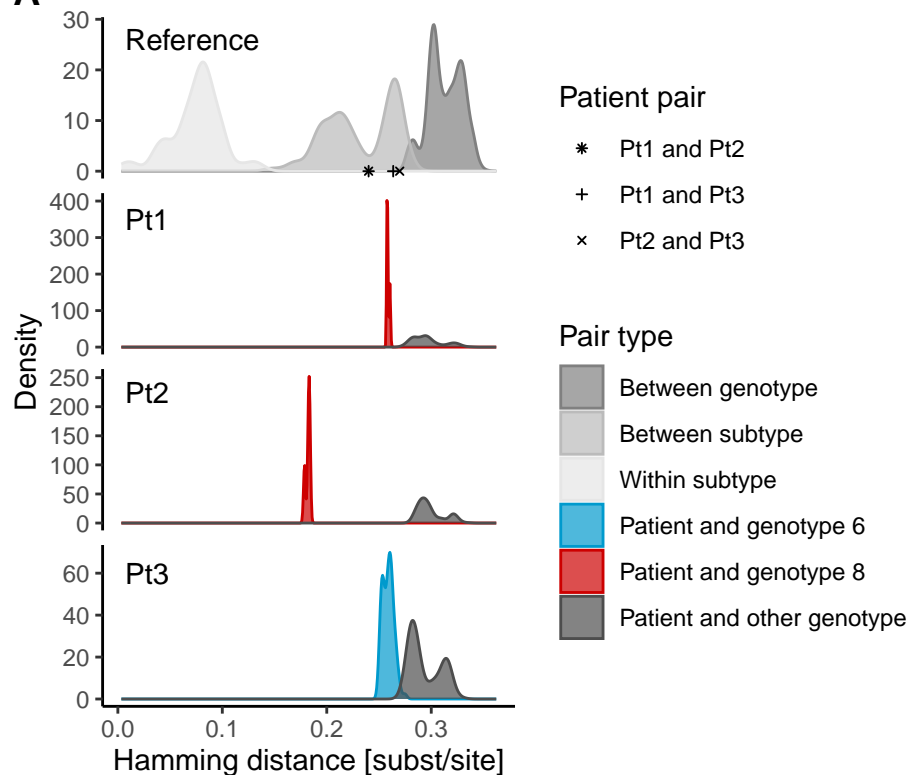
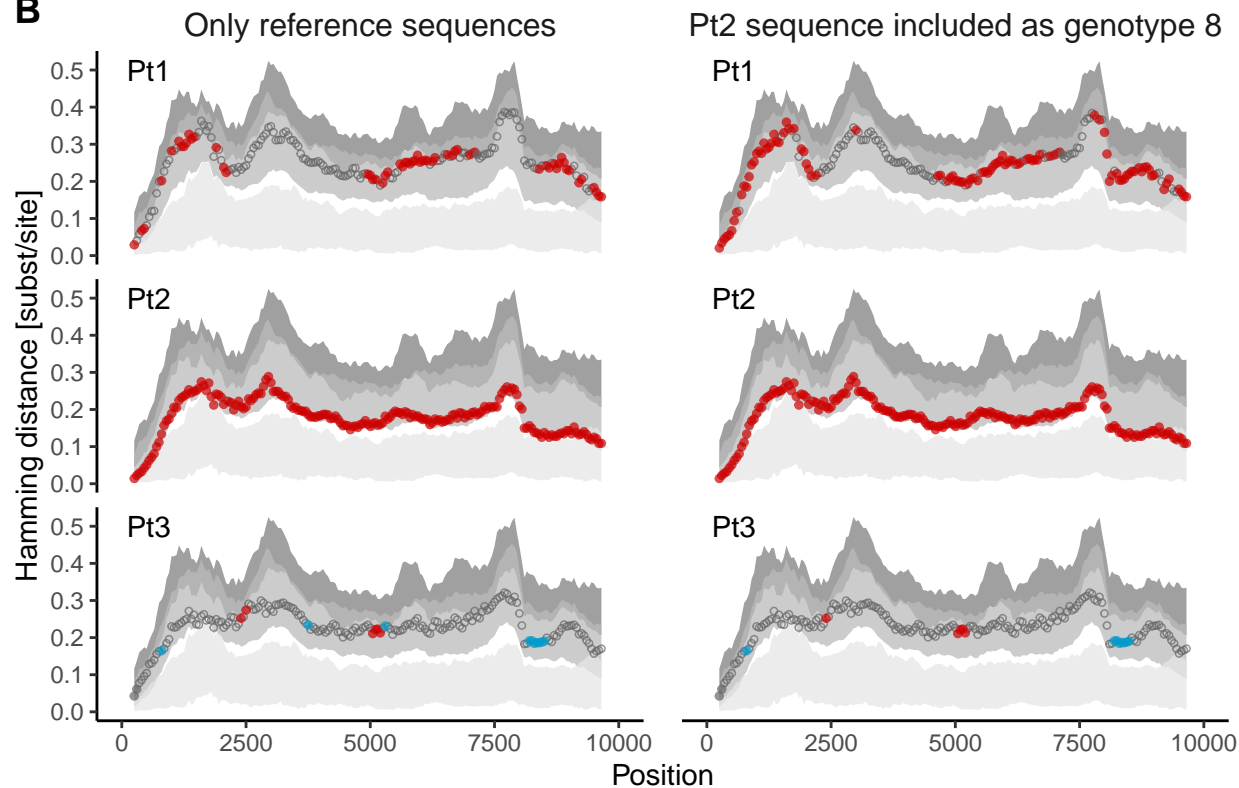


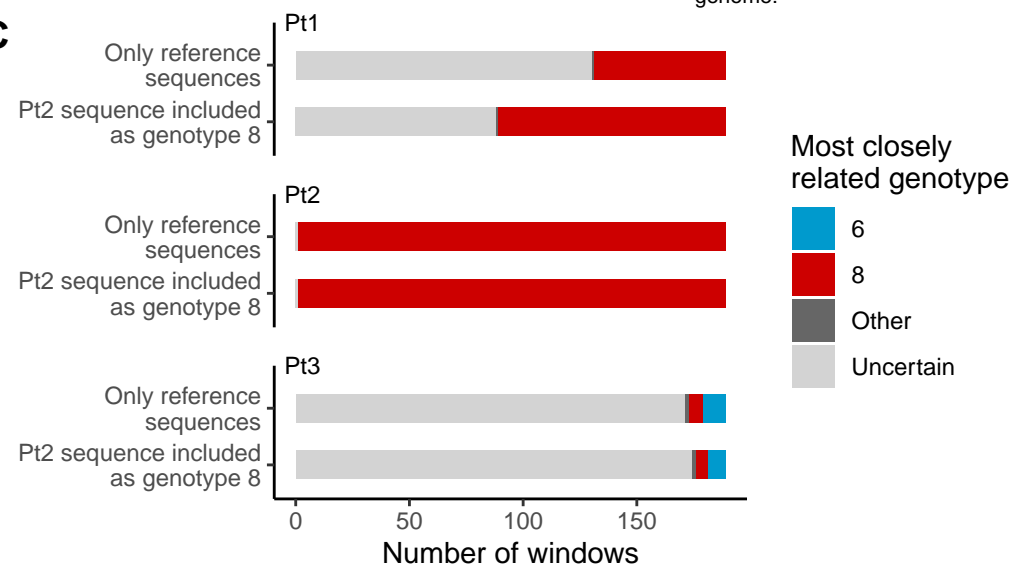
A**B**

Most closely related genotype

- 6
- 8
- Other
- Uncertain

95% CI

- Between genotype
- Between subtype
- Within subtype

C**D**