

# 13<sup>th</sup> EUROPEAN PATHOLOGY CONGRESS

August 02-03, 2017 Milan, Italy

## Genomic drivers of human cancers: Diagnostics, prognostics and therapeutics

Accurate prediction of clinical courses of human cancers remains elusive. In recent studies, we performed whole genome analysis on prostate and liver cancers. Our result showed that combination of genome copy number variance and novel fusion transcripts specific for cancer achieved high accuracy in predicting clinical outcomes of these cancers. Interestingly, some of these fusion genes

### Notes: