Strain: VH7209

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| Genotype: | taap-1(hd7209[LoxP + myo-2p::GFP::unc-54 3' UTR + rps-27p::neoR::unc-54 3' UTR + LoxP]) II. |
| Type: | Homozygous |
| QC Results: | PPPP |
| Deletion: | 1080 bp |
| Flanks: | TTTTTAATTTGAAGTTTTCCATTAATTCCA - [deletion] - GGGTTTTTCTTCGCAAGATTCACGTTTCGG |
| DNA Files: | taap-1\_Deletion.ape, taap-1\_Wildtype.ape |

**Primer location and sequences**

Primer 1-4 coordinates are from the Deletion ApE file. Primer 5 coordinates are from the Wildtype ApE file.

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| --- | --- | --- | --- |
| **No** | **Position** | **Name** | **Sequence (5’ to 3’)** |
| 1 | 1495-1516 | taap-1-1 SV F | AGAATGACTCCTGGATTGCTCG |
| 2 | 3060-3083 | pMYO-2 SEC Insertion Primer 2 | CAGAACAATGAGTACAGTGTGTGC |
| 3 | 6595-6616 | NeoR SEC Insertion Primer 2 | ATCTCCTGTCATCTCACCTTGC |
| 4 | 9362-9385 | taap-1-1 SV R | GCCATCCTAATATGGACTTCCAGG |
| 5 | 2667-2687 | taap-1-1 WT R | ACCAACATCGTGTGACCTACG |

**PCR products**

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| --- | --- | --- | --- |
| **Product (see figure)** | **Primer** | **Product Size** | **QC Result** |
| {US} - Upstream insertion site | 1+2 | 1589 bp | Pass (Band present) |
| {DS} - Downstream insertion site | 3+4 | 2791 bp | Pass (Band present) |
| {WoM} – WT PCR on mutant | 1+5 | no product, if WT sequence is deleted | Pass (Band absent) |
| {WoM} – WT PCR on wildtype | 1+5 | 1193 bp | Pass (Band present) |

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| **PCR Program:**  (1) 94˚C - 30 sec  (2) 94˚C - 30 sec  (3) 60˚C - 30 sec  (4) 72˚C - 3 min  (5) To step (2) x 34 times  (6) 72˚C - 5 min  (7) 4˚C - hold | A diagram of a gene deletion  Description automatically generated |