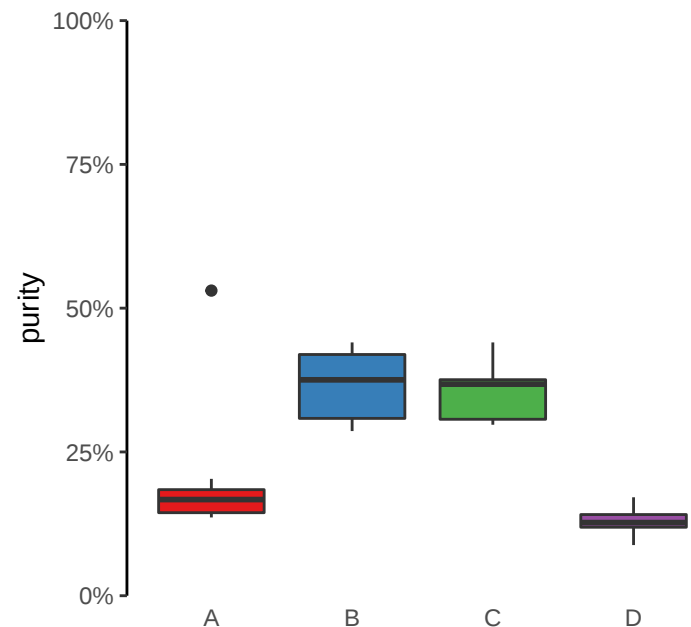
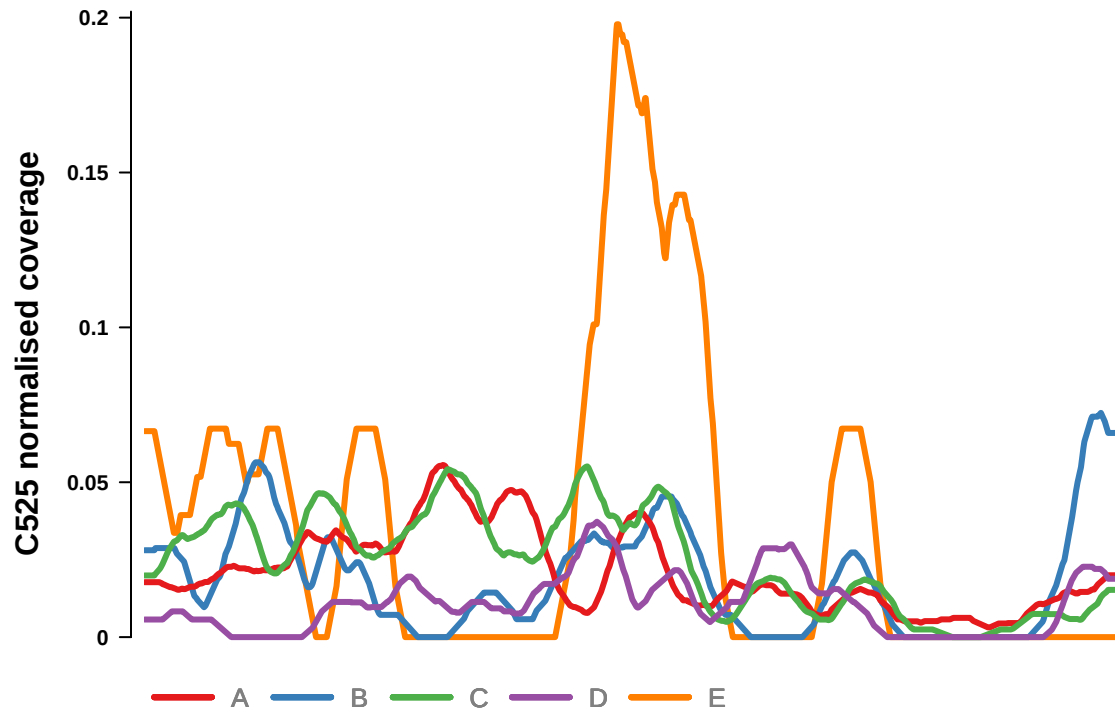


AKR1B1 promoter loss chr7:134458962-134459462

Putatively Subclonal (logFC = -0.86)

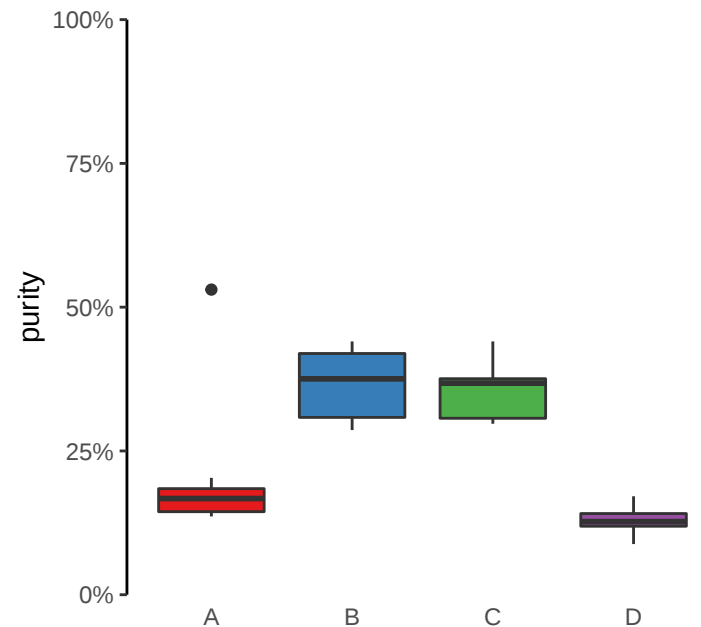
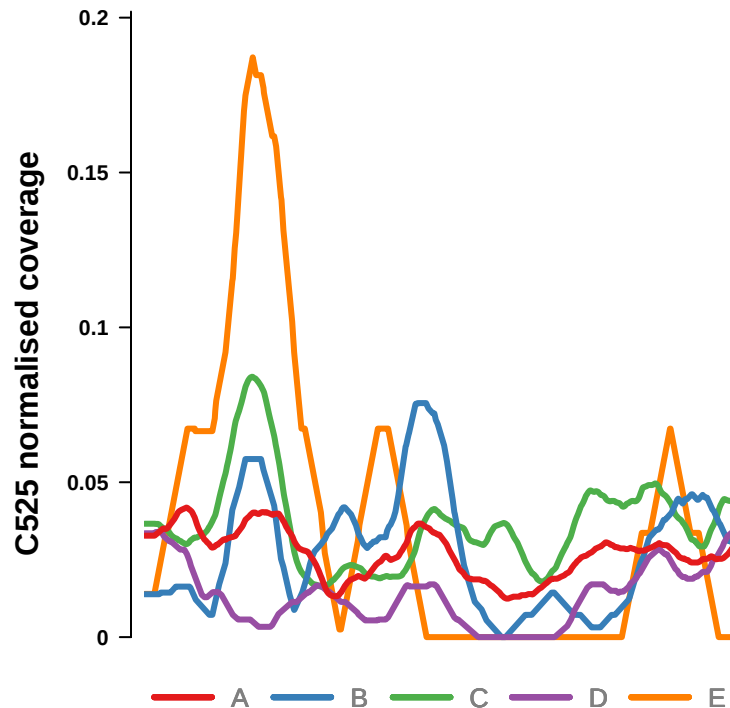
	P.	Adjusted P.
<i>~region vs. ~</i>	2.77e-03 **	1.04e-02 *
<i>~purity vs ~</i>	4.77e-01	8.63e-01
<i>~purity+region vs ~purity</i>	6.74e-03 **	2.37e-02 *
<i>Carcinoma vs. normal</i>	NA	NA



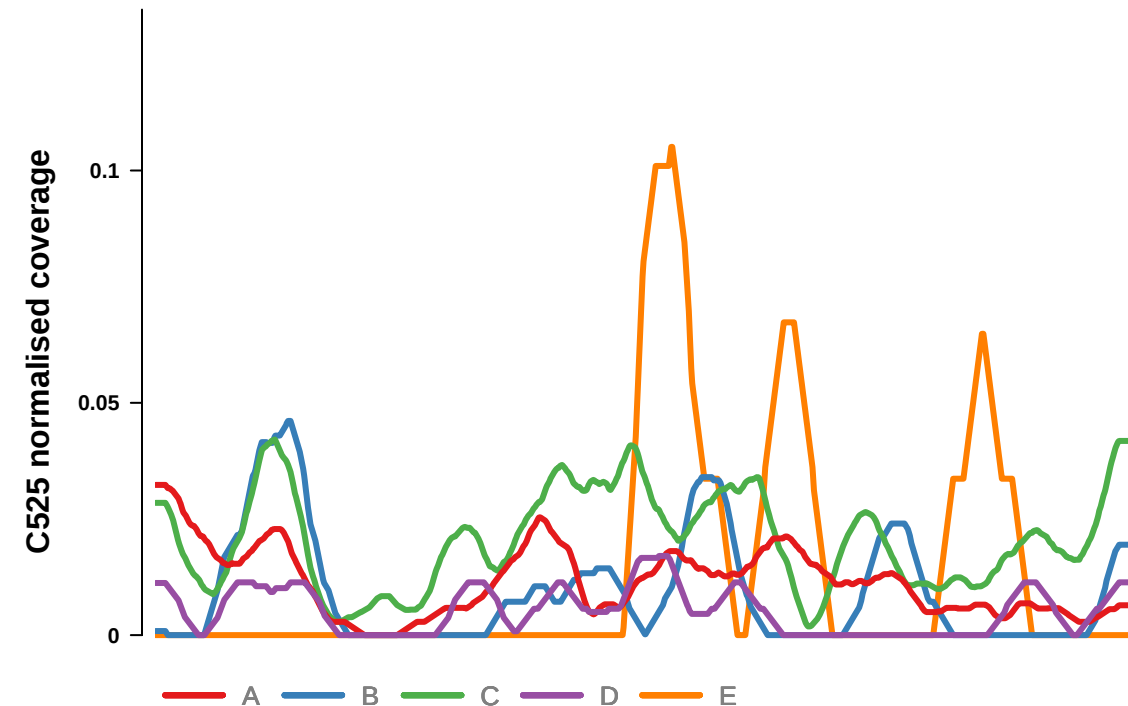
EPS8L1 promoter loss
chr19:55086973–55087473

Putatively Subclonal (logFC = -1.42)

	P.	Adjusted P.
<i>~region vs. ~</i>	1.14e-04 ***	9.8e-04 ***
<i>~purity vs ~</i>	7.1e-02	7.18e-01
<i>~purity+region vs ~purity</i>	1.22e-03 **	7.66e-03 **
<i>Carcinoma vs. normal</i>	2.39e-02 *	2.39e-02 *

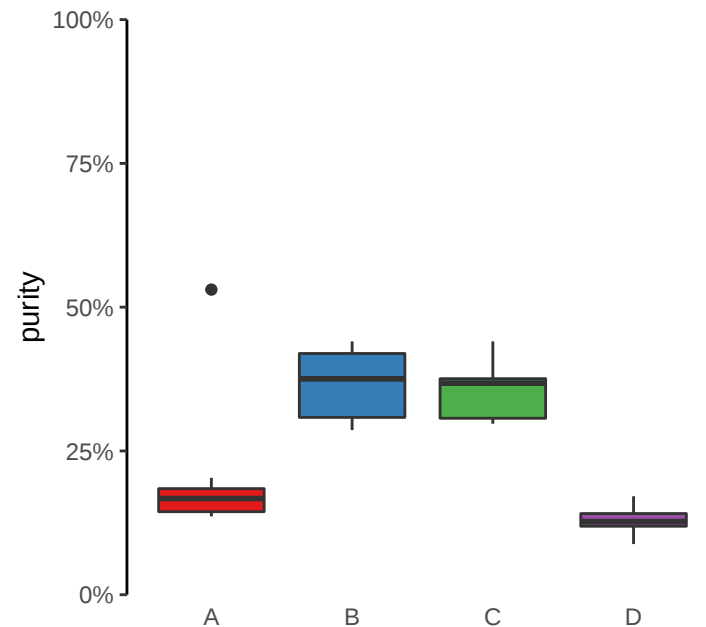


LETM1 promoter loss chr4:1839336-1839836

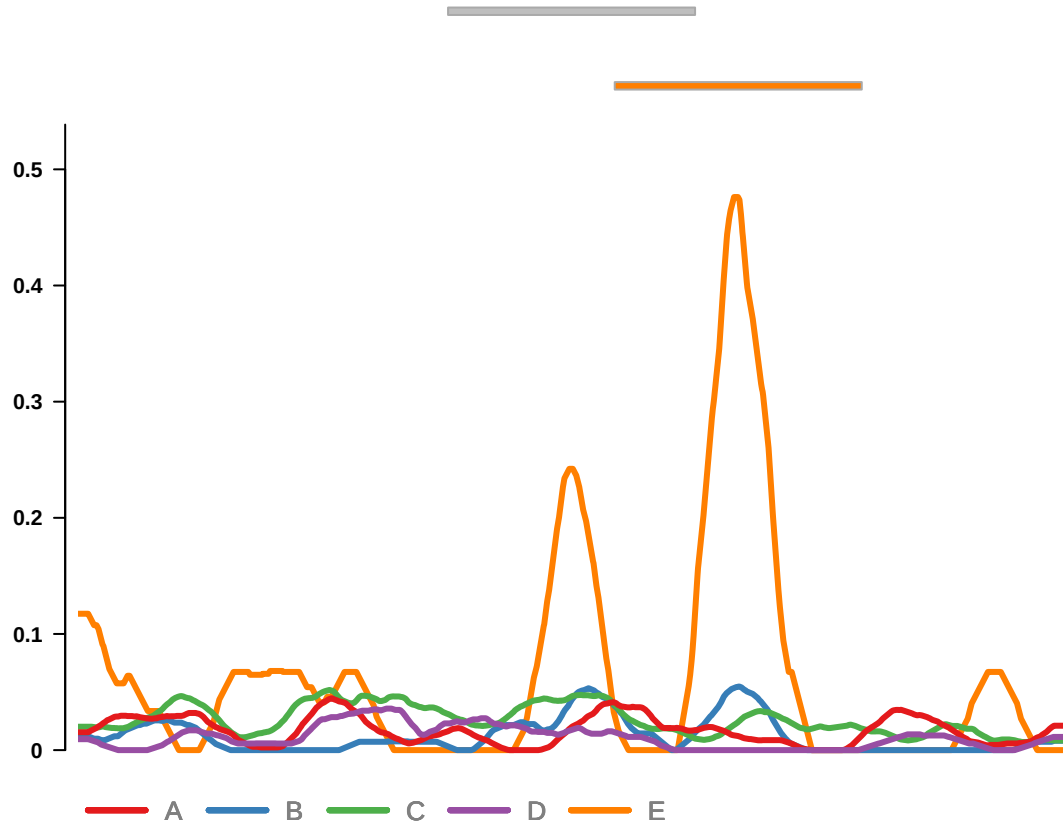


Putatively Subclonal (logFC = -0.47)

	P.	Adjusted P.
<i>~region vs. ~</i>	1.64e-05 ****	4.7e-04 ***
<i>~purity vs ~</i>	4.86e-02 *	6.12e-01
<i>~purity+region vs ~purity</i>	4.79e-04 ***	3.9e-03 **
<i>Carcinoma vs. normal</i>	NA	NA

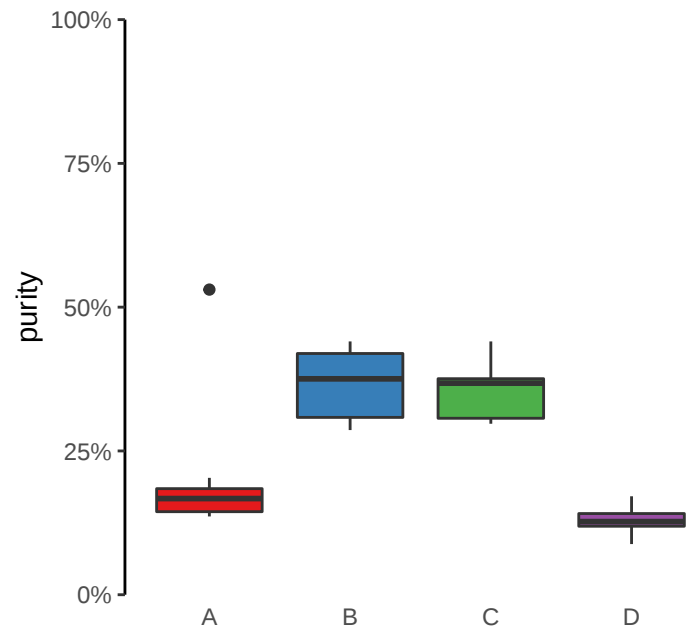


TIAM1 promoter loss
chr21:31558930–31559430



Putatively Subclonal (logFC = -1.93)

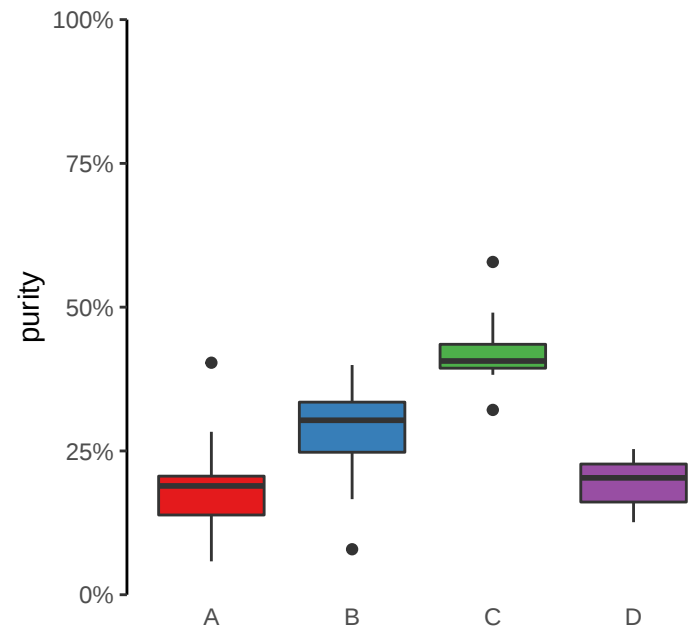
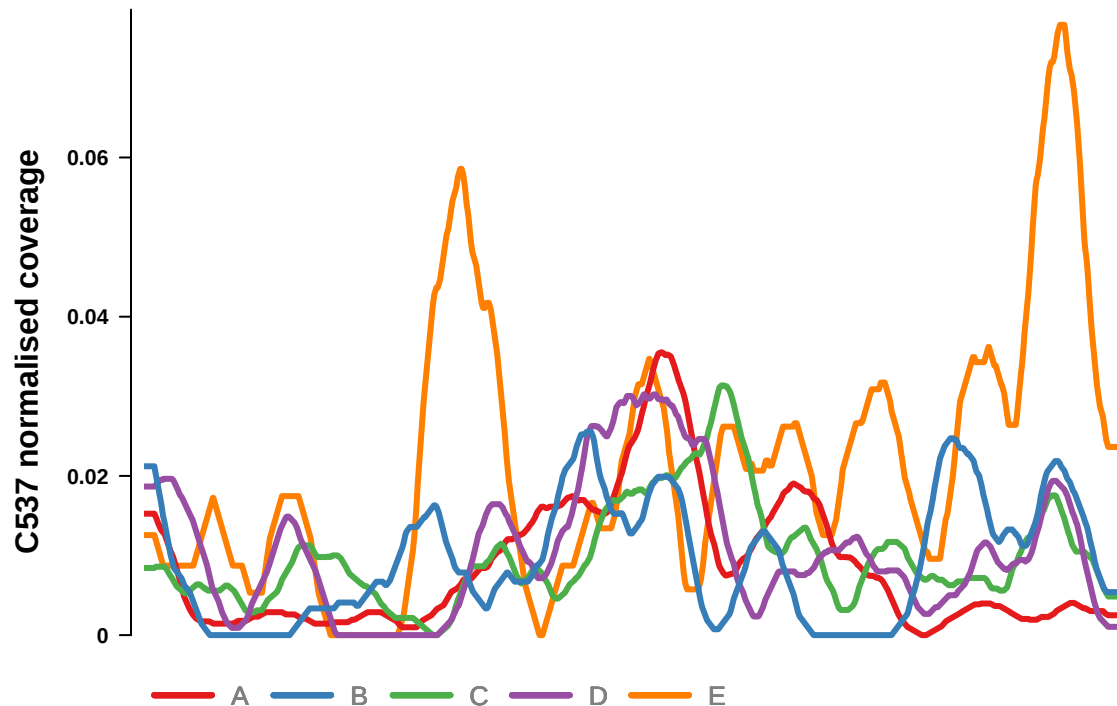
	P.	Adjusted P.
<i>~region vs. ~</i>	1.77e-03 **	7.63e-03 **
<i>~purity vs ~</i>	3.22e-01	7.48e-01
<i>~purity+region vs ~purity</i>	5.79e-03 **	2.21e-02 *
<i>Carcinoma vs. normal</i>	1.45e-03 **	1.45e-03 **



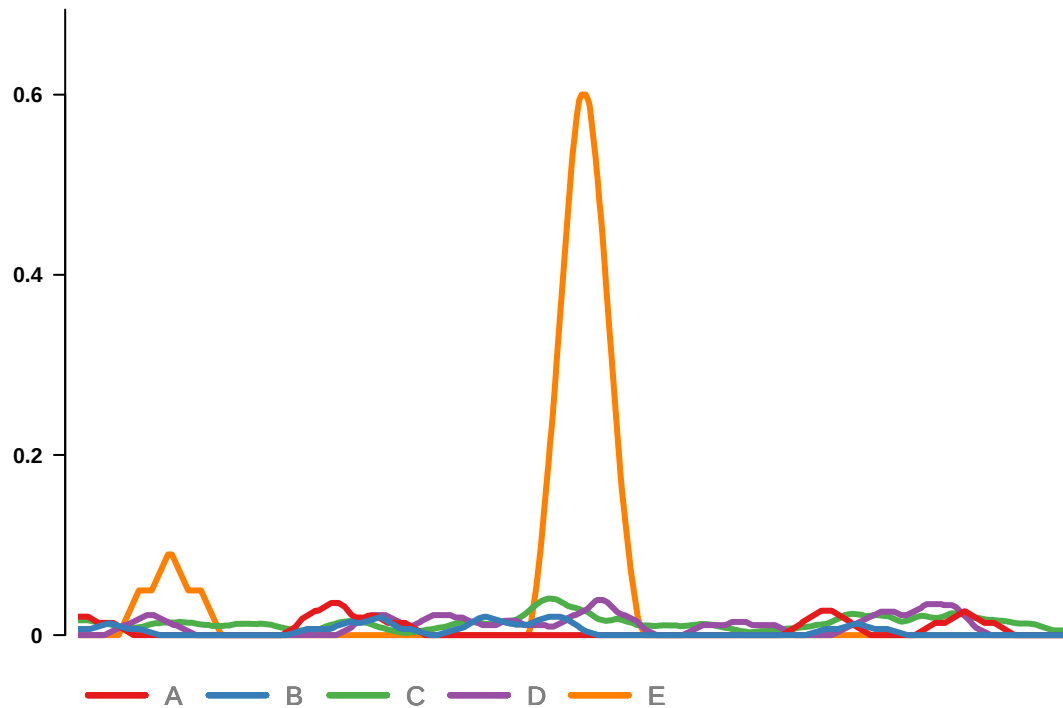
LPCAT4 promoter loss
chr15:34362318–34362818

Putatively Subclonal (logFC = -0.69)

	P.	Adjusted P.
<i>~region vs. ~</i>	1.44e-02 *	6.02e-02
<i>~purity vs ~</i>	7.33e-01	8.29e-01
<i>~purity+region vs ~purity</i>	3.01e-03 **	3.96e-02 *
<i>Carcinoma vs. normal</i>	NA	NA

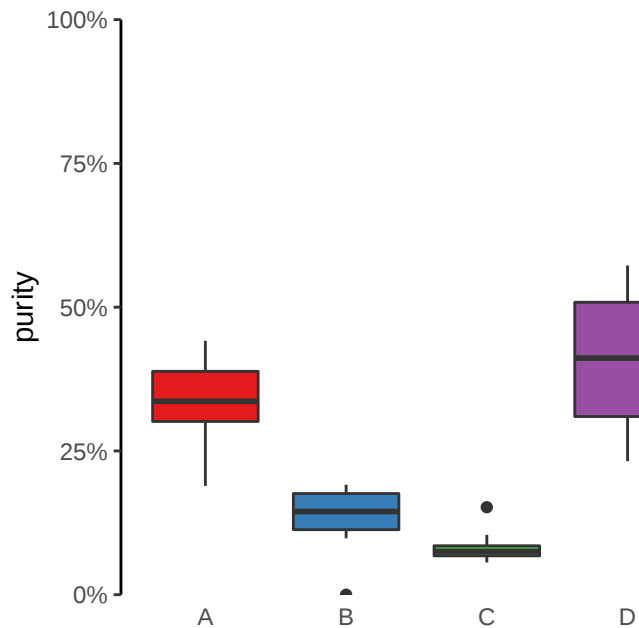


R3HDM1 promoter loss chr2:135633520–135634020

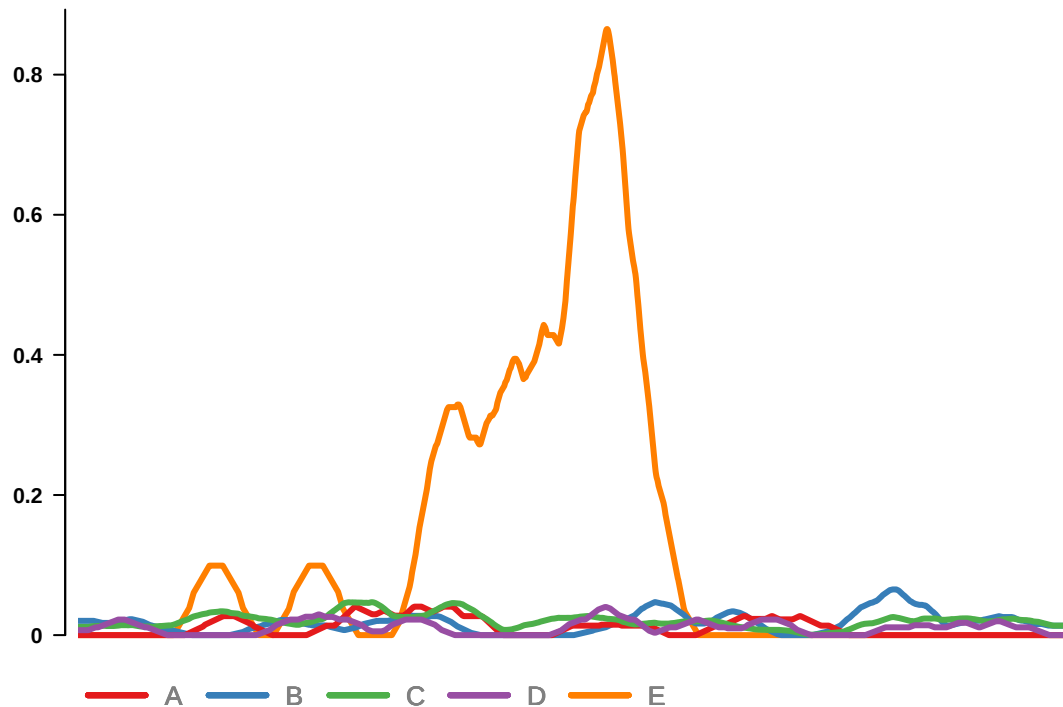


Putatively Subclonal (logFC = -0.57)

	P.	Adjusted P.
<i>~region vs. ~</i>	4.47e-05 ****	2.62e-04 ***
<i>~purity vs ~</i>	2.84e-03 **	6.41e-03 **
<i>~purity+region vs ~purity</i>	8.54e-03 **	4.22e-02 *
<i>Carcinoma vs. normal</i>	NA	NA

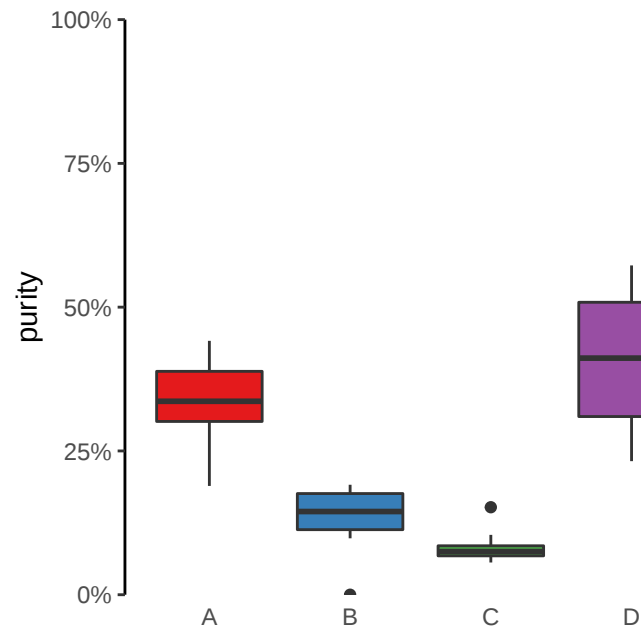


SMAD3 promoter loss
chr15:67127194–67127694



Putatively Subclonal (logFC = -1.32)

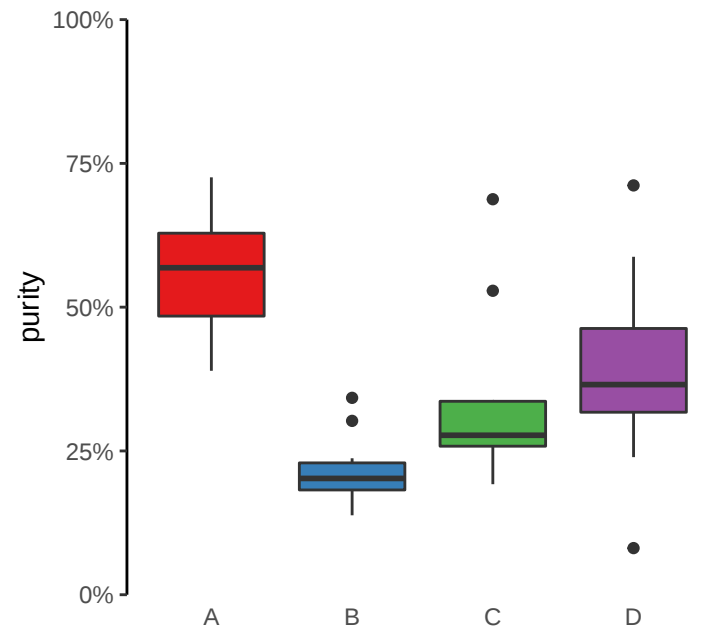
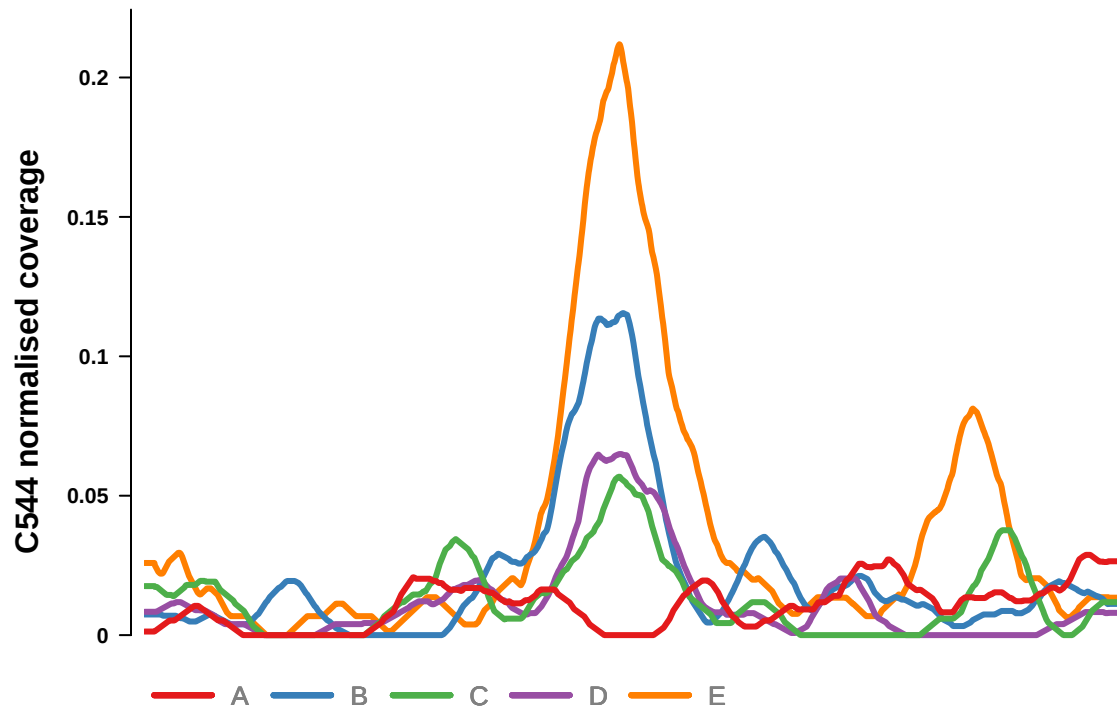
	P.	Adjusted P.
<i>~region vs. ~</i>	1.42e-07 ****	4.16e-06 ****
<i>~purity vs ~</i>	7.03e-05 ****	6.48e-04 ***
<i>~purity+region vs ~purity</i>	3.12e-04 ***	4.93e-03 **
<i>Carcinoma vs. normal</i>	1.05e-01	1.05e-01



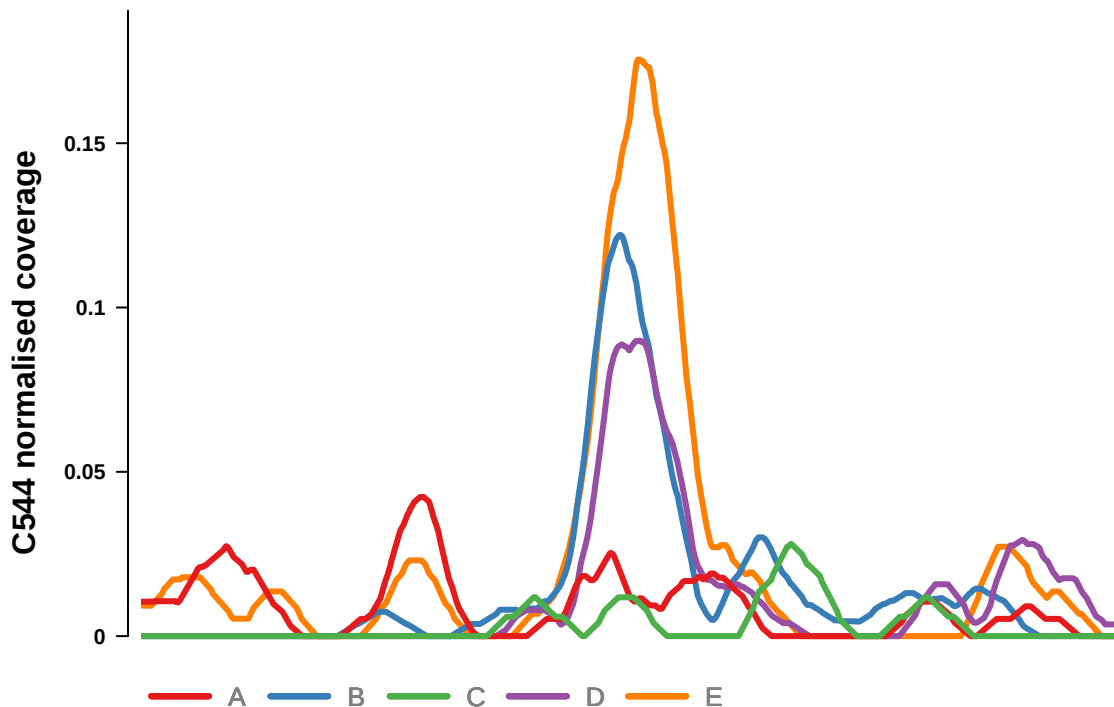
LPCAT4 promoter loss
chr15:34362318–34362818

Putatively Subclonal (logFC = -1.32)

	P.	Adjusted P.
<i>~region vs. ~</i>	1.89e-05 ****	3.48e-04 ***
<i>~purity vs ~</i>	3.26e-03 **	2.16e-02 *
<i>~purity+region vs ~purity</i>	1.8e-03 **	1.59e-02 *
<i>Carcinoma vs. normal</i>	2.91e-02 *	2.91e-02 *

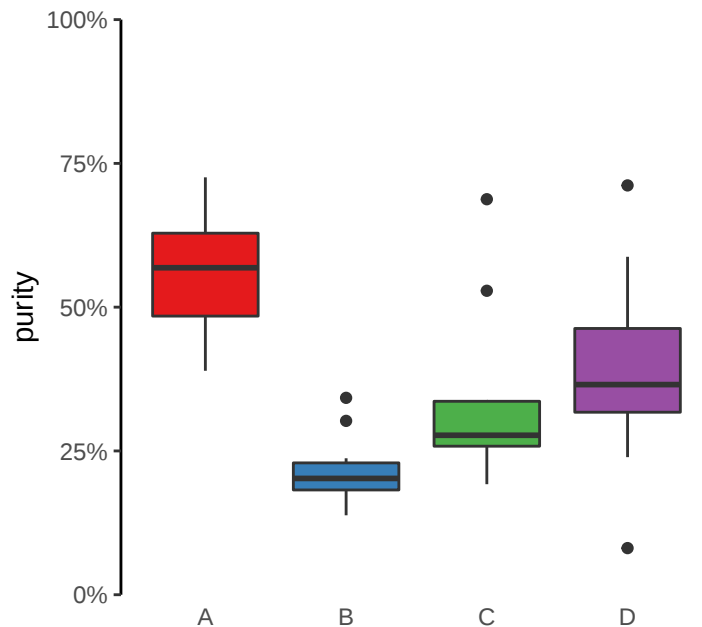


R3HDM1 promoter loss chr2:135633520–135634020

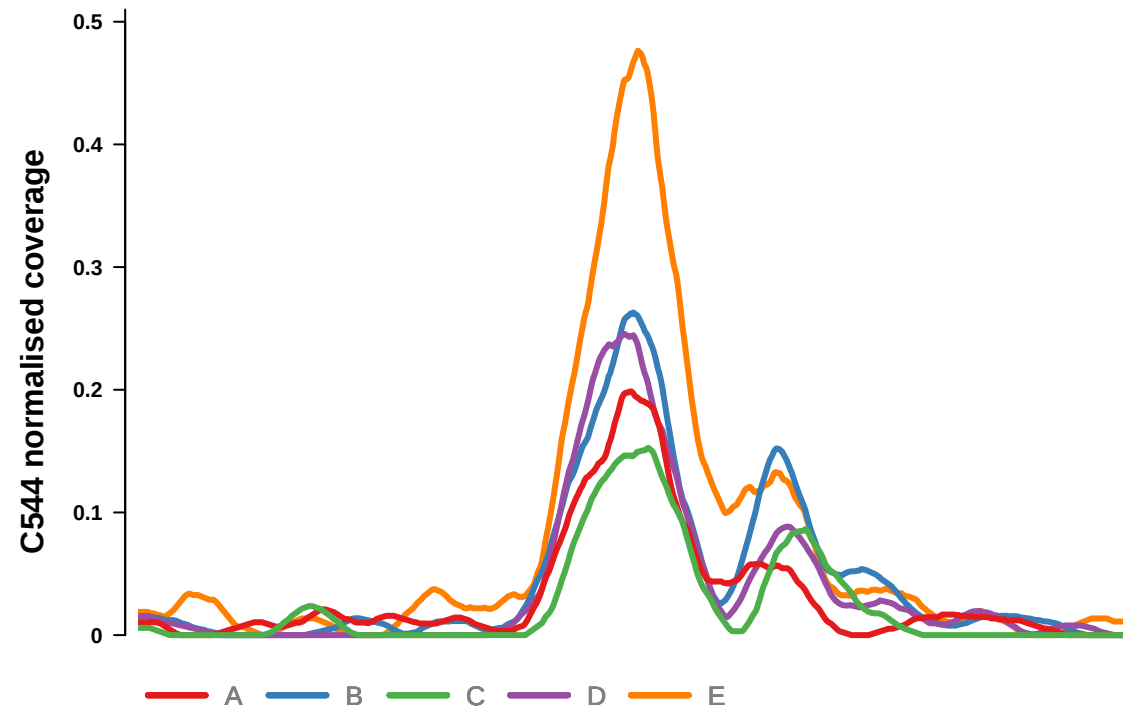


Putatively Subclonal (logFC = -1.24)

	P.	Adjusted P.
<i>~region vs. ~</i>	6.97e-05 ****	6.14e-04 ***
<i>~purity vs ~</i>	3.73e-02 *	9.93e-02
<i>~purity+region vs ~purity</i>	9.16e-04 ***	1.17e-02 *
<i>Carcinoma vs. normal</i>	5.05e-02	5.05e-02

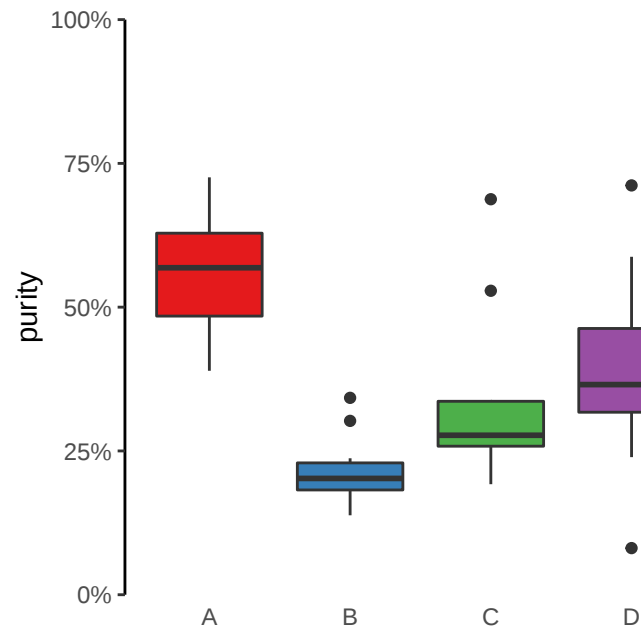


SPAG16 promoter loss
chr2:213284089–213284589

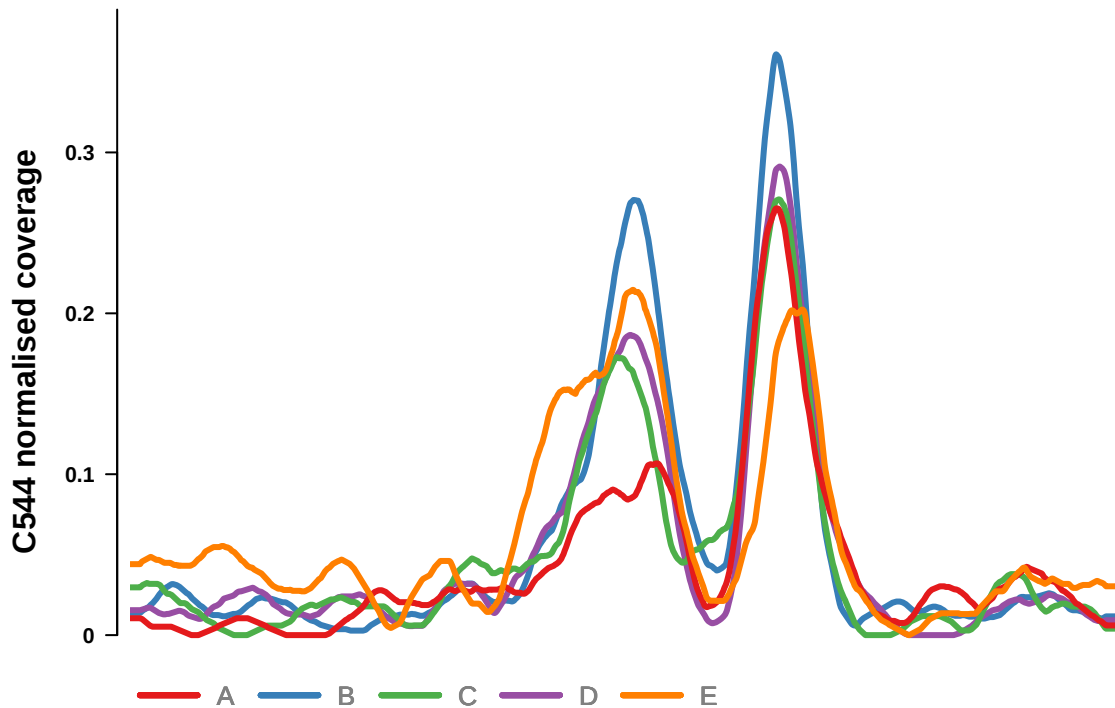


Putatively Subclonal (logFC = -0.12)

	P.	Adjusted P.
<i>~region vs. ~</i>	4.67e-03 **	1.96e-02 *
<i>~purity vs ~</i>	7.21e-01	8e-01
<i>~purity+region vs ~purity</i>	5.32e-03 **	2.83e-02 *
<i>Carcinoma vs. normal</i>	NA	NA

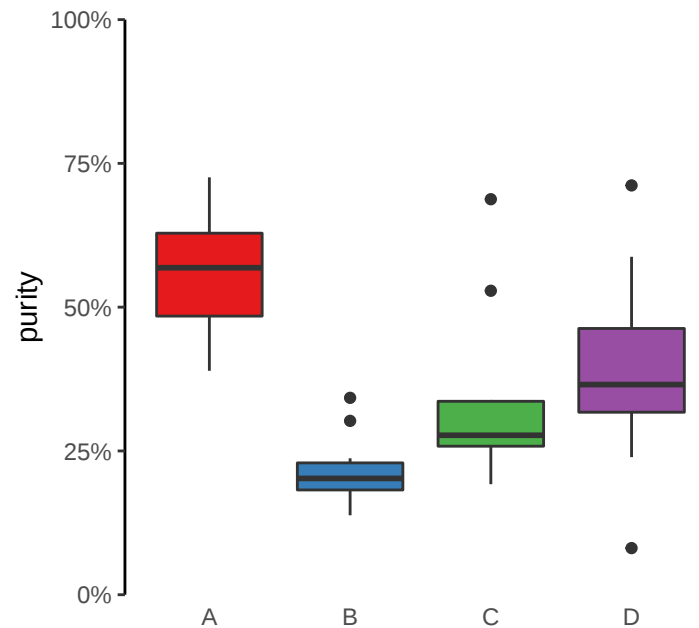


TIAM1 promoter loss
chr21:31558930–31559430

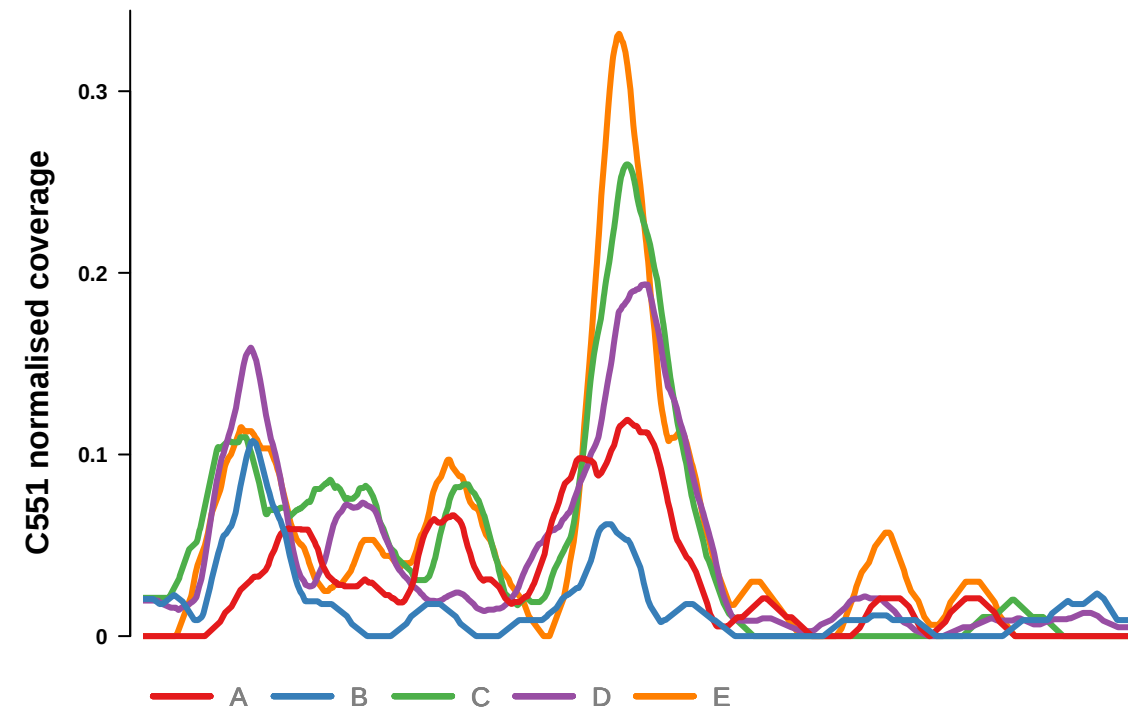


Putatively Subclonal (logFC = -0.63)

	P.	Adjusted P.
<i>~region vs. ~</i>	1.93e-04 ***	1.55e-03 **
<i>~purity vs ~</i>	3.51e-01	5.18e-01
<i>~purity+region vs ~purity</i>	3.65e-04 ***	9.1e-03 **
<i>Carcinoma vs. normal</i>	NA	NA

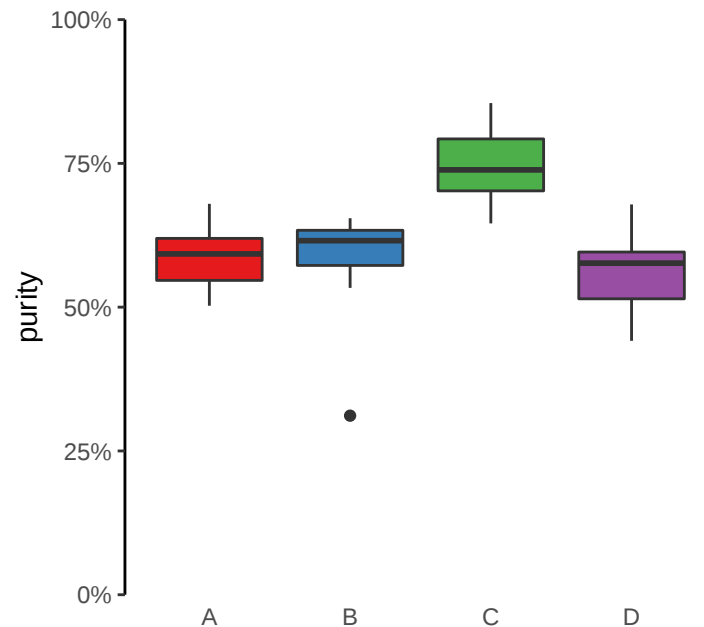


AKR1B1 promoter loss
chr7:134458962–134459462

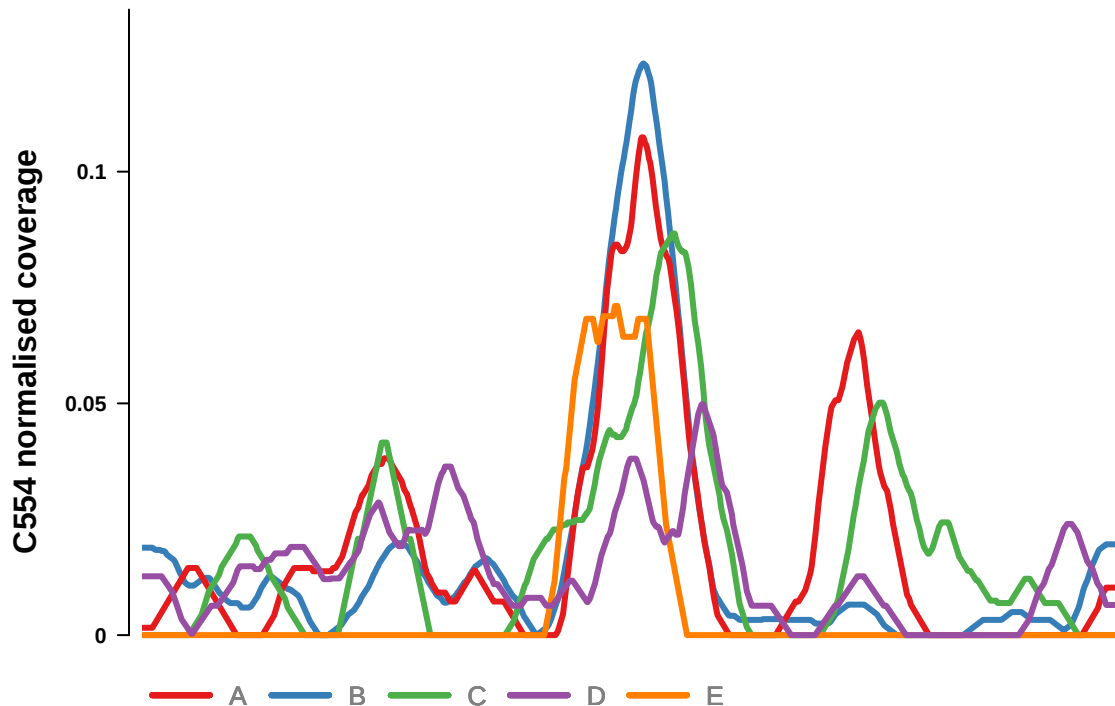


Putatively Subclonal (logFC = -0.24)

	P.	Adjusted P.
<i>~region vs. ~</i>	1.84e-04 ***	8.09e-03 **
<i>~purity vs ~</i>	7.62e-01	9.44e-01
<i>~purity+region vs ~purity</i>	6.86e-05 ****	6.03e-03 **
<i>Carcinoma vs. normal</i>	NA	NA

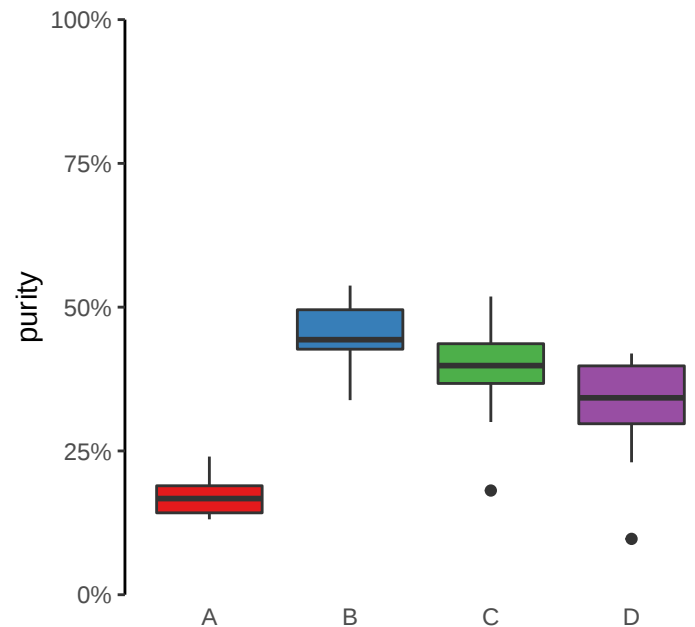


AMBRA1 promoter loss
chr11:46492426–46492926

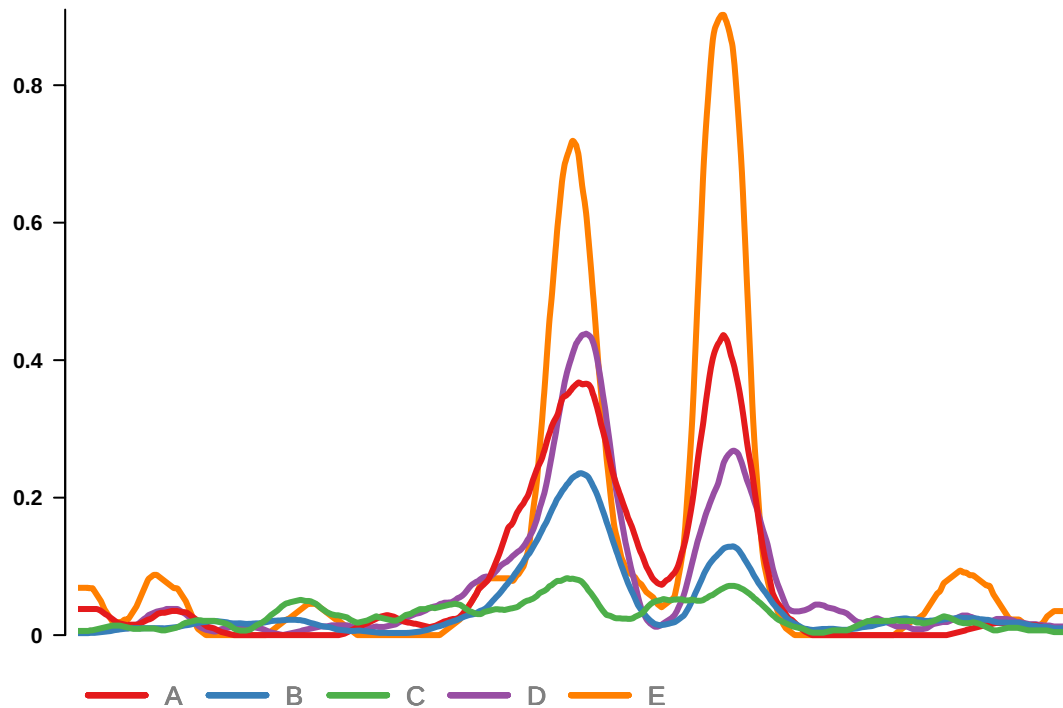


Putatively Subclonal (logFC = -0.2)

	P.	Adjusted P.
<i>~region vs. ~</i>	8.53e-04 ***	4.37e-03 **
<i>~purity vs ~</i>	7.87e-02	2.22e-01
<i>~purity+region vs ~purity</i>	5.97e-03 **	3.52e-02 *
<i>Carcinoma vs. normal</i>	NA	NA



TIAM1 promoter loss
chr21:31558930–31559430



Putatively Subclonal (logFC = -0.43)

	P.	Adjusted P.
<i>~region vs. ~</i>	2.82e-01	3.63e-01
<i>~purity vs ~</i>	8.9e-01	9.43e-01
<i>~purity+region vs ~purity</i>	1.24e-02 *	4.38e-02 *
<i>Carcinoma vs. normal</i>	NA	NA

