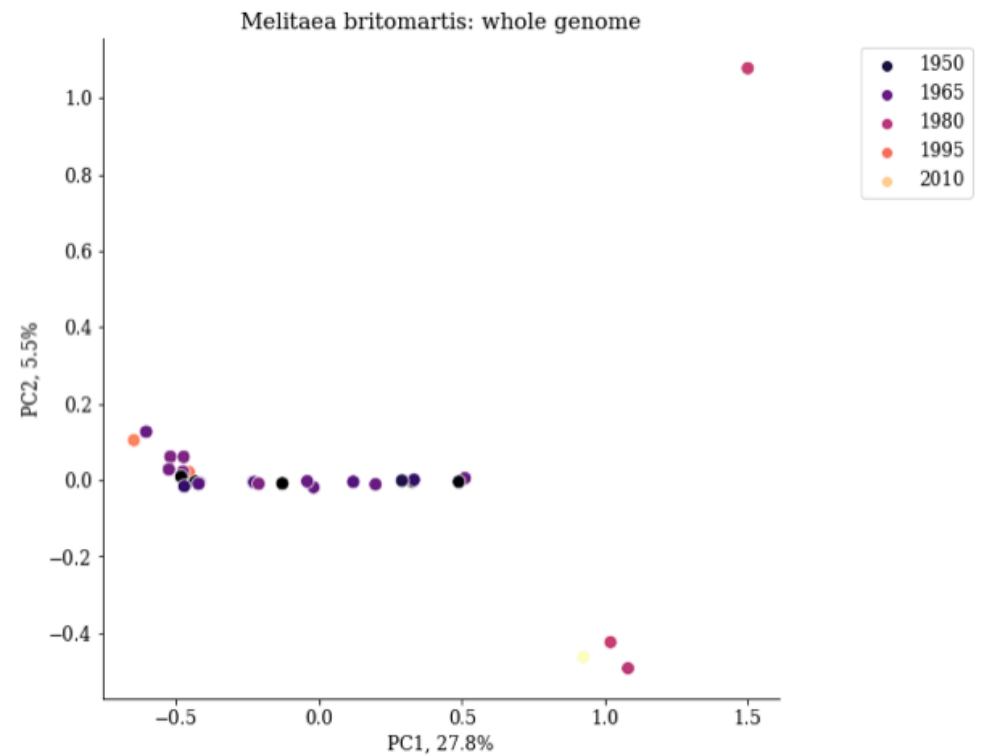
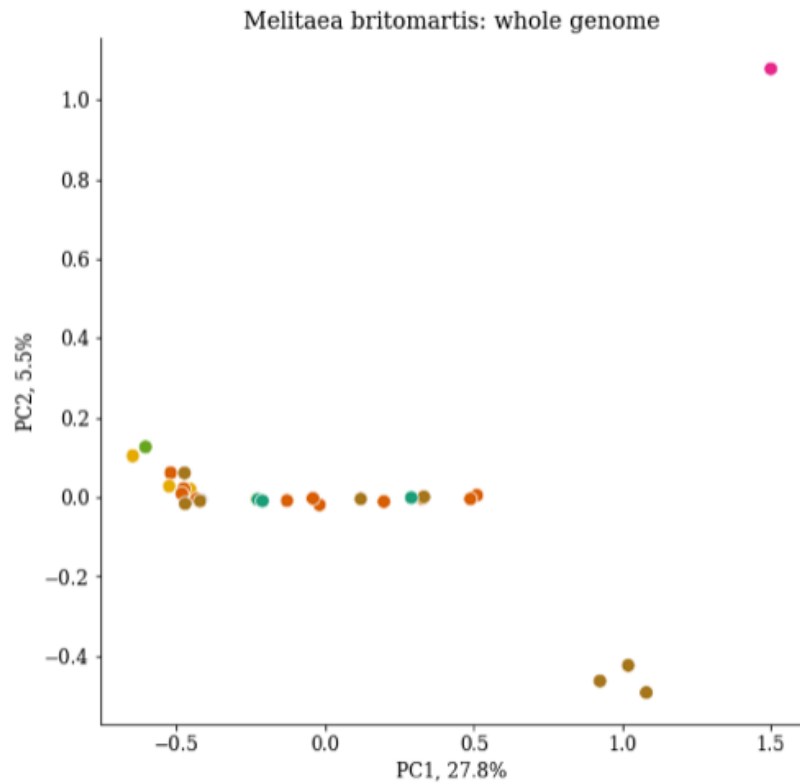


Swedish *Melitaea britomartis* population structure

ANGSD approach



Swedish *Melitaea britomartis* population structure

ANGSD approach

Filters:

MAF > 0.05

Minimum mapping quality 20 -

Base quality score 20

5 < Depth < 30

Minimum N of individuals = 10

Stats:

Wateson's theta (mean over per chromosome estimate) = 39.16

Pairwise theta (mean over per chromosome estimate) = 35.08

Number of sites = 5342

pi (from Waterson) = 0.28

pi (from pairwise) = 0.25

Tajimas'D = -0.35

***Melitaea britomartis* population structure (autosomes)**

ANGSD approach

