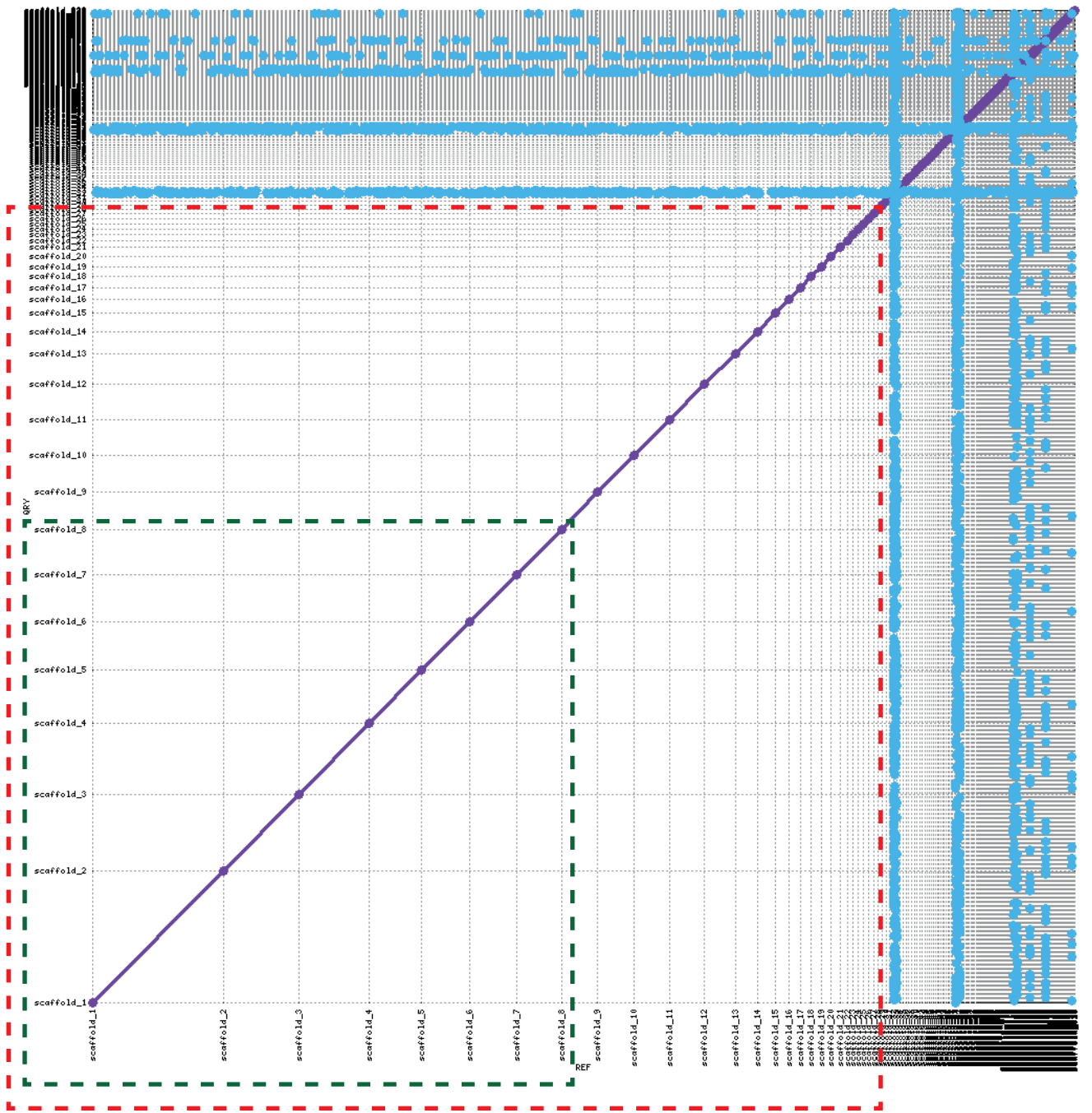


*P. dumerilii* self-self alignment map



- 50% of assembly in 8 scaffolds
- 80% of assembly in 28 scaffolds

● Probable repeat elements/centromeric and/or telomeric genomic sequences