**Figure 3-source data 6. *DOG1* PCR primers used in the complementation experiment.** We designed an Eden-2 specific forward primer because of structural variants at the 3’ end of *DOG1*. The underlined sequences correspond to the flanking sequences of the EcoRI restriction site in the pGreenII 0029 that was used for transformation.

|  |  |  |  |
| --- | --- | --- | --- |
| **ID** | **Name** | **Forward** | **Reverse** |
| 992 | Ale-Stenar-44-4 | CGGGCTGCAGGAATTGGGGCATCTAGAATTGTGTCAT | GCTTGATATCGAATTTGACTTCGGCTTCTTTTTAACC |
| 1002 | Ale-Stenar-64-24 | CGGGCTGCAGGAATTGGGGCATCTAGAATTGTGTCAT | GCTTGATATCGAATTTGACTTCGGCTTCTTTTTAACC |
| 6913 | Eden-2 | CGGGCTGCAGGAATTGCTTACTAATCGCGATCCAAAC | GCTTGATATCGAATTTGACTTCGGCTTCTTTTTAACC |