



Figure S3. Cumulative distribution function (CDF) of the P -values of 48 nuclear receptors.

The CDF represents associations between the individual NR gene expression and survival from univariate Cox models from MDACC lung cancer cohort. Each dot represents a P -value for one NR gene and the solid red line represents the expected CDF for randomly picked genes. The dashed green line corresponds to $P=0.05$. 37 NR genes have P -values smaller than 0.05. SHP and PR genes are indicated in the plot.