**Altered in 246 (97.23%) of 253 samples.**

TMB axis: 1772 ... 0

| Gene | Percentage |
|------|-----------|
| TP53 | 63% |
| TTN | 61% |
| CSMD3 | 50% |
| MUC16 | 49% |
| RYR2 | 43% |
| LRP1B | 43% |
| ZFHX4 | 42% |
| USH2A | 39% |
| XIRP2 | 32% |
| KRAS | 32% |
| FLG | 30% |
| SPTA1 | 30% |
| COL11A1 | 26% |
| MUC17 | 26% |
| NAV3 | 26% |
| RYR3 | 25% |
| CSMD1 | 24% |
| FAT3 | 24% |
| PCDH15 | 24% |
| ANK2 | 23% |

No. of samples: 0 ... 160

Legend:
- Missense_Mutation
- Nonsense_Mutation
- Splice_Site
- Frame_Shift_Del
- Frame_Shift_Ins
- In_Frame_Del
- Multi_Hit