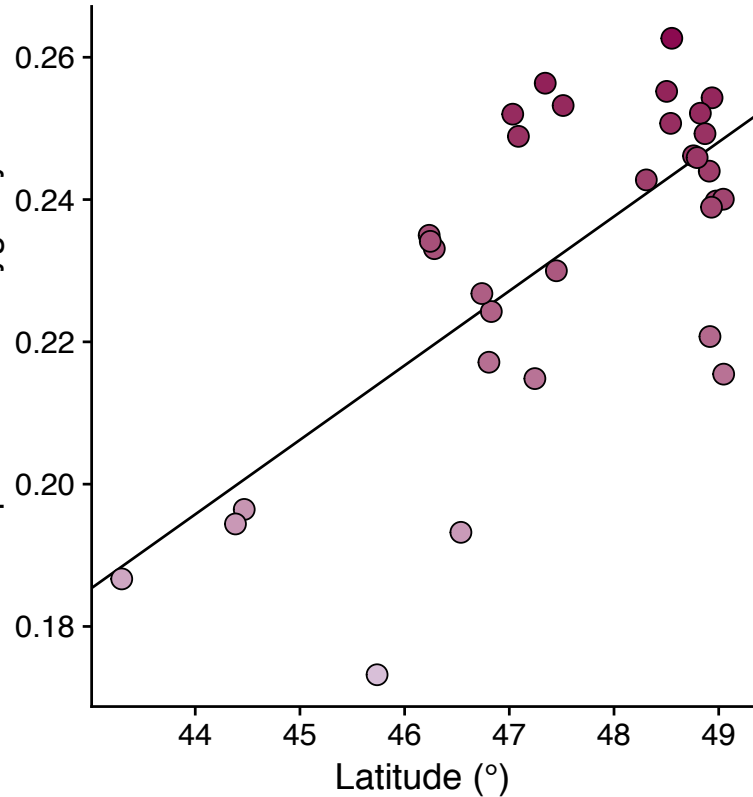
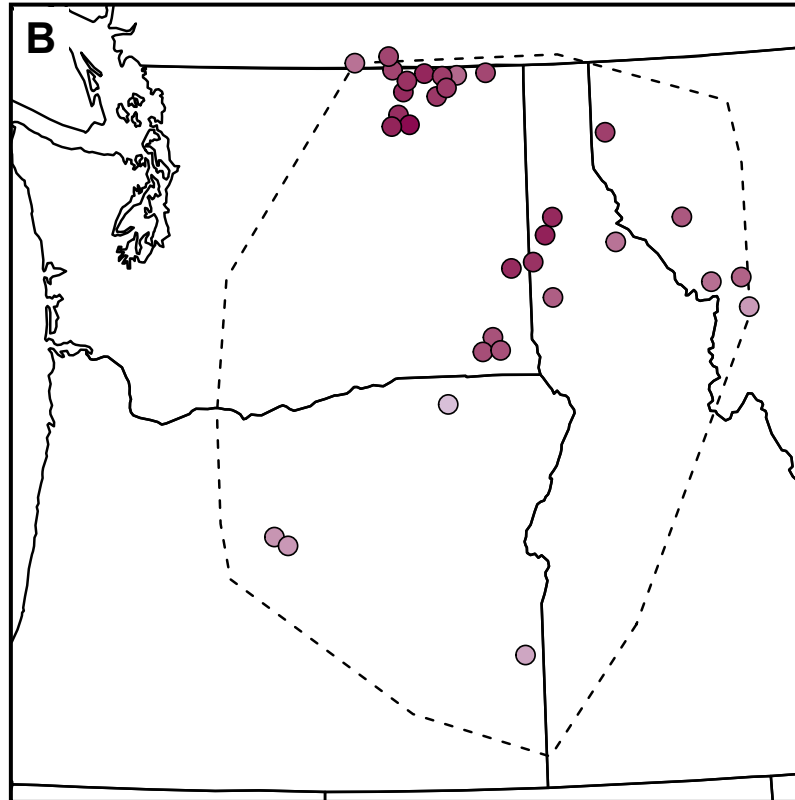


A

Expected heterozygosity

**B**

Expected heterozygosity

0.26

0.24

0.22

0.20

0.18