Table 1:

Distance-based analysis of hybrid effects on predicted weight loss similarity (all mice: n = 335)

| Predictor Variable | Coefficient | 95% CI | P-value | *1* |
| --- | --- | --- | --- | --- |
| Subspecies genetic distance | 0.0068 | (-0.0098, 0.0234) | 0.421 |  |
| **Hybridization distance (hHe)** | **0.0288** | **(0.0026, 0.0550)** | **0.031** | **\*** |
| **Mean hybridization level** | **-0.0292** | **(-0.0459, -0.0125)** | **< 0.001** | **\*\*\*** |
| Sex difference | 0.0061 | (-0.0002, 0.0123) | 0.059 | . |
| Subspecies genetic distance | -0.0409 | (-0.0916, 0.0099) | 0.114 |  |
| **Infection difference effect** | **-0.0141** | **(-0.0204, -0.0078)** | **< 0.001** | **\*\*\*** |
| *1*Significance: \*\*\* p < 0.001, \*\* p < 0.01, \* p < 0.05, . p < 0.1 | | | | |