|  |  |  |  |  |
| --- | --- | --- | --- | --- |
| **Supplementary Table 2.** | | |  |  |
|  | | | | |
| **SNPID** | **Scaffold** | **Position** | **Major Allele** | **Minor Allele** |
| G01 | GL429767 | 16839644 | T | A |
| G02 | GL429767 | 22649683 | T | A |
| G03 | GL429767 | 46534125 | C | T |
| G04 | GL429768 | 3921142 | G | C |
| G05 | GL429768 | 29962249 | A | G |
| G06 | GL429768 | 9041952 | A | G |
| G07 | GL429770 | 8507307 | A | C |
| G08 | GL429772 | 14993622 | T | C |
| G09 | GL429773 | 7204887 | C | T |
| G10 | GL429775 | 5556309 | A | C |
| G11 | GL429775 | 17831035 | C | T |
| G12 | GL429778 | 3561000 | G | A |
| G13 | GL429781 | 7594948 | A | T |
| G14 | GL429783 | 665407 | C | T |
| G15 | GL429792 | 753145 | C | A |
| G16 | GL429801 | 6986210 | A | T |
| G17 | GL429802 | 4091453 | C | T |
| G18 | GL429806 | 3311579 | T | C |
| G19 | GL429816 | 3229647 | G | A |
| G20 | GL429819 | 365354 | G | A |
| G21 | GL429835 | 2966433 | A | G |
| G22 | GL429841 | 3522450 | A | G |
| G23 | GL429841 | 612532 | G | A |
| G24 | GL429842 | 4124241 | A | G |
| G25 | GL429845 | 2091046 | C | T |
| G26 | GL429855 | 4401152 | G | C |
| G27 | GL429859 | 300331 | T | C |
| G28 | GL429861 | 293372 | A | G |
| G29 | GL429861 | 3065399 | A | G |
| G30 | GL429867 | 2575249 | A | G |
| G31 | GL429873 | 2056629 | T | A |
| G32 | GL429879 | 384336 | G | A |
| G33 | GL429885 | 3445677 | T | A |
| G34 | GL429885 | 617822 | C | T |
| G35 | GL429885 | 2043530 | T | C |
| G36 | GL429888 | 3508435 | A | T |
| G37 | GL429898 | 676714 | T | C |
| G38 | GL429910 | 2414078 | A | C |
| G39 | GL429927 | 1241213 | A | G |
| G40 | GL429929 | 2943933 | G | T |
| G41 | GL429930 | 2586279 | C | T |
| G42 | GL429955 | 1484586 | G | A |
| G43 | GL429962 | 689293 | A | C |
| G44 | GL429965 | 1956978 | G | A |
| G45 | GL429998 | 99324 | C | G |
| G46 | GL430008 | 1396110 | C | T |
| G47 | GL430018 | 180278 | T | A |
| G48 | GL430029 | 1661413 | A | C |
| G49 | GL430036 | 626424 | G | T |
| G50 | GL430101 | 496908 | T | C |
| G51 | GL430121 | 936584 | T | C |
| G52 | GL430156 | 916661 | C | T |
| G53 | GL430158 | 593841 | T | G |
| G54 | GL430166 | 11932 | C | T |
| G55 | GL430239 | 60325 | G | A |
| G56 | GL430239 | 60326 | T | C |
| G57 | GL430271 | 525210 | C | T |
| G58 | GL430281 | 514067 | T | C |
| G59 | GL430338 | 350547 | T | C |
| G60 | GL430482 | 173352 | C | T |
| G61 | GL430686 | 91034 | C | T |
| G62 | GL430746 | 9663 | C | T |
| G63 | GL432768 | 965 | C | T |

**Supplementary Table 2.**

SNPs that were identified by Gignoux-Wolfsohn et al (2021) as putatively under selection due to white-nose syndrome and that were included in the target SNP eQTL analysis. Scaffold and position refer to the Myoluc 2.0 genome publicly available from ENSEMBL and NCBI. The red colored allele of the major and minor alleles is the one found to be positively selected in Gignoux-Wolfsohn et al (2021).