|  |  |  |
| --- | --- | --- |
| Oligo | | DNA |
| Name | | Sequence (5'-3') |
| T318N fs\*26 | F | CCA GCA CGC ACT CAA CCT CTG TGA GCA GAG |
| T318N fs\*26 | R | CTC TGC TCA CAG AGG TTG AGT GCG TGC TGG |
| S319L fs\*2 | F | CAG CAC GCA CTC ACT CTG TGA GCA GAG G |
| S319L fs\*2 | R | CCT CTG CTC ACA GAG TGA GTG CGT GCT G |
| S324F fs\*21 | F | GTG AGC AGA GGG TTC CTC CAG CCT CAA GAT C |
| S324F fs\*21 | R | GAT CTT GAG GCT GGA GGA ACC CTC TGC TCA C |
| S324P fs\*42 | F | CTC TGT GAG CAG AGG TCC AGC CTC AAG ATC |
| S324P fs\*42 | R | GAT CTT GAG GCT GGA CCT CTG CTC ACA GAG |
| R322Q fs\*22 | F | CAC CTC TGT GAG CCA GAG GGT CCA GCC |
| R322Q fs\*22 | R | GGC TGG ACC CTC TGG CTC ACA GAG GTG |
| L326Q fs\*17 | F | GCA GAG GGT CCA GCC AAG ATC CTC TCC AAA G |
| L326Q fs\*17 | R | CTT TGG AGA GGA TCT TGG CTG GAC CCT CTG C |
| K327R fs\*17 | F | GAG GGT CCA GCC TCA GAG ATC CTC TCC AAA G |
| K327R fs\*17 | R | CTT TGG AGA GGA TCT CTG AGG CTG GAC CCT C |
| V340L fs\*27 | F | GGT GGA CAT TCA TCT CTG TTT CCA CTG AGT C |
| V340L fs\*27 | R | GAC TCA GTG GAA ACA GAG ATG AAT GTC CAC C |
| S346\* | F | CTG AGT CTG AGT AAG TTT TCA CTC C |
| S346\* | R | GGA GTG AAA ACT TAC TCA GAC TCA G |
| S339L\*27 | F | GAGGTGGACATTCTCTGTTTCCACTG |
| S339L\*27 | R | CAGTGGAAACAGAGAATGTCCACCTC |