



Supplementary Figure 2. Expression of *ARRB1* and *ARRB2* transcripts in lung ADC, lung SCC and non-tumour samples. Analysis of RNAseq (ILLUMINAHiSeq) datasets downloaded from TCGA Data Portal. Whiskers delimit the minimal and maximal values. n=515 ADC; n= 502 SCC and n= 110 non-tumoral lung tissue. Statistical analysis by one-way ANOVA with Tukey post-hoc analysis *** P ≤ 0.001 .