



Corrigendum

# RNAIndel: discovering somatic coding indels from tumor RNA-Seq data

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Bioinformatics (2019) doi: 10.1093/bioinformatics/btz753

In the above paper, Table 4 has been corrected as follows.

Original:

NSCLC (n=90)	PON	TPR				#False Positives			
		Mutect2		RNAIndel		Mutect2		RNAIndel	
		+	-	+	-	+	-	+	-
<i>s</i>		0.464	0.485	0.887	0.887	94	50	6	5
<i>m</i>		0.603	0.603	0.941	0.941	26	16	7	7

Corrected version:

NSCLC (n=90)	PON	TPR				Median #FP/Sample			
		Mutect2		RNAIndel		Mutect2		RNAIndel	
		+	-	+	-	+	-	+	-
<i>s</i>		0.464	0.485	0.887	0.887	50	94	5	6
<i>m</i>		0.603	0.603	0.941	0.941	16	26	7	7