Table S2. Primer sequences of HCC prognostic genes and internal reference gene beta Actin (ACTB)

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| Primer name | Sense | Anti-sense |
| PPIA | GGCAAATGCTGGACCCAACACA | TGCTGGTCTTGCCATTCCTGGA |
| ALG3 | CTTTGCTGTGCTCTACCTGGCT | CGCAGCACAAAGATGGAGTGGA |
| CTSA | GCTTCGTGAAGGAGTTCTCCCA | CTGTGGTCATCAGTATGGCTGC |
| CAD | TAGTCCTTGGCTCTGGCGTCTA | TAGTCGGTGCTGACTGTCTCTG |
| B3GAT3 | GCCCTTGCTGTTAGATAAGCCC | GCTTCTCTGTCCGAGTATGCCA |
| TRAPPC3 | CTGGTCACCCAGCTATGTAAGG | ACCTCCCAACATTTGACCGAGC |
| HSP90AA1 | TCTGCCTCTGGTGATGAGATGG | CGTTCCACAAAGGCTGAGTTAGC |
| SRD5A3 | TCTTCCAGGACCTGATCCGCTA | CAGGAAGCCATTCCACAGCACT |
| BAG2 | TAGCCAGGACATGAGGCAGATC | GCTTTAGGGATTCTTGCTGCTGG |
| DNAJC1 | GGACTCCAGGTCGATGGGAAAA | GGAGCAGGTCACTGAATCCTTC |
| ADAMTS5 | CCTGGTCCAAATGCACTTCAGC | TCGTAGGTCTGTCCTGGGAGTT |
| PLOD2 | GACAGCGTTCTCTTCGTCCTCA | CTCCAGCCTTTTCGTGGTGACT |
| DYNC1LI1 | GCAAAGCAACCACCAACTGCAG | ATGGGTGACACGCTGGCAACAT |
| TUBULIN | TTGGGAGGTCATCAGCGATGAG | AGGCTCCAGATCCACCAGGATG |
| ST6GALNAC4 | ACACCTTCACGGAGCGCATGAT | CCATAGACCACGATCTCCTCAC |
| ACTB | CACCATTGGCAATGAGCGGTTC | AGGTCTTTGCGGATGTCCACGT |