Table 1:

Testing infection-dependent hybrid effects in infected mice only (n = 133)

| Predictor Variable | Coefficient | 95% CI | P-value | *1* |
| --- | --- | --- | --- | --- |
| **Subspecies genetic distance** | **0.0233** | **(0.0059, 0.0407)** | **0.009** | **\*\*** |
| **Hybridization distance (hHe)** | **0.0670** | **(0.0397, 0.0943)** | **< 0.001** | **\*\*\*** |
| **Mean hybridization level** | **0.0557** | **(0.0373, 0.0740)** | **< 0.001** | **\*\*\*** |
| Sex difference | -0.0014 | (-0.0078, 0.0049) | 0.654 |  |
| **Subspecies genetic distance** | **-0.0877** | **(-0.1428, -0.0326)** | **0.002** | **\*\*** |
| *1*Significance: \*\*\* p < 0.001, \*\* p < 0.01, \* p < 0.05, . p < 0.1 | | | | |