|  |  |  |  |  |
| --- | --- | --- | --- | --- |
| **Target species** | **Gene** | **Primer** | **Sequence 5’-3’** | **Amplicon**  **size (bp)** |
| ***E. faecalis*** | *acuI* | Ecfaeca\_acuI-F | TCATTTCAAGCATTTACGTTAAGAGA | 73 |
| Ecfaeca\_acuI-R | CGTCTAAAGTAATGGTTTCTAGTTGA |
| g3060 | Ecfaeca\_g3060-F | CCCTCTTTAACATTAATTGGACTTGC | 74 |
| Ecfaeca\_g3060-R | TGATTTCTTAATTTATAGGCTTGGCT |
| ***E. faecium*** | *cysS* | Ecfaeci\_cysS-F | GCAGCCACCAATTTACAACGA | 86 |
| Ecfaeci\_cysS-R | TCATCTGCCAAATTCTCTGAGG |
| *purD* | Ecfaeci\_purD-F | GGAATGAAAGAACTAGGACGCT | 71 |
| Ecfaeci\_purD-R | GGTCCTTCCTTCGTAGCAAT |
| ***P. acidilactici*** | *asnS* | Pdacidi\_asnS-F | GCCATCCACGAATTCCTCCA | 80 |
| Pdacidi\_asnS-R | GCACCTTCCGTATCACTAGAAGT |
| g1164 | Pdacidi\_g1164-F | TTTAGGAGCAATCATCGCAATG | 75 |
| Pdacidi\_g1164-R | CCAACTAATACCGCGCTACC |
| ***P. pentosaceus*** | *nagK* | Pdpento\_nagK-F | GGGTGTCACTAACGGAGCAA | 107 |
| Pdpento\_nagK-R | CAATTCCGCCCGCTTGTAAG |
| g4364 | Pdpento\_g4364-F | TGCTTACCACCAACCTGCTT | 76 |
| Pdpento\_g4364-R | TGGACGTTCTAGGTTTCCGT |