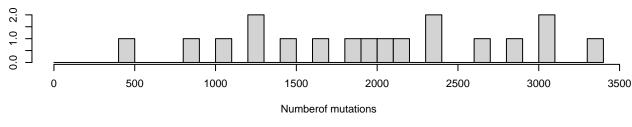
Frequency

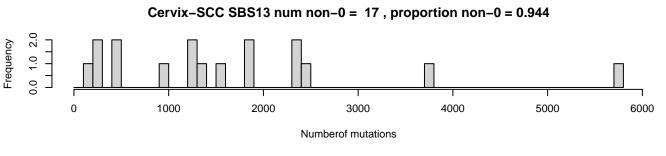
Frequency

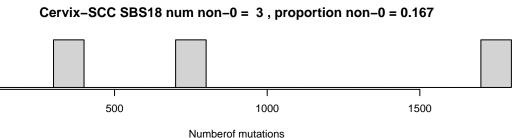
9.0 0.0

0









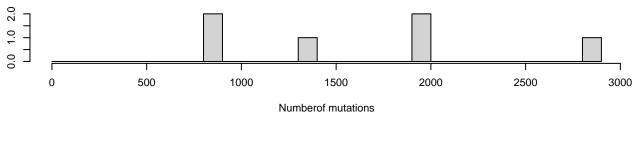
Cervix-SCC SBS40 num non-0 = 6, proportion non-0 = 0.333

Frequency

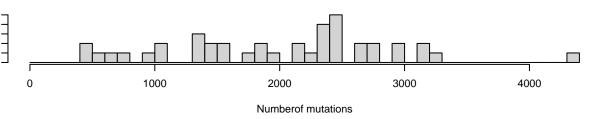
Frequency

Frequency

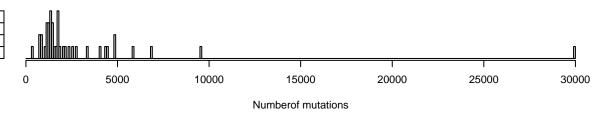
Frequency



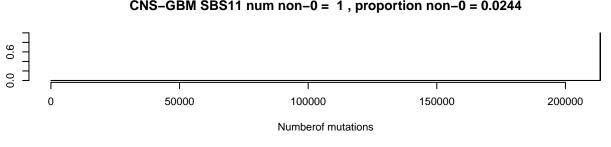
CNS-GBM SBS1 num non-0 = 41, proportion non-0 = 1



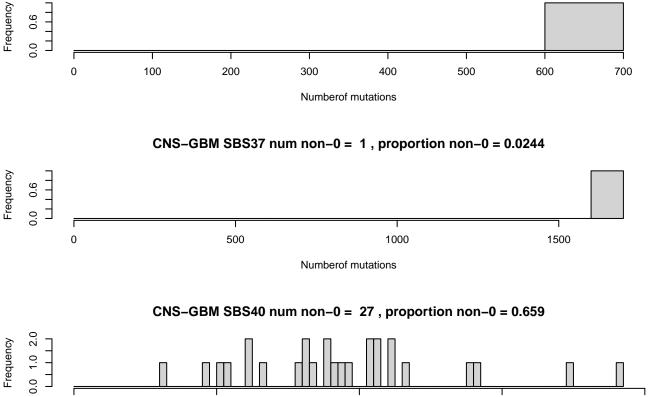
CNS-GBM SBS5 num non-0 = 41, proportion non-0 = 1

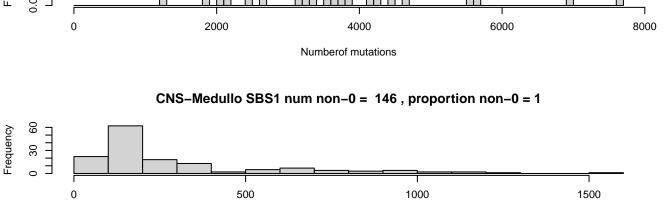


CNS-GBM SBS11 num non-0 = 1, proportion non-0 = 0.0244



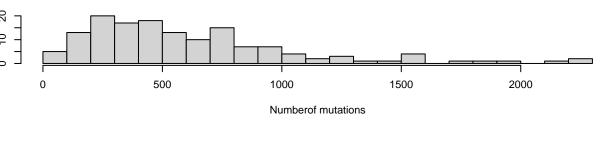
CNS-GBM SBS30 num non-0 = 1 , proportion non-0 = 0.0244





Number of mutations

CNS-Medullo SBS5 num non-0 = 146, proportion non-0 = 1



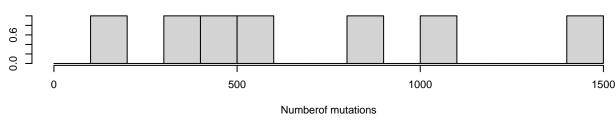
Frequency

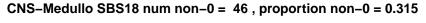
Frequency

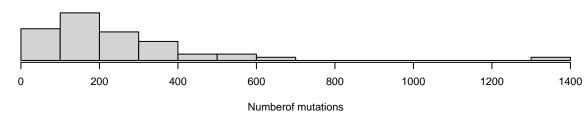
Frequency

Frequency

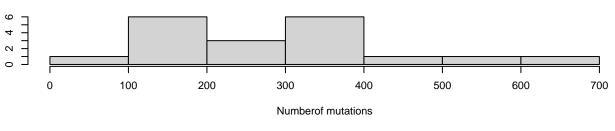
CNS-Medullo SBS8 num non-0 = 7, proportion non-0 = 0.0479



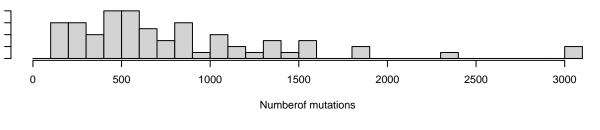




CNS-Medullo SBS39 num non-0 = 19 , proportion non-0 = 0.13



CNS-Medullo SBS40 num non-0 = 66, proportion non-0 = 0.452



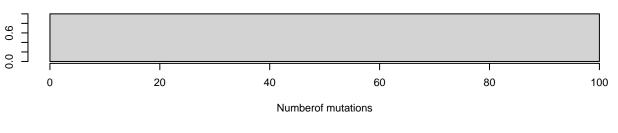
Frequency

Frequency

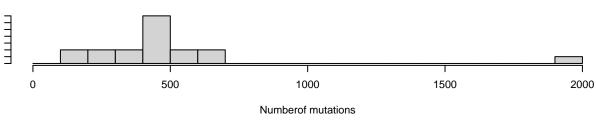
-requency

Frequency

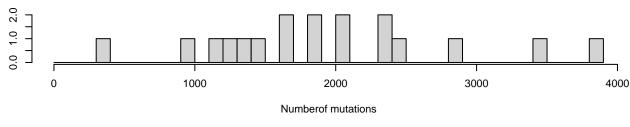
CNS-Medullo SBS60 num non-0 = 1, proportion non-0 = 0.00685



CNS-Oligo SBS1 num non-0 = 18, proportion non-0 = 1



CNS-Oligo SBS5 num non-0 = 18, proportion non-0 = 1



CNS-Oligo SBS8 num non-0 = 1, proportion non-0 = 0.0556

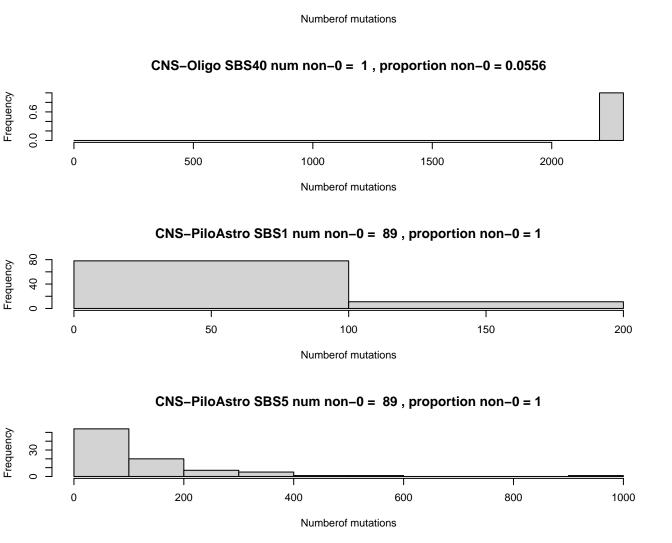
1000

1500

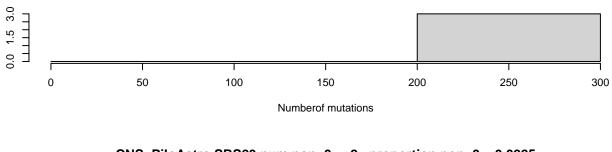
500

0.0 0.6

0



CNS-PiloAstro SBS19 num non-0 = 3, proportion non-0 = 0.0337



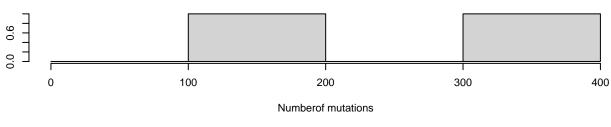
Frequency

Frequency

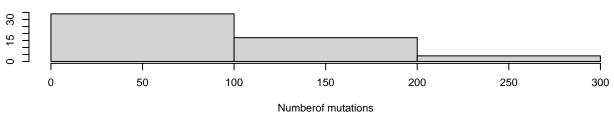
Frequency

Frequency

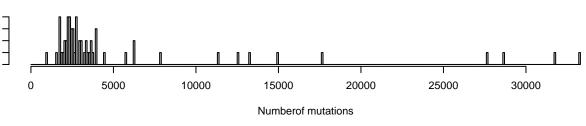
CNS-PiloAstro SBS23 num non-0 = 2, proportion non-0 = 0.0225



CNS-PiloAstro SBS40 num non-0 = 55, proportion non-0 = 0.618



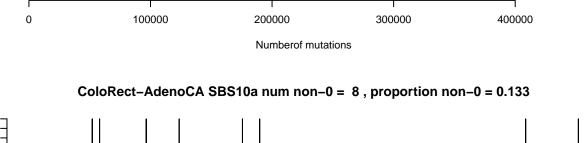
ColoRect-AdenoCA SBS1 num non-0 = 60 , proportion non-0 = 1

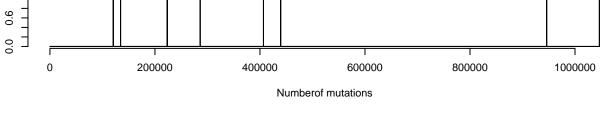


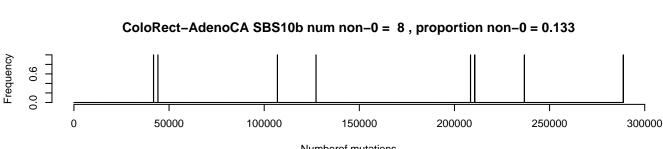
ColoRect-AdenoCA SBS5 num non-0 = 57, proportion non-0 = 0.95

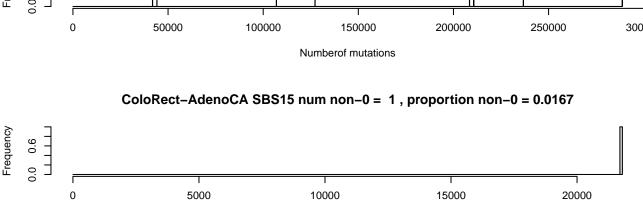
Frequency 0.0 1.0 2.0

Frequency









Number of mutations

ColoRect-AdenoCA SBS17a num non-0 = 4, proportion non-0 = 0.0667 500 1000 1500 Number of mutations ColoRect-AdenoCA SBS17b num non-0 = 4, proportion non-0 = 0.0667 1000 2000 3000 4000 5000 Number of mutations ColoRect-AdenoCA SBS18 num non-0 = 35, proportion non-0 = 0.583 2000 8000 4000 6000 10000 Number of mutations ColoRect-AdenoCA SBS28 num non-0 = 13, proportion non-0 = 0.217

Frequency 0.0 1.0 2.0

Frequency

9.0 0.0

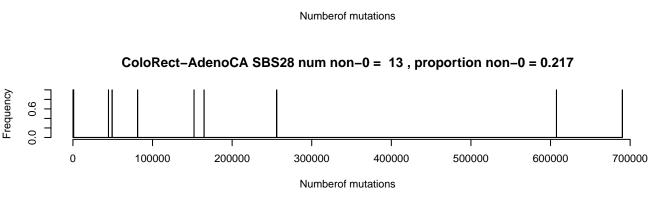
0.0 1.5 3.0

Frequency

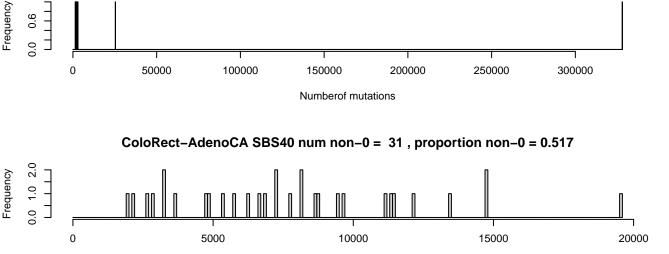
0

0

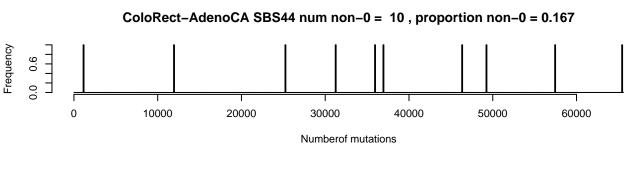
0

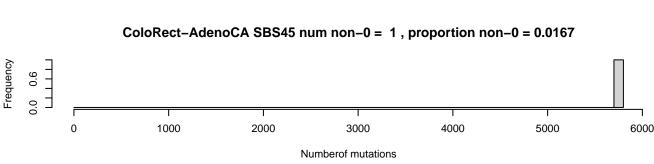


ColoRect-AdenoCA SBS37 num non-0 = 8, proportion non-0 = 0.133

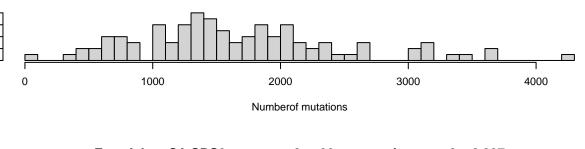


Number of mutations





Eso-AdenoCA SBS1 num non-0 = 98, proportion non-0 = 1

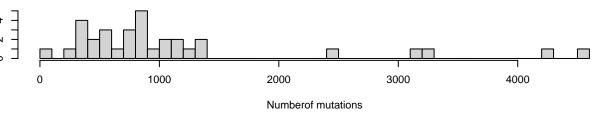


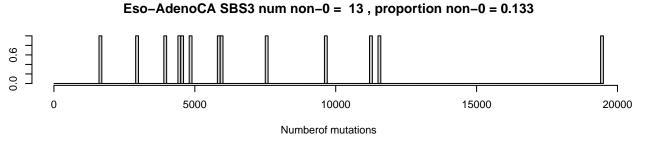
Frequency

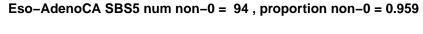
Frequency

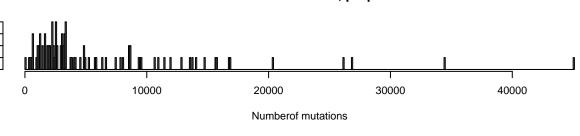
Frequency

Eso-AdenoCA SBS2 num non-0 = 33, proportion non-0 = 0.337

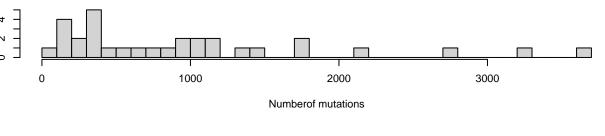








Eso-AdenoCA SBS13 num non-0 = 31 , proportion non-0 = 0.316



Frequency

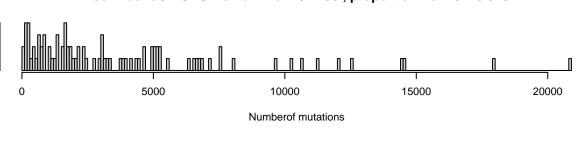
requency-

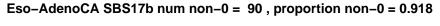
0.0 1.5 3.0

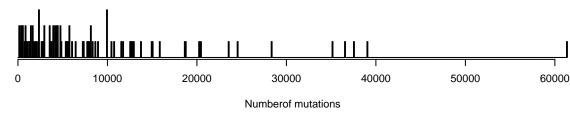
Frequency

-requency

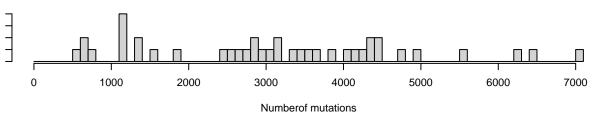
Eso-AdenoCA SBS17a num non-0 = 90, proportion non-0 = 0.918



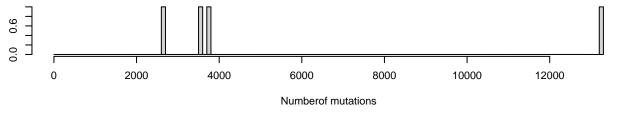




Eso-AdenoCA SBS18 num non-0 = 40 , proportion non-0 = 0.408



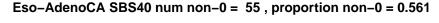
Eso-AdenoCA SBS28 num non-0 = 4, proportion non-0 = 0.0408

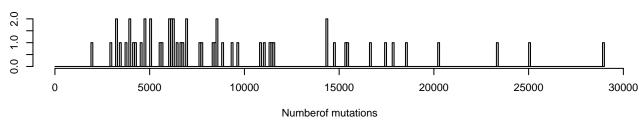


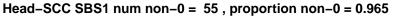
Frequency

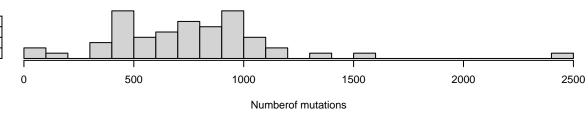
Frequency

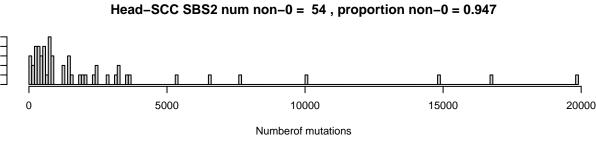
-requency











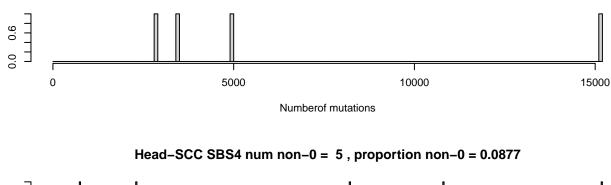
Head-SCC SBS3 num non-0 = 4, proportion non-0 = 0.0702

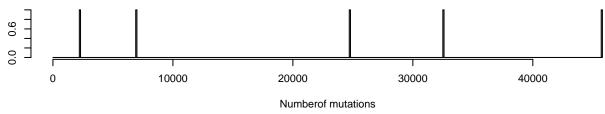
Frequency

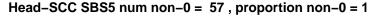
Frequency

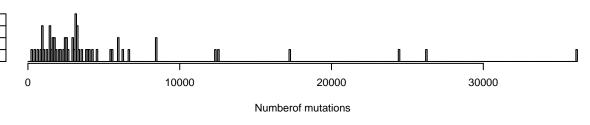
Frequency

Frequency

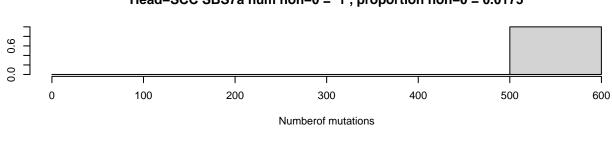




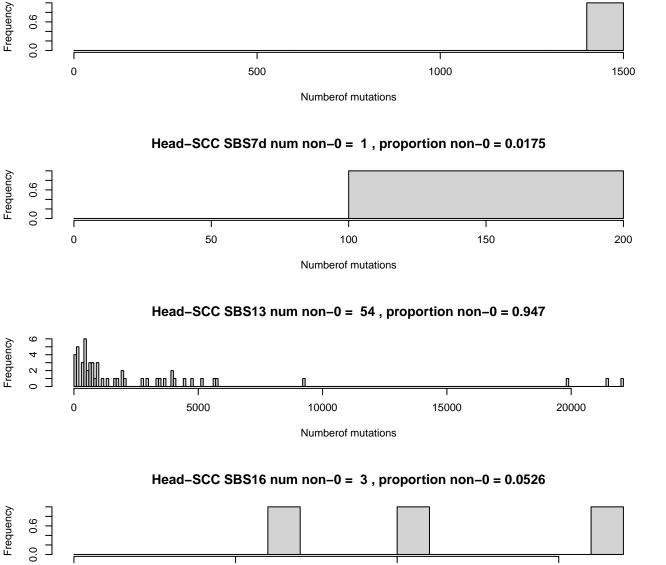




Head-SCC SBS7a num non-0 = 1, proportion non-0 = 0.0175

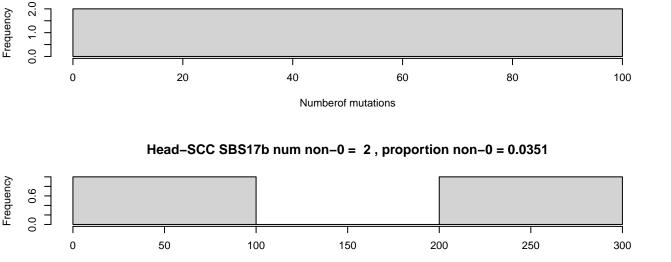


Head–SCC SBS7b num non–0 = 1, proportion non–0 = 0.0175

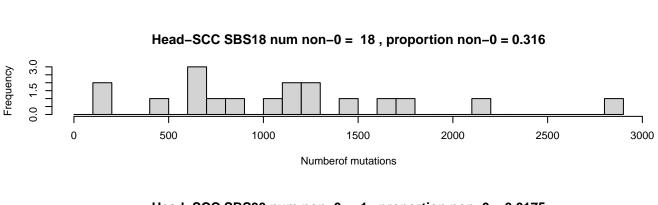


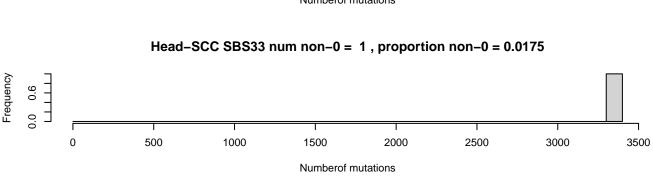
Number of mutations

Head-SCC SBS17a num non-0 = 2, proportion non-0 = 0.0351



Number of mutations



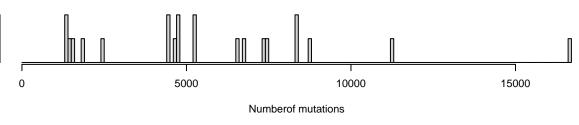


Head-SCC SBS40 num non-0 = 22, proportion non-0 = 0.386

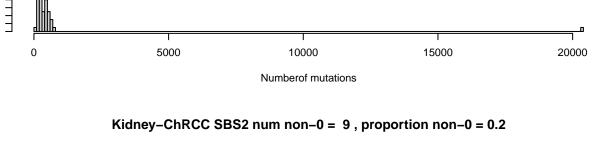
0.0 1.0 2.0 Frequency

Frequency

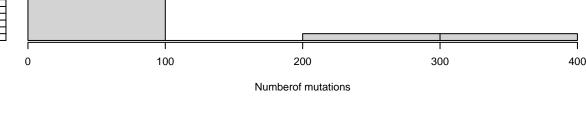
Frequency

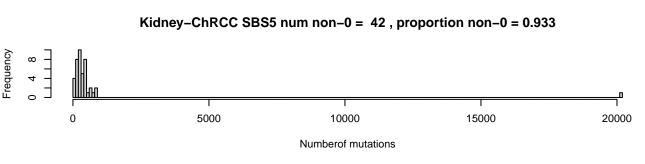


Kidney-ChRCC SBS1 num non-0 = 45, proportion non-0 = 1



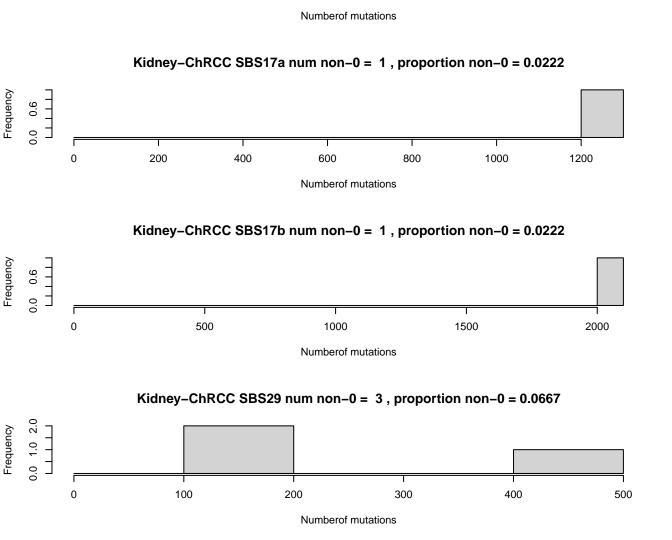




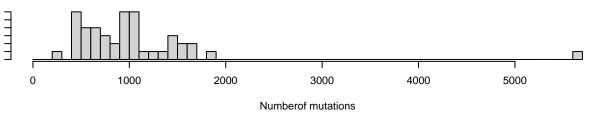


Kidney-ChRCC SBS13 num non-0 = 8, proportion non-0 = 0.178

Frequency



Kidney-ChRCC SBS40 num non-0 = 44 , proportion non-0 = 0.978

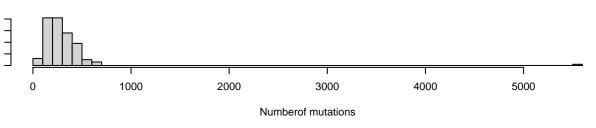


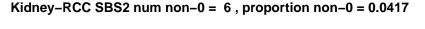
Frequency

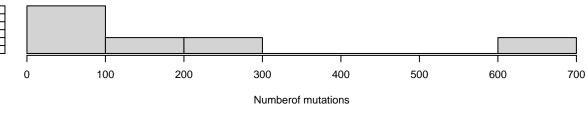
Frequency 0.0 1.5 3.0

-requency

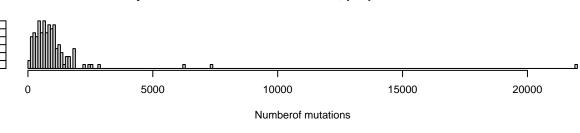
Kidney-RCC SBS1 num non-0 = 144, proportion non-0 = 1



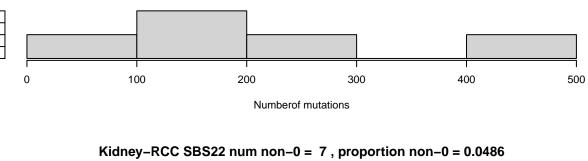




Kidney-RCC SBS5 num non-0 = 135, proportion non-0 = 0.938



Kidney-RCC SBS13 num non-0 = 5, proportion non-0 = 0.0347

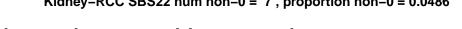


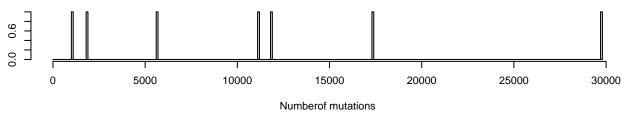
Frequency 0.0 1.0 2.0

Frequency

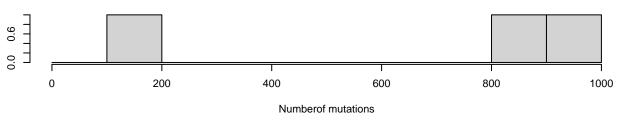
Frequency

-requency

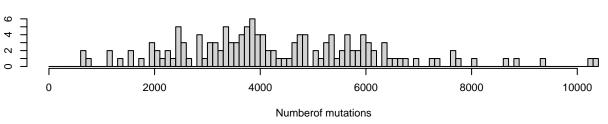




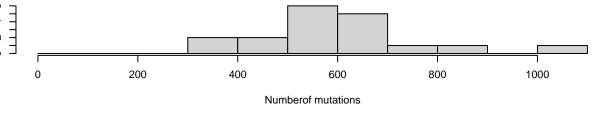
Kidney-RCC SBS29 num non-0 = 3, proportion non-0 = 0.0208



Kidney-RCC SBS40 num non-0 = 141 , proportion non-0 = 0.979



Kidney-RCC SBS41 num non-0 = 18, proportion non-0 = 0.125



Frequency

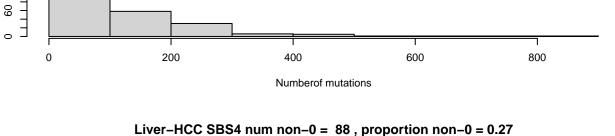
Frequency

-requency

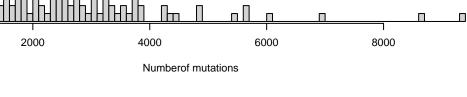
Frequency

0

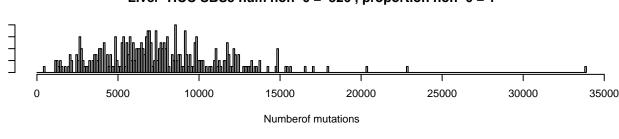
Liver-HCC SBS1 num non-0 = 213, proportion non-0 = 0.653



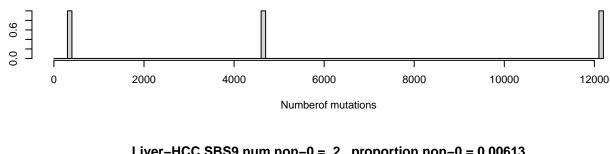




Liver-HCC SBS5 num non-0 = 326, proportion non-0 = 1



Liver-HCC SBS6 num non-0 = 3, proportion non-0 = 0.0092



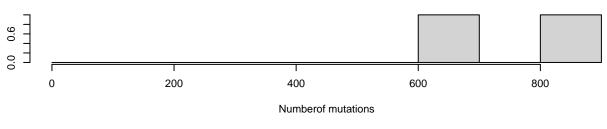
Frequency

Frequency

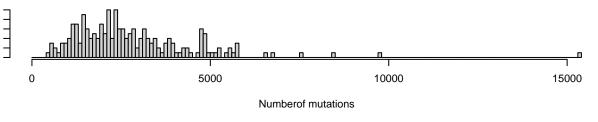
Frequency

Frequency

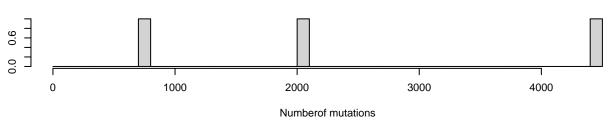
Liver-HCC SBS9 num non-0 = 2, proportion non-0 = 0.00613



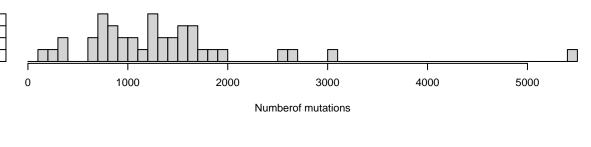
Liver-HCC SBS12 num non-0 = 198, proportion non-0 = 0.607



Liver-HCC SBS14 num non-0 = 3, proportion non-0 = 0.0092



Liver-HCC SBS16 num non-0 = 39, proportion non-0 = 0.12

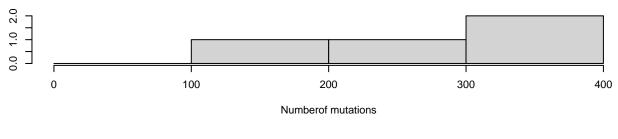


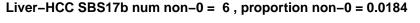
Frequency

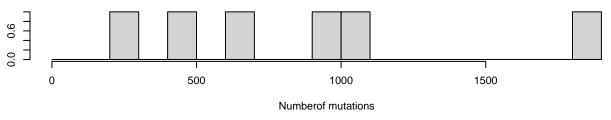
Frequency

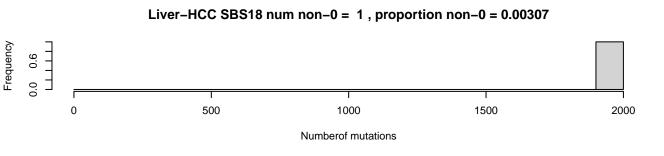
Frequency

Liver-HCC SBS17a num non-0 = 4, proportion non-0 = 0.0123





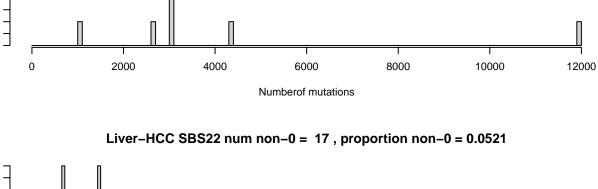


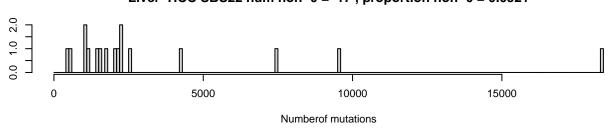


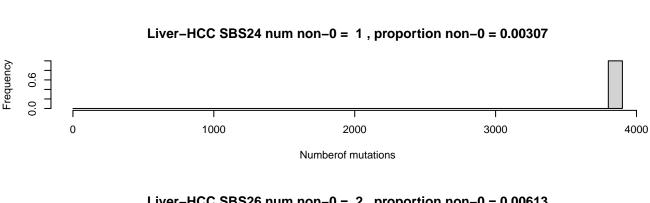
Liver-HCC SBS19 num non-0 = 6, proportion non-0 = 0.0184

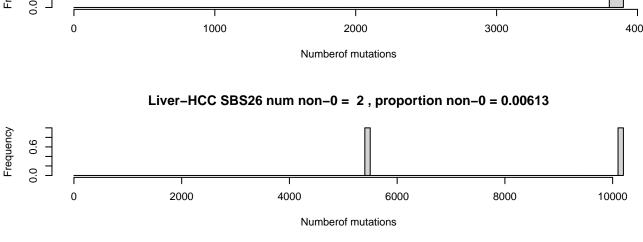
Frequency 0.0 1.0 2.0

Frequency

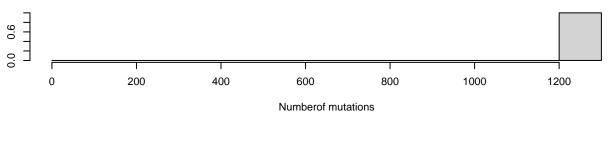








Liver-HCC SBS28 num non-0 = 1, proportion non-0 = 0.00307



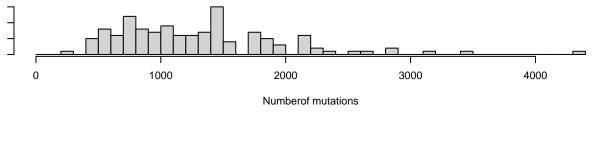
Frequency

Frequency

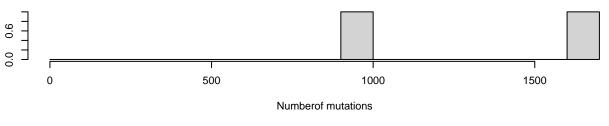
Frequency

Frequency

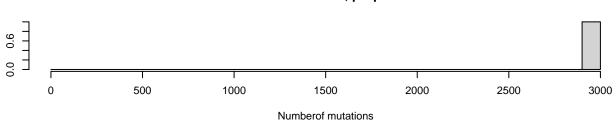
Liver-HCC SBS29 num non-0 = 125, proportion non-0 = 0.383



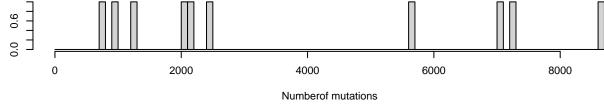
Liver-HCC SBS30 num non-0 = 2, proportion non-0 = 0.00613



Liver-HCC SBS31 num non-0 = 1, proportion non-0 = 0.00307



Liver–HCC SBS35 num non–0 = 10 , proportion non–0 = 0.0307



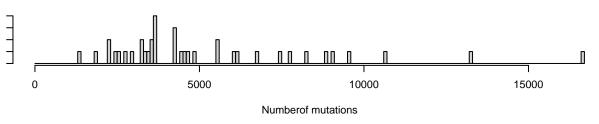
Frequency

Frequency

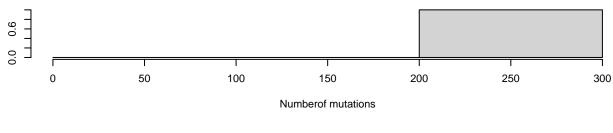
Frequency

Frequency

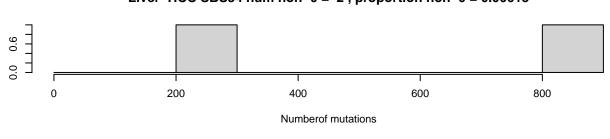
Liver-HCC SBS40 num non-0 = 39, proportion non-0 = 0.12



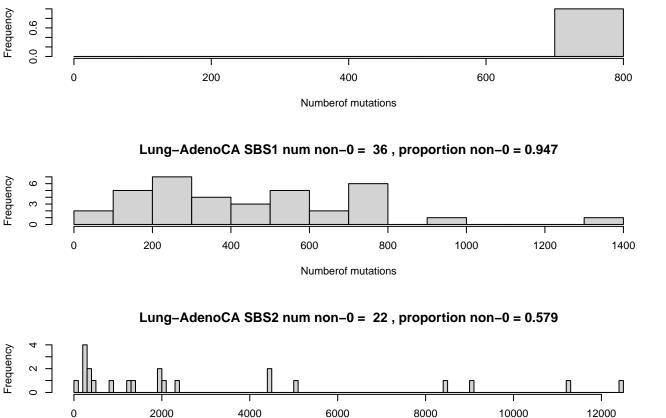
Liver-HCC SBS53 num non-0 = 1, proportion non-0 = 0.00307

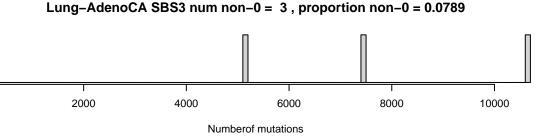


Liver-HCC SBS54 num non-0 = 2, proportion non-0 = 0.00613



Liver-HCC SBS56 num non-0 = 1, proportion non-0 = 0.00307





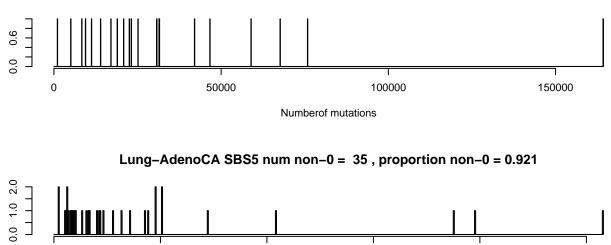
Number of mutations

0.0 0.6

0

Frequency

Lung-AdenoCA SBS4 num non-0 = 21 , proportion non-0 = 0.553



Frequency

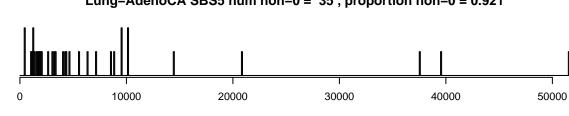
Frequency

Frequency

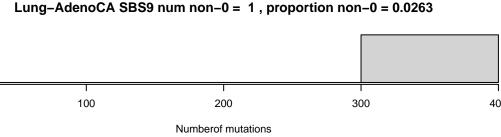
-requency

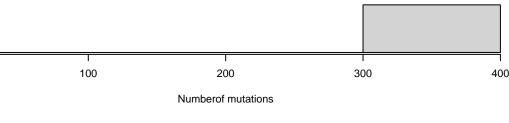
0.0 0.6

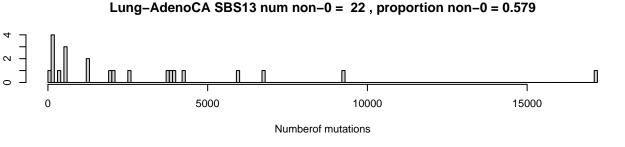
0



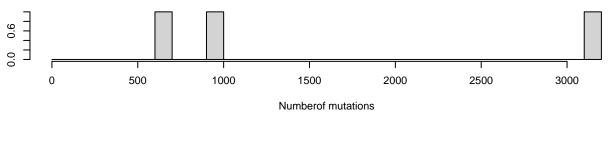
Number of mutations







Lung-AdenoCA SBS17a num non-0 = 3 , proportion non-0 = 0.0789



Frequency

Frequency

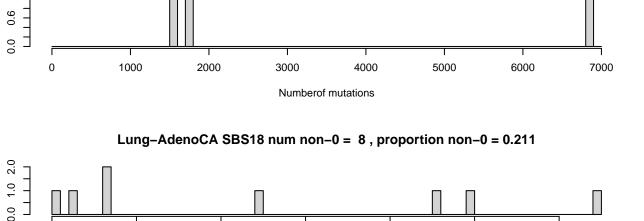
Frequency

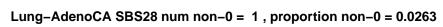
Frequency

1000

2000

Lung-AdenoCA SBS17b num non-0 = 3, proportion non-0 = 0.0789





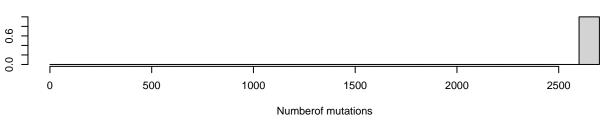
Number of mutations

4000

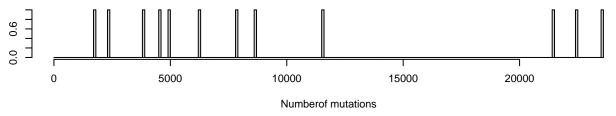
5000

6000

3000



Lung-AdenoCA SBS40 num non-0 = 12 , proportion non-0 = 0.316

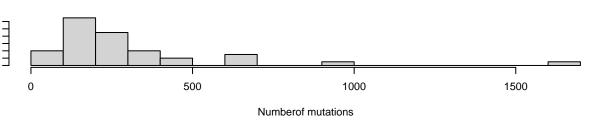


Frequency

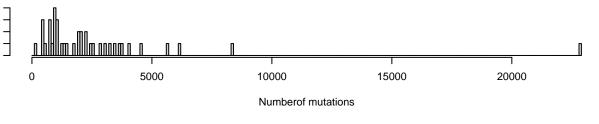
Frequency

Frequency

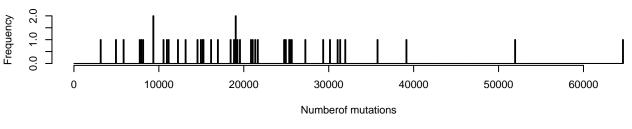
Lung-SCC SBS1 num non-0 = 37, proportion non-0 = 0.771



Lung-SCC SBS2 num non-0 = 40 , proportion non-0 = 0.833



Lung-SCC SBS4 num non-0 = 44, proportion non-0 = 0.917



Lung-SCC SBS5 num non-0 = 48, proportion non-0 = 1

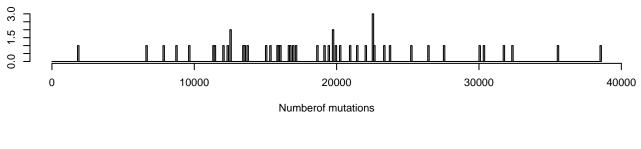
Frequency

Frequency

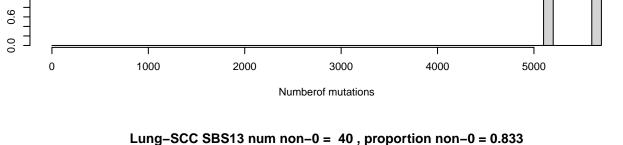
-requency

-requency

0



Lung-SCC SBS8 num non-0 = 2, proportion non-0 = 0.0417

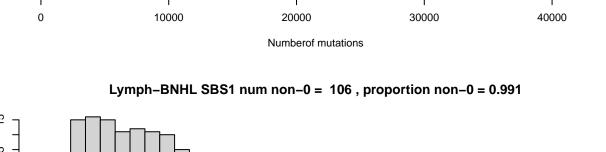




2000

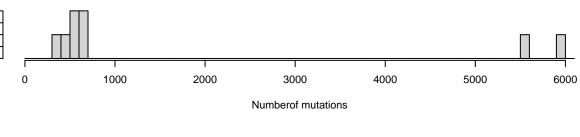
Number of mutations

3000



1000

Lymph-BNHL SBS2 num non-0 = 8, proportion non-0 = 0.0748

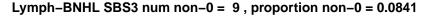


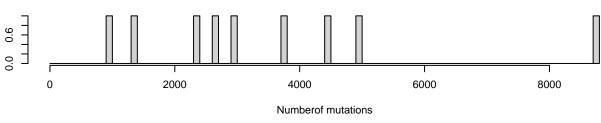
Frequency 0.0 1.0 2.0

Frequency

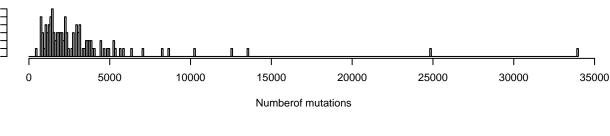
-requency

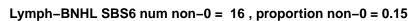
Frequency

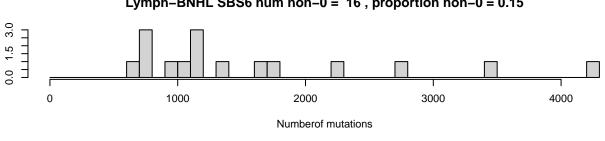


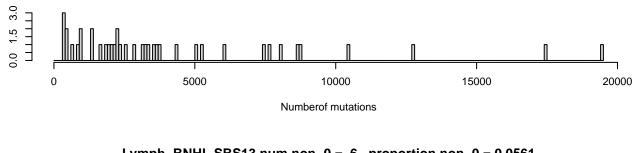


Lymph–BNHL SBS5 num non–0 = 106, proportion non–0 = 0.991

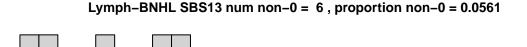








Lymph-BNHL SBS9 num non-0 = 40, proportion non-0 = 0.374

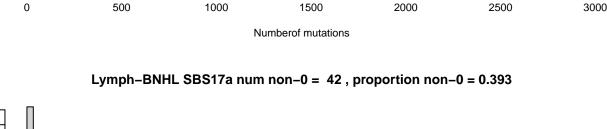


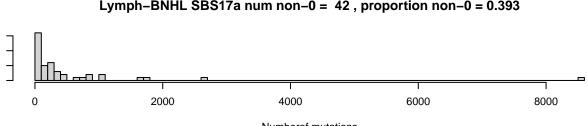
Frequency

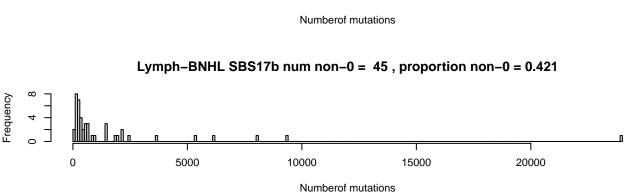
Frequency

-requency

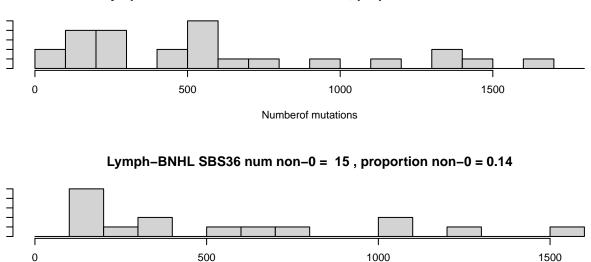
0.0 0.6







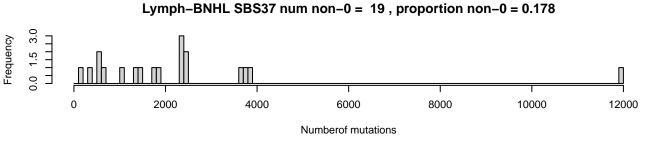
Lymph–BNHL SBS34 num non–0 = 25, proportion non–0 = 0.234



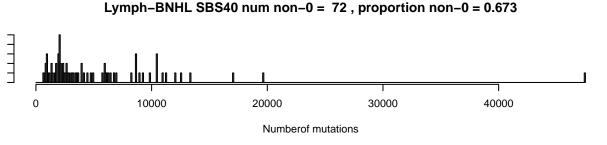
Frequency

Frequency

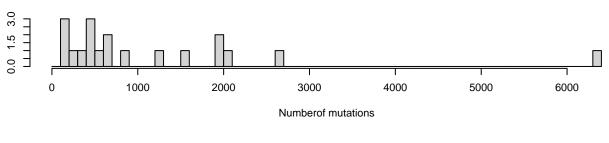
Frequency



Number of mutations



Lymph-BNHL SBS56 num non-0 = 19, proportion non-0 = 0.178



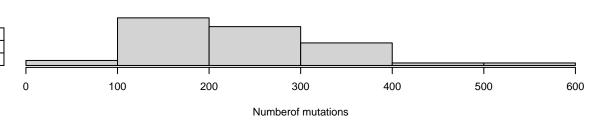
Frequency

Frequency

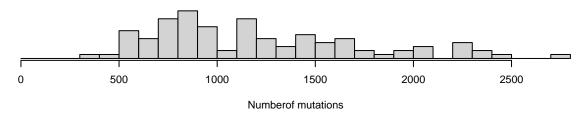
-requency

-requency

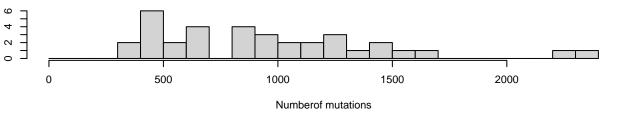
Lymph-CLL SBS1 num non-0 = 95, proportion non-0 = 1



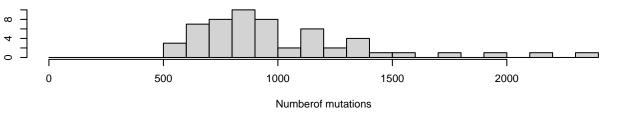
Lymph–CLL SBS5 num non–0 = 95, proportion non–0 = 1



Lymph-CLL SBS9 num non-0 = 35, proportion non-0 = 0.368



Lymph–CLL SBS40 num non–0 = 56, proportion non–0 = 0.589



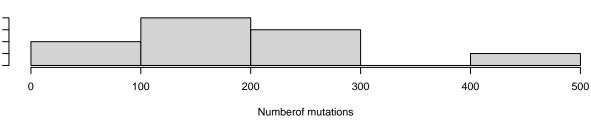
Frequency

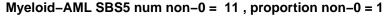
Frequency

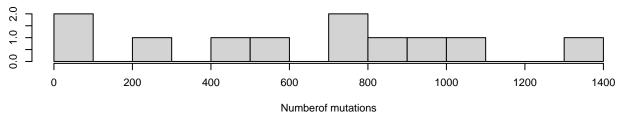
Frequency

Frequency

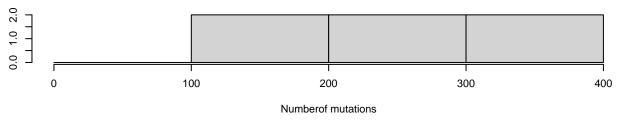
Myeloid-AML SBS1 num non-0 = 10, proportion non-0 = 0.909







Myeloid-AML SBS18 num non-0 = 6, proportion non-0 = 0.545



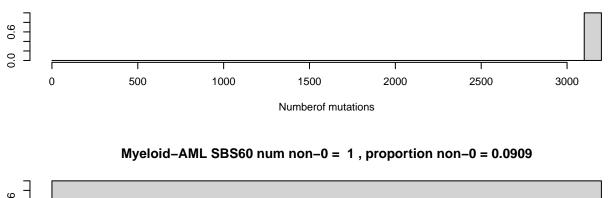
Myeloid-AML SBS31 num non-0 = 1 , proportion non-0 = 0.0909

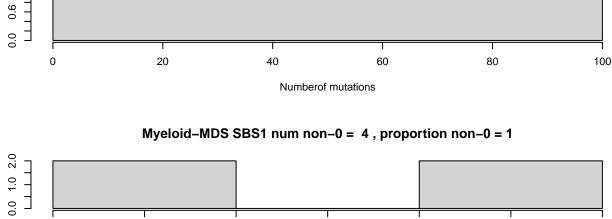
Frequency

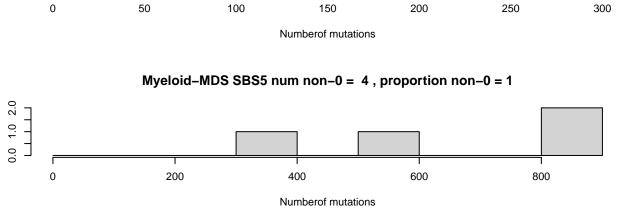
Frequency

Frequency

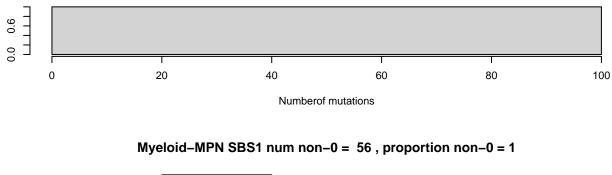
Frequency







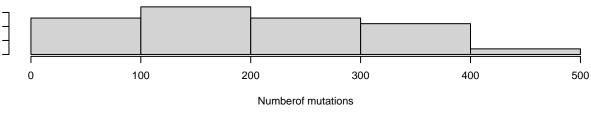
Myeloid-MDS SBS23 num non-0 = 1 , proportion non-0 = 0.25

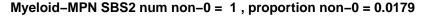


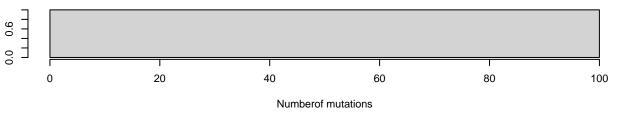
Frequency

-requency

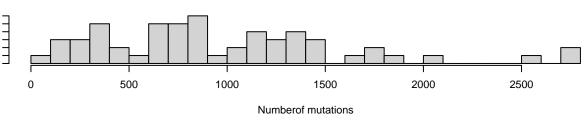
Frequency





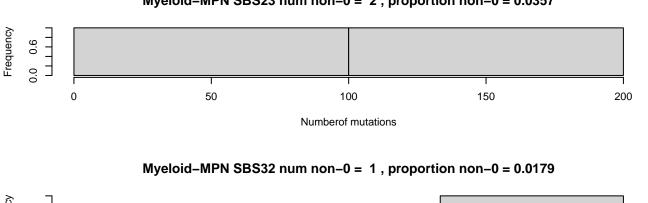


Myeloid-MPN SBS5 num non-0 = 56, proportion non-0 = 1



Myeloid-MPN SBS19 num non-0 = 3 , proportion non-0 = 0.0536 100 200 300 Number of mutations Myeloid-MPN SBS23 num non-0 = 2 , proportion non-0 = 0.0357

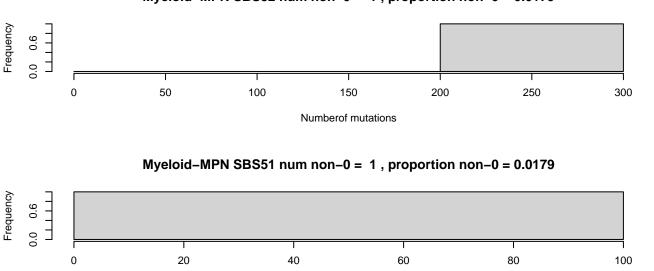
400



Frequency

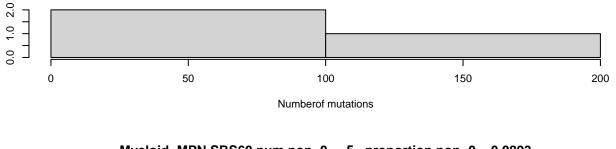
0.0 0.6

0



Number of mutations

Myeloid-MPN SBS58 num non-0 = 3, proportion non-0 = 0.0536



Frequency

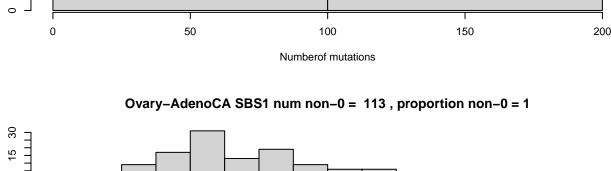
-requency

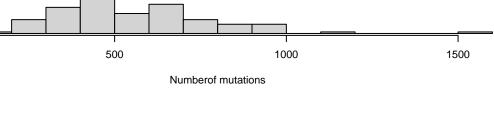
Frequency

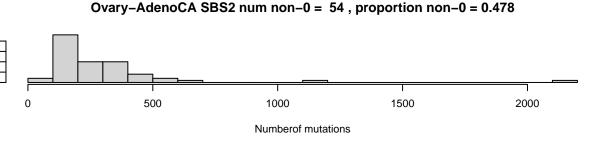
Frequency

0

Myeloid-MPN SBS60 num non-0 = 5, proportion non-0 = 0.0893

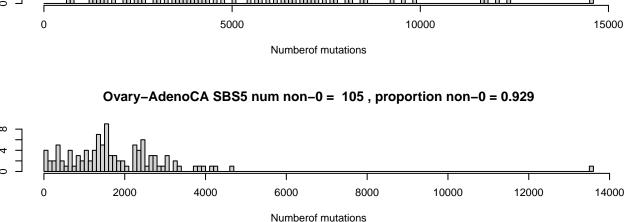


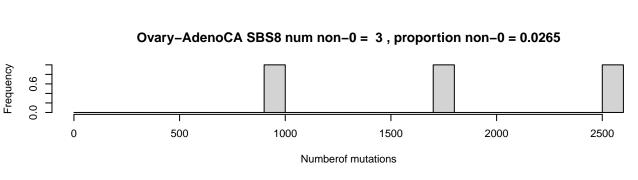


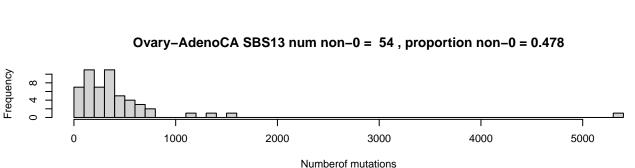


Ovary-AdenoCA SBS3 num non-0 = 106, proportion non-0 = 0.938

Frequency







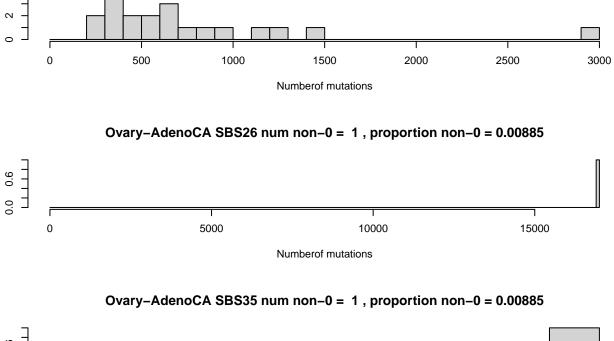
Ovary-AdenoCA SBS18 num non-0 = 20 , proportion non-0 = 0.177

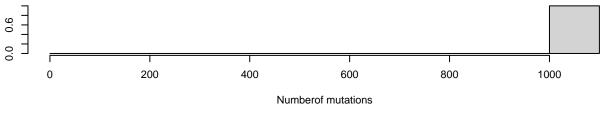
Frequency

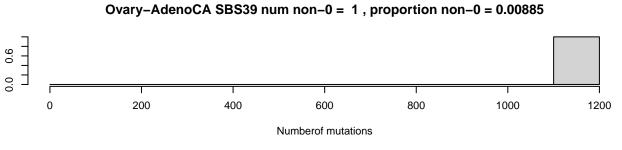
Frequency

Frequency

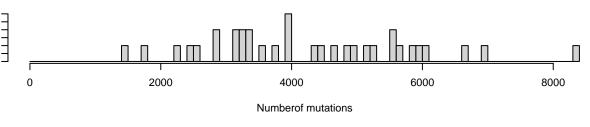
Frequency







Ovary-AdenoCA SBS40 num non-0 = 34 , proportion non-0 = 0.301



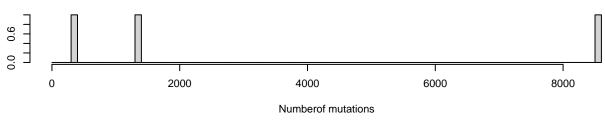
Frequency 0.0 1.5 3.0

Frequency

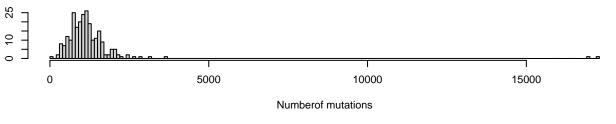
-requency

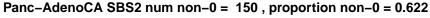
Frequency

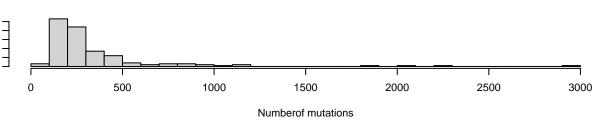
Ovary-AdenoCA SBS41 num non-0 = 3, proportion non-0 = 0.0265



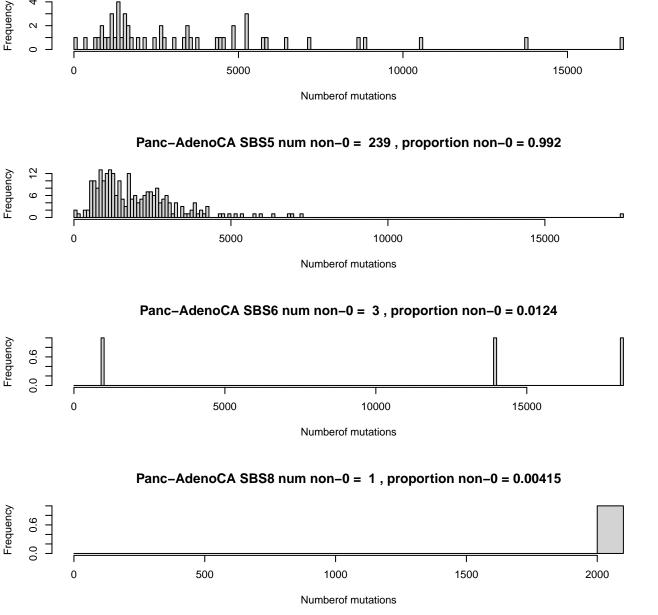
Panc-AdenoCA SBS1 num non-0 = 241, proportion non-0 = 1



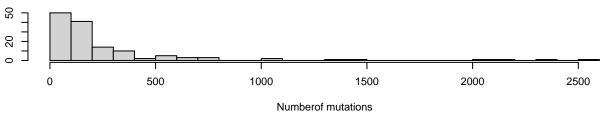




Panc-AdenoCA SBS3 num non-0 = 52, proportion non-0 = 0.216



Panc-AdenoCA SBS13 num non-0 = 136, proportion non-0 = 0.564

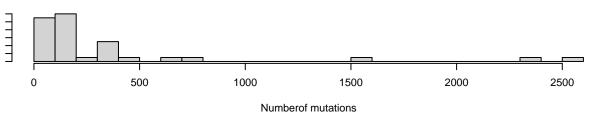


Frequency

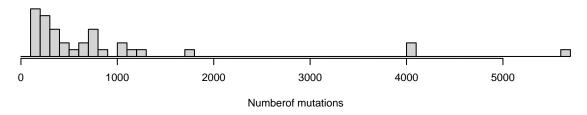
Frequency

Frequency

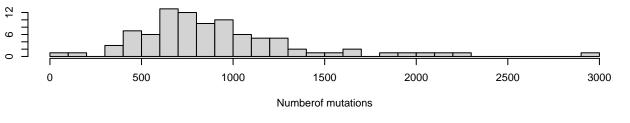
Panc-AdenoCA SBS17a num non-0 = 35, proportion non-0 = 0.145



Panc-AdenoCA SBS17b num non-0 = 35, proportion non-0 = 0.145



Panc-AdenoCA SBS18 num non-0 = 90, proportion non-0 = 0.373

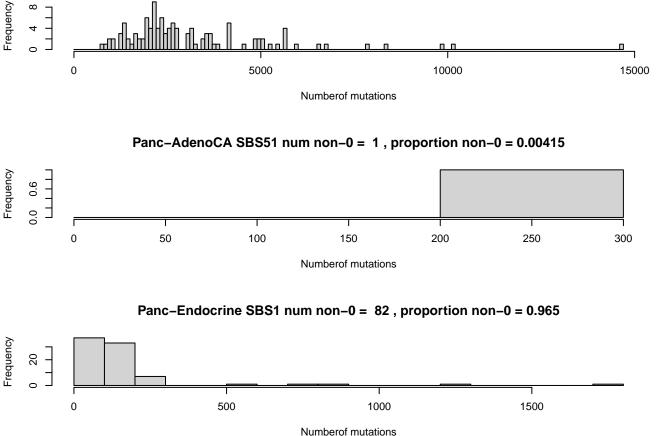


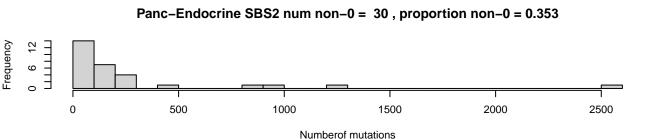
0.0 0.6 Frequency Number of mutations Panc-AdenoCA SBS26 num non-0 = 2, proportion non-0 = 0.0083 Frequency 0.0 0.6 Number of mutations Panc-AdenoCA SBS28 num non-0 = 5, proportion non-0 = 0.0207 Frequency 0.0 0.6 Number of mutations Panc-AdenoCA SBS30 num non-0 = 11, proportion non-0 = 0.0456 0.0 1.5 3.0 -requency

Number of mutations

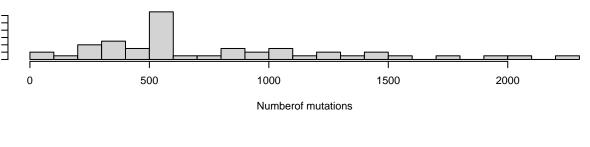
Panc-AdenoCA SBS20 num non-0 = 3, proportion non-0 = 0.0124

Panc-AdenoCA SBS40 num non-0 = 113, proportion non-0 = 0.469





Panc-Endocrine SBS3 num non-0 = 49, proportion non-0 = 0.576



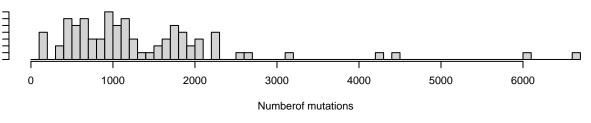
Frequency

Frequency

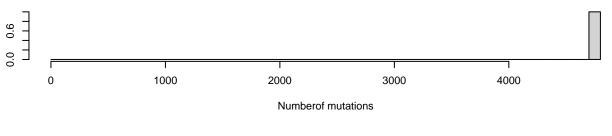
Frequency

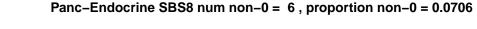
Frequency

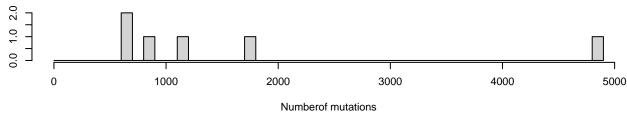
Panc-Endocrine SBS5 num non-0 = 82, proportion non-0 = 0.965



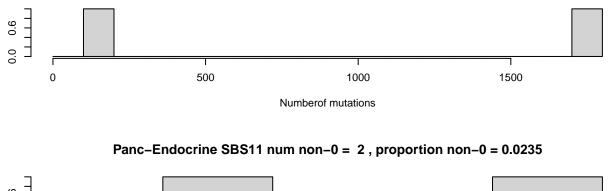
Panc-Endocrine SBS6 num non-0 = 1, proportion non-0 = 0.0118







Panc-Endocrine SBS9 num non-0 = 2, proportion non-0 = 0.0235

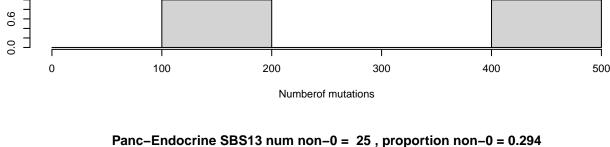


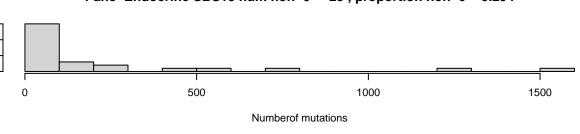
Frequency

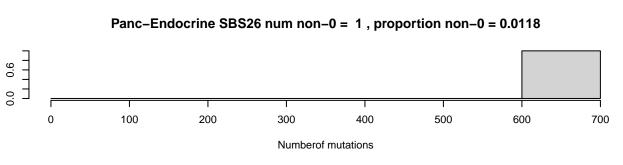
Frequency

Frequency

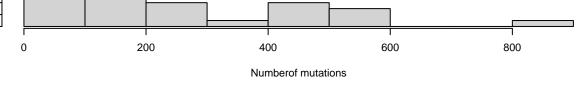
Frequency





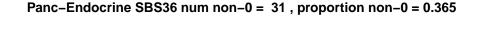


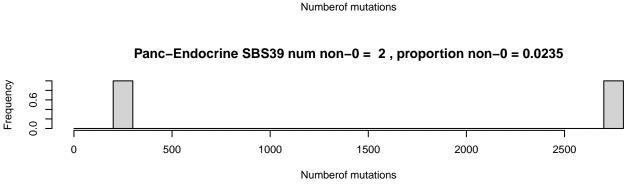
Panc-Endocrine SBS30 num non-0 = 26, proportion non-0 = 0.306

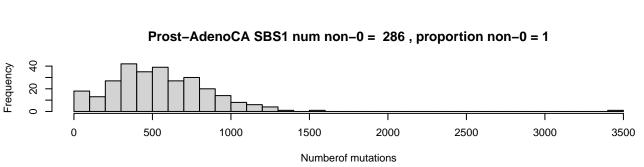


Frequency

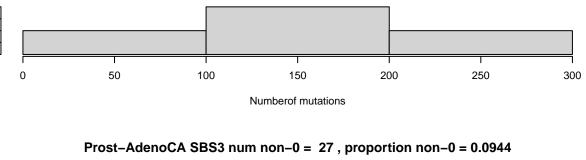
Frequency







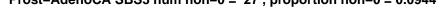
Prost-AdenoCA SBS2 num non-0 = 8, proportion non-0 = 0.028

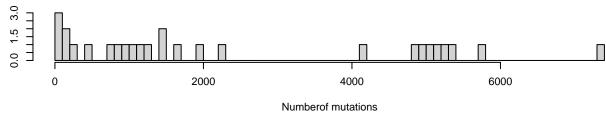


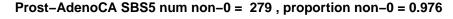
Frequency

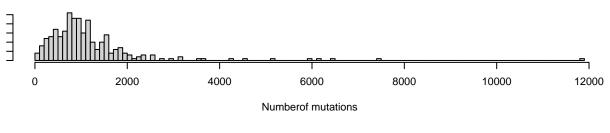
Frequency

Frequency

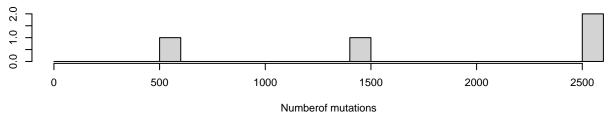




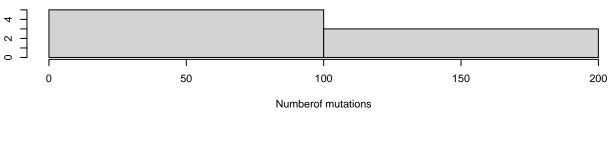




Prost-AdenoCA SBS8 num non-0 = 4, proportion non-0 = 0.014



Prost-AdenoCA SBS13 num non-0 = 8, proportion non-0 = 0.028



Frequency

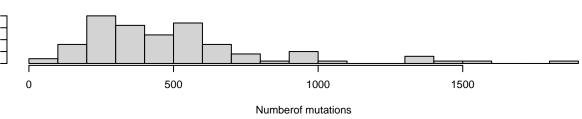
Frequency

Frequency

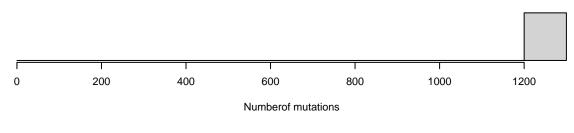
Frequency

0.0 0.6

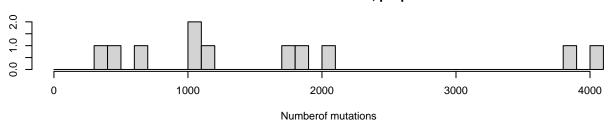
Prost-AdenoCA SBS18 num non-0 = 100, proportion non-0 = 0.35



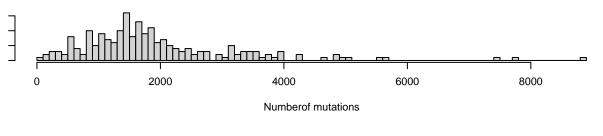
Prost-AdenoCA SBS33 num non-0 = 1, proportion non-0 = 0.0035



Prost-AdenoCA SBS37 num non-0 = 11 , proportion non-0 = 0.0385



Prost-AdenoCA SBS40 num non-0 = 204, proportion non-0 = 0.713



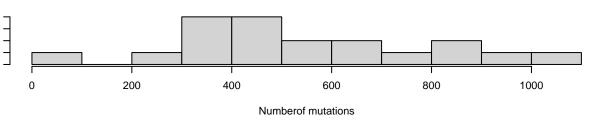
Frequency

-requency

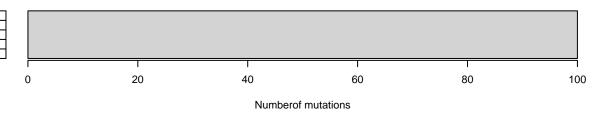
Frequency 0.0 0.6

Frequency

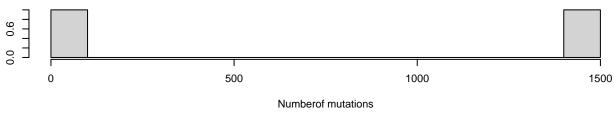
Prost-AdenoCA SBS41 num non-0 = 19 , proportion non-0 = 0.0664



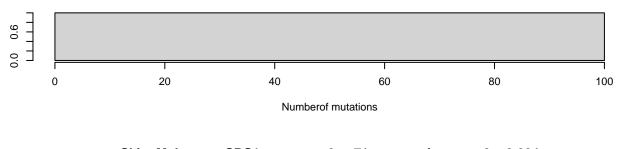
Prost-AdenoCA SBS45 num non-0 = 1, proportion non-0 = 0.0035



Prost-AdenoCA SBS52 num non-0 = 2, proportion non-0 = 0.00699



Prost-AdenoCA SBS58 num non-0 = 1 , proportion non-0 = 0.0035



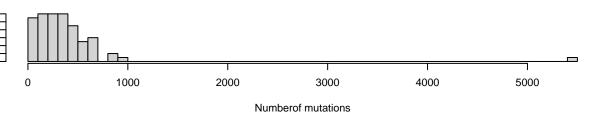
Frequency

Frequency

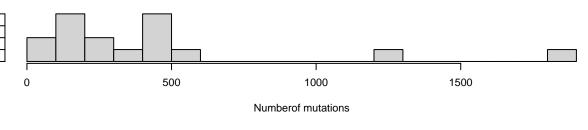
Frequency

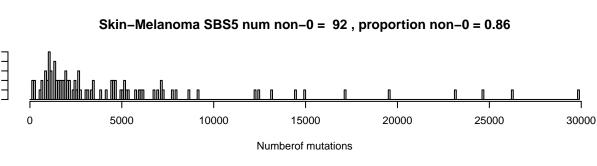
Frequency

Skin-Melanoma SBS1 num non-0 = 71, proportion non-0 = 0.664

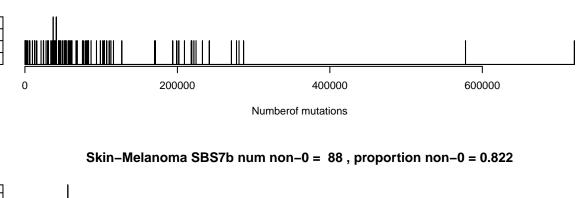


Skin-Melanoma SBS2 num non-0 = 16, proportion non-0 = 0.15





Skin-Melanoma SBS7a num non-0 = 88 , proportion non-0 = 0.822



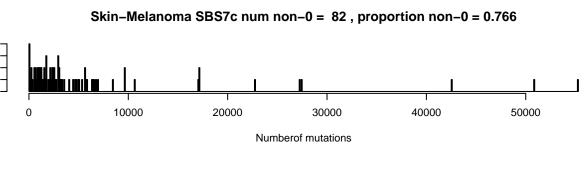
Frequency 0.0 1.0 2.0

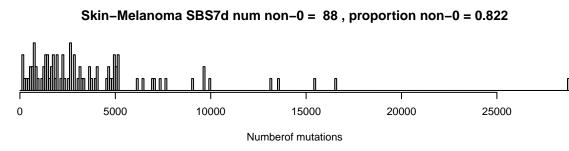
Frequency

-requency

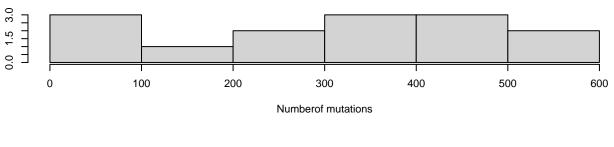
-requency







Skin-Melanoma SBS13 num non-0 = 14 , proportion non-0 = 0.131



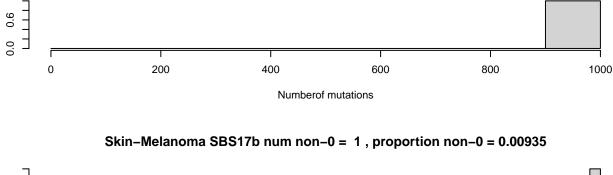
Frequency

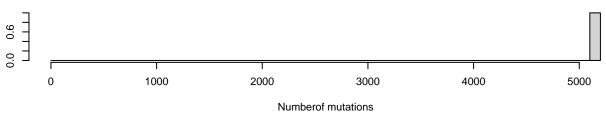
Frequency

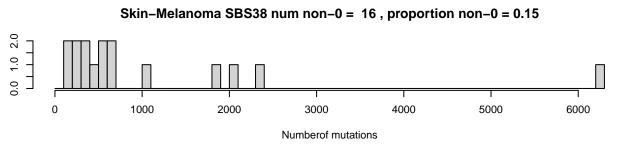
Frequency

Frequency

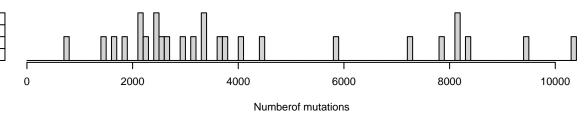
Skin-Melanoma SBS17a num non-0 = 1, proportion non-0 = 0.00935







Skin-Melanoma SBS40 num non-0 = 27 , proportion non-0 = 0.252



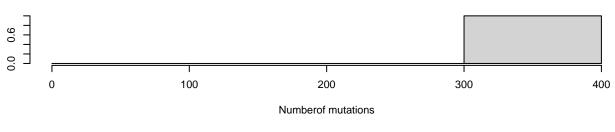
Frequency 0.0 1.0 2.0

Frequency

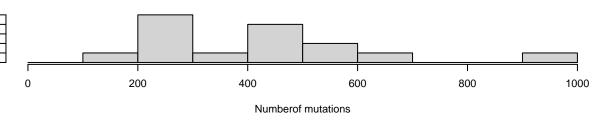
-requency

Frequency

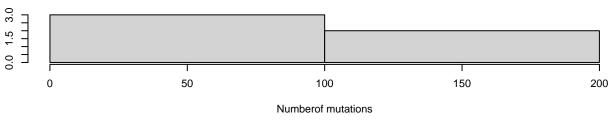
Skin-Melanoma SBS58 num non-0 = 1 , proportion non-0 = 0.00935



SoftTissue-Leiomyo SBS1 num non-0 = 15, proportion non-0 = 1



SoftTissue-Leiomyo SBS2 num non-0 = 5, proportion non-0 = 0.333



SoftTissue-Leiomyo SBS5 num non-0 = 15, proportion non-0 = 1 500 1500 1000 2000 Number of mutations SoftTissue-Leiomyo SBS13 num non-0 = 6, proportion non-0 = 0.4100 200 300 400 500 Number of mutations SoftTissue-Leiomyo SBS17a num non-0 = 1, proportion non-0 = 0.0667 100 150 50 200 Number of mutations SoftTissue-Leiomyo SBS17b num non-0 = 1, proportion non-0 = 0.0667

0.0 1.0 2.0 Frequency

0.0 1.5 3.0

0.0 0.6

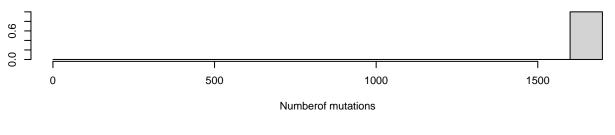
Frequency

Frequency

0

0 0.0 0.6 Frequency 100 200 0 300 400 Number of mutations

SoftTissue-Leiomyo SBS30 num non-0 = 1, proportion non-0 = 0.0667



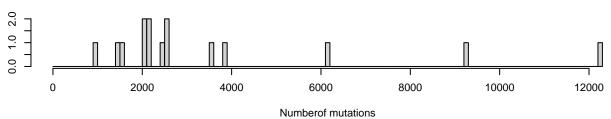
Frequency

Frequency

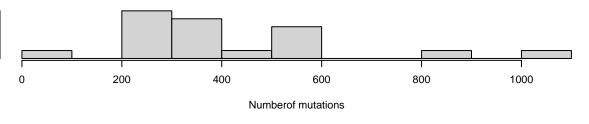
-requency

Frequency

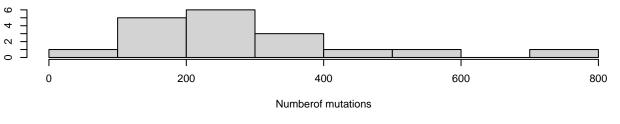
SoftTissue-Leiomyo SBS40 num non-0 = 15, proportion non-0 = 1



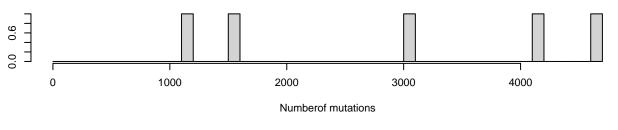
SoftTissue-Liposarc SBS1 num non-0 = 19, proportion non-0 = 1



SoftTissue-Liposarc SBS2 num non-0 = 18, proportion non-0 = 0.947



SoftTissue-Liposarc SBS3 num non-0 = 5, proportion non-0 = 0.263



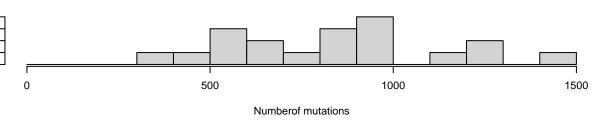
Frequency

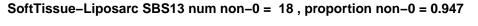
-requency

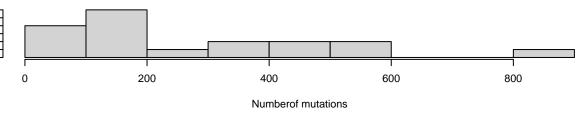
Frequency

Frequency

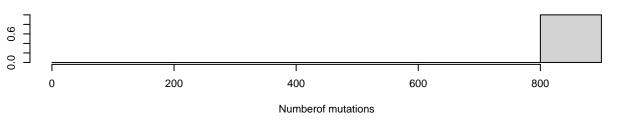
SoftTissue-Liposarc SBS5 num non-0 = 19, proportion non-0 = 1



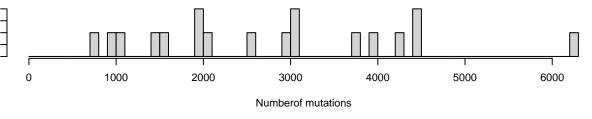




SoftTissue-Liposarc SBS37 num non-0 = 1, proportion non-0 = 0.0526

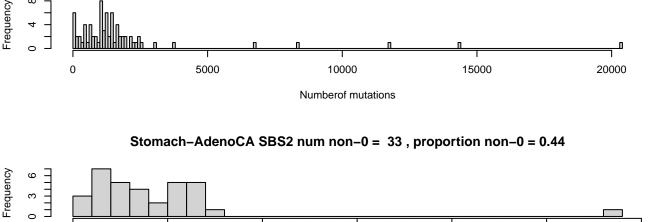


SoftTissue-Liposarc SBS40 num non-0 = 18, proportion non-0 = 0.947



Frequency 0.0 1.0 2.0

Stomach-AdenoCA SBS1 num non-0 = 75, proportion non-0 = 1



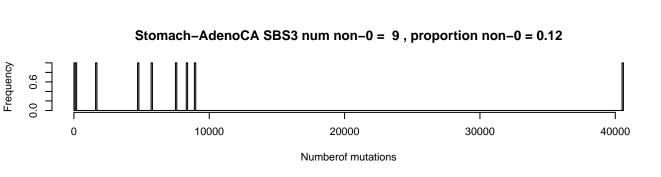
1500

Number of mutations

2000

2500

3000

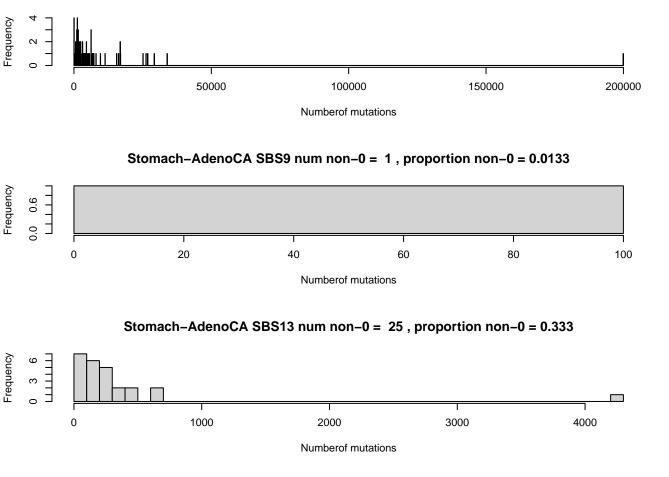


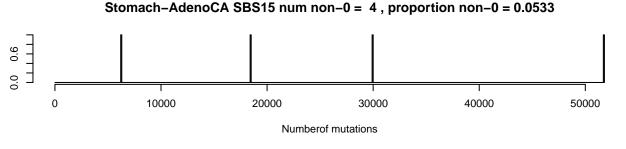
500

0

1000

Stomach-AdenoCA SBS5 num non-0 = 72 , proportion non-0 = 0.96

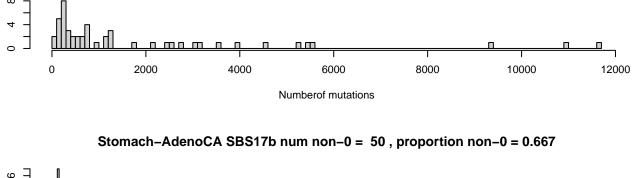


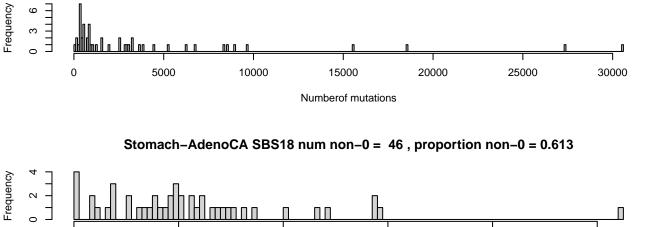


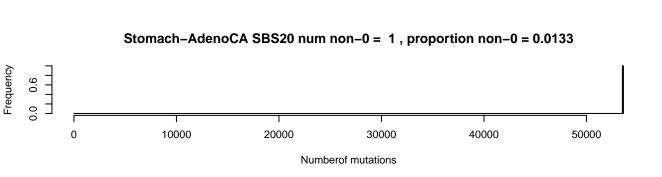
Frequency

Stomach–AdenoCA SBS17a num non–0 = 50 , proportion non–0 = 0.667

Frequency



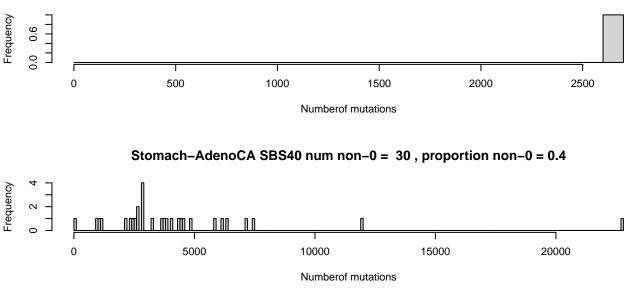




Number of mutations

Frequency 0.0 0.6 20000 60000 40000 80000 Number of mutations Stomach-AdenoCA SBS26 num non-0 = 3, proportion non-0 = 0.04 Frequency 9.0 0.0 5000 10000 15000 0 20000 Number of mutations Stomach-AdenoCA SBS28 num non-0 = 1, proportion non-0 = 0.0133 0.0 0.6 0 500 1000 1500 2000 2500 Number of mutations

Stomach-AdenoCA SBS21 num non-0 = 2, proportion non-0 = 0.0267



Frequency 0.0 0.6 4000 10000 6000 8000 2000 Number of mutations Stomach-AdenoCA SBS43 num non-0 = 1, proportion non-0 = 0.0133 Frequency 9.0 0.0 20 80 40 60 100 Number of mutations Stomach-AdenoCA SBS44 num non-0 = 5, proportion non-0 = 0.0667 0.0 0.6 Frequency 10000 0 20000 30000 Number of mutations Stomach-AdenoCA SBS51 num non-0 = 1, proportion non-0 = 0.0133 0.0 0.6 Frequency 20 40 60 80 0 100 Number of mutations

Stomach-AdenoCA SBS41 num non-0 = 11, proportion non-0 = 0.147

Stomach-AdenoCA SBS58 num non-0 = 1, proportion non-0 = 0.0133

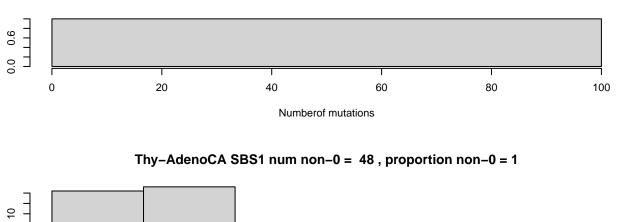
Frequency

-requency

Frequency

Frequency

0



300

400

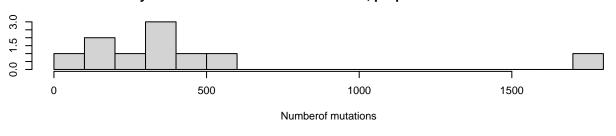
500

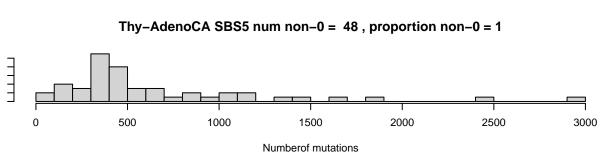
600



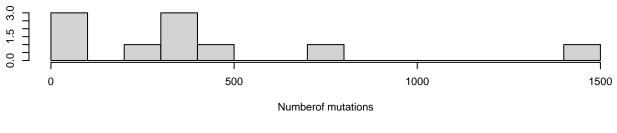
200

100





Thy-AdenoCA SBS13 num non-0 = 10, proportion non-0 = 0.208



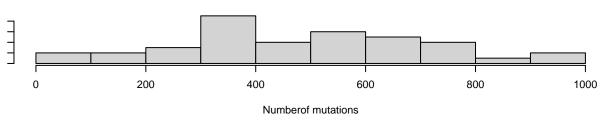
Frequency

Frequency

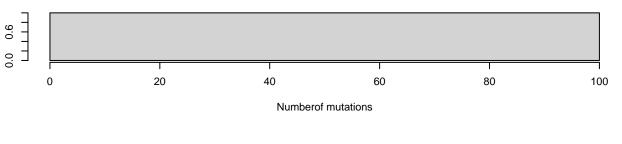
Frequency

-requency

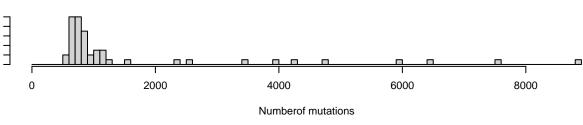
Thy-AdenoCA SBS40 num non-0 = 38, proportion non-0 = 0.792



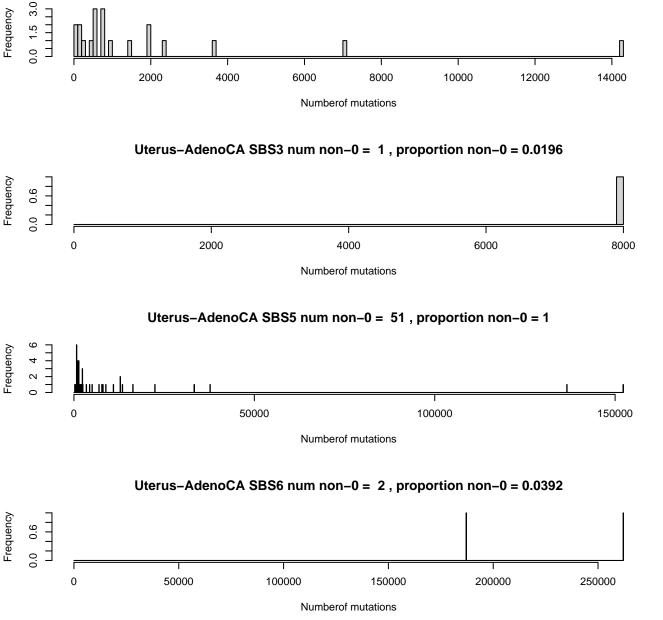
Thy-AdenoCA SBS58 num non-0 = 1, proportion non-0 = 0.0208



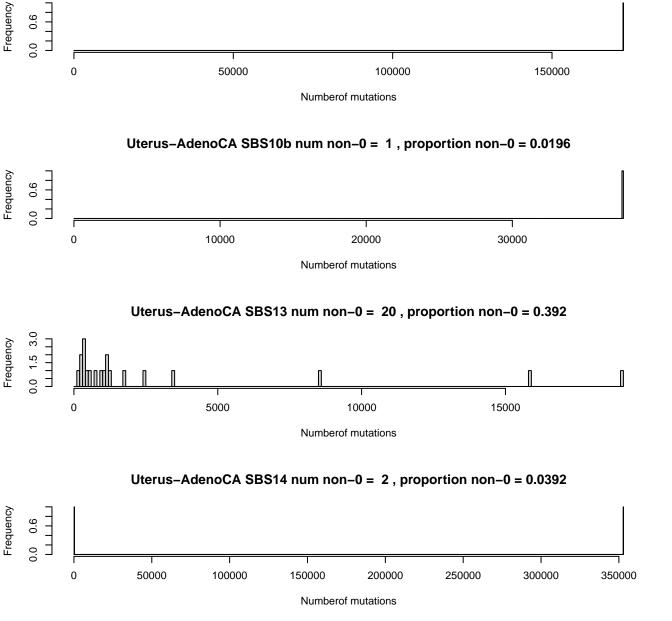
Uterus-AdenoCA SBS1 num non-0 = 49, proportion non-0 = 0.961



Uterus-AdenoCA SBS2 num non-0 = 20 , proportion non-0 = 0.392



Uterus-AdenoCA SBS10a num non-0 = 1 , proportion non-0 = 0.0196



Uterus-AdenoCA SBS15 num non-0 = 1, proportion non-0 = 0.0196

100000

Number of mutations

150000

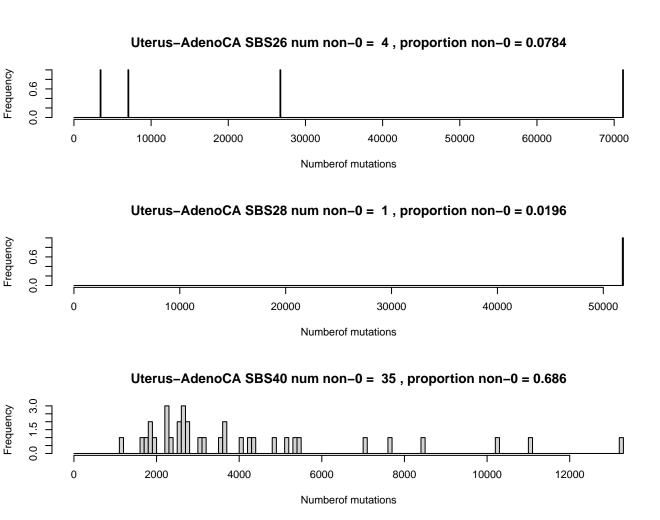
200000

50000

0.0 0.6

0

Frequency



Uterus-AdenoCA SBS44 num non-0 = 9, proportion non-0 = 0.176

