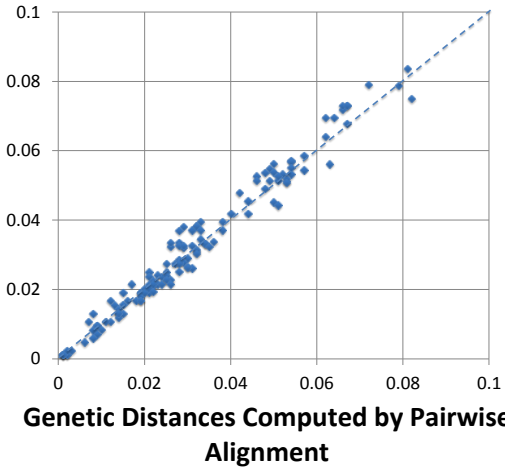


A
Genetic Distances Computed by
Multiple Sequence Alignment



B

