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| **genomic variant** | **Primers sequence** \* |
| NG\_005905.2: g.163181\_169408del6228 (exon 21-22 del) | F: TGCAAGATTTCACACTTACCTGT  R: ATGGAAGCCATTGTCCTCTG |
| NG\_005905.2: g.160396\_164568del4173 (exon 20 del) | F: GGTCTCAAGCTGTCCTCCTG  R: GTGCCACTCTCACTACCTGT |
| NG\_005905.2:g.145185\_151339del6155 (exon 16-17 del) | F: GGTTGTTGATGTGGAGC  R: AACCTCAGCCTCCAGAGTTC |

\* F: forward strand; R: reverse strand