Source Data for Figure 1—figure supplement 2F

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| Genotype | Marker | Number of values | Mean (number of cells) | Std. Deviation | Std. Error of Mean | p value |
| Control | Kr+ Zfh2- | 31 | 2.000 | 0.000 | 0.000 | NA |
| NB3-1>Hb | Kr+ Zfh2- | 60 | 5.633 | 1.461 | 0.1886 | <0.0001\* |
|  |  |  |  |  |  |  |
| Control | Kr+ Zfh2+ | 31 | 1.000 | 0.000 | 0.000 | NA |
| NB3-1>Hb | Kr+ Zfh2+ | 90 | 0.1167 | 0.3237 | 0.04179 | <0.0001\* |
|  |  |  |  |  |  |  |
| Control | Hb- Cut+ | 31 | 1.000 | 0.000 | 0.000 | NA |
| NB3-1>Hb | Hb- Cut+ | 60 | 0.111 | 0.3160 | 0.03331 | <0.0001\* |

\* t test, with Welch’s correction (for un-equal Std. Deviation)