**Table 2. Mean and median scaffold *F*ST between morphologically defined taxonomy (Figure 1) and genomic clusters (Figure 2).**

|  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- |
| **Morphological taxa** | | | | | | | |
|  | *M. suaveolens* | *M. spicata* | | *M.* x *rotundifolia* | | | |
| *M. longifolia* | 0.141/0.116 | 0.043/0.029 | | 0.039/0.024 | | | |
| *M. suaveolens* | - | 0.104/0.082 | | 0.074/0.049 | | | |
| *M. spicata* |  | - | | 0.037/0.023 | | | |
| *M.* x *rotundifolia* |  |  | | - | | | |
| **Genomic clusters** | | | | | | | |
|  | sua | spi1 | spi2 | rot1 | rot2 | rot3 | rot4 |
| long | 0.098/0.070 | 0.092/0.059 | 0.039/0.021 | 0.118/0.076 | 0.076/0.025 | 0.035/0.017 | 0.080/0.014 |
| sua | - | 0.117/0.086 | 0.100/0.075 | 0.103/0.056 | 0.087/0.042 | 0.089/0.062 | 0.142/0.096 |
| spi1 |  | - | 0.086/0.057 | 0.146/0.111 | 0.102/0.065 | 0.084/0.056 | 0.109/0.065 |
| spi2 |  |  | - | 0.119/0.079 | 0.076/0.038 | 0.046/0.028 | 0.071/0.019 |
| rot1 |  |  |  | - | 0.100/0.056 | 0.094/0.057 | 0.153/0.115 |
| rot2 |  |  |  |  | - | 0.056/0.021 | 0.095/0.036 |
| rot3 |  |  |  |  |  | - | 0.082/0.033 |
| rot4 |  |  |  |  |  |  | - |