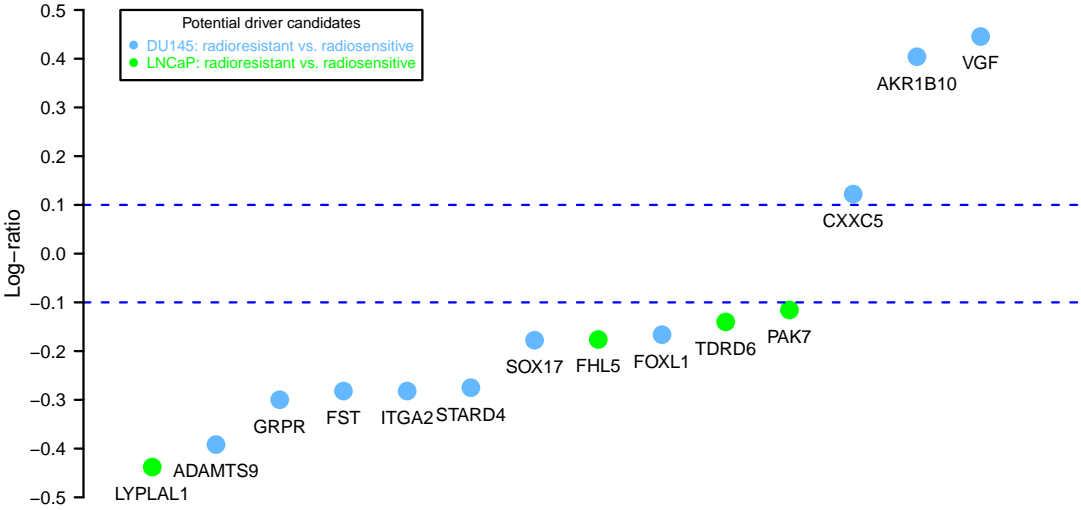


**S6 Figure: Copy number alteration levels of driver candidate genes.**



**S6 Figure:** Measured copy number alteration levels of predicted driver candidate genes with strong impact on the expression of radioresistance marker genes. Log<sub>2</sub>-ratios of copy number alteration levels comparing radioresistant to radiosensitive cell lines of DU145 or LNCaP are shown on the y-axis. The genes are sorted according to increasing log-ratios and visualized by dots that are colored according to the cell line in which a candidate gene has been predicted. The two blue dashed horizontal lines mark the log-ratio cutoff that was used to select these genes from our aCGH data.