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 | 0.0028 | (-0.0135, 0.0192) | 0.734 |  |
| Hybridization distance (hHe) | 0.0060 | (-0.0199, 0.0320) | 0.649 |  |
| Mean hybridization level | -0.0117 | (-0.0279, 0.0046) | 0.158 |  |
| Sex difference | 0.0030 | (-0.0032, 0.0092) | 0.344 |  |
| Subspecies genetic distance | -0.0184 | (-0.0686, 0.0317) | 0.471 |  |
| **Infection difference effect** | **-0.0205** | **(-0.0268, -0.0143)** | **< 0.001** | **\*\*\*** |
| *1*Significance: \*\*\* p < 0.001, \*\* p < 0.01, \* p < 0.05, . p < 0.1 | | | | |