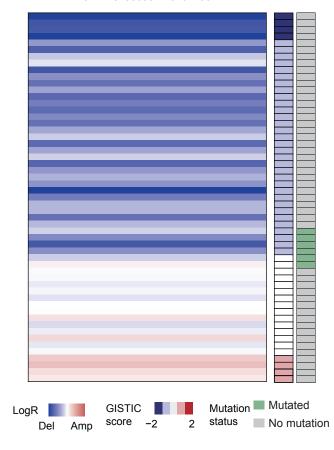
FAT1 chr4:187508937-187644987



Supplementary information, Figure S8 Recurrent *FAT1* mutations in SCCE. Left: heat map of copy number alteration for *FAT1*. Each row depicts one patient. Middle column, copy number value of each patient. Values are provided by GISTIC2.0. 2 represents high copy number gain (amplification), 1 represents lower copy number gain, -1 represents low copy number loss and -2 represents higher copy number loss (deletion). Right, somatic mutation status for each patient.