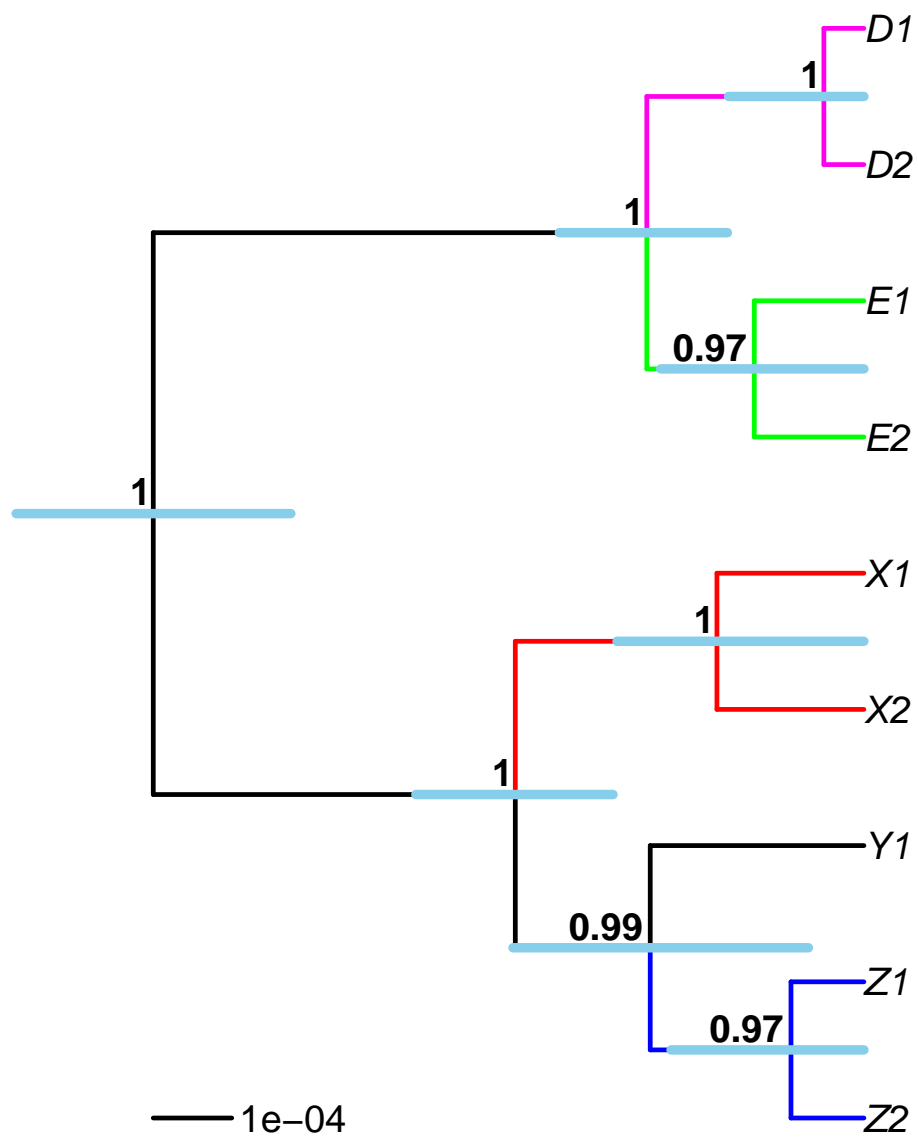
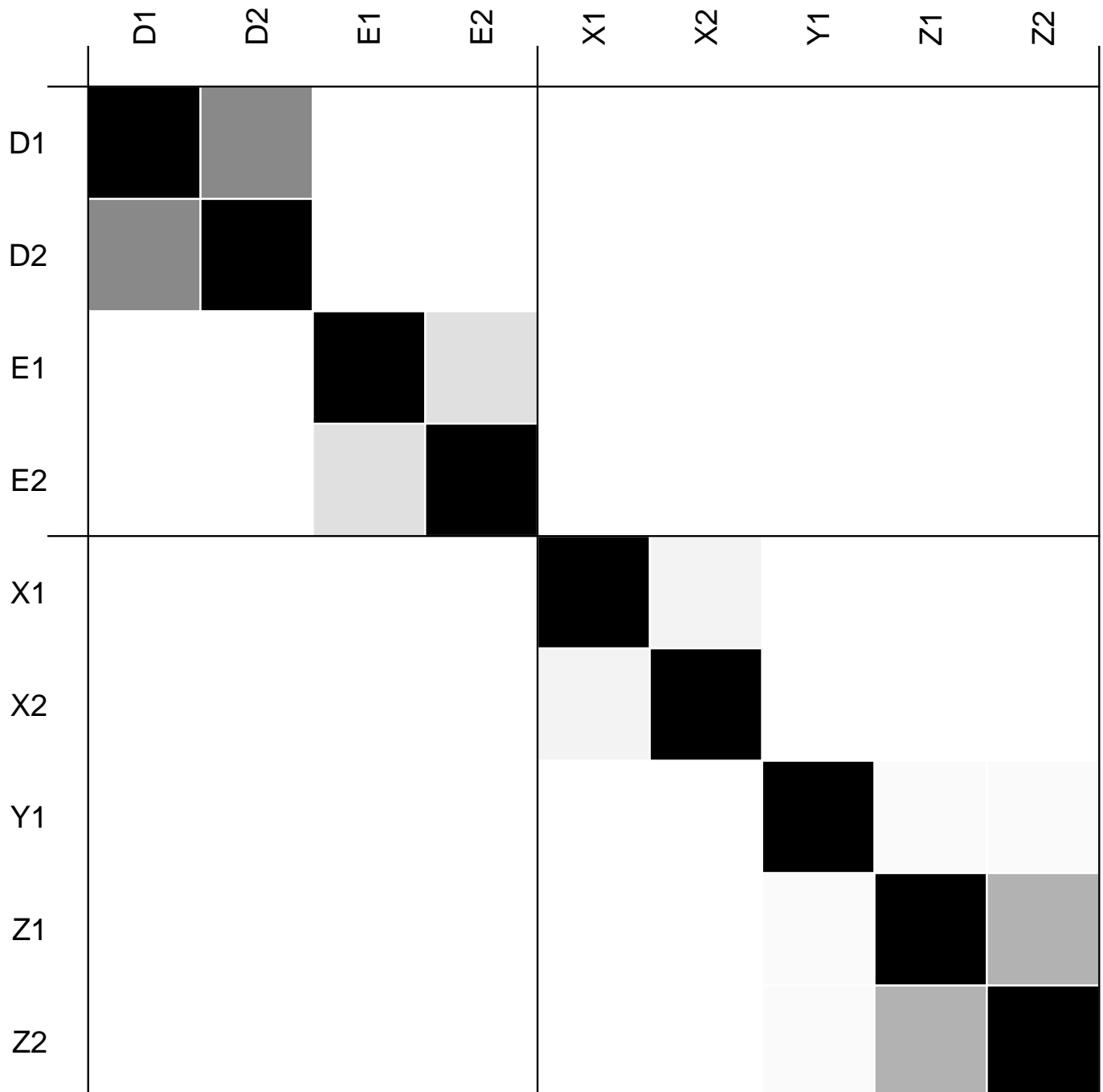


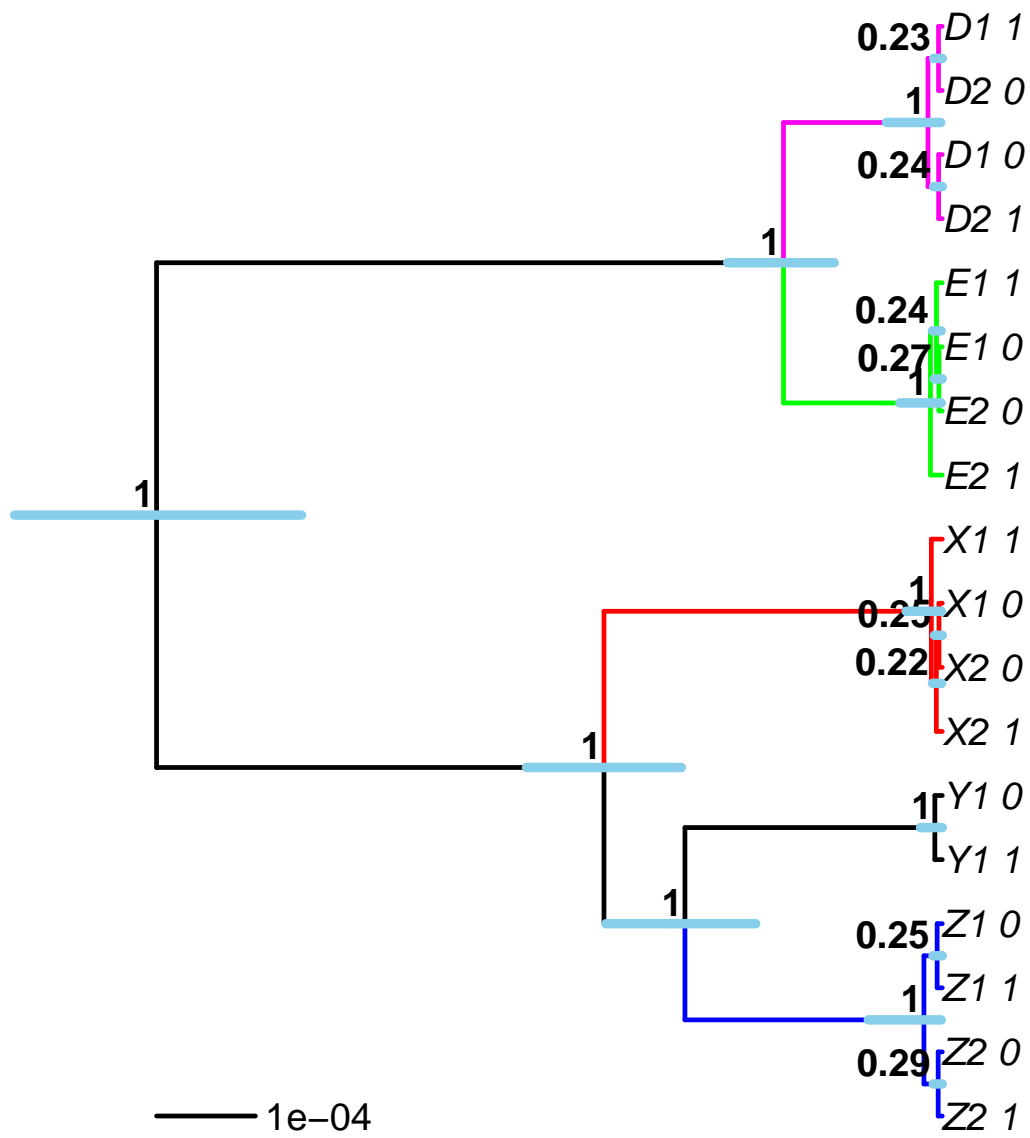
Sim-rep 1: Contig sequence alignments (n=150), STACEY species tree



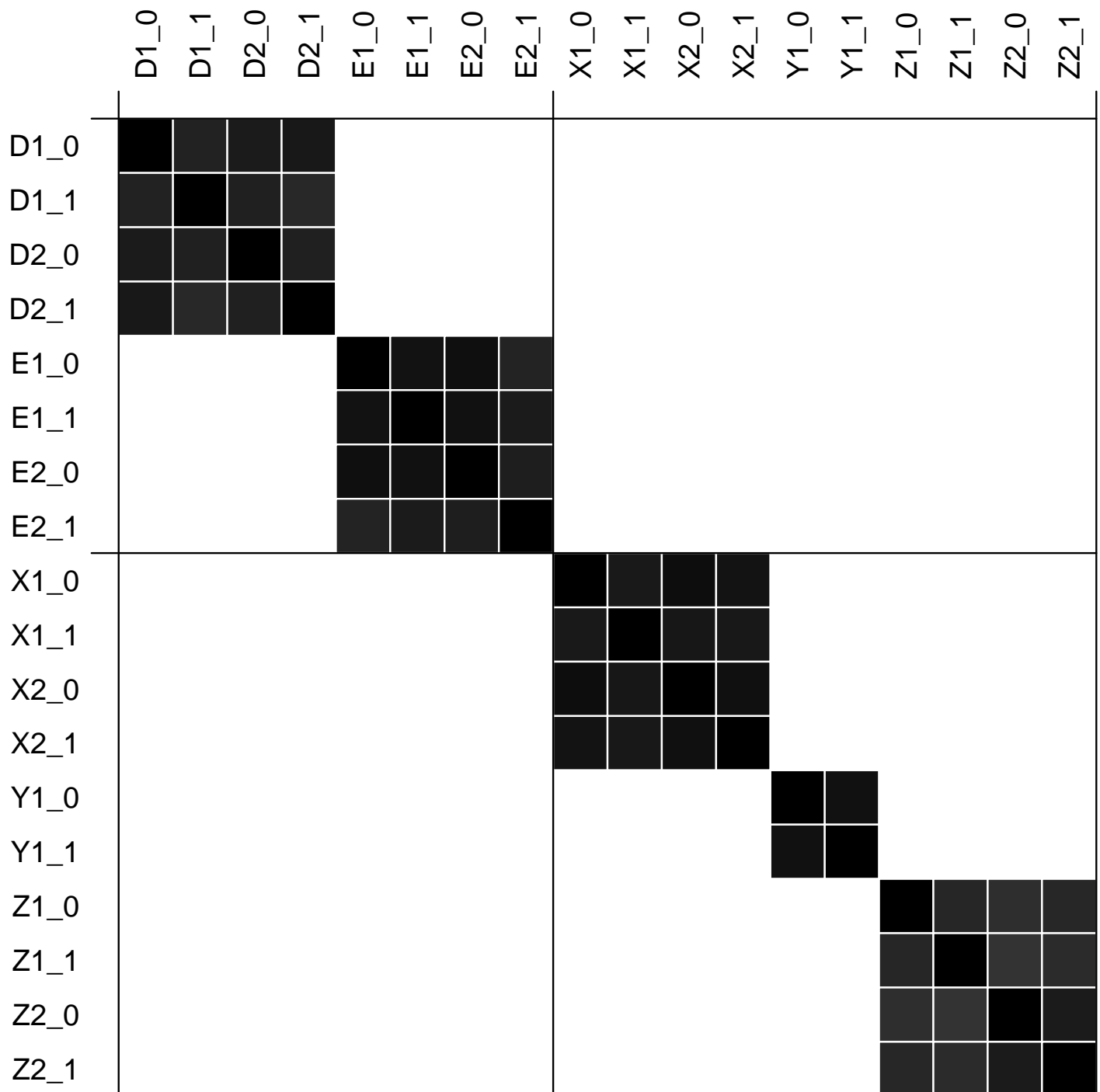
Sim-rep 1: Contig sequence alignments (n=150), STACEY similarity matrix



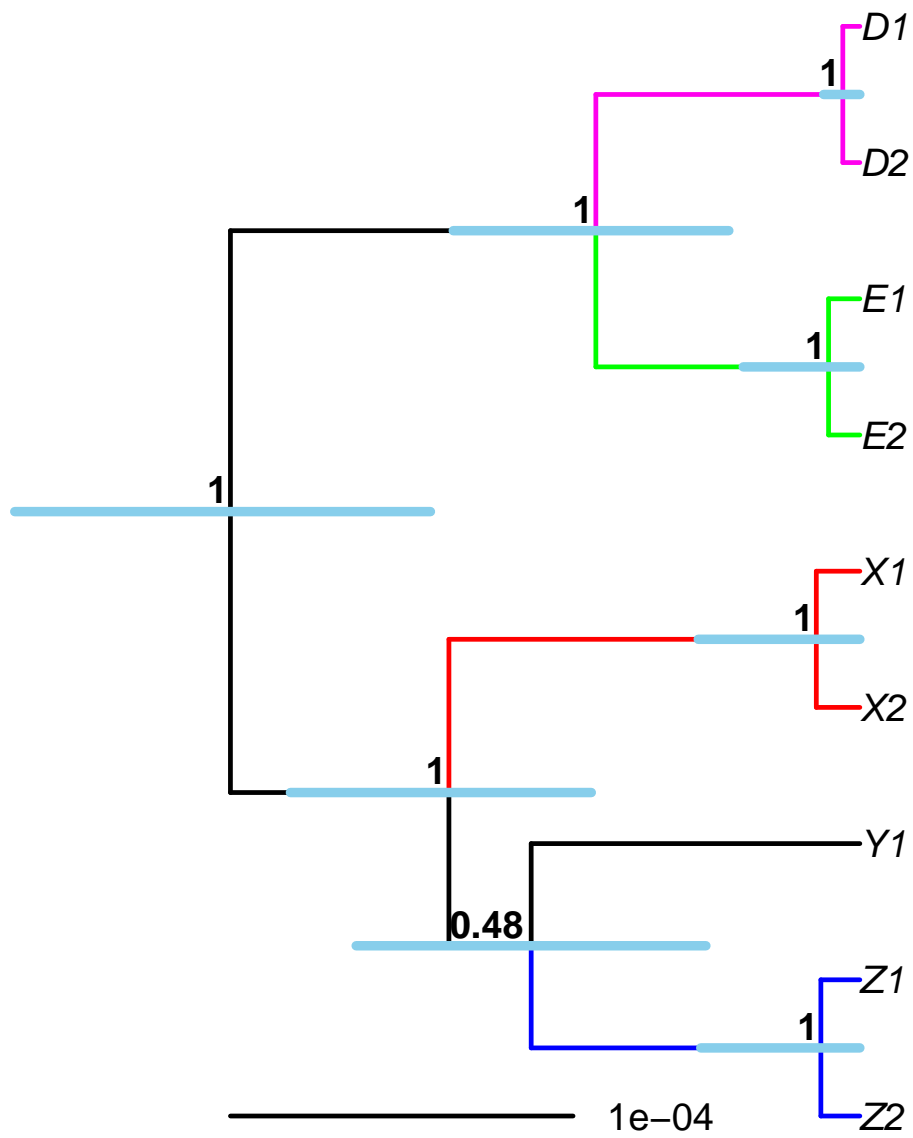
Sim-rep 1: Phased allele alignments (n=150), STACEY species tree



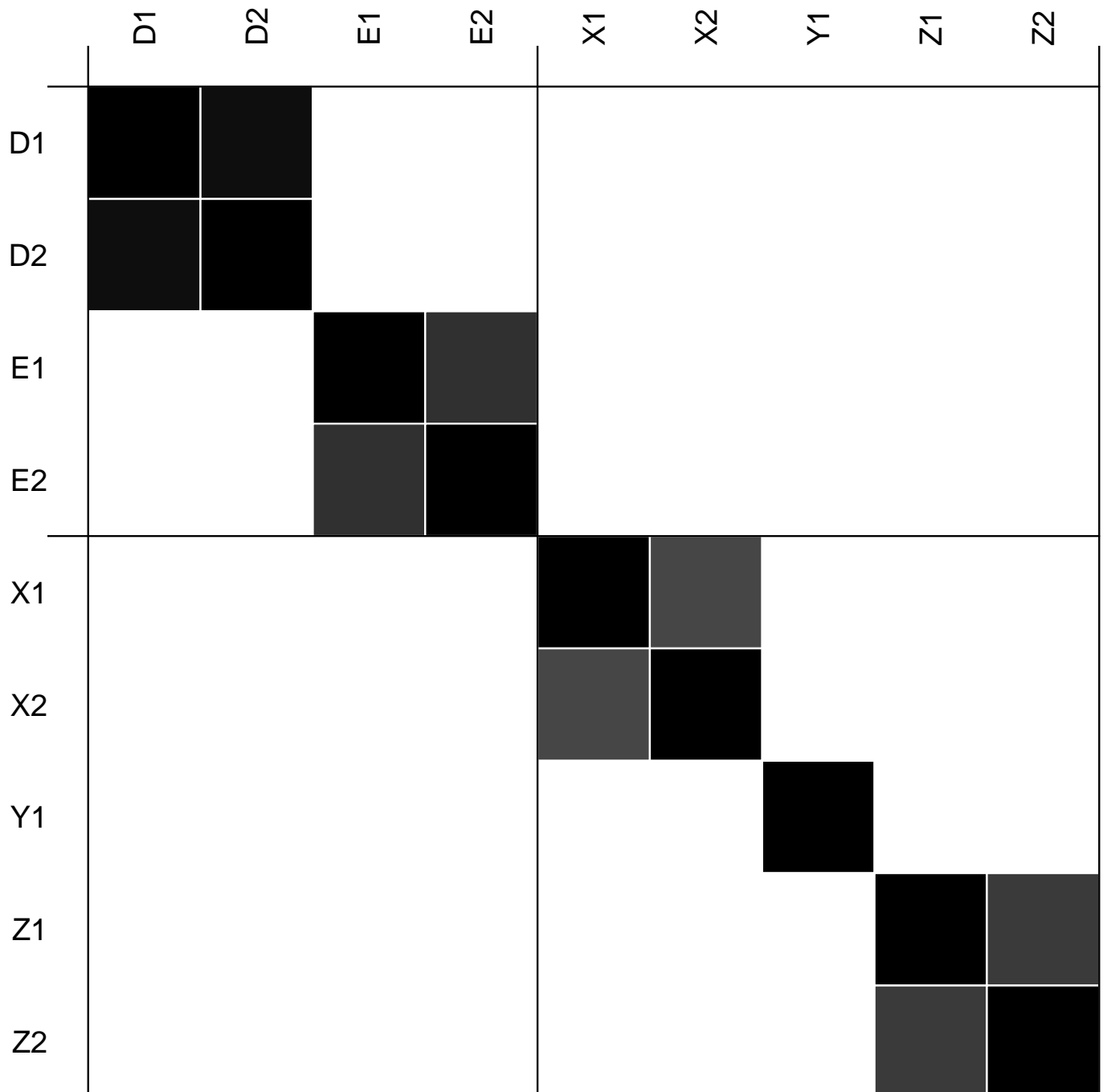
Sim-rep 1: Phased allele alignments (n=150), STACEY similarity matrix



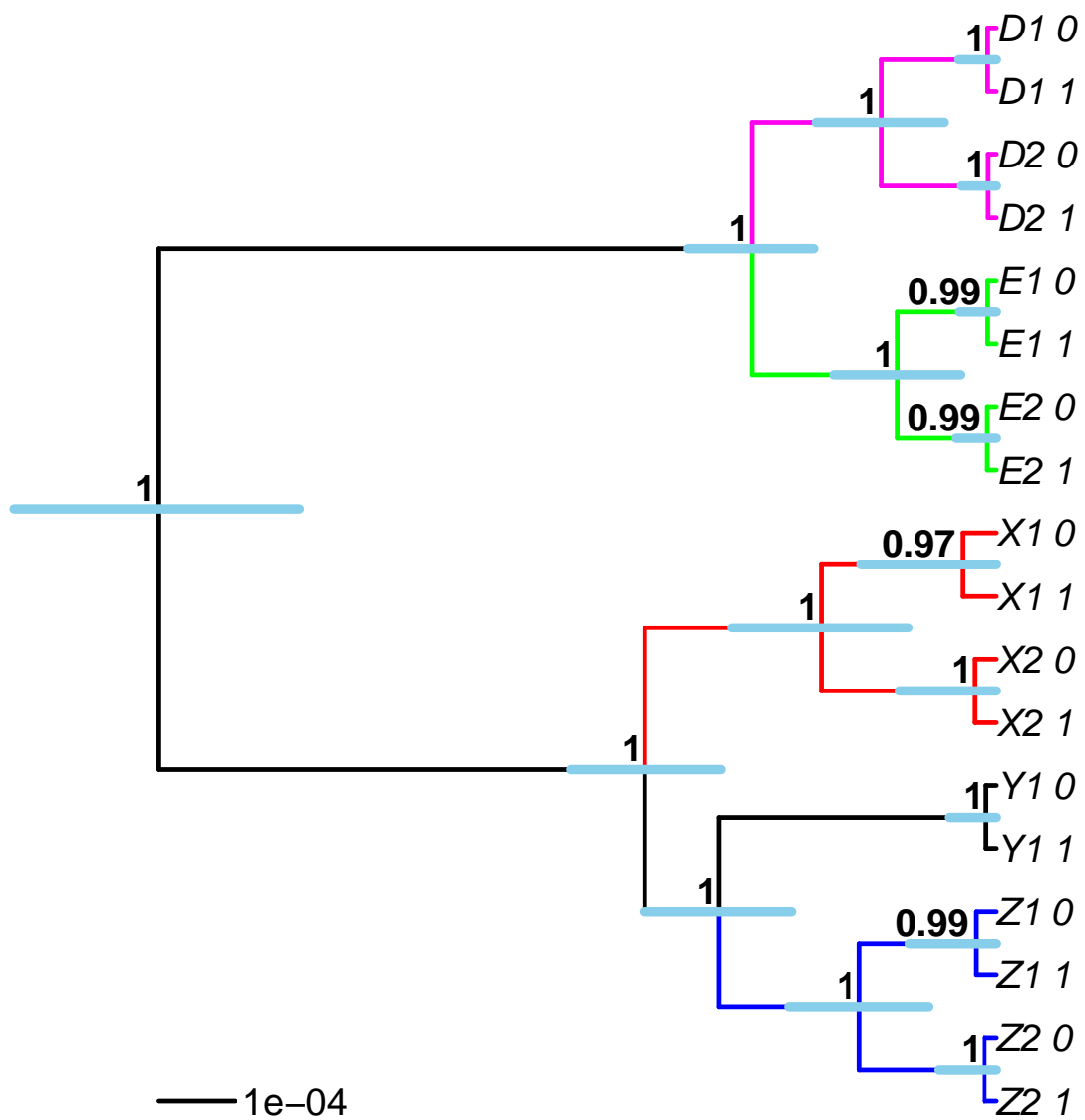
Sim-rep 1: IUPAC consensus alignments (n=150), STACEY species tree



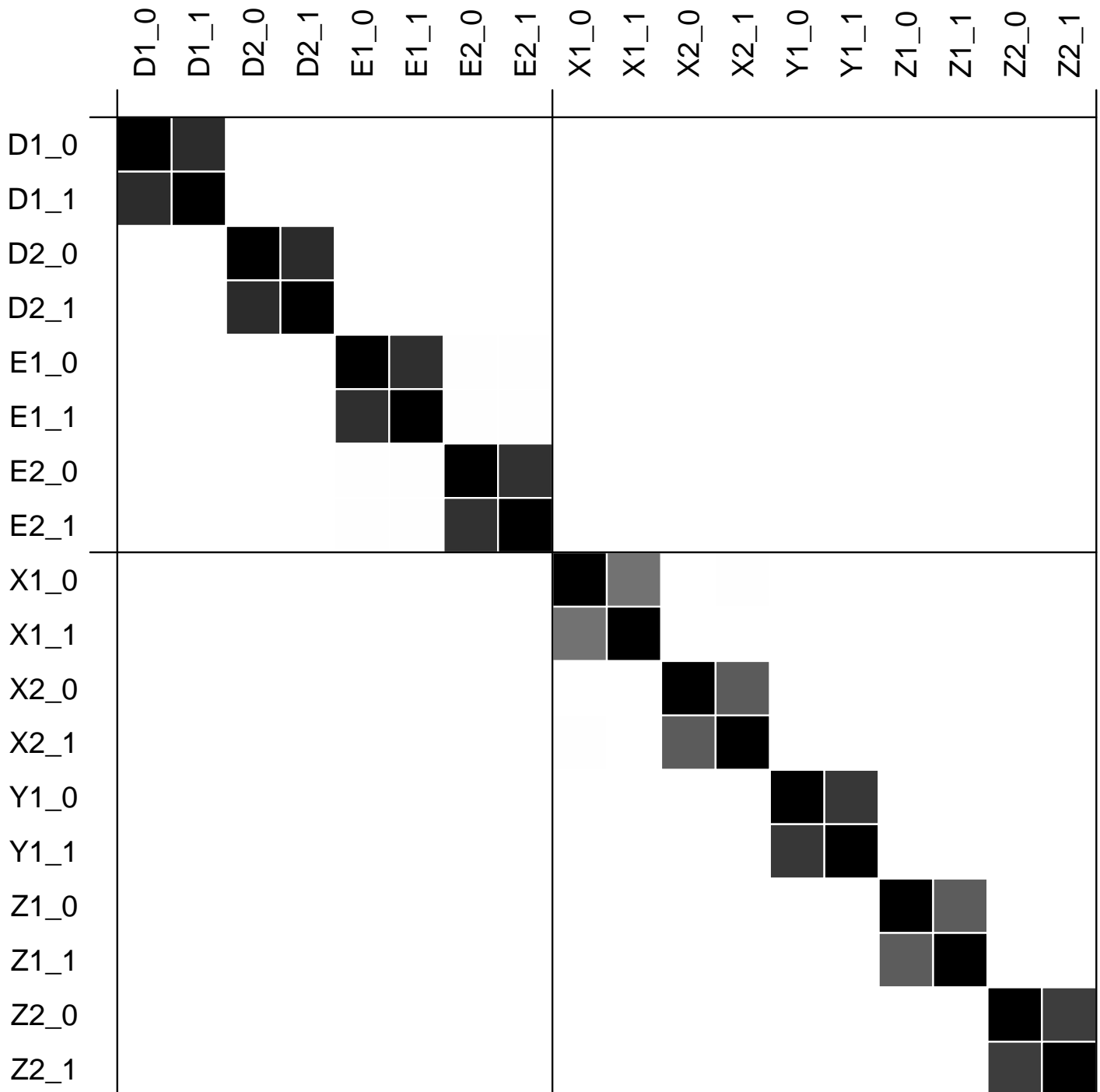
Sim-rep 1: IUPAC consensus alignments (n=150), STACEY similarity matrix



Sim-rep 1: Chimeric allele alignments (n=150), STACEY species tree

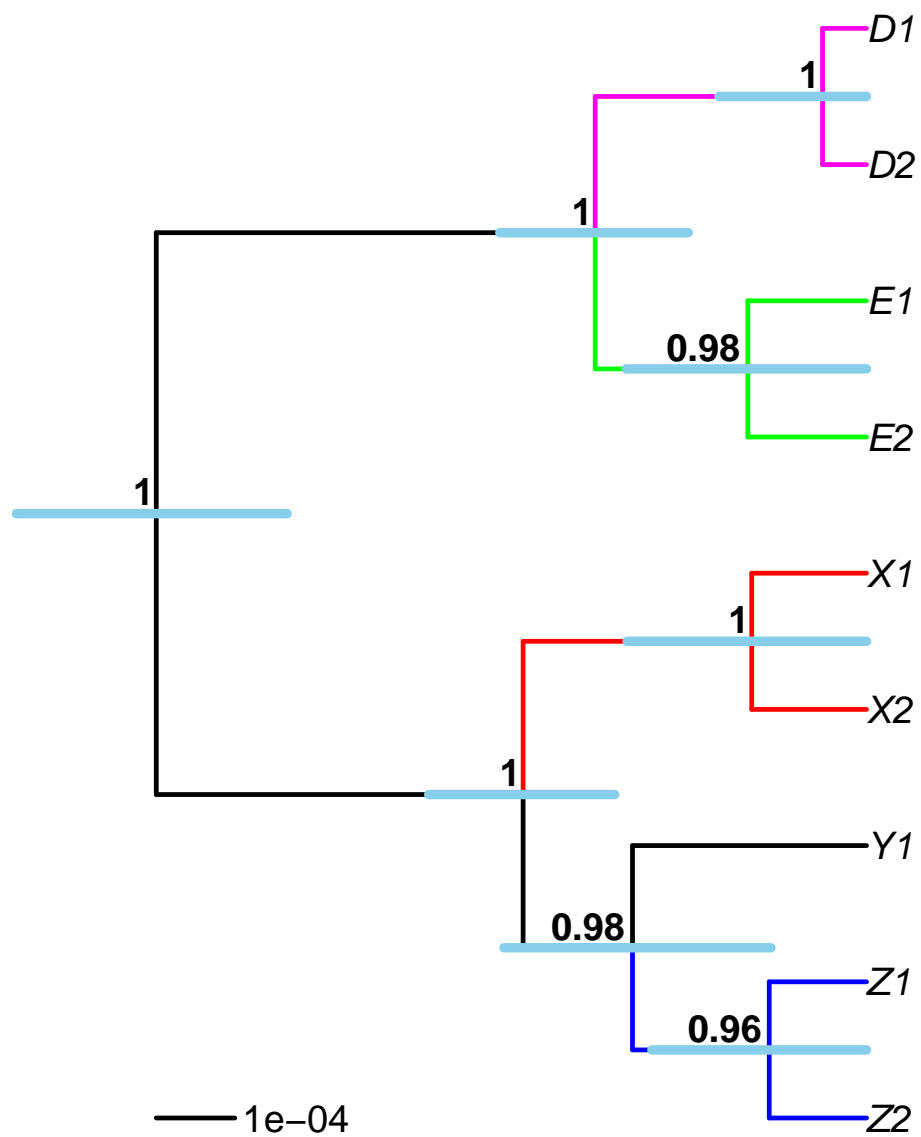


Sim-rep 1: Chimeric allele alignments (n=150), STACEY similarity matrix

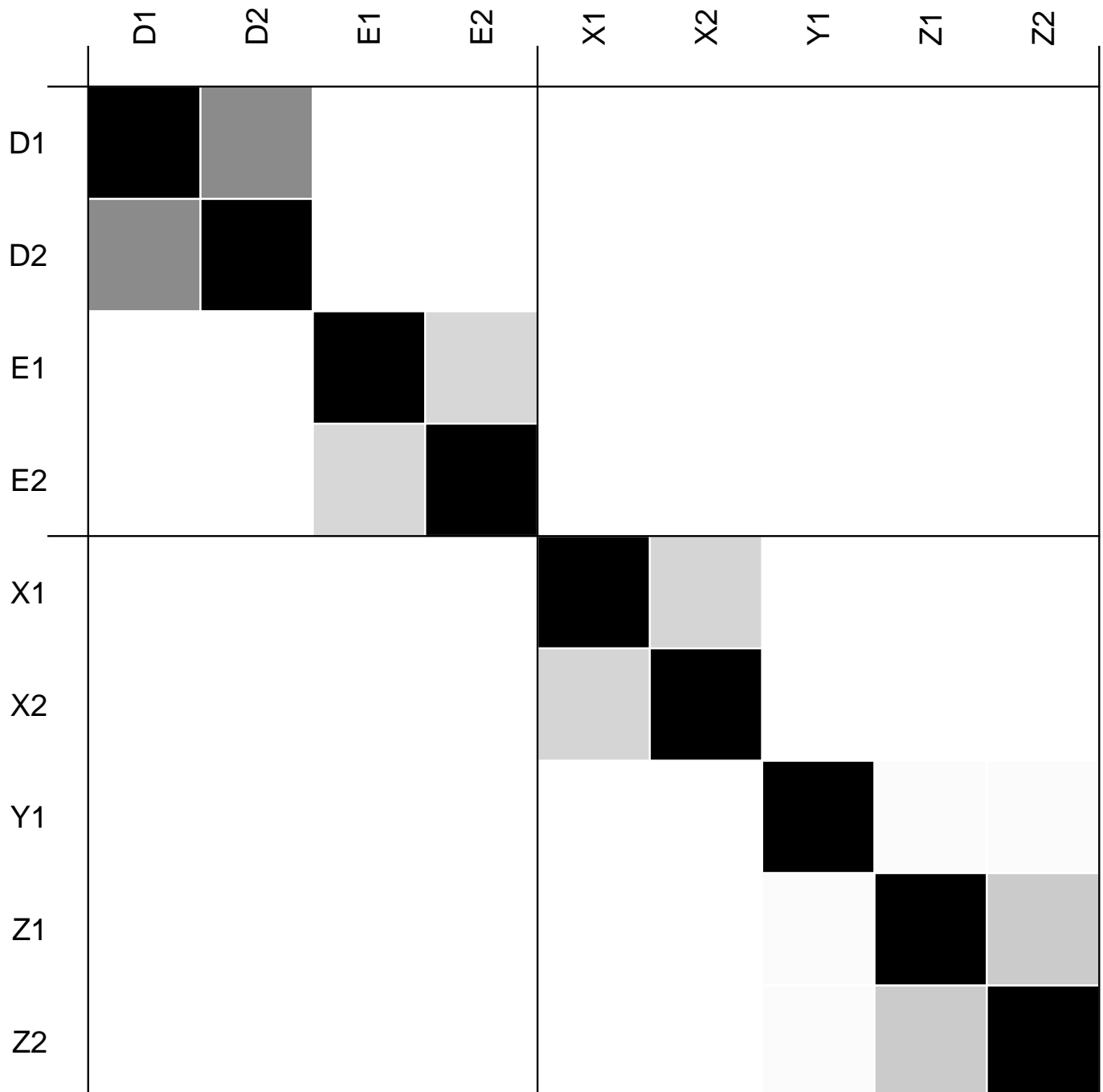




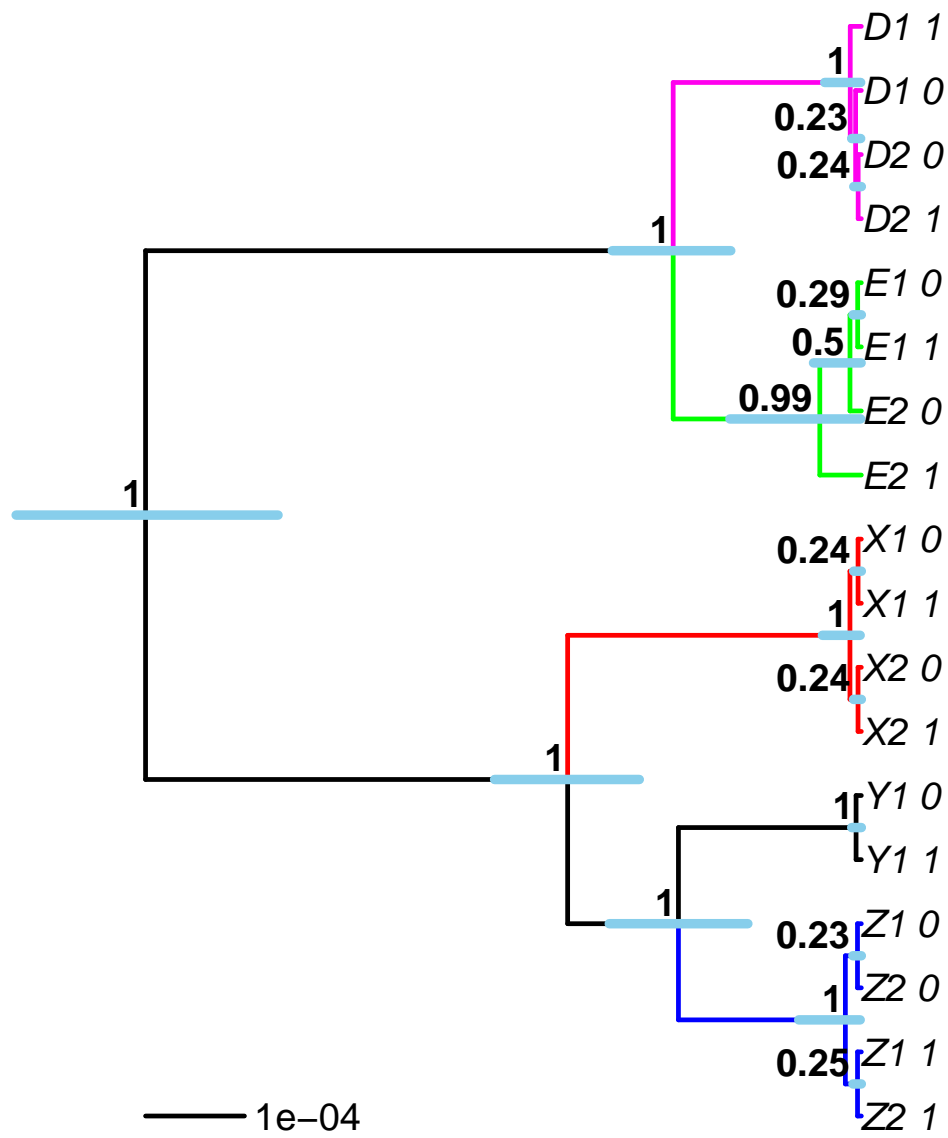
Sim-rep 2: Contig sequence alignments (n=150), STACEY species tree



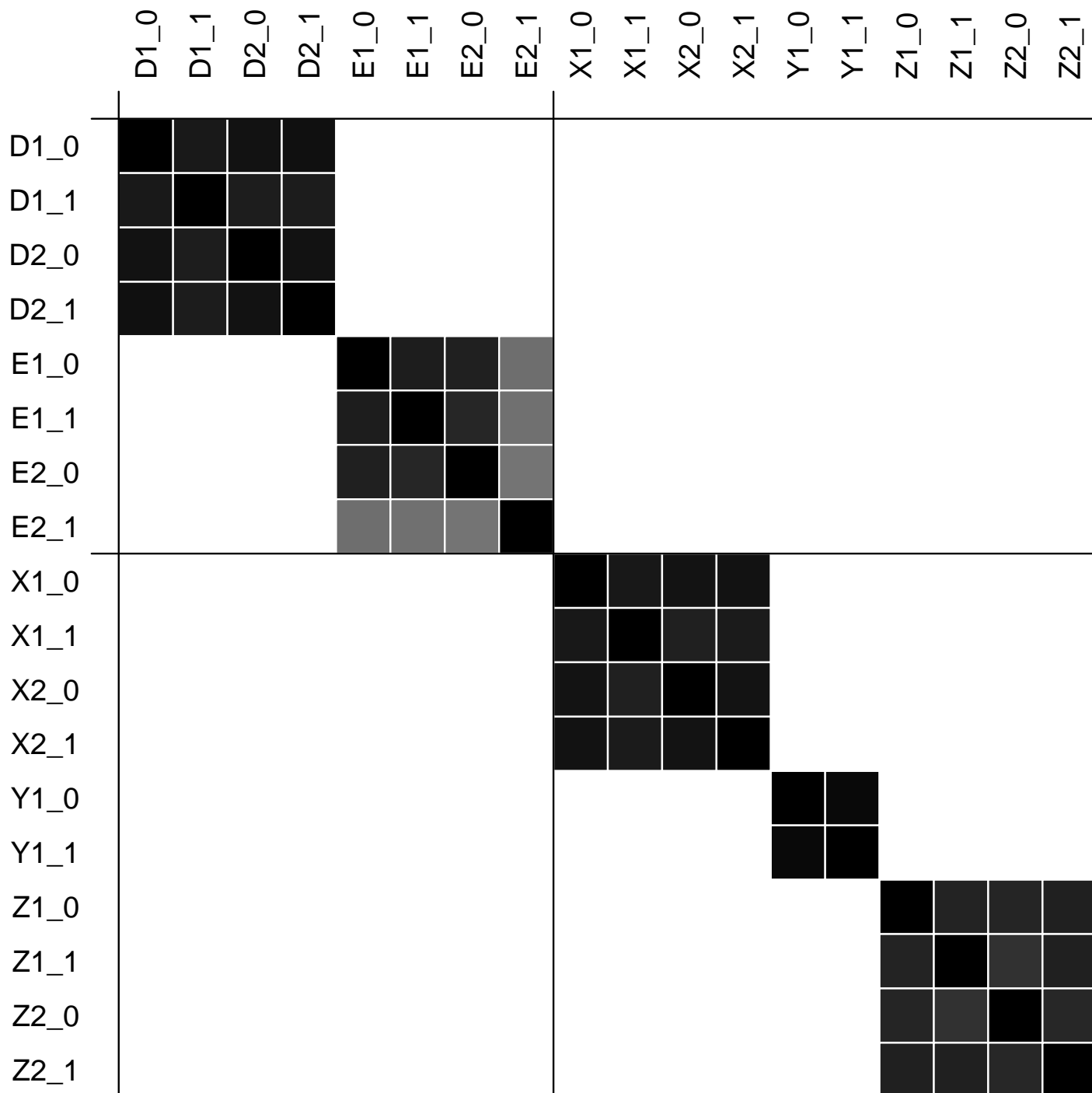
Sim-rep 2: Contig sequence alignments (n=150), STACEY similarity matrix



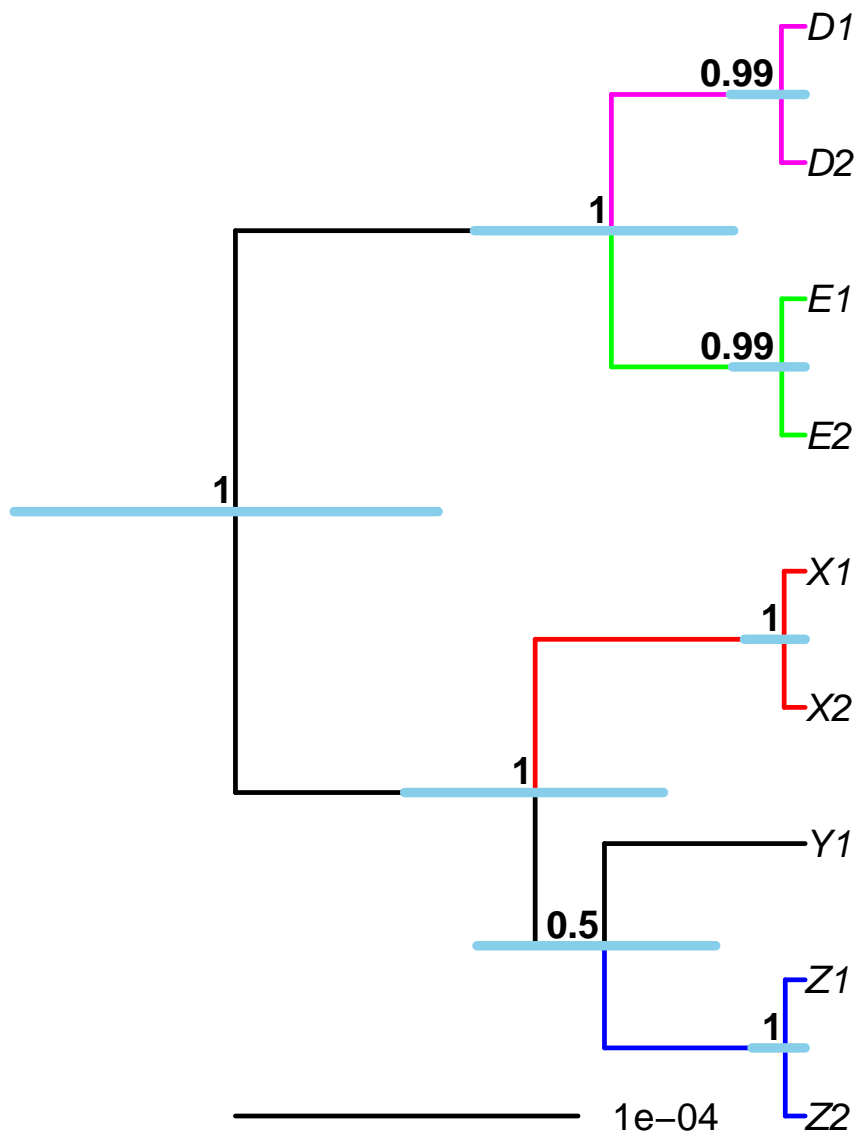
Sim-rep 2: Phased allele alignments (n=150), STACEY species tree



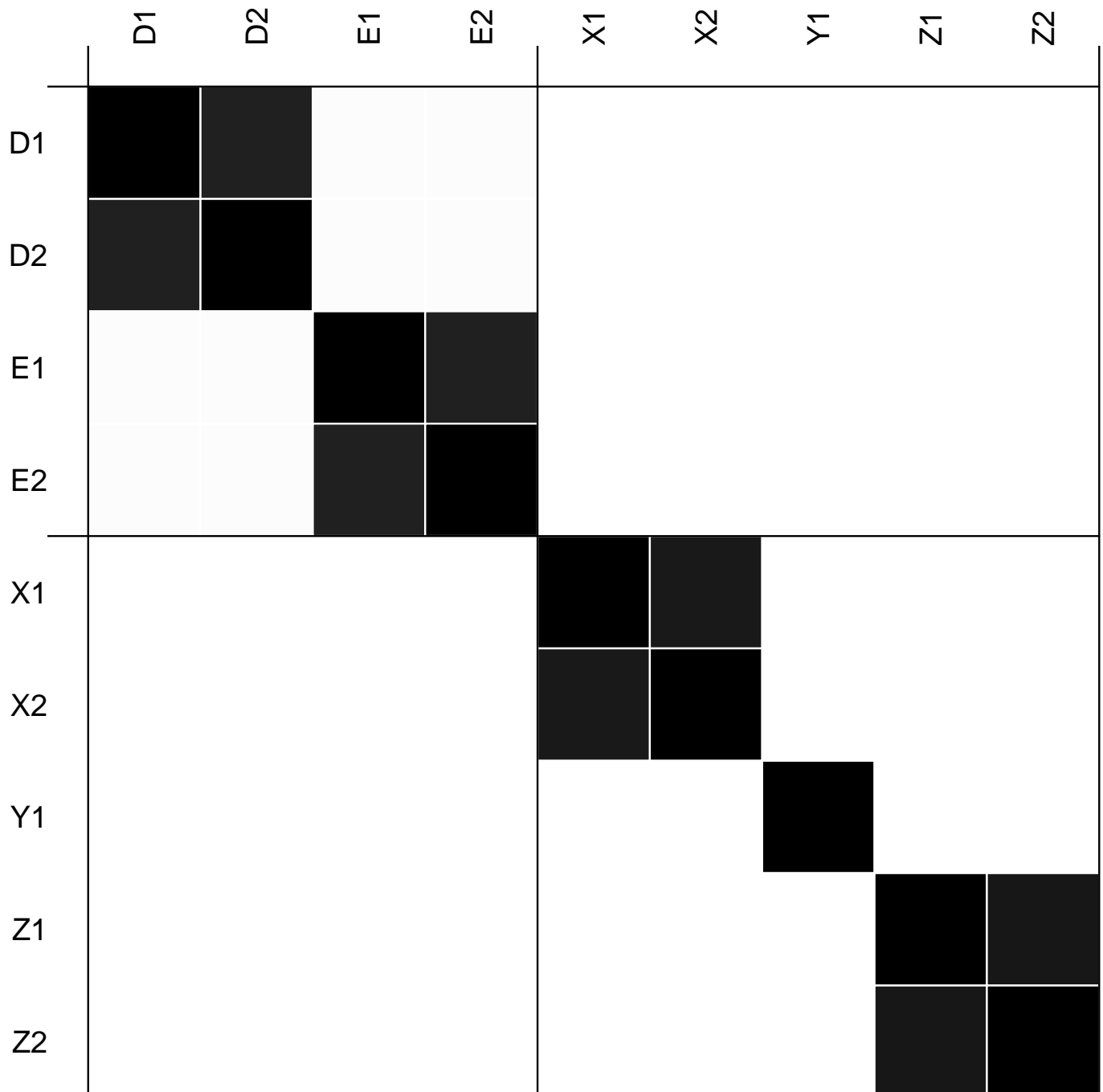
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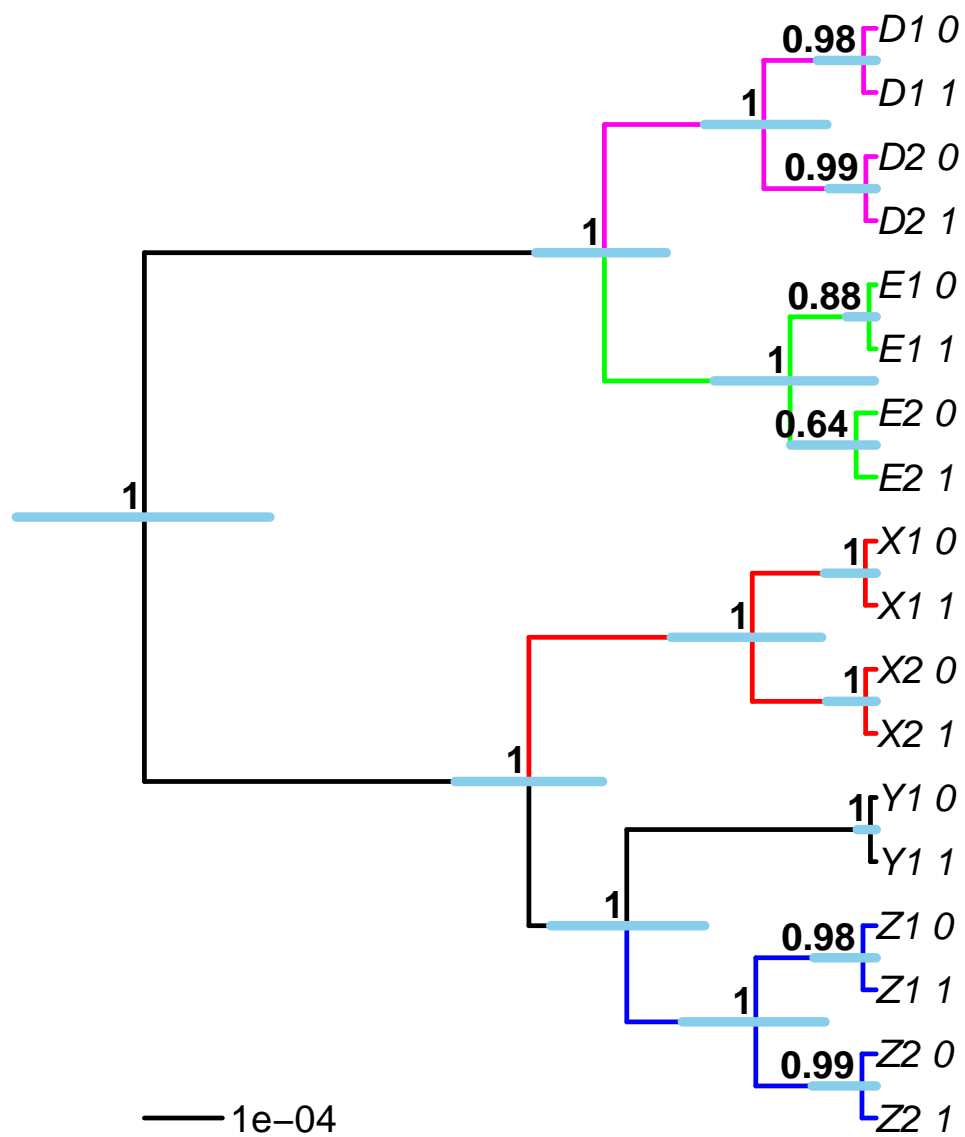
Sim-rep 2: IUPAC consensus alignments (n=150), STACEY species tree



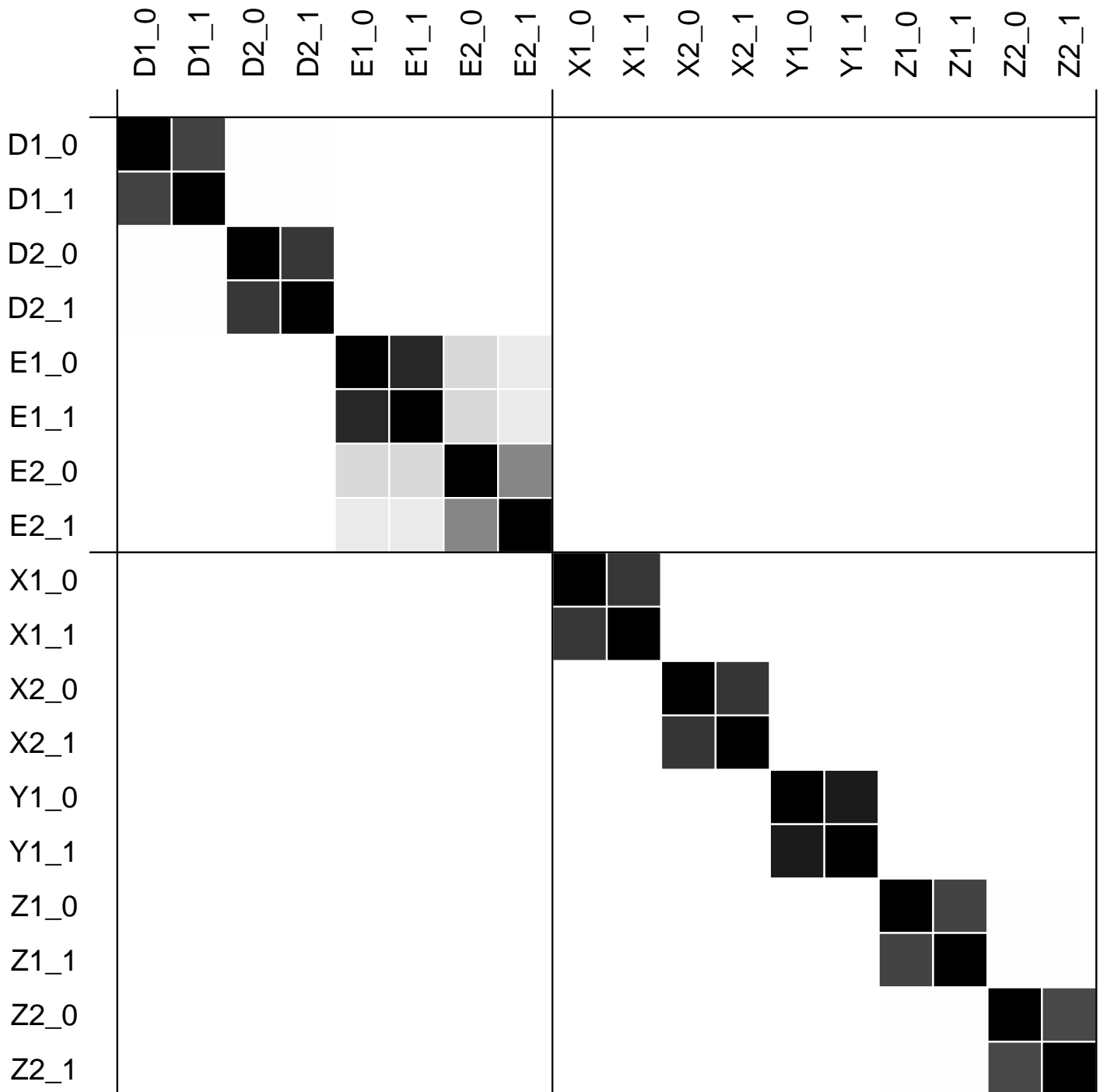
Sim-rep 2: IUPAC consensus alignments (n=150), STACEY similarity matrix



Sim-rep 2: Chimeric allele alignments (n=150), STACEY species tree

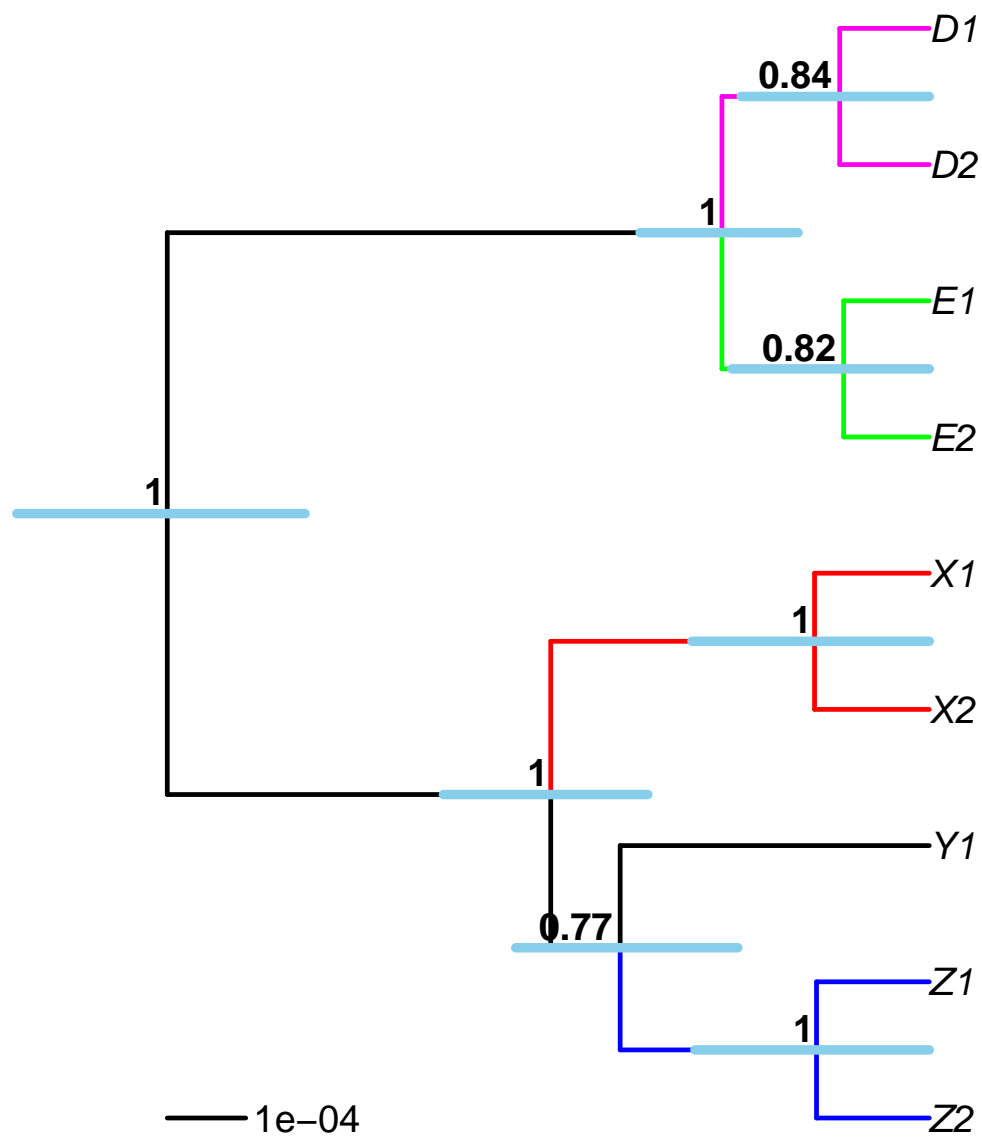


Sim-rep 2: Chimeric allele alignments (n=150), STACEY similarity matrix

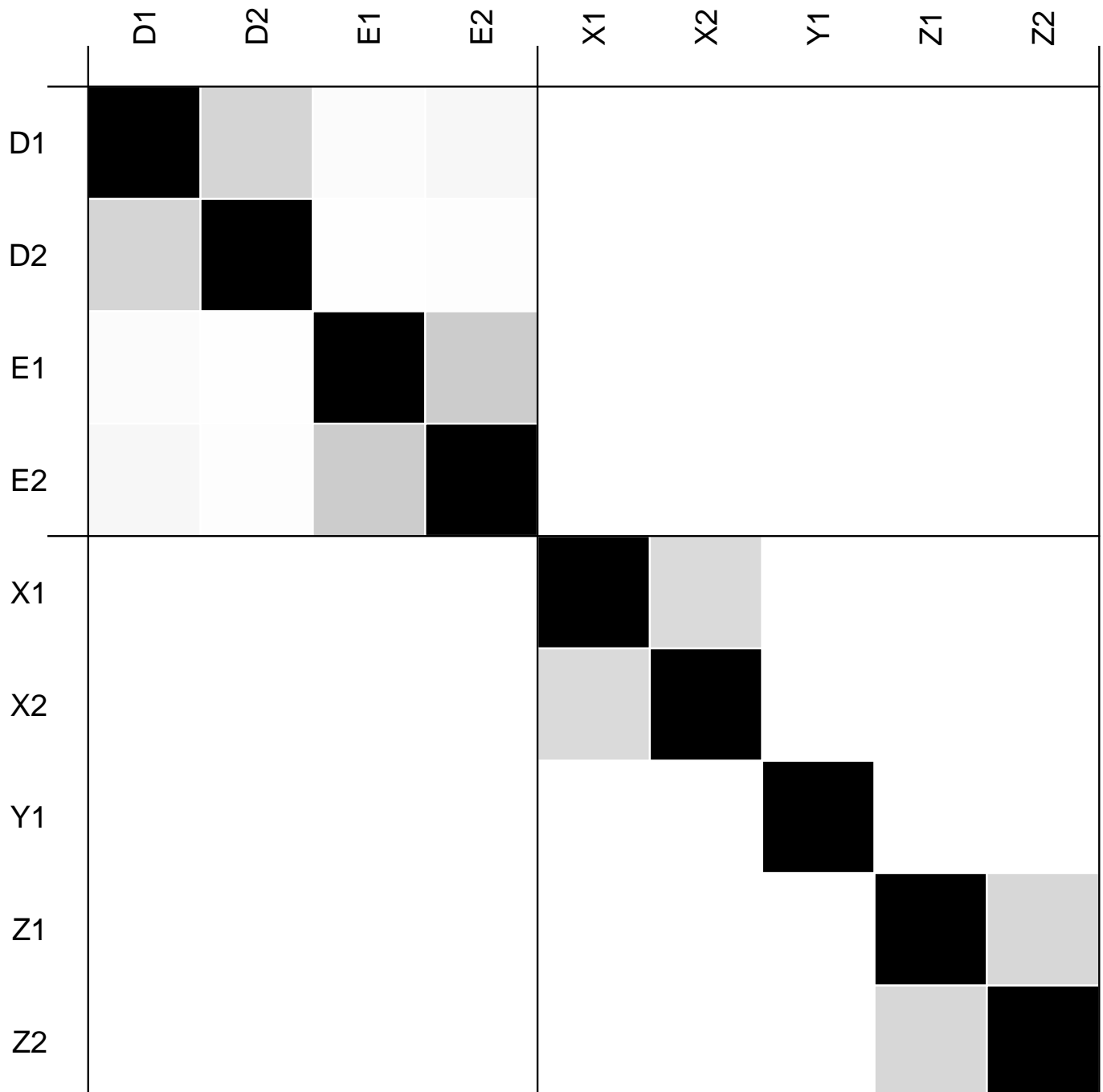




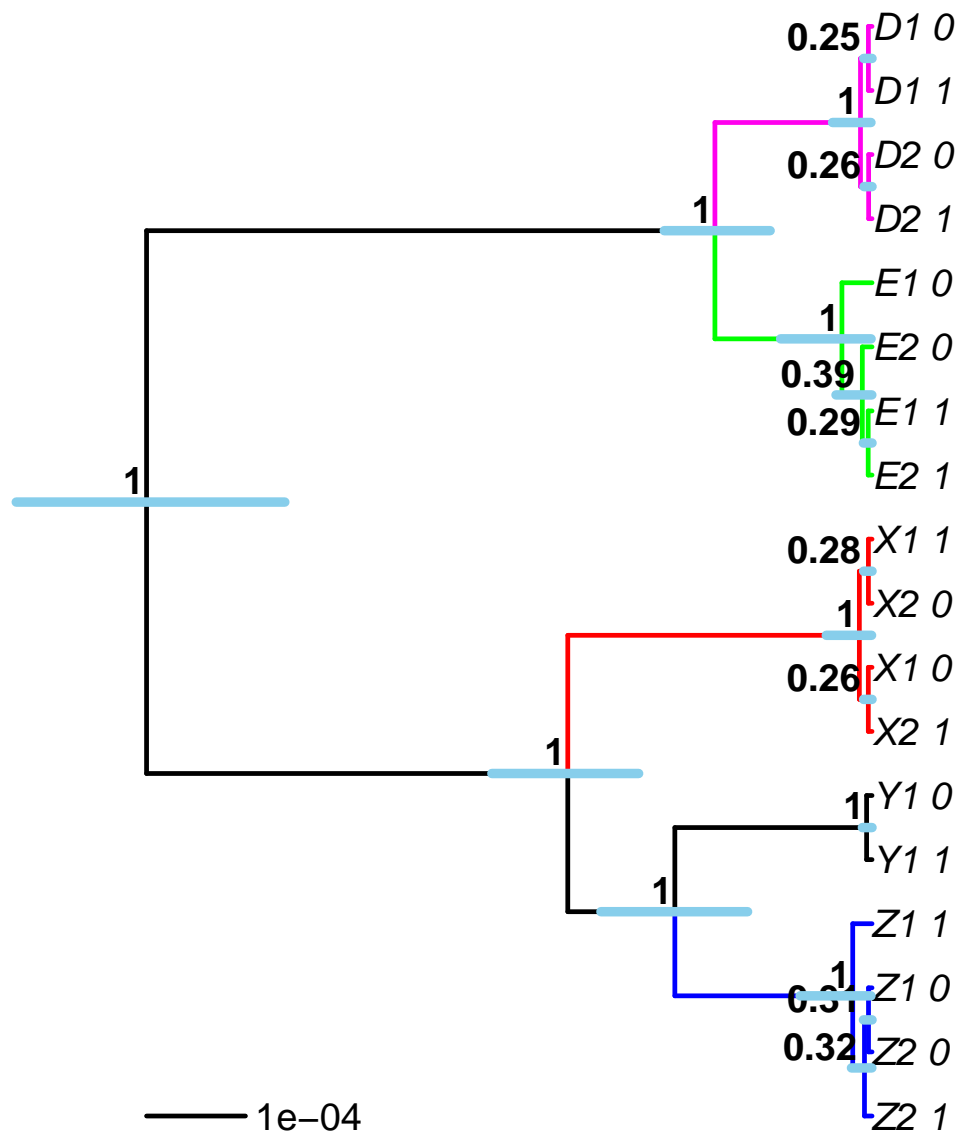
Sim-rep 3: Contig sequence alignments (n=150), STACEY species tree



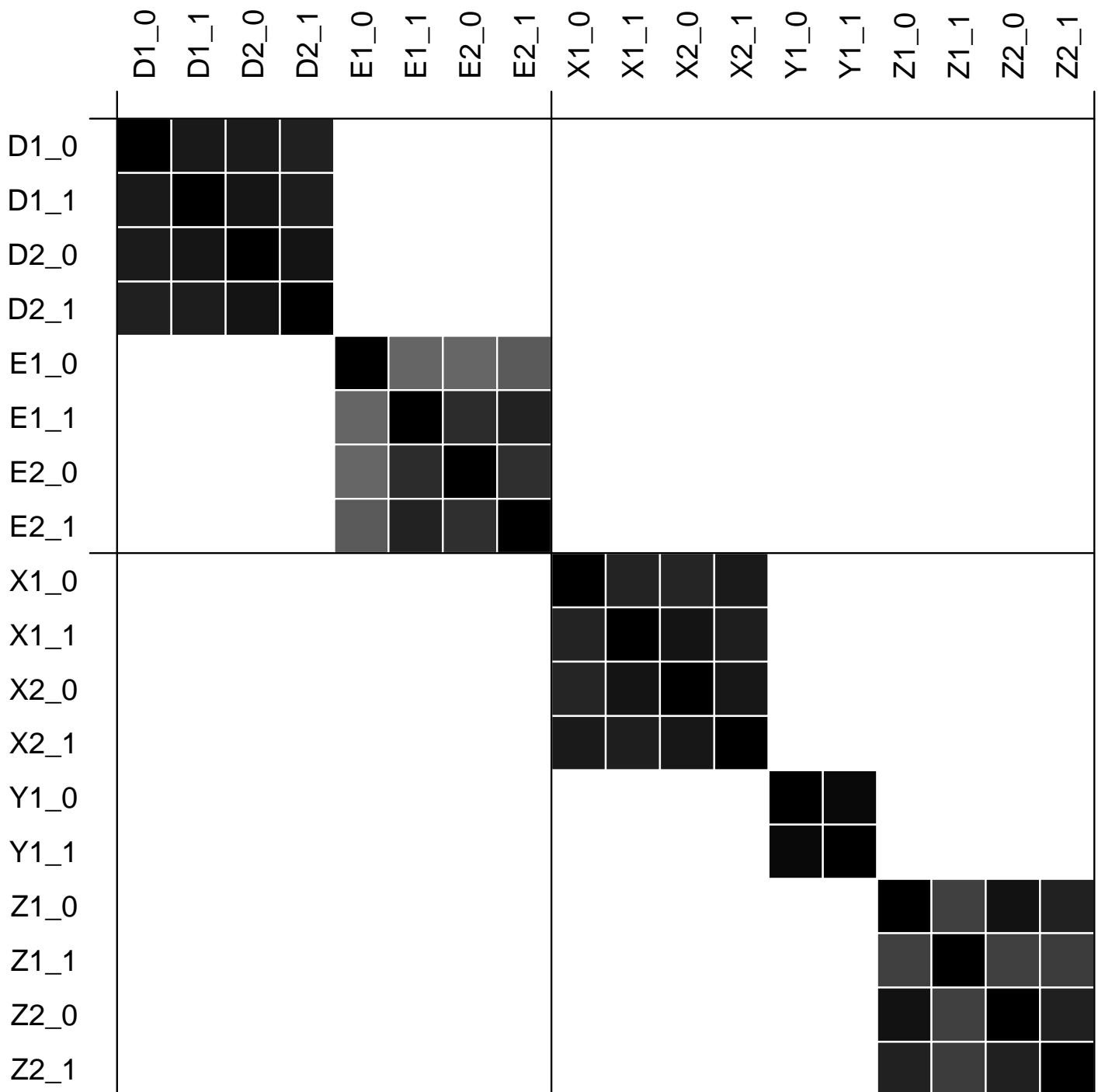
Sim-rep 3: Contig sequence alignments (n=150), STACEY similarity matrix



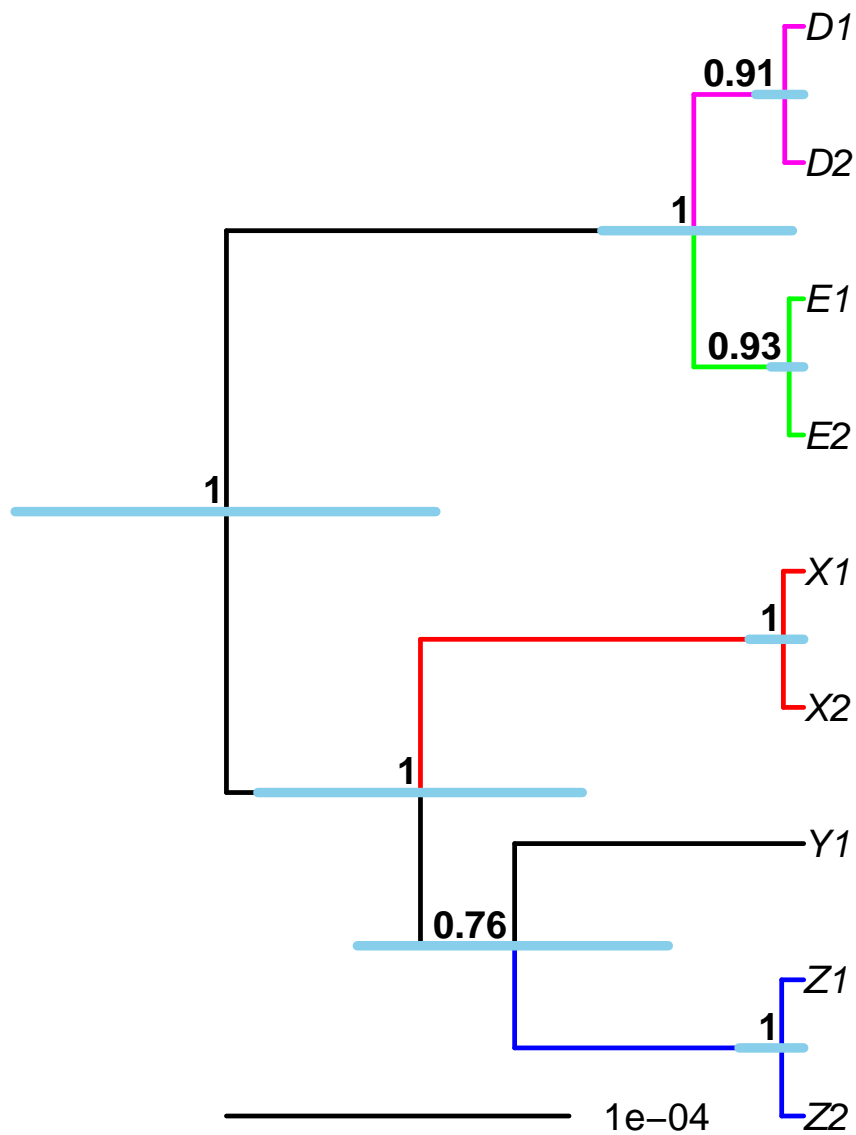
Sim-rep 3: Phased allele alignments (n=150), STACEY species tree



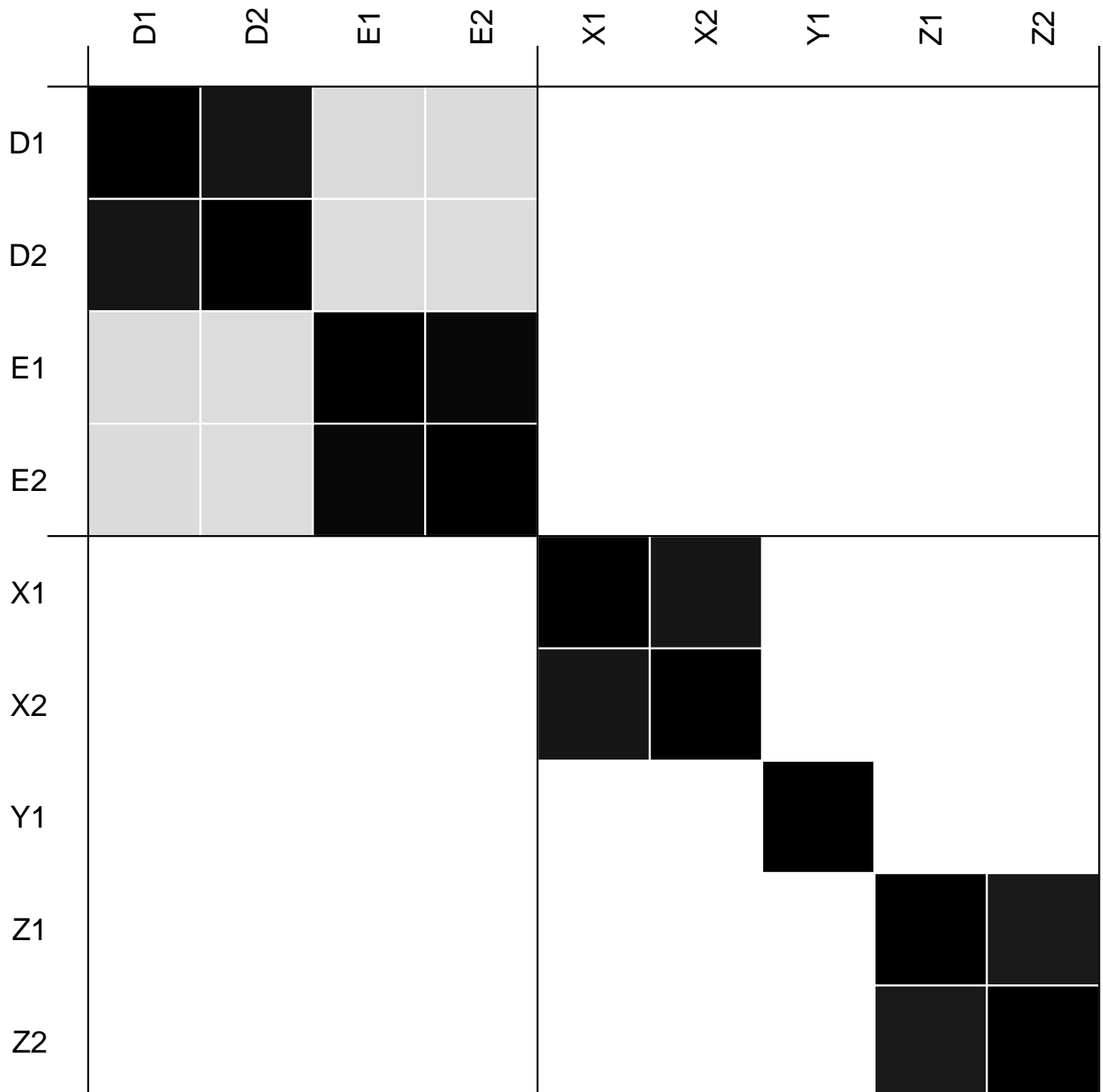
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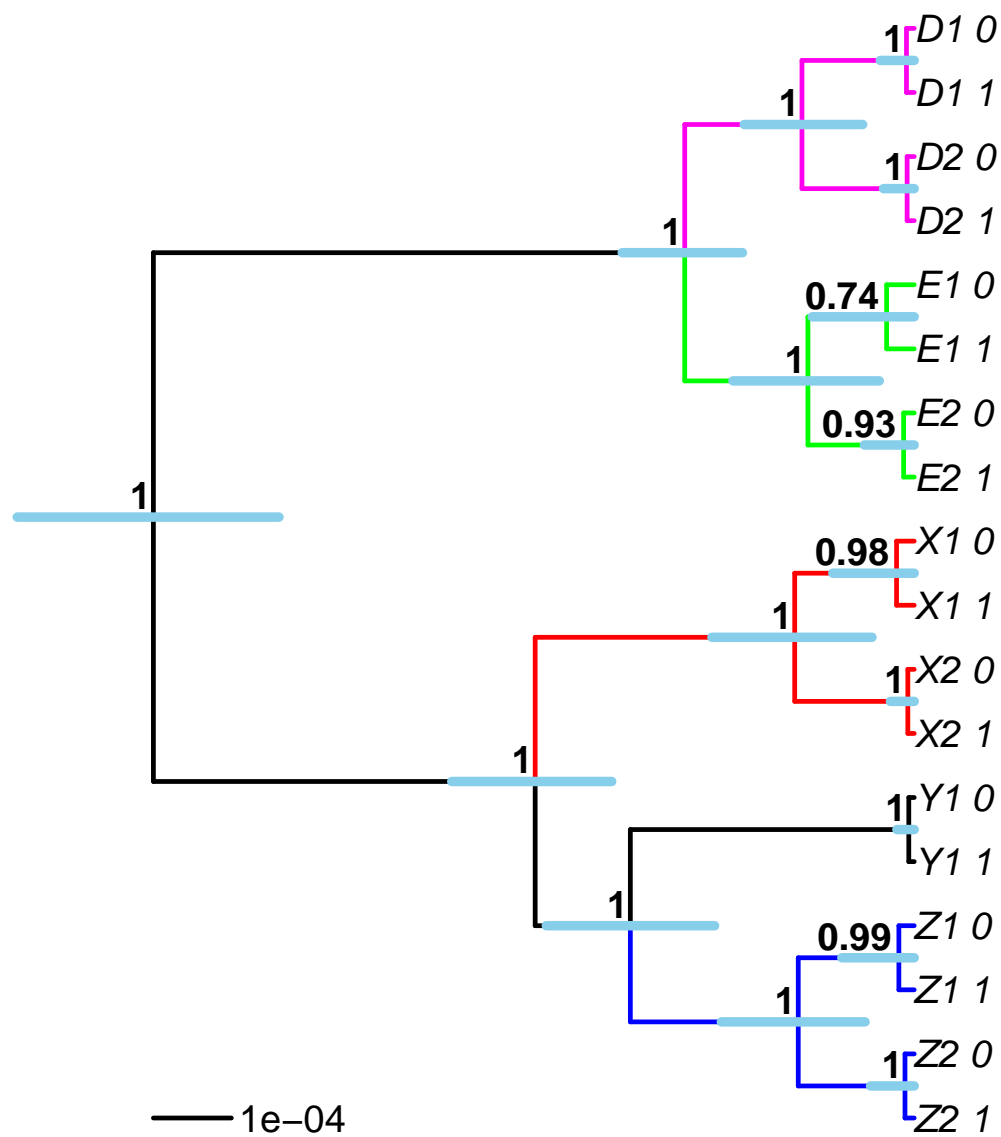
Sim-rep 3: IUPAC consensus alignments (n=150), STACEY species tree



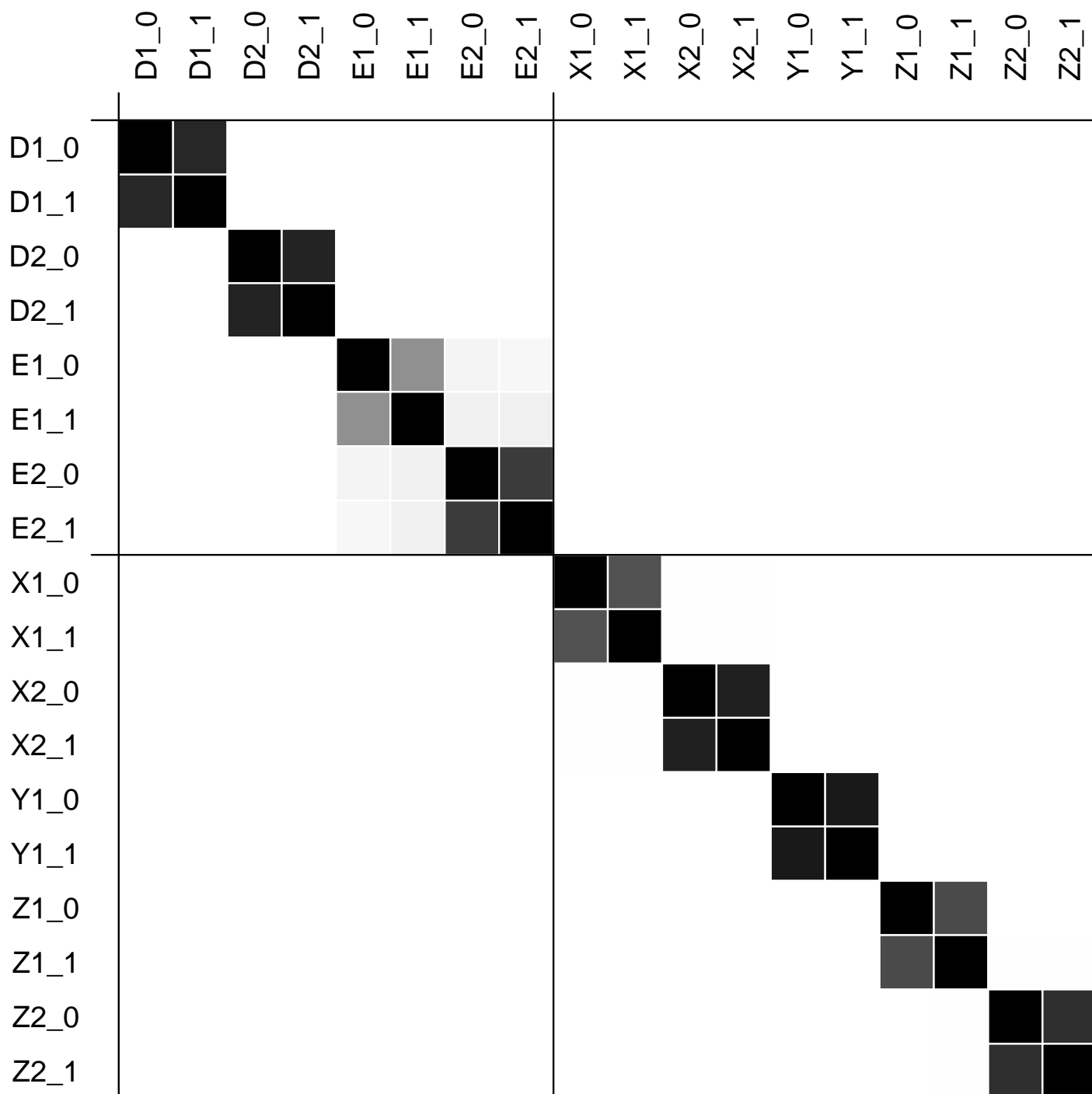
Sim-rep 3: IUPAC consensus alignments (n=150), STACEY similarity matrix



Sim-rep 3: Chimeric allele alignments (n=150), STACEY species tree

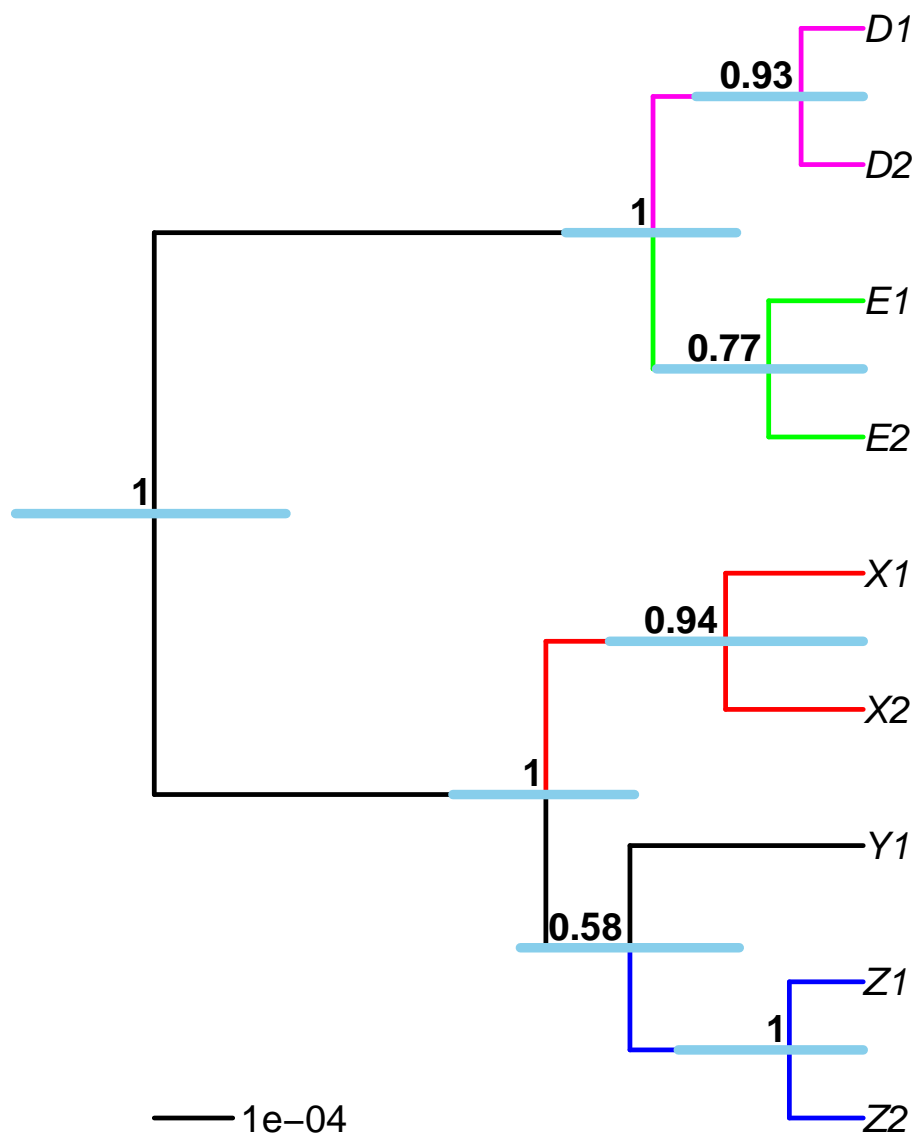


Sim-rep 3: Chimeric allele alignments (n=150), STACEY similarity matrix

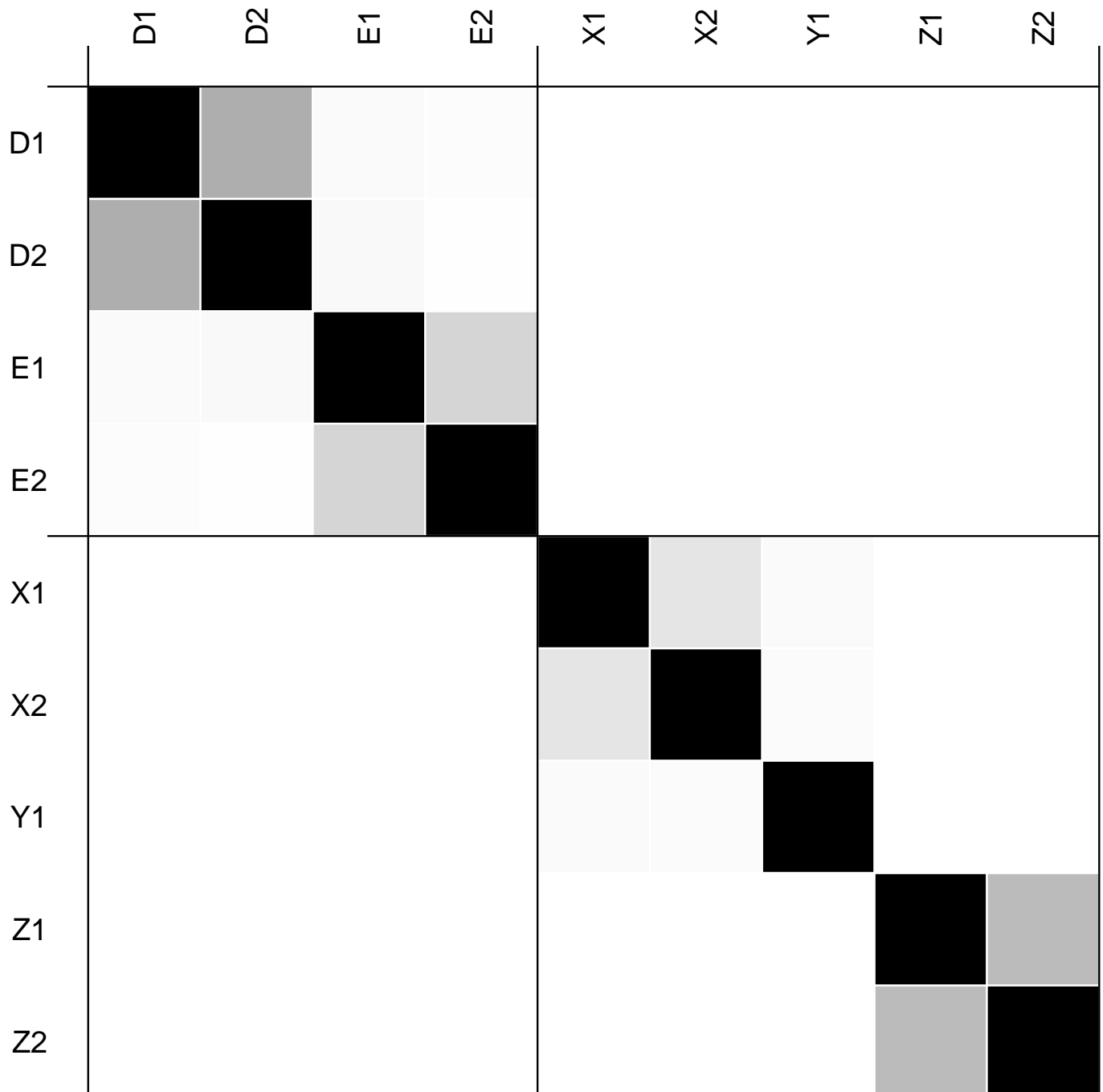




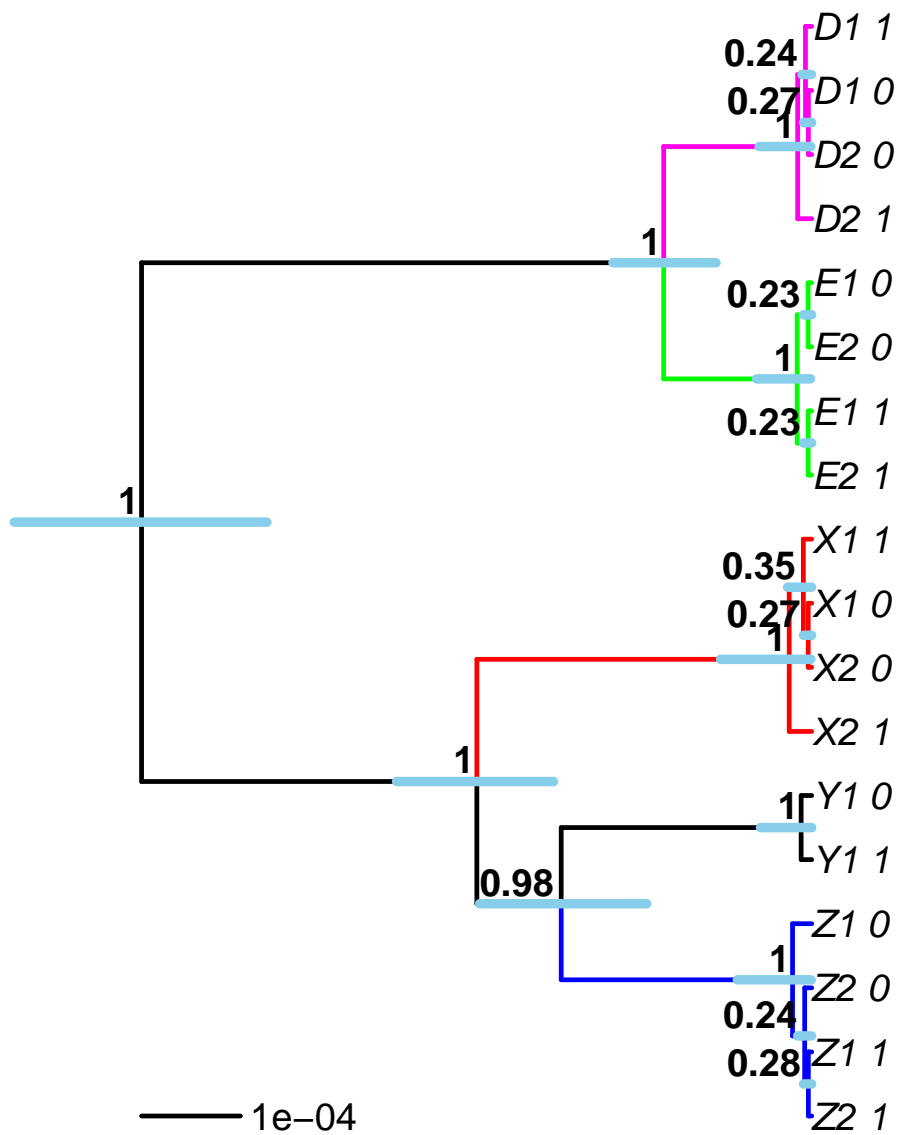
Sim-rep 4: Contig sequence alignments (n=150), STACEY species tree



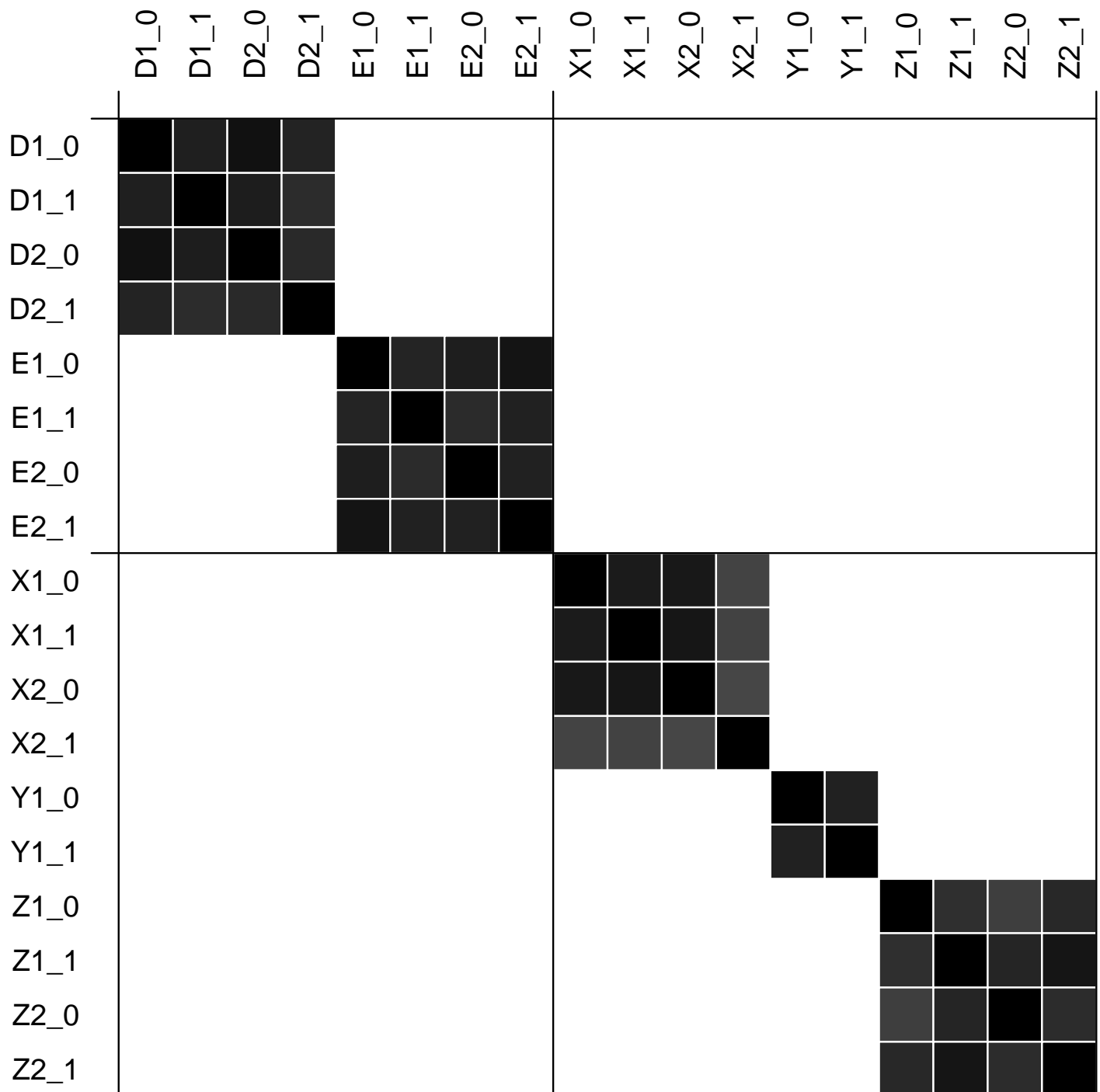
Sim-rep 4: Contig sequence alignments (n=150), STACEY similarity matrix



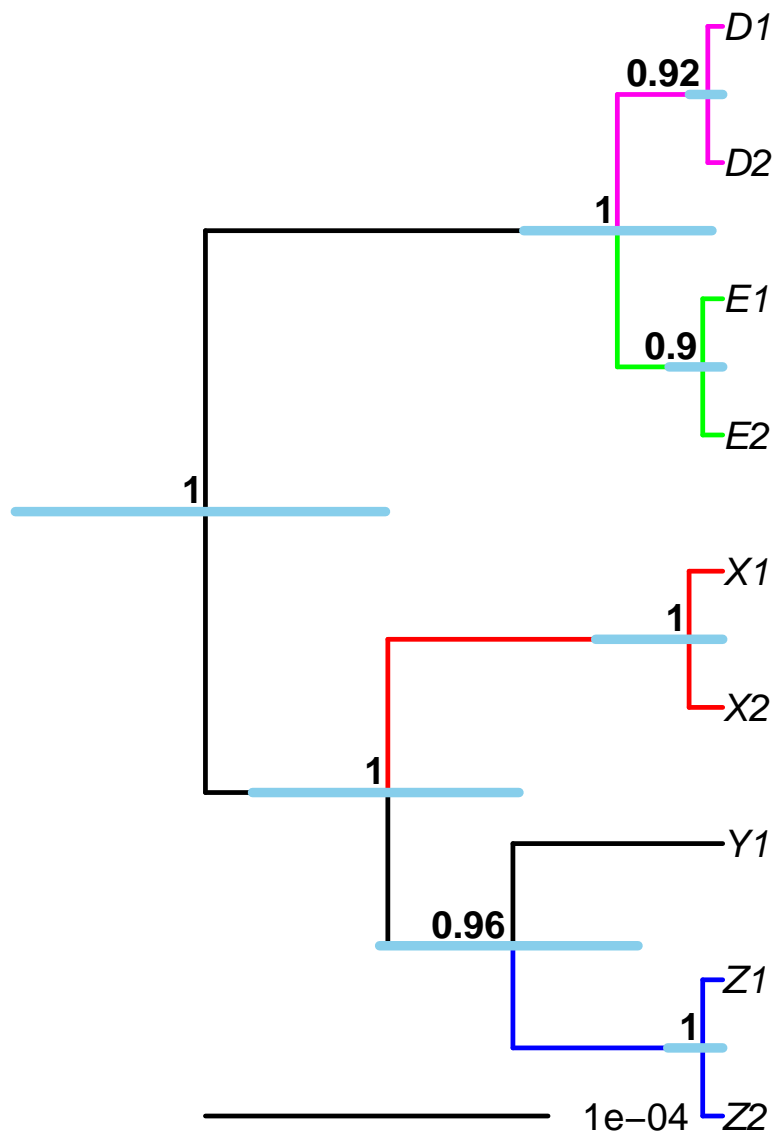
Sim-rep 4: Phased allele alignments (n=150), STACEY species tree



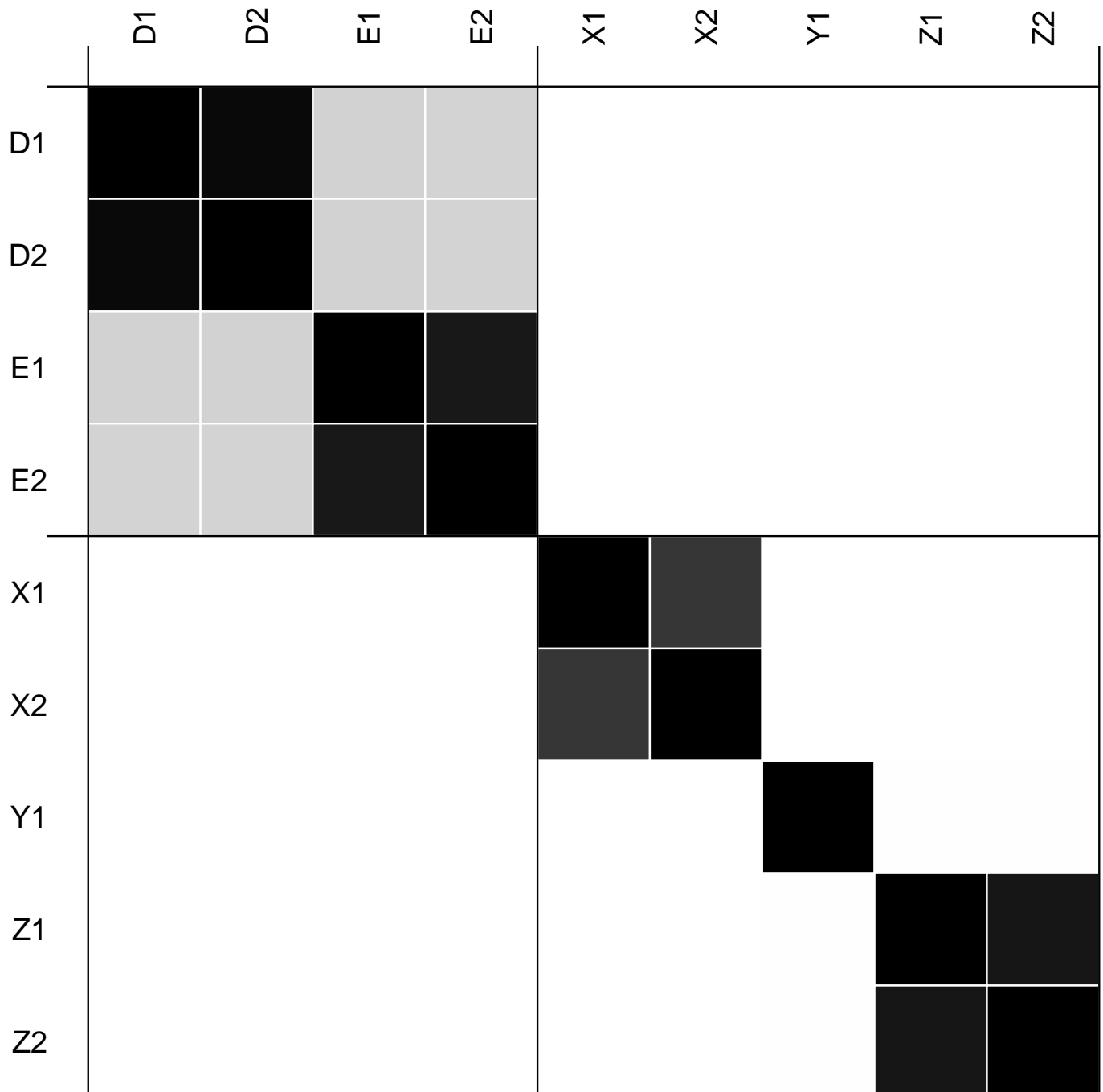
Sim-rep 4: Phased allele alignments (n=150), STACEY similarity matrix



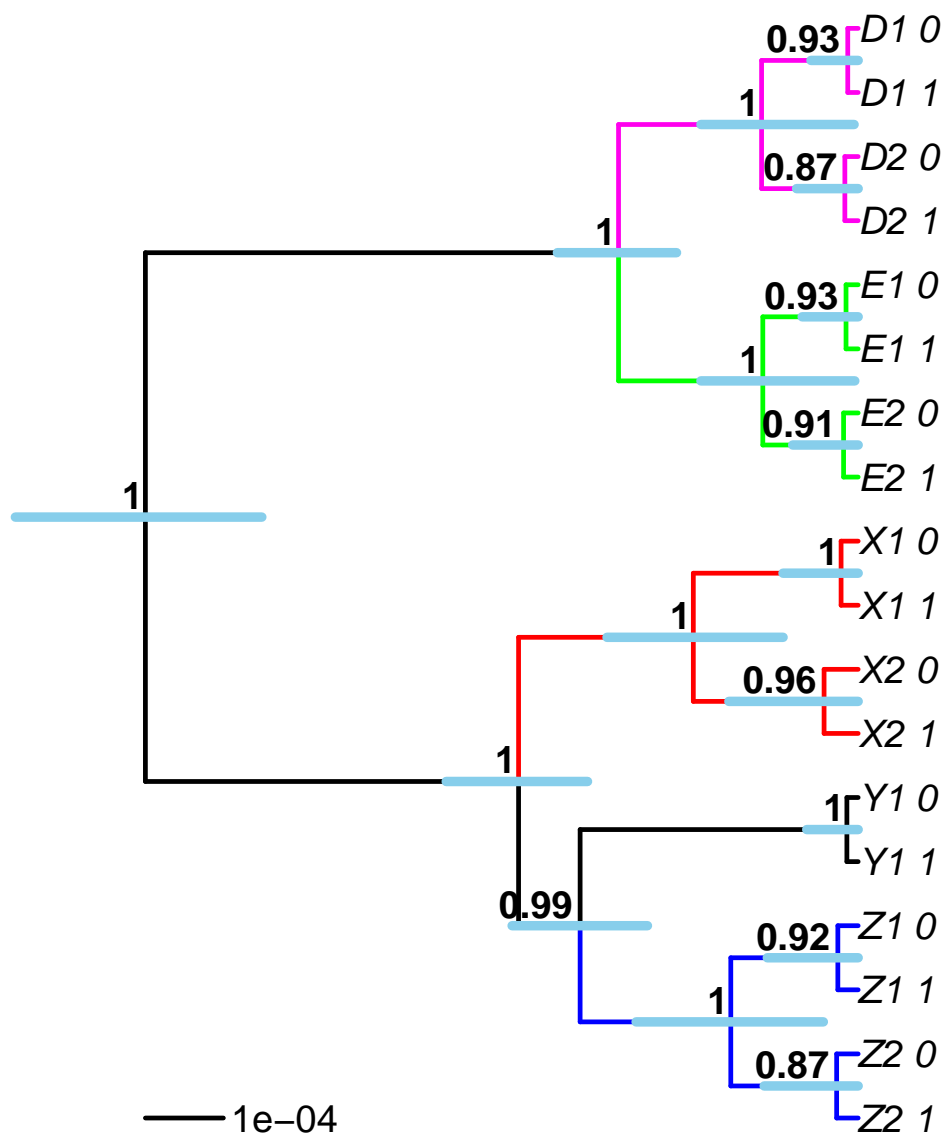
Sim-rep 4: IUPAC consensus alignments (n=150), STACEY species tree



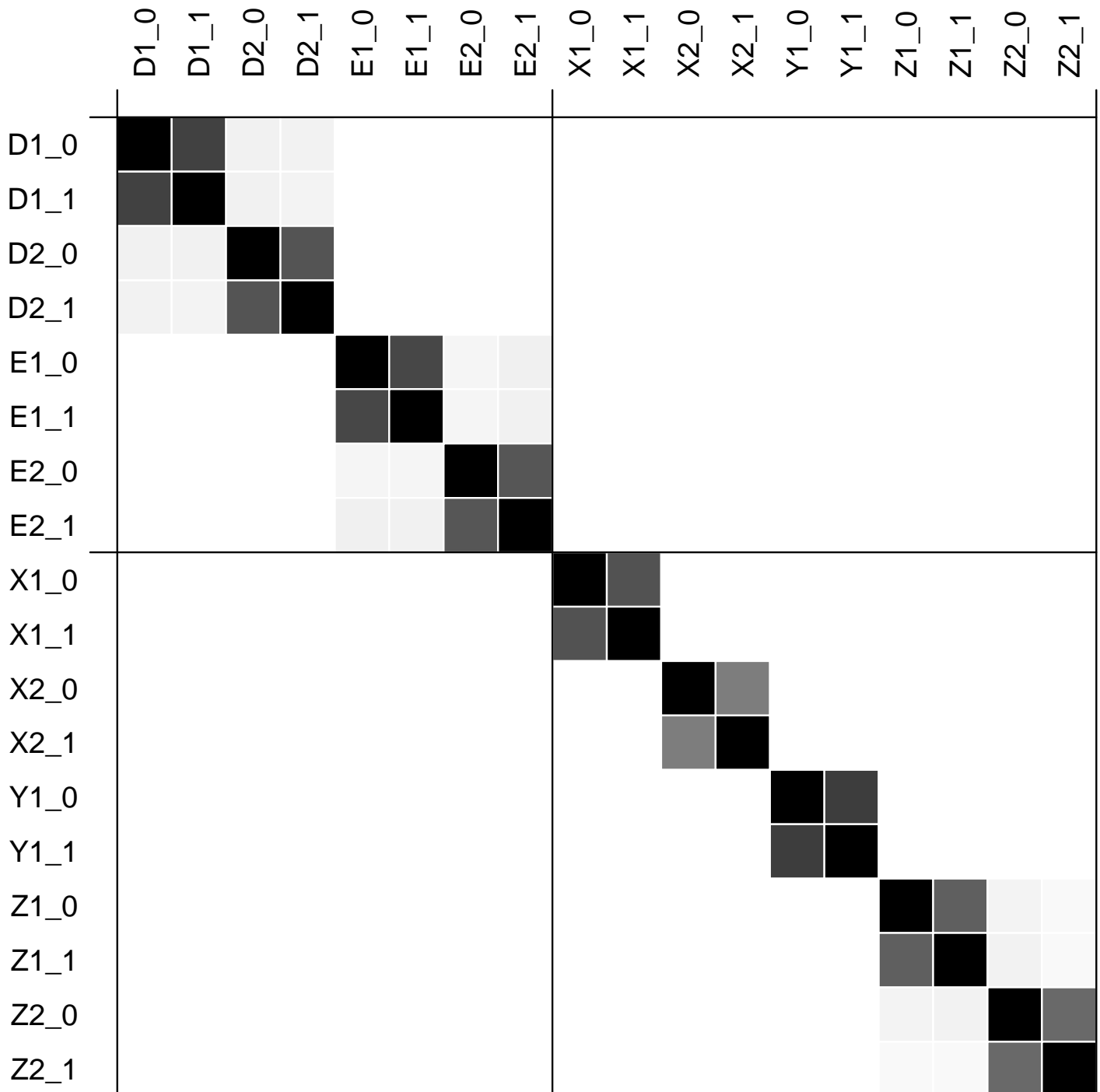
Sim-rep 4: IUPAC consensus alignments (n=150), STACEY similarity matrix



Sim-rep 4: Chimeric allele alignments (n=150), STACEY species tree

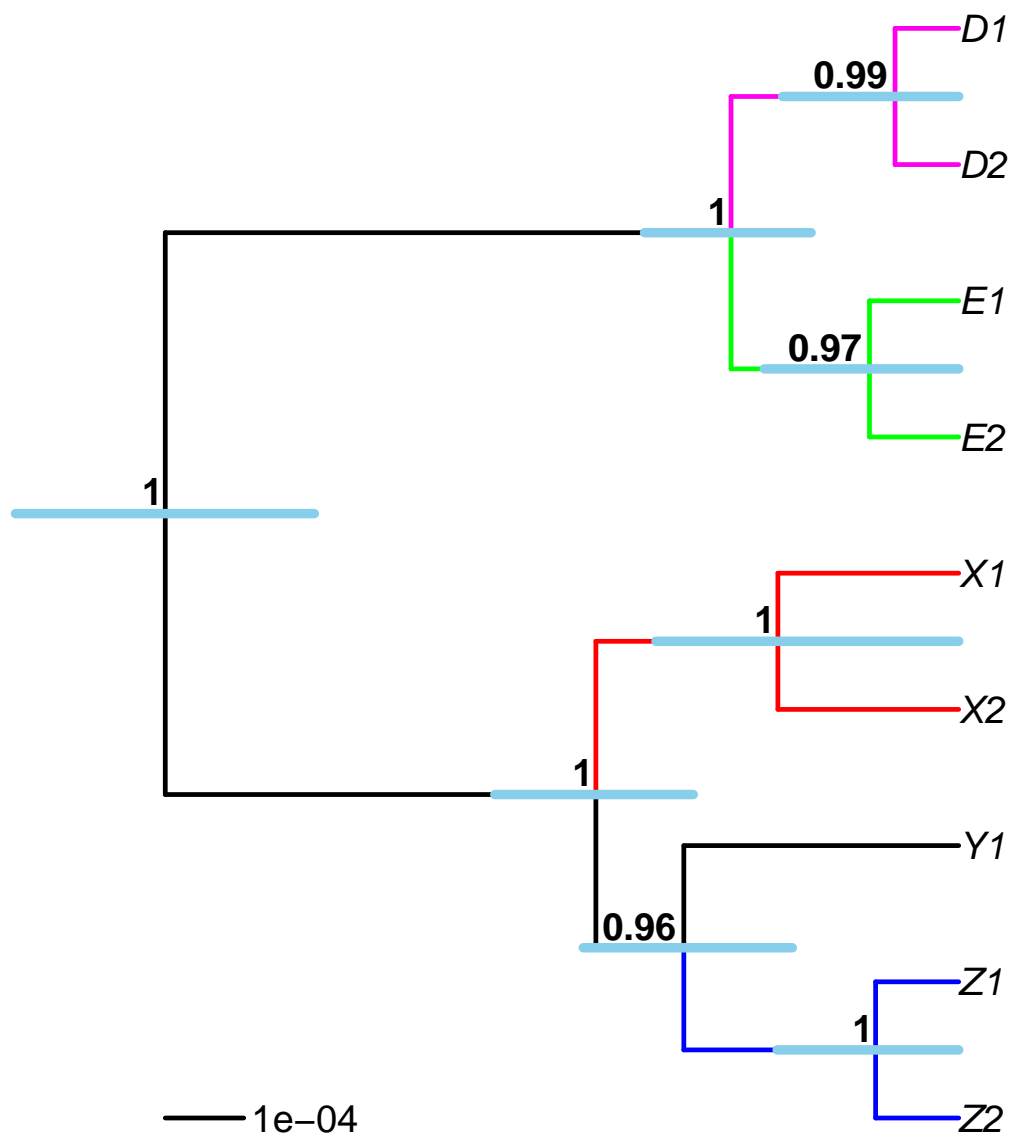


Sim-rep 4: Chimeric allele alignments (n=150), STACEY similarity matrix

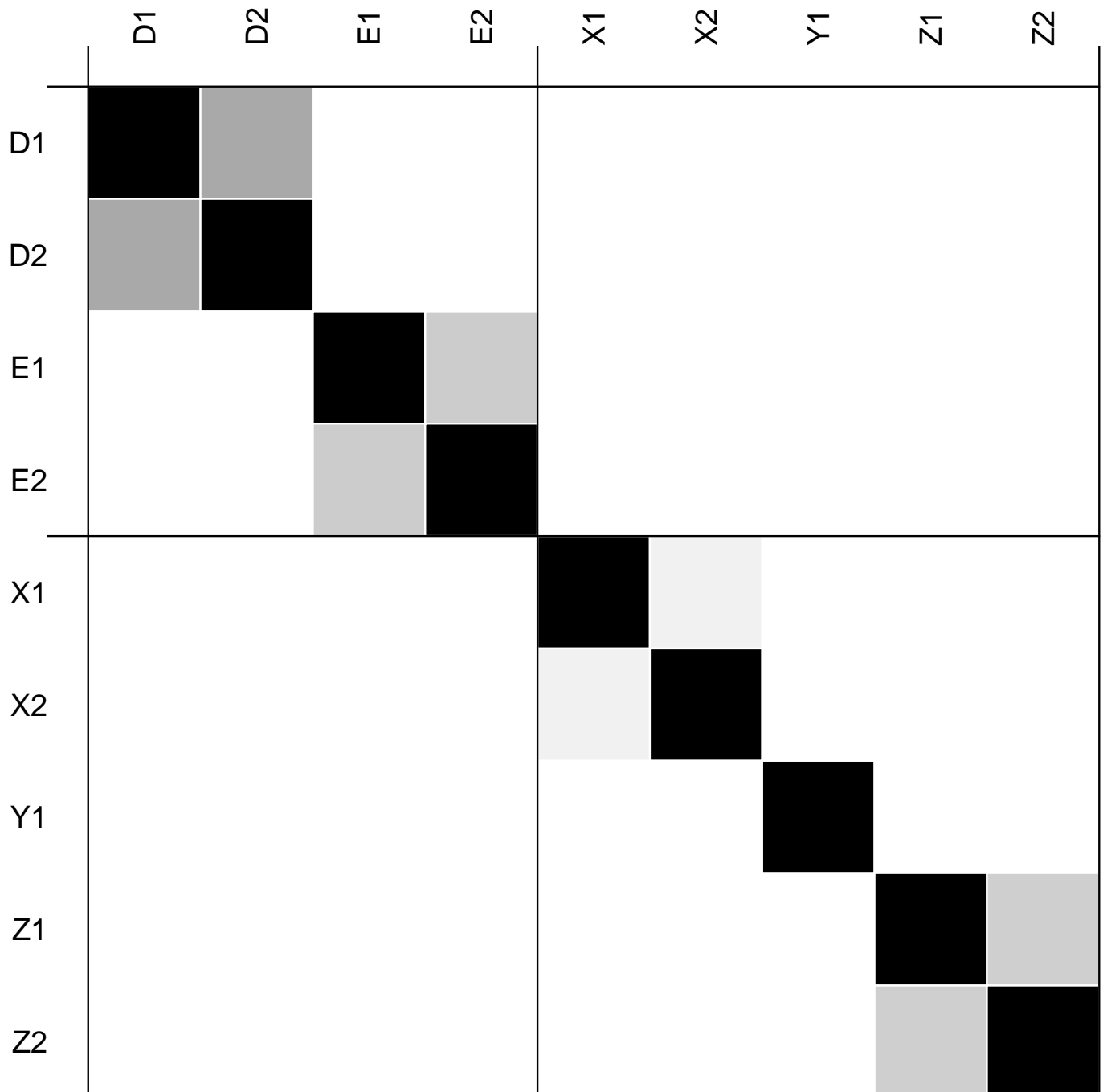




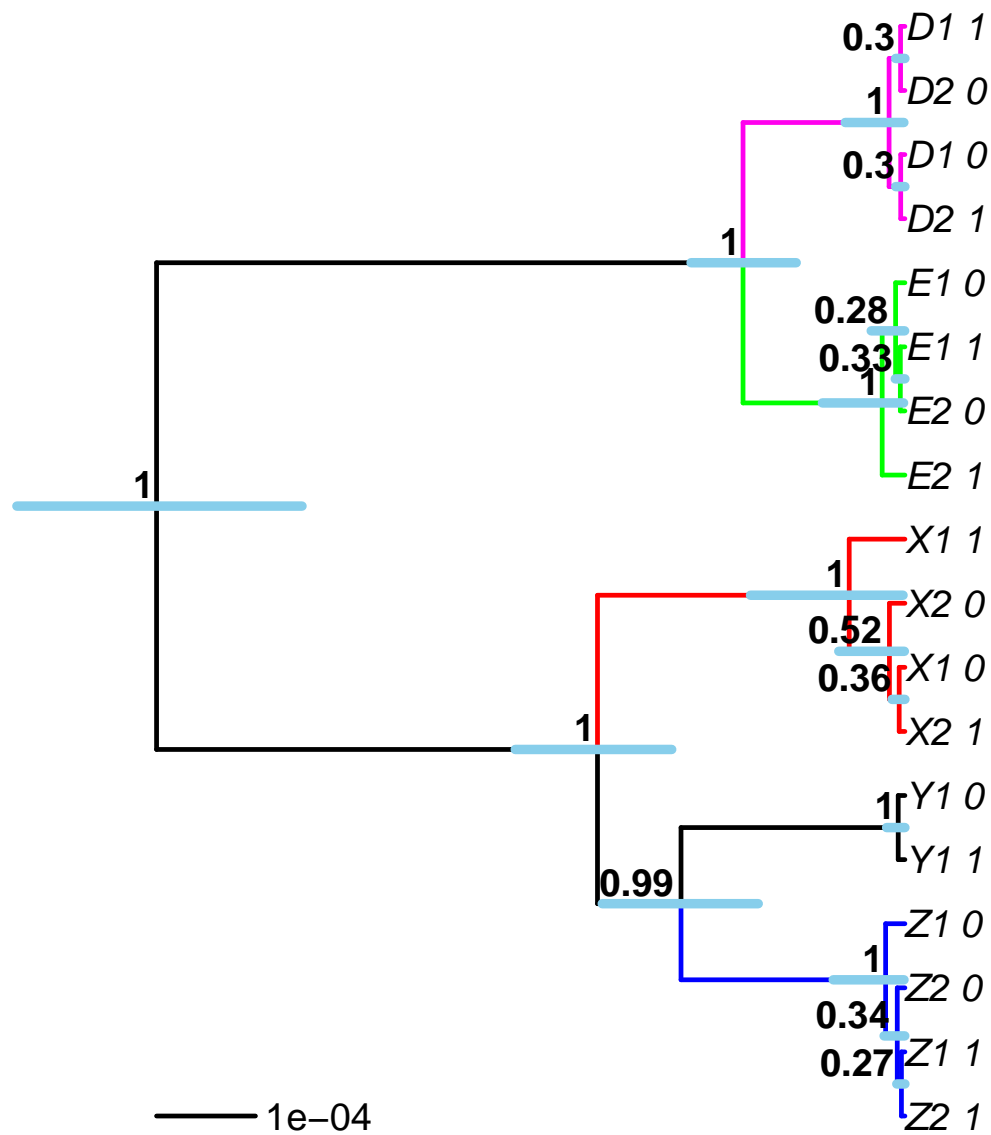
Sim-rep 5: Contig sequence alignments (n=150), STACEY species tree



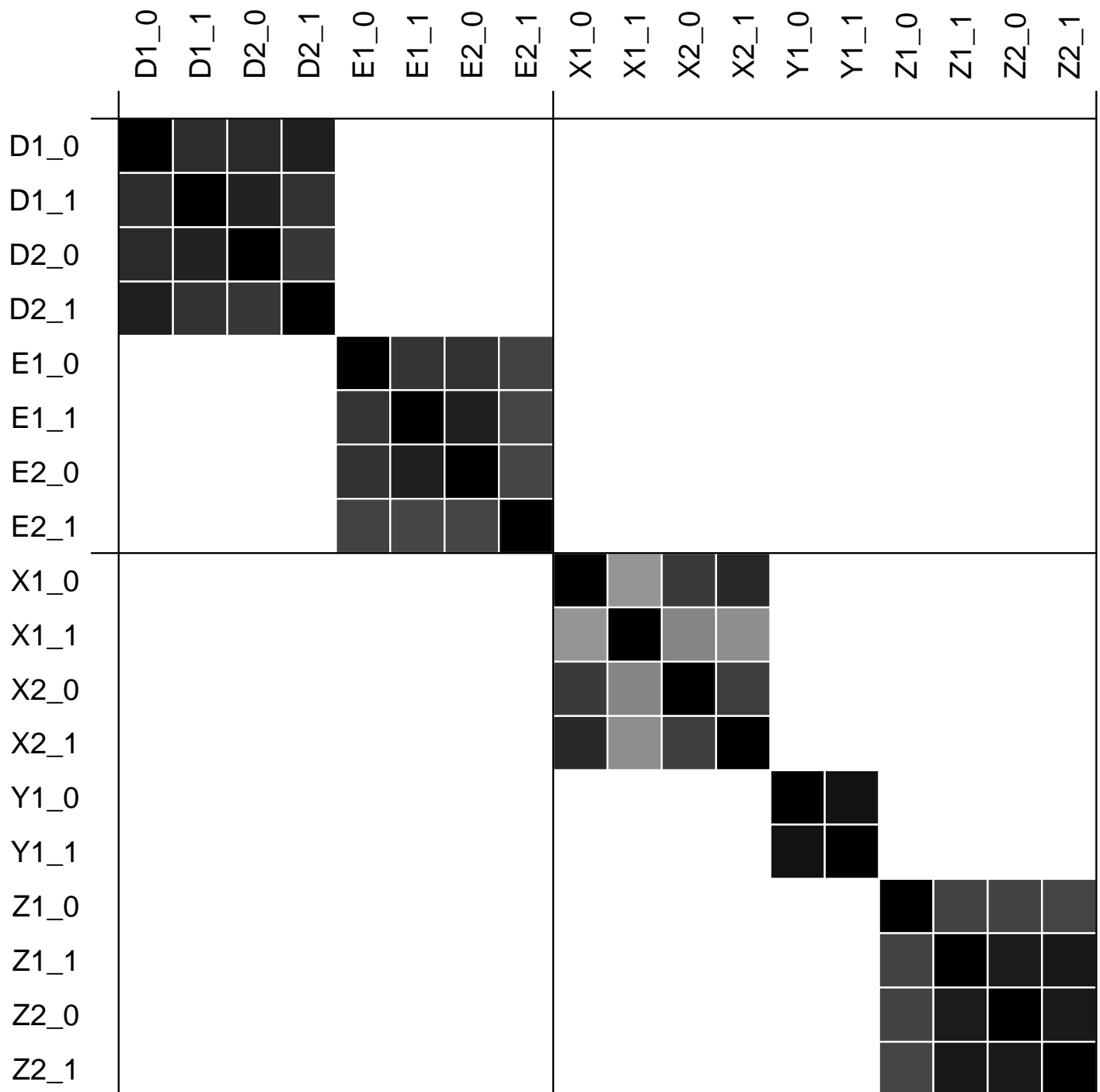
Sim-rep 5: Contig sequence alignments (n=150), STACEY similarity matrix



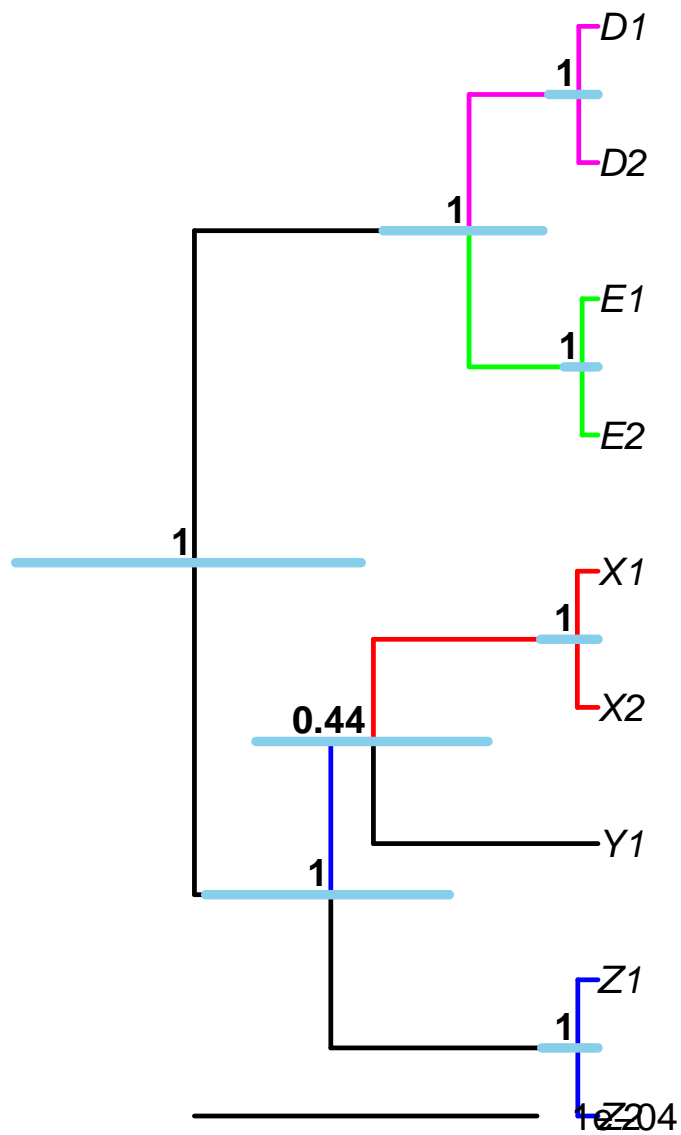
Sim-rep 5: Phased allele alignments (n=150), STACEY species tree



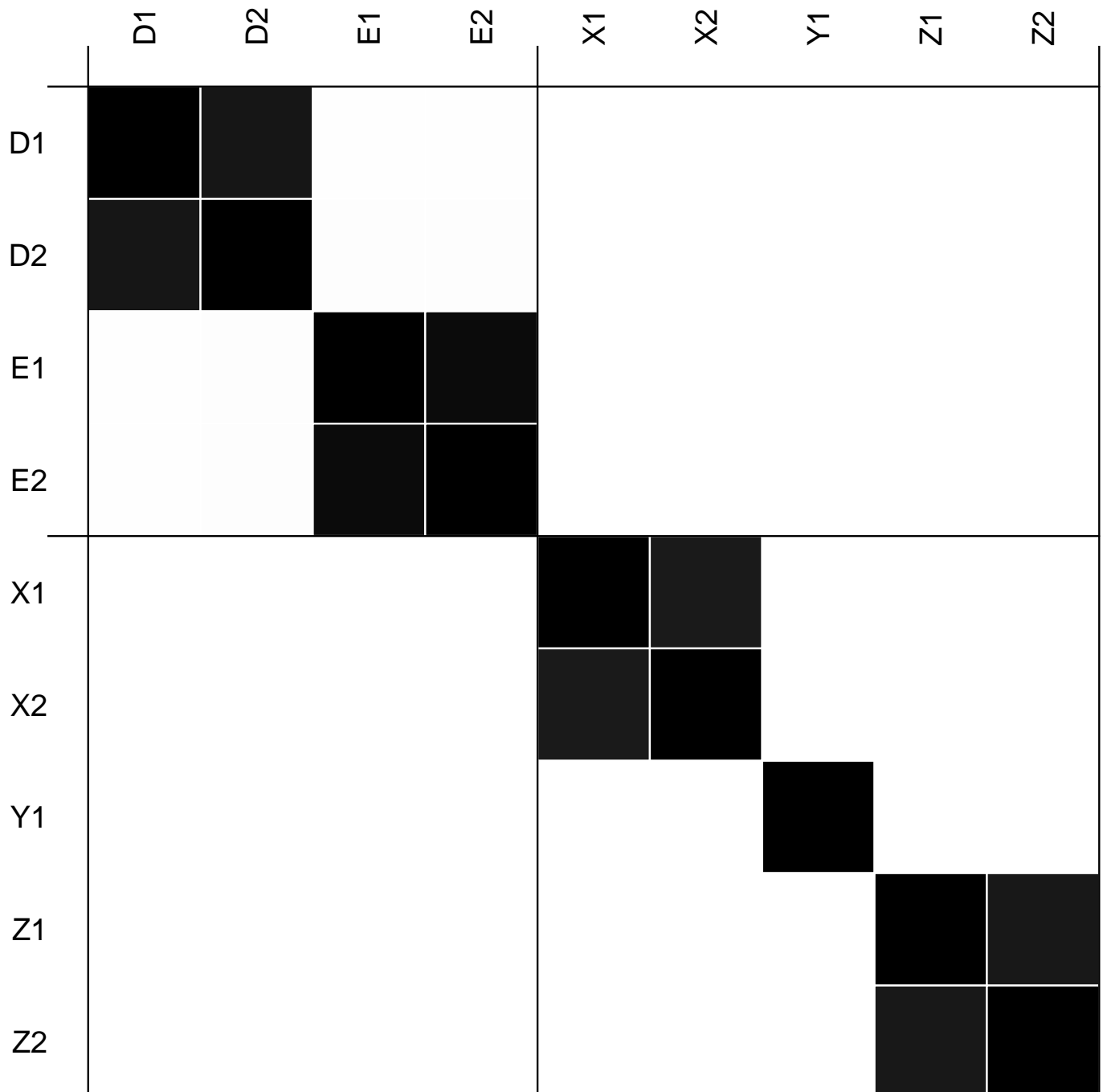
Sim-rep 5: Phased allele alignments (n=150), STACEY similarity matrix



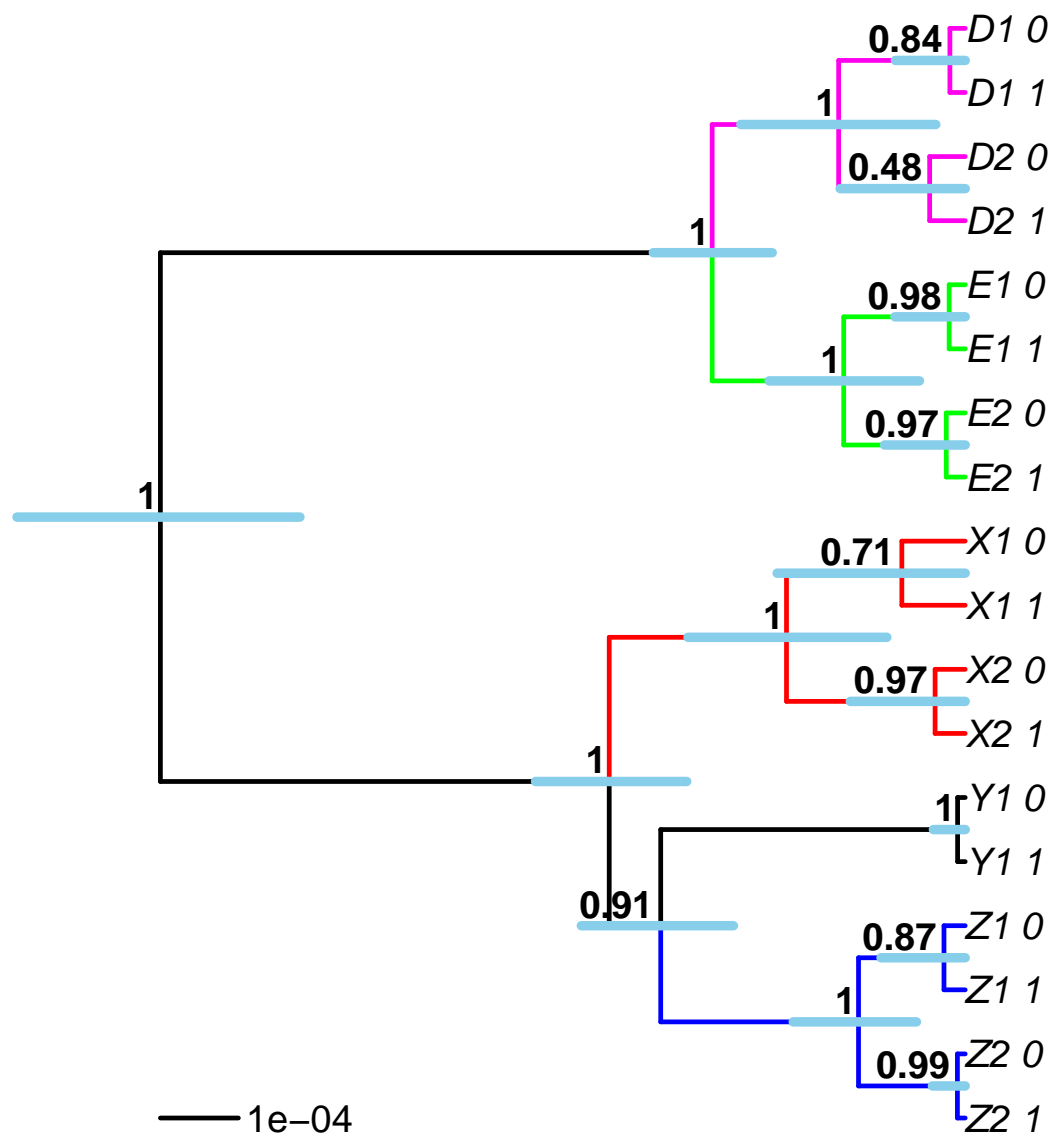
Sim-rep 5: IUPAC consensus alignments (n=150), STACEY species tree



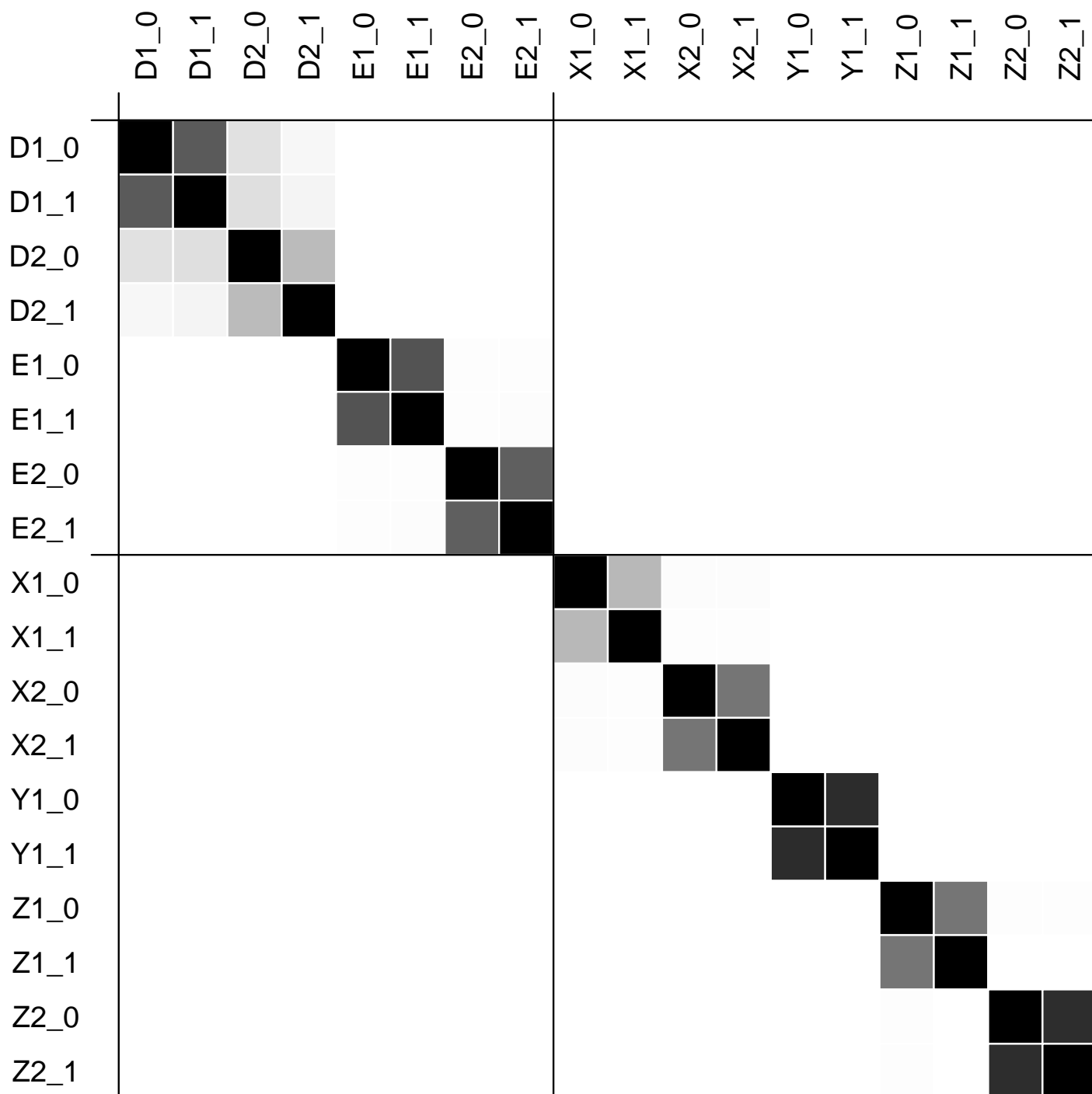
Sim-rep 5: IUPAC consensus alignments (n=150), STACEY similarity matrix



Sim-rep 5: Chimeric allele alignments (n=150), STACEY species tree

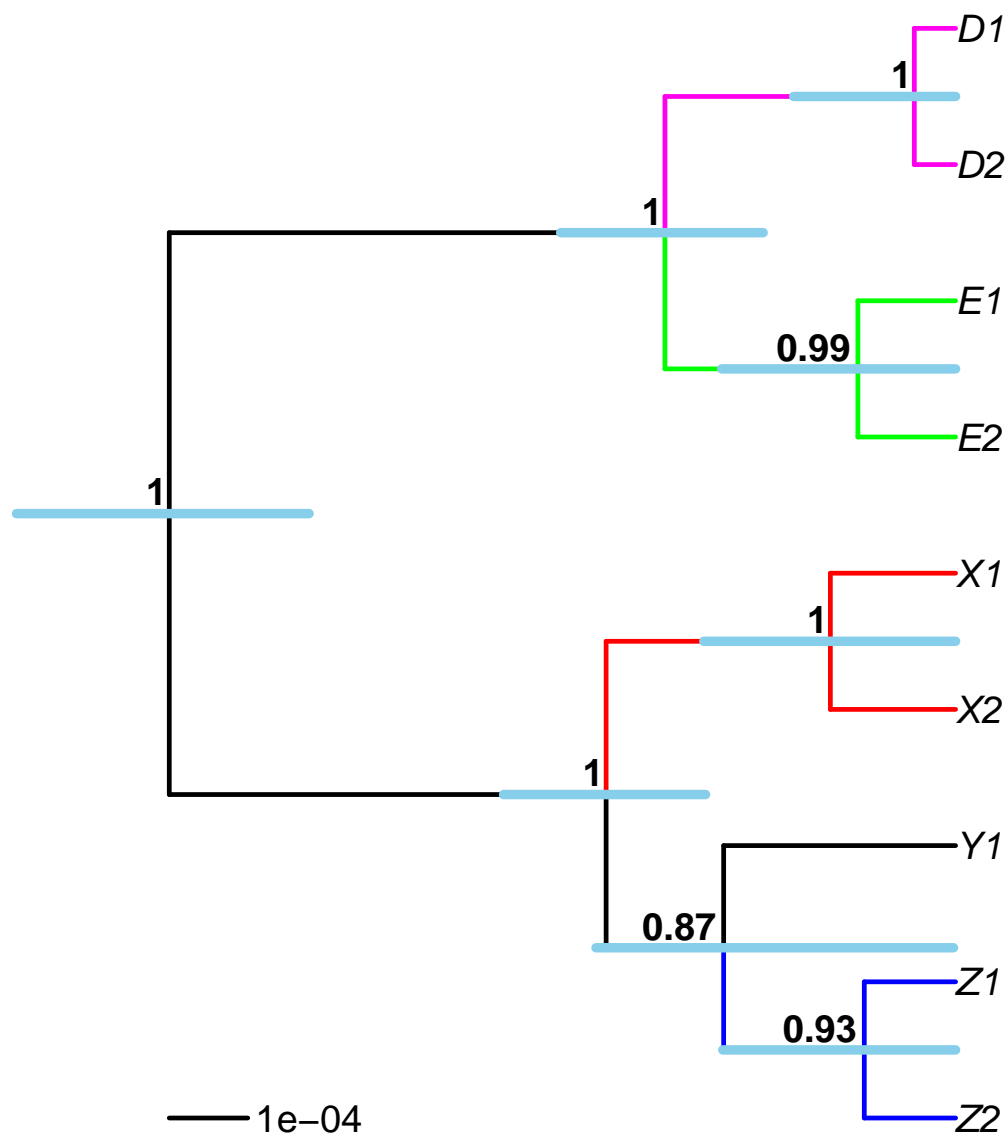


Sim-rep 5: Chimeric allele alignments (n=150), STACEY similarity matrix

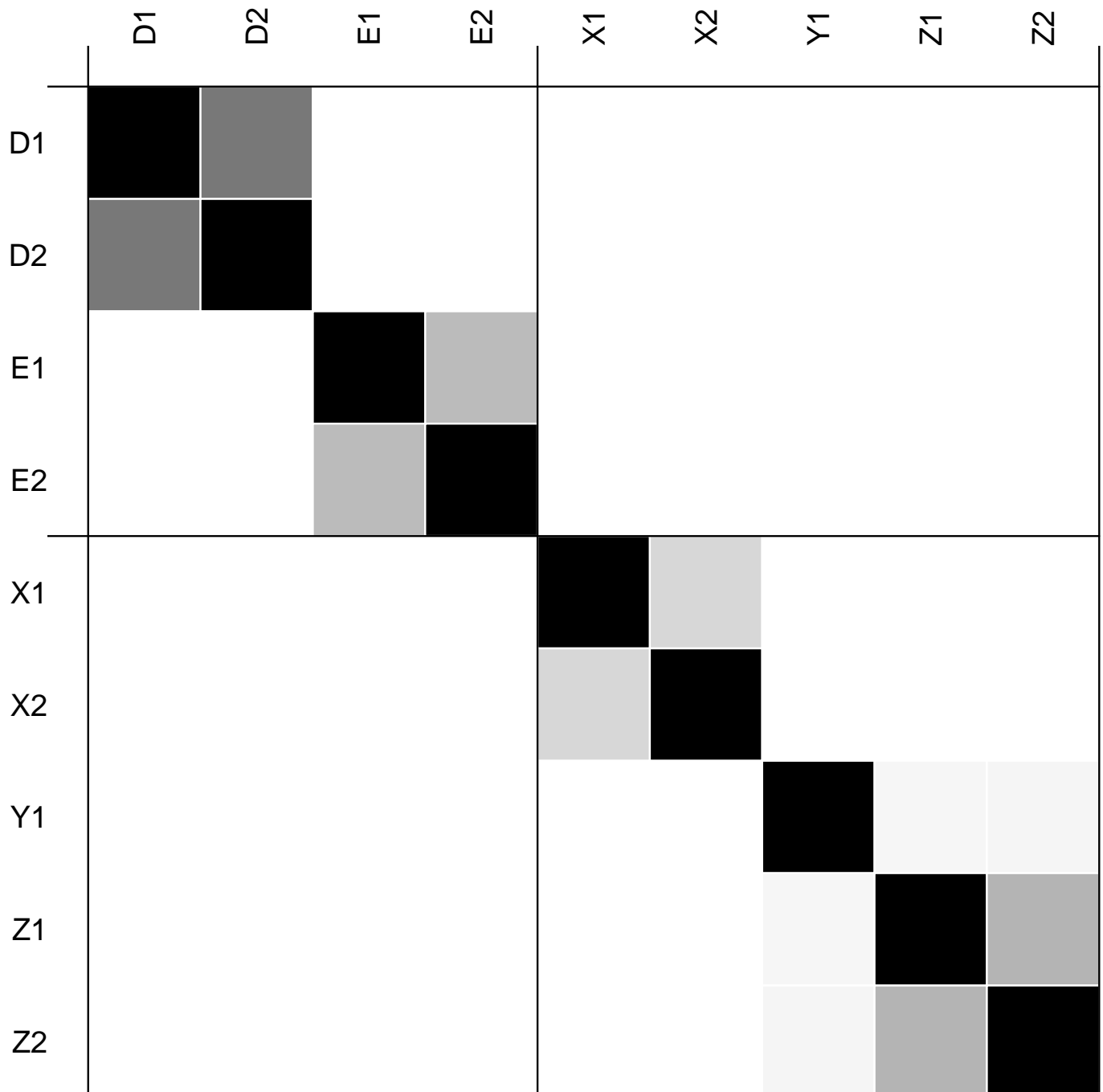




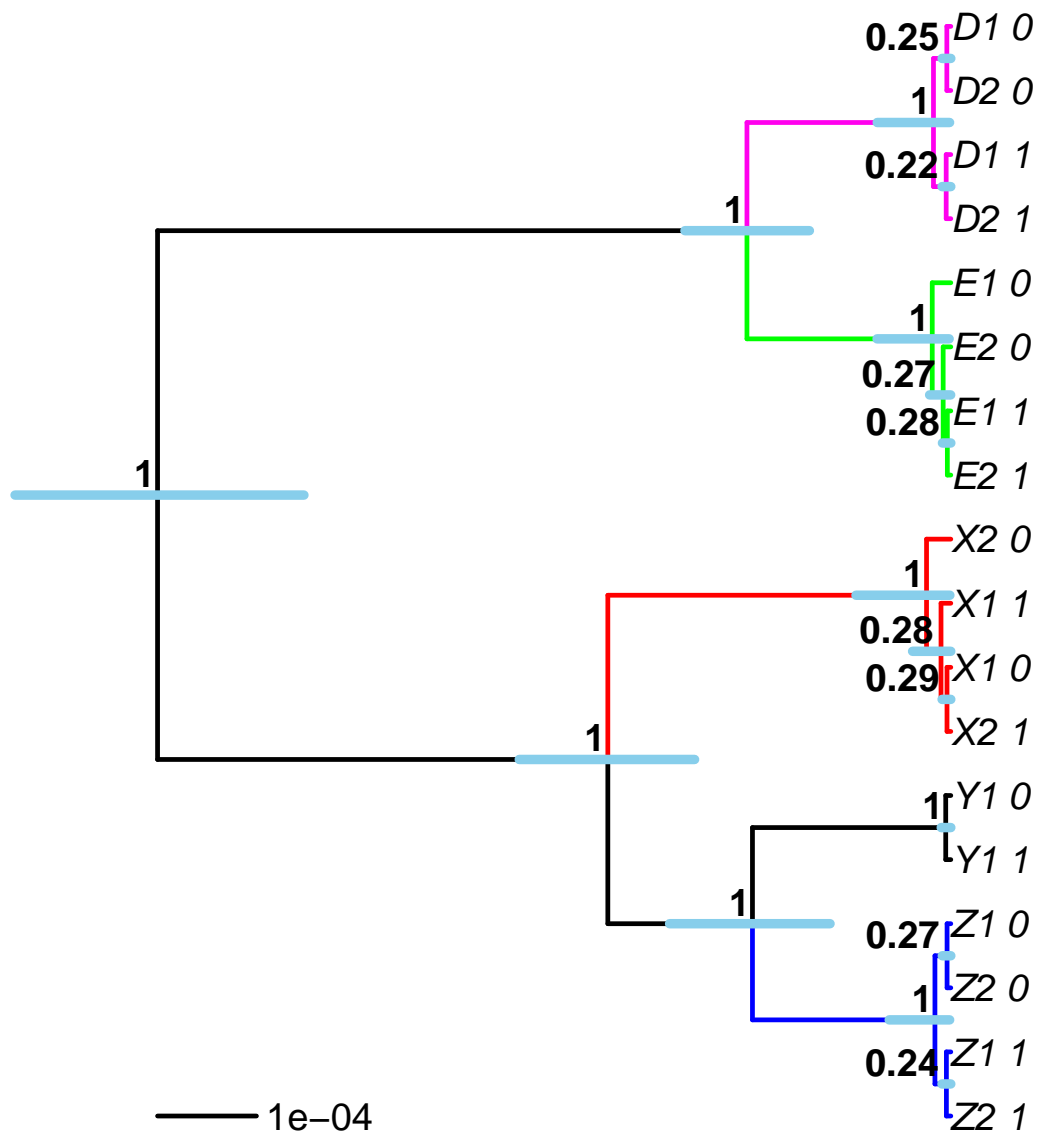
Sim-rep 6: Contig sequence alignments (n=150), STACEY species tree



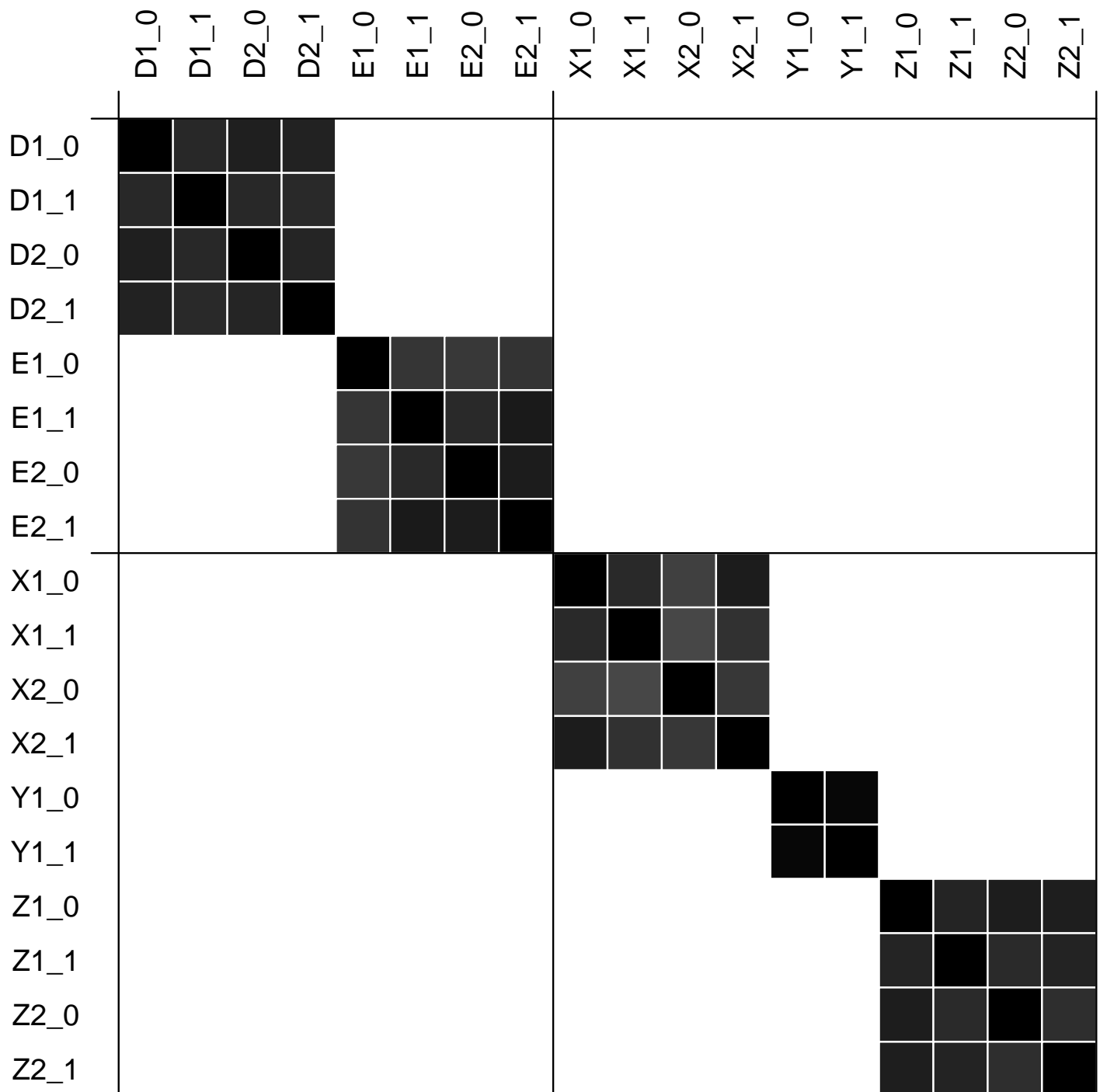
Sim-rep 6: Contig sequence alignments (n=150), STACEY similarity matrix



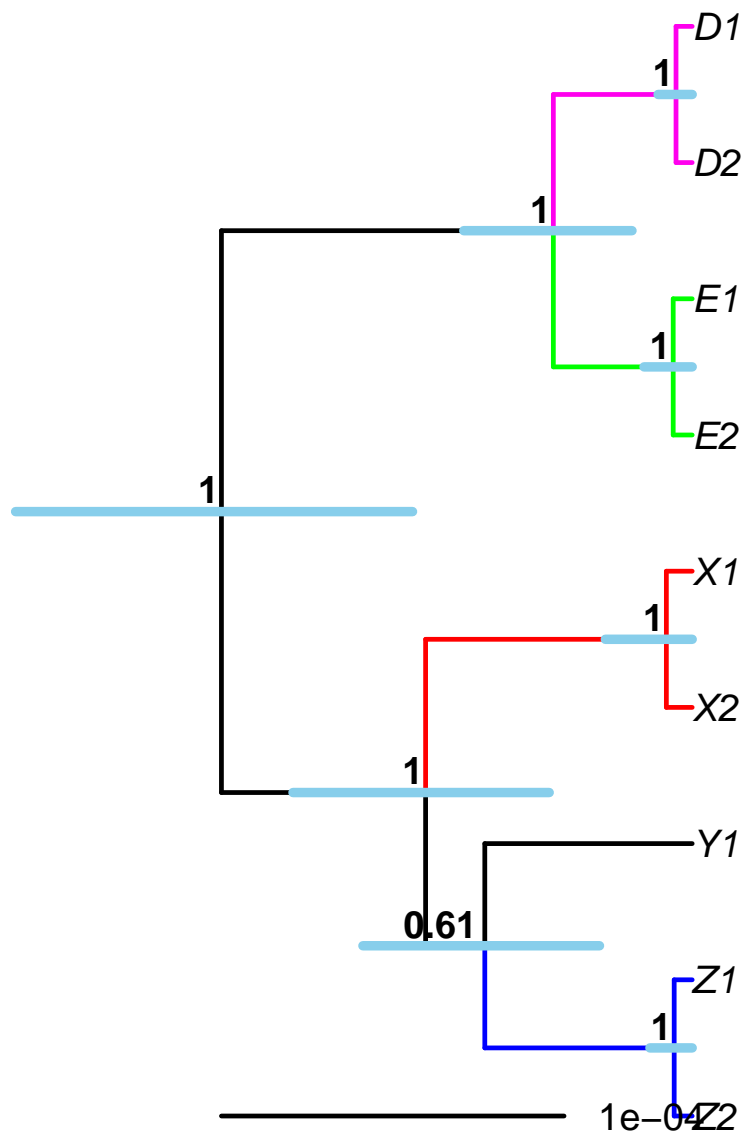
Sim-rep 6: Phased allele alignments (n=150), STACEY species tree



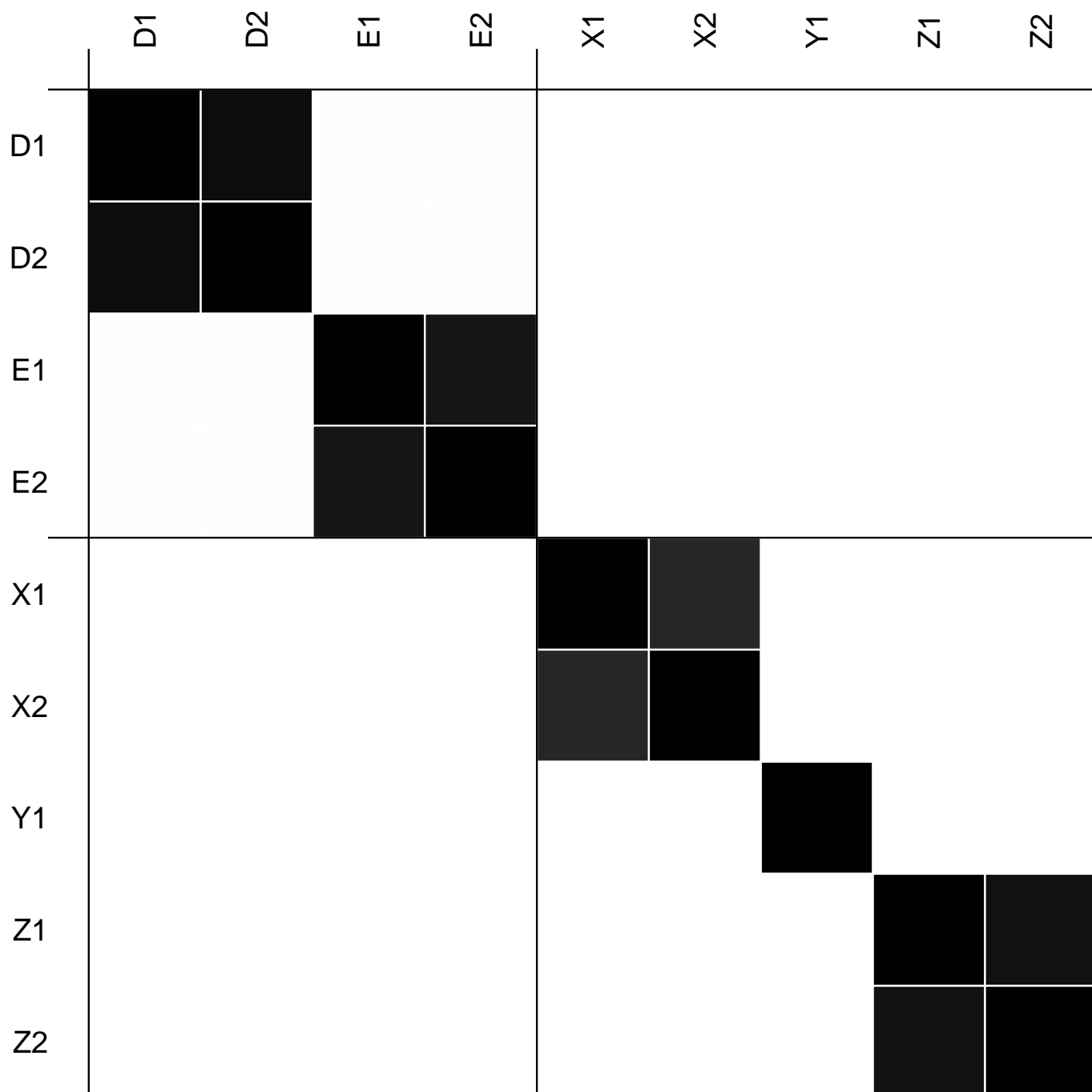
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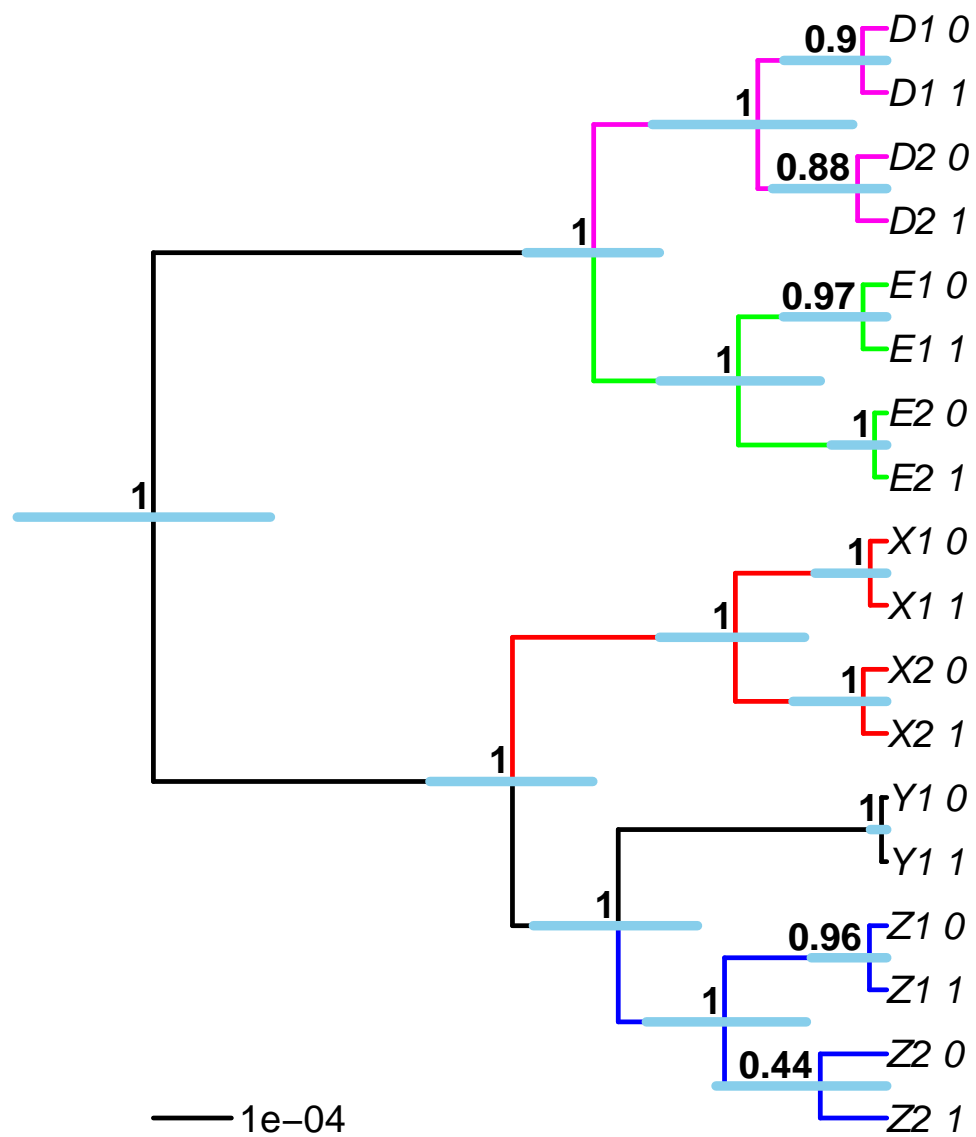
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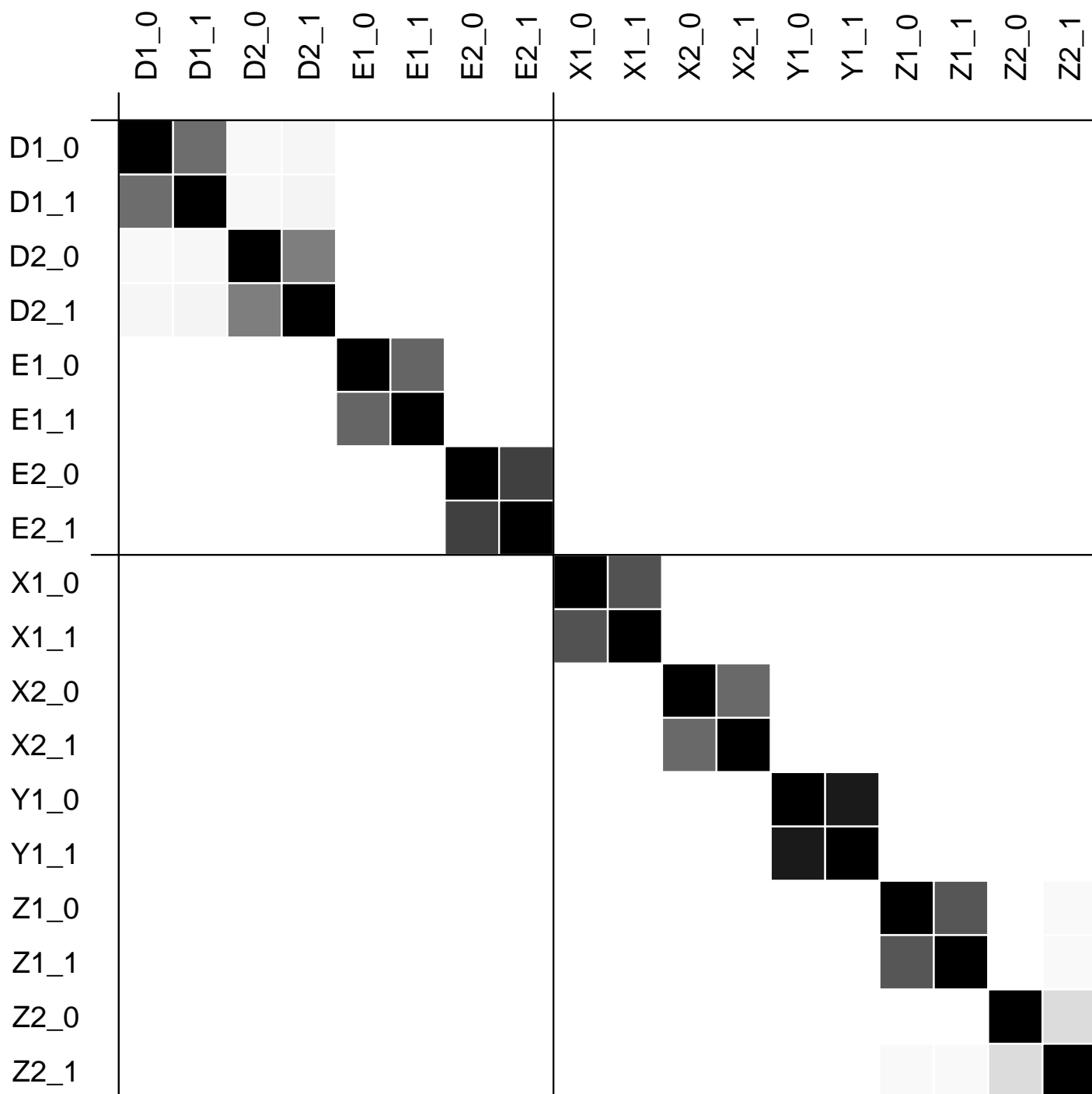
Sim-rep 6: IUPAC consensus alignments (n=150), STACEY similarity matrix



Sim-rep 6: Chimeric allele alignments (n=150), STACEY species tree

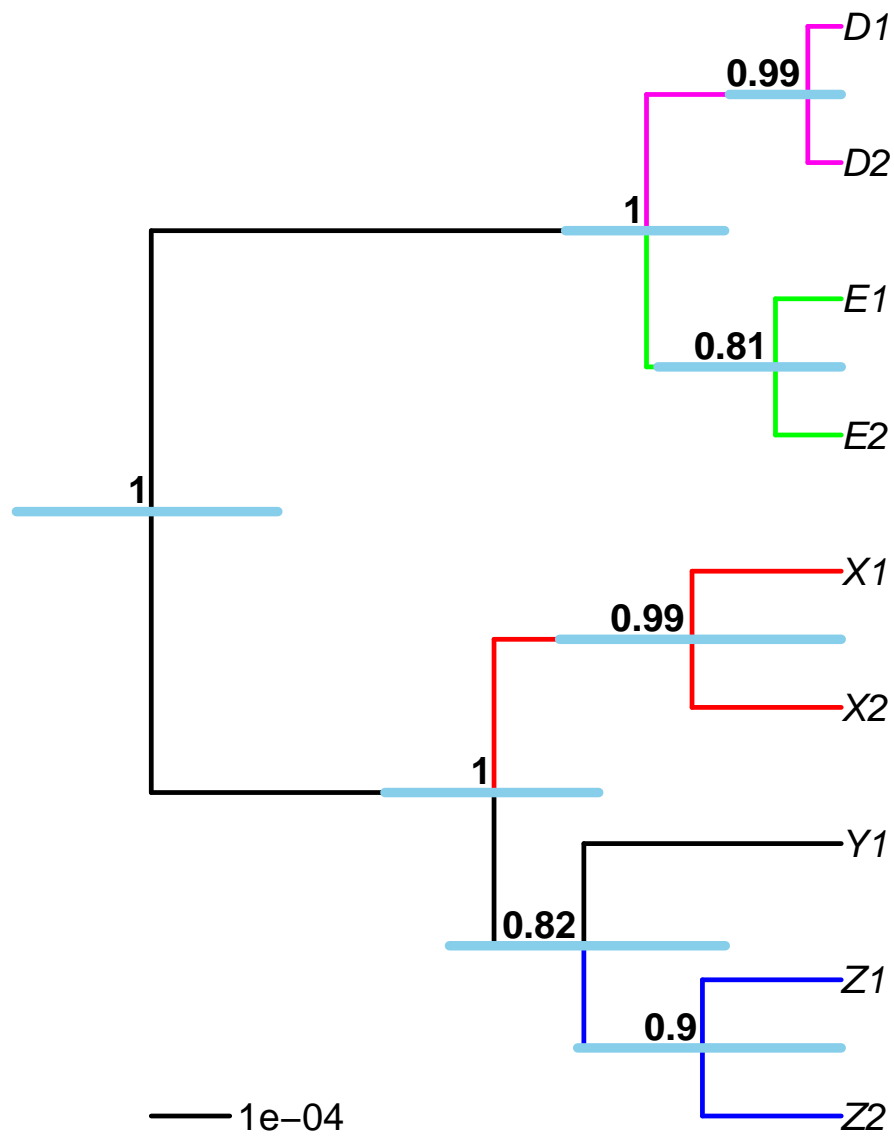


Sim-rep 6: Chimeric allele alignments (n=150), STACEY similarity matrix

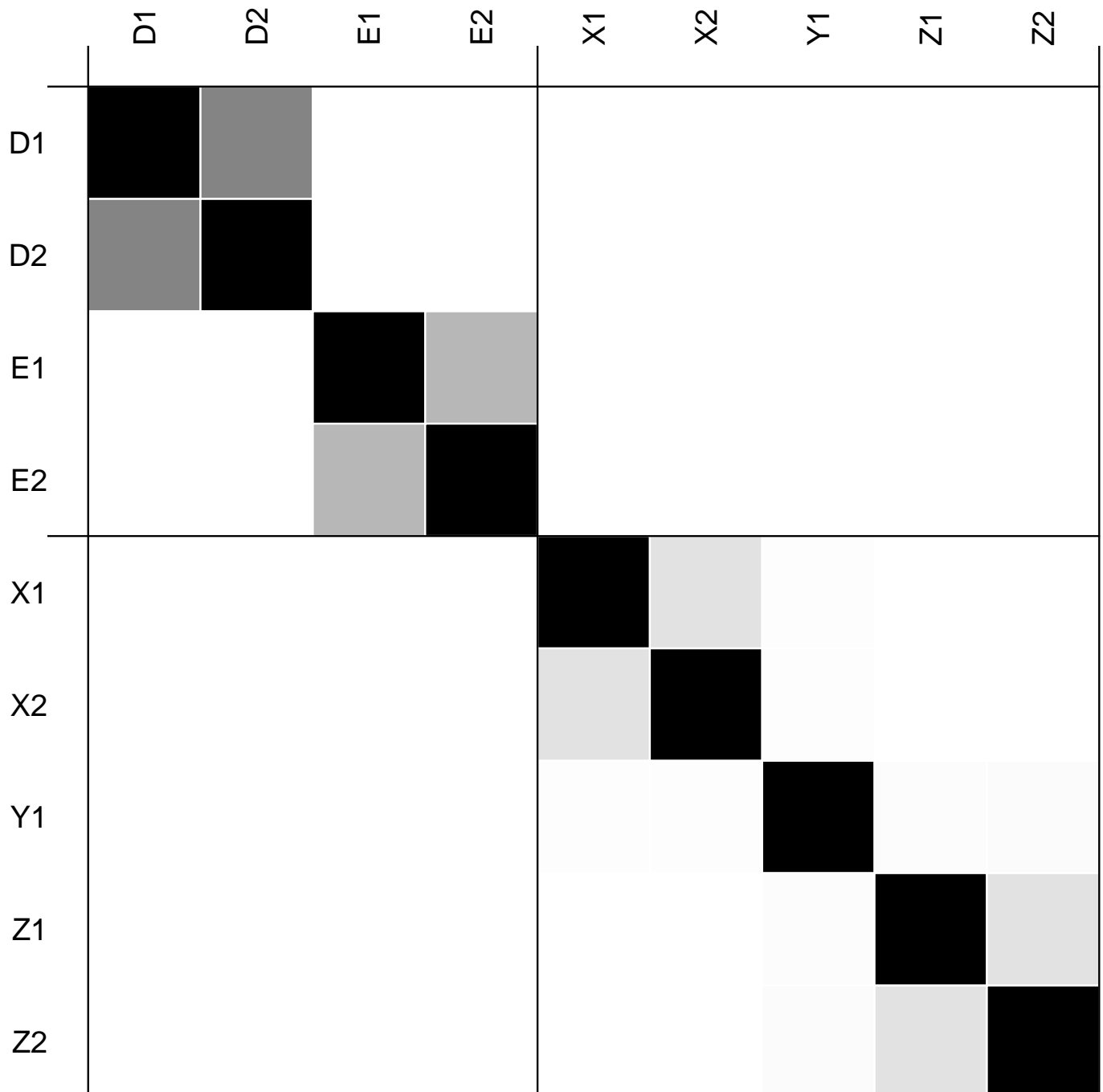




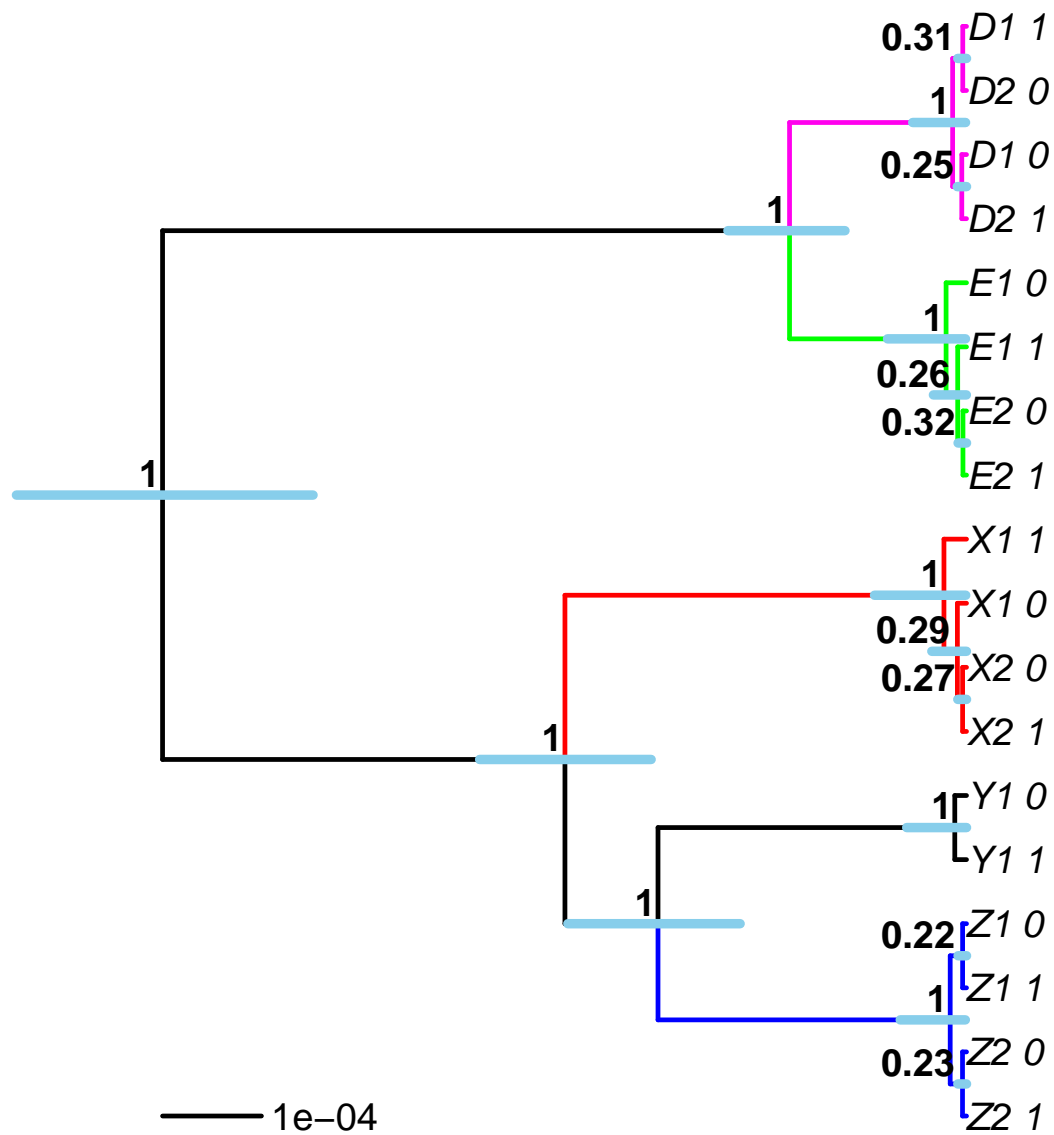
Sim-rep 7: Contig sequence alignments (n=150), STACEY species tree



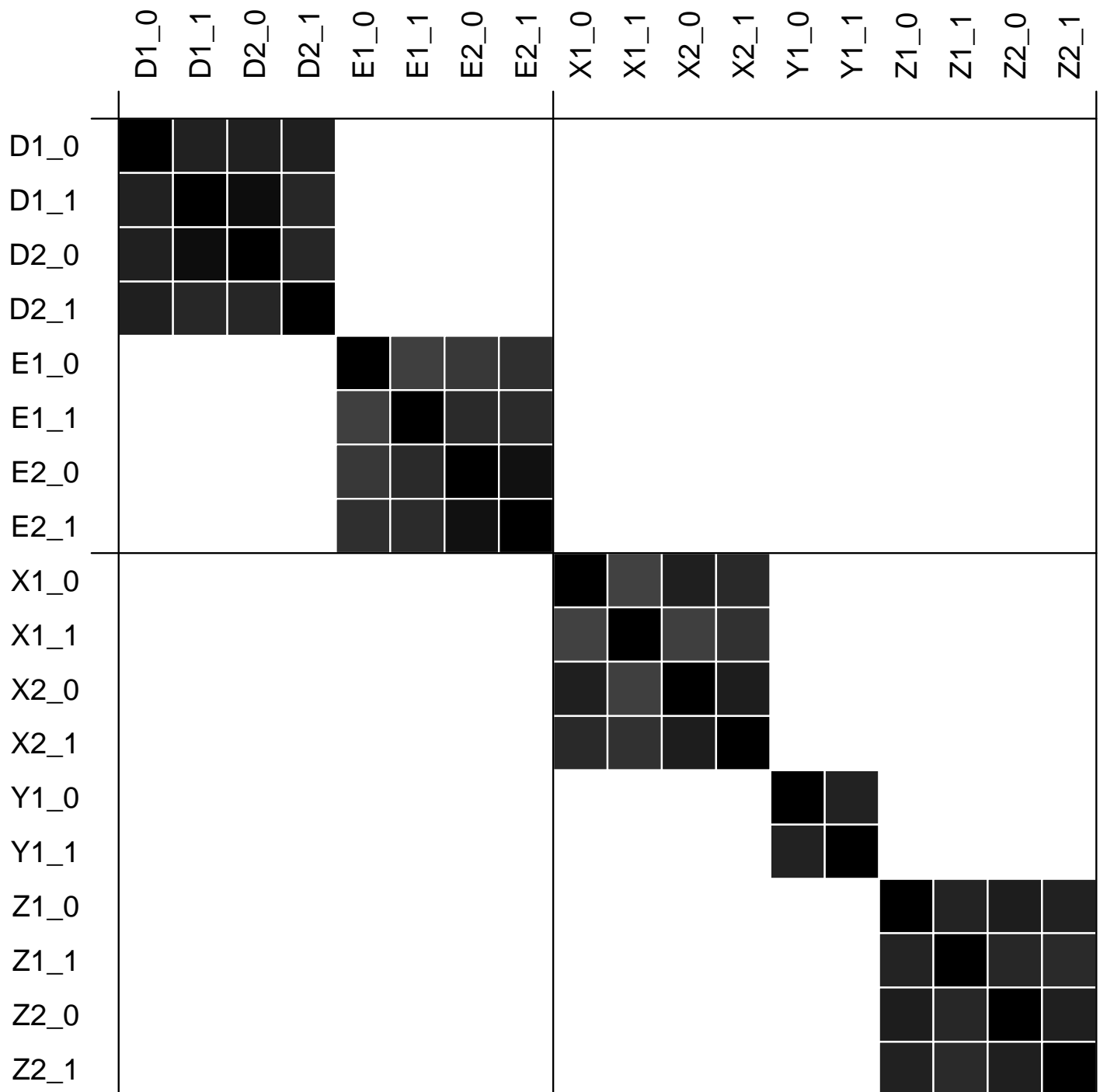
Sim-rep 7: Contig sequence alignments (n=150), STACEY similarity matrix



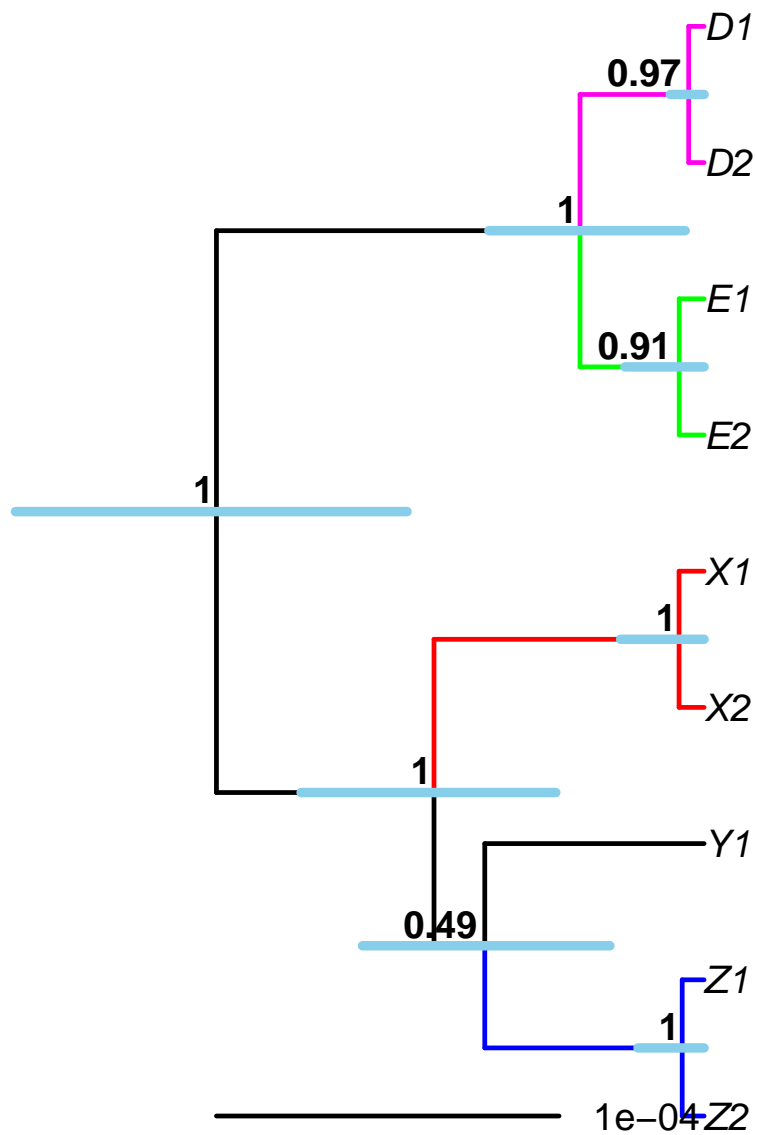
Sim-rep 7: Phased allele alignments (n=150), STACEY species tree



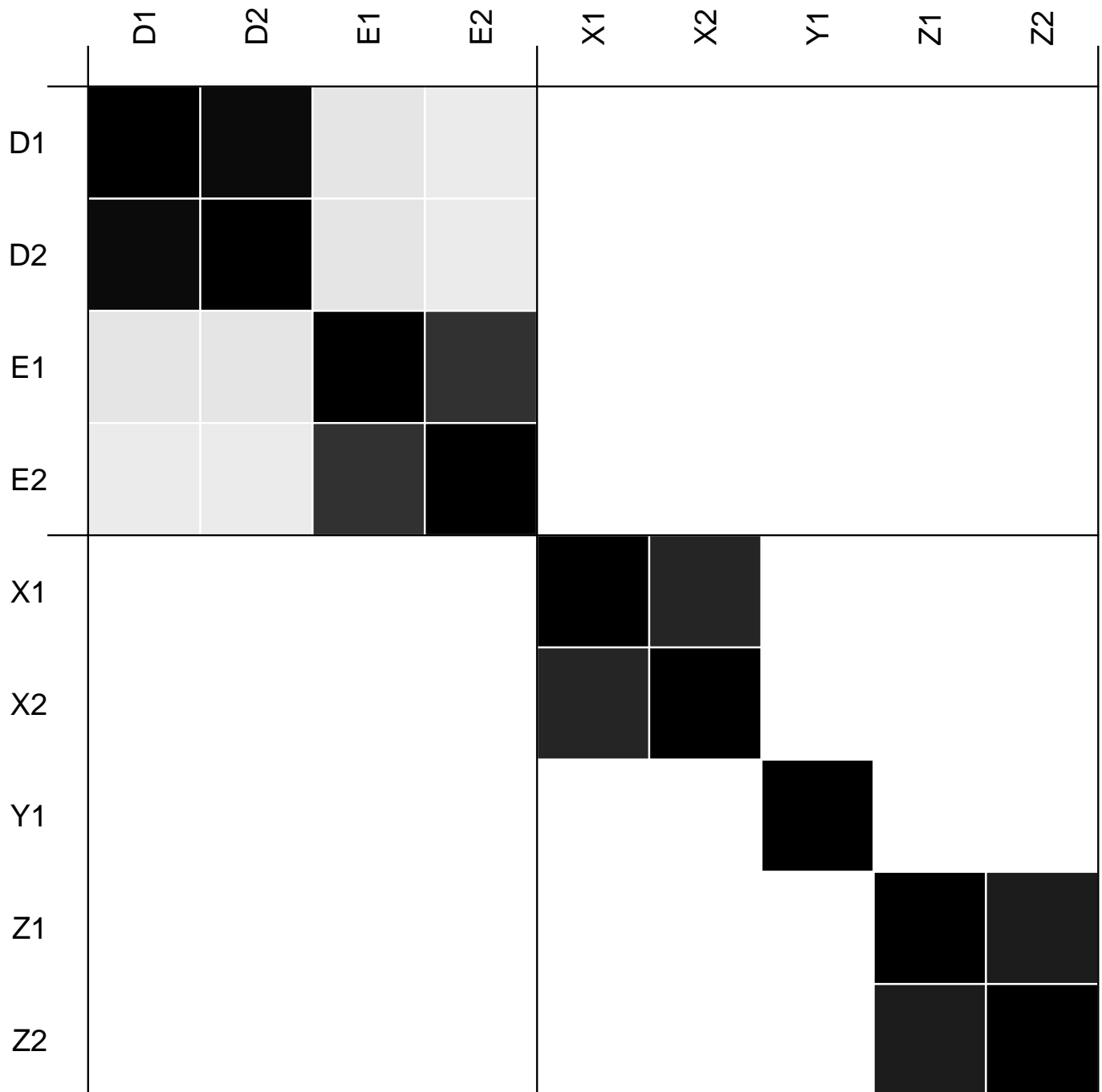
Sim-rep 7: Phased allele alignments (n=150), STACEY similarity matrix



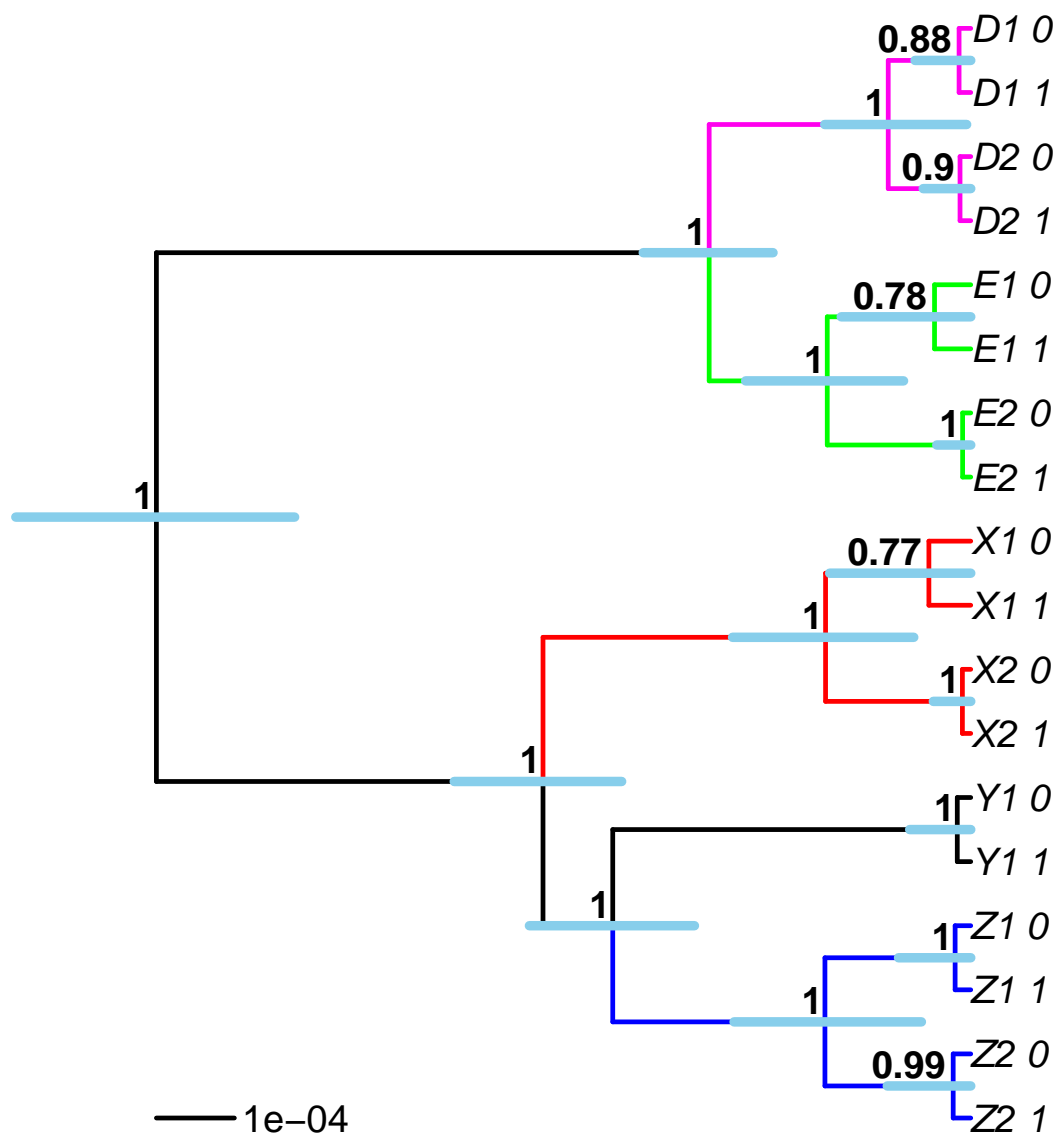
Sim-rep 7: IUPAC consensus alignments (n=150), STACEY species tree



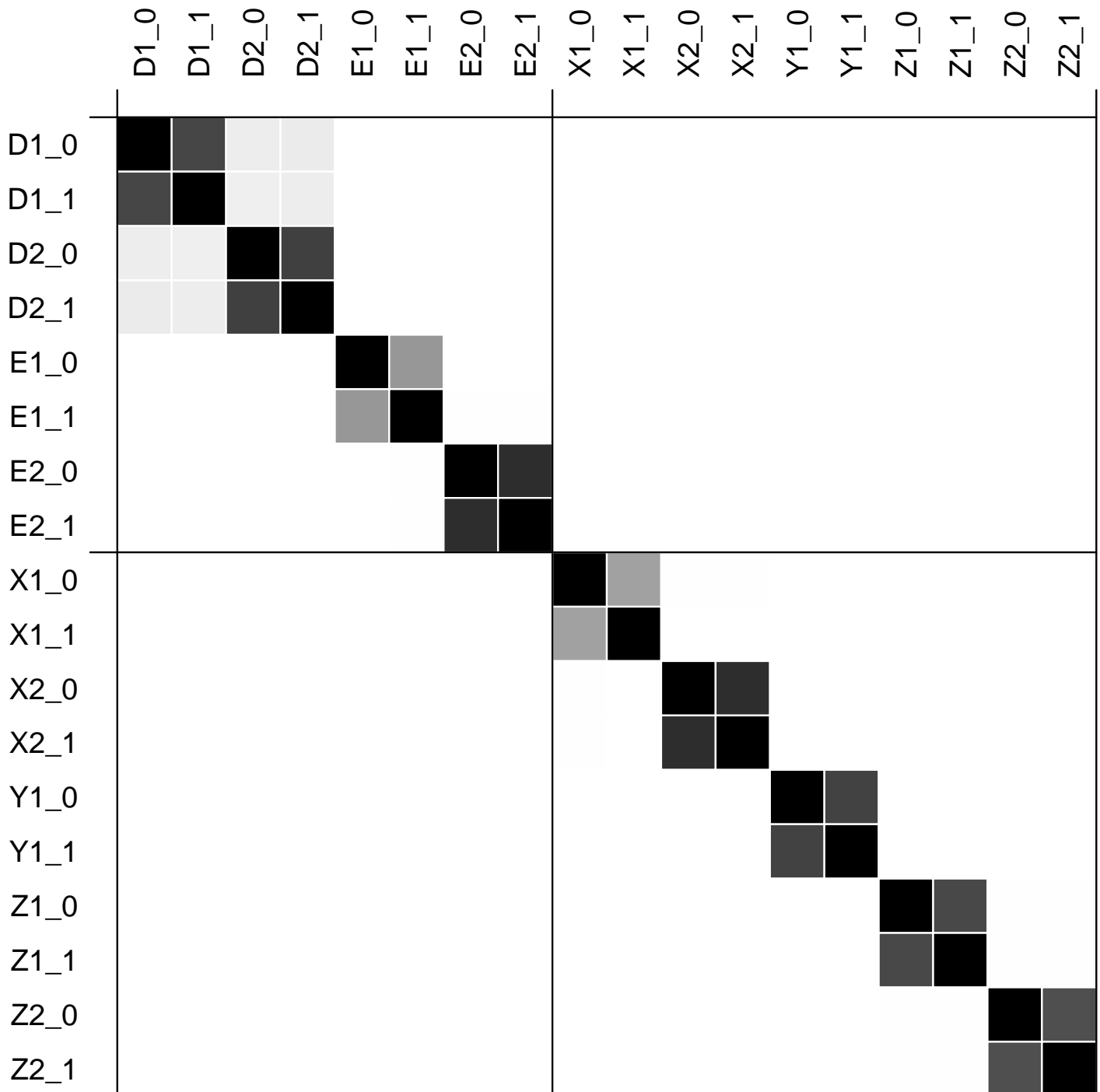
Sim-rep 7: IUPAC consensus alignments (n=150), STACEY similarity matrix



Sim-rep 7: Chimeric allele alignments (n=150), STACEY species tree

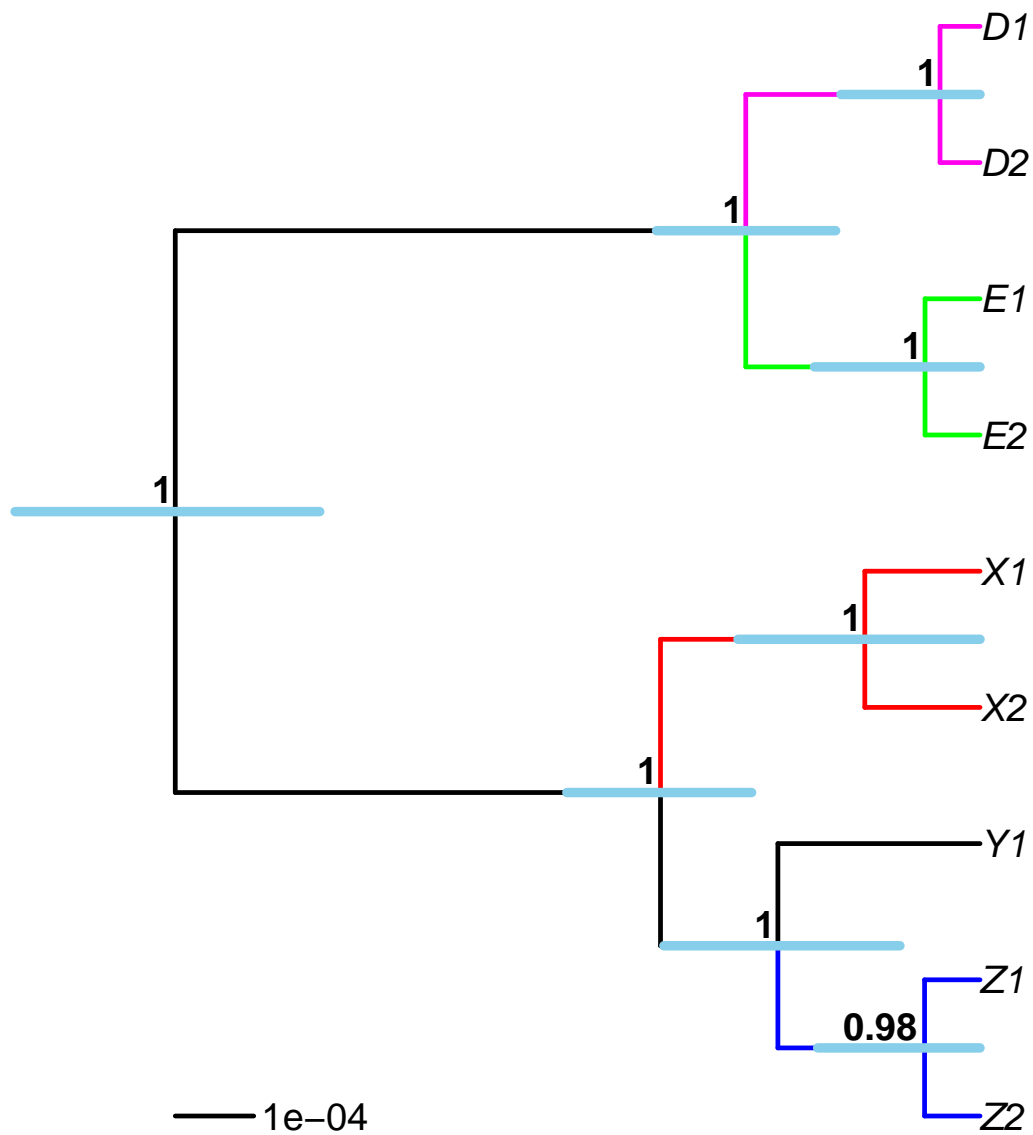


Sim-rep 7: Chimeric allele alignments (n=150), STACEY similarity matrix

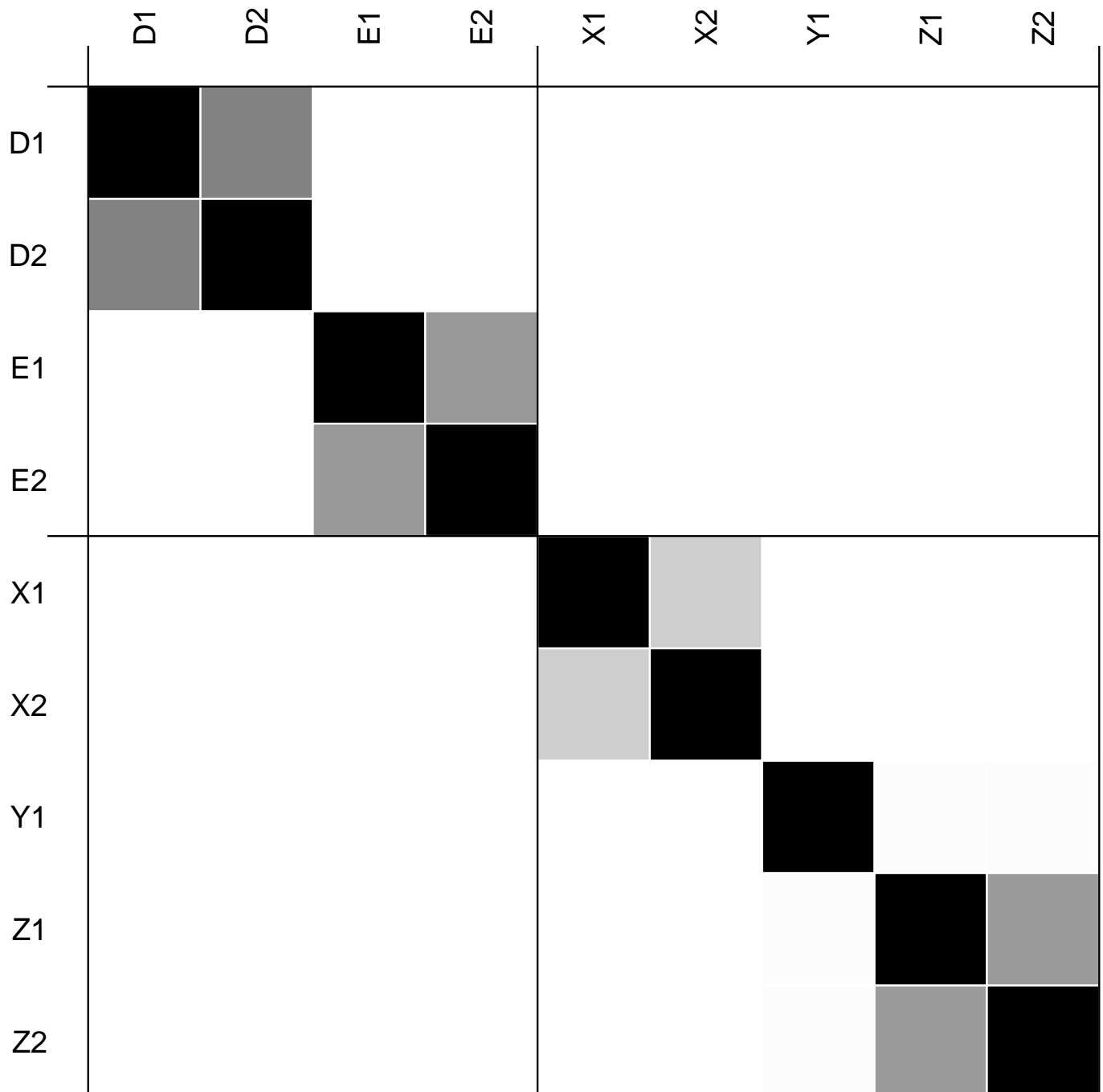




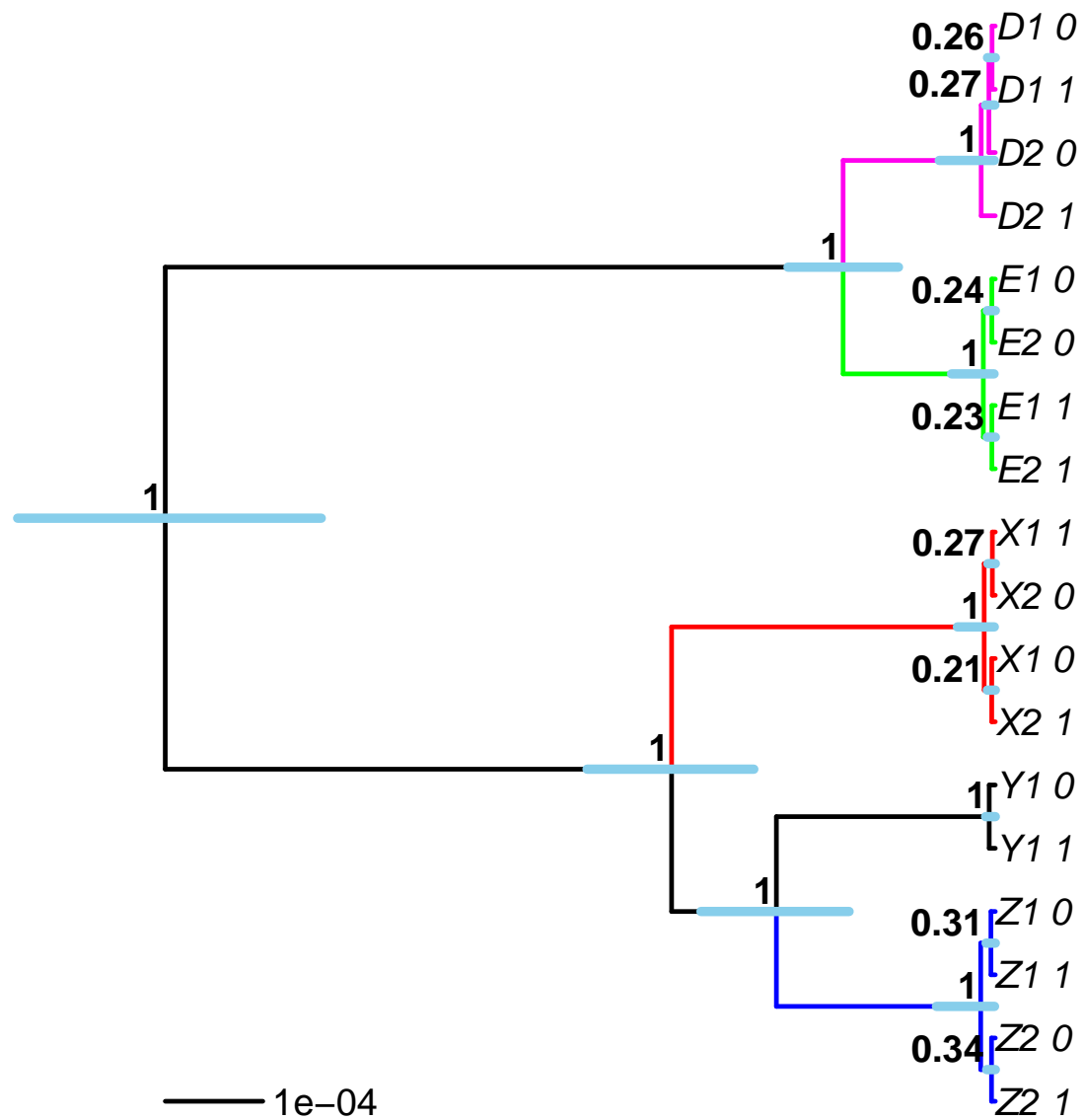
Sim-rep 8: Contig sequence alignments (n=150), STACEY species tree



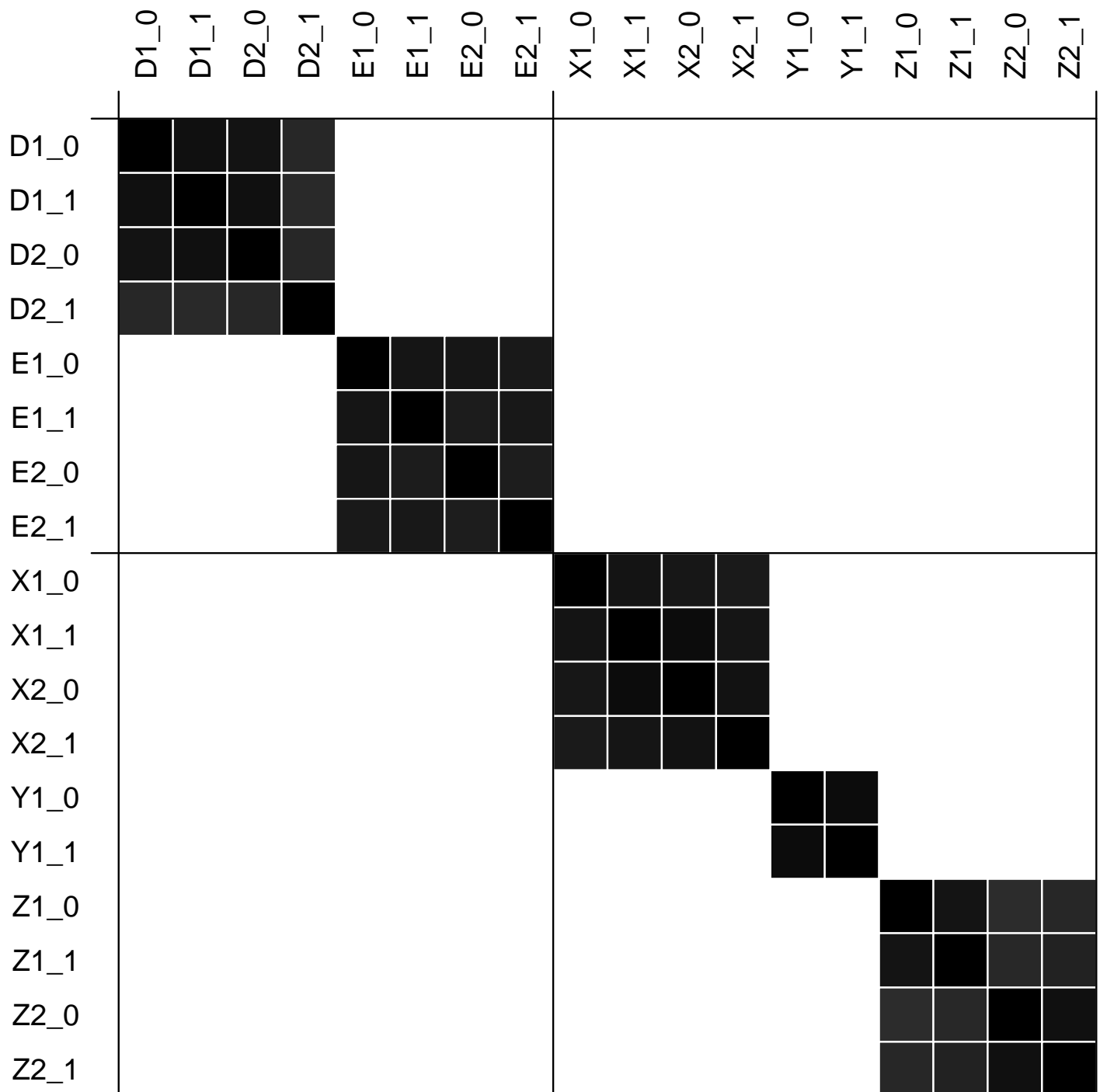
Sim-rep 8: Contig sequence alignments (n=150), STACEY similarity matrix



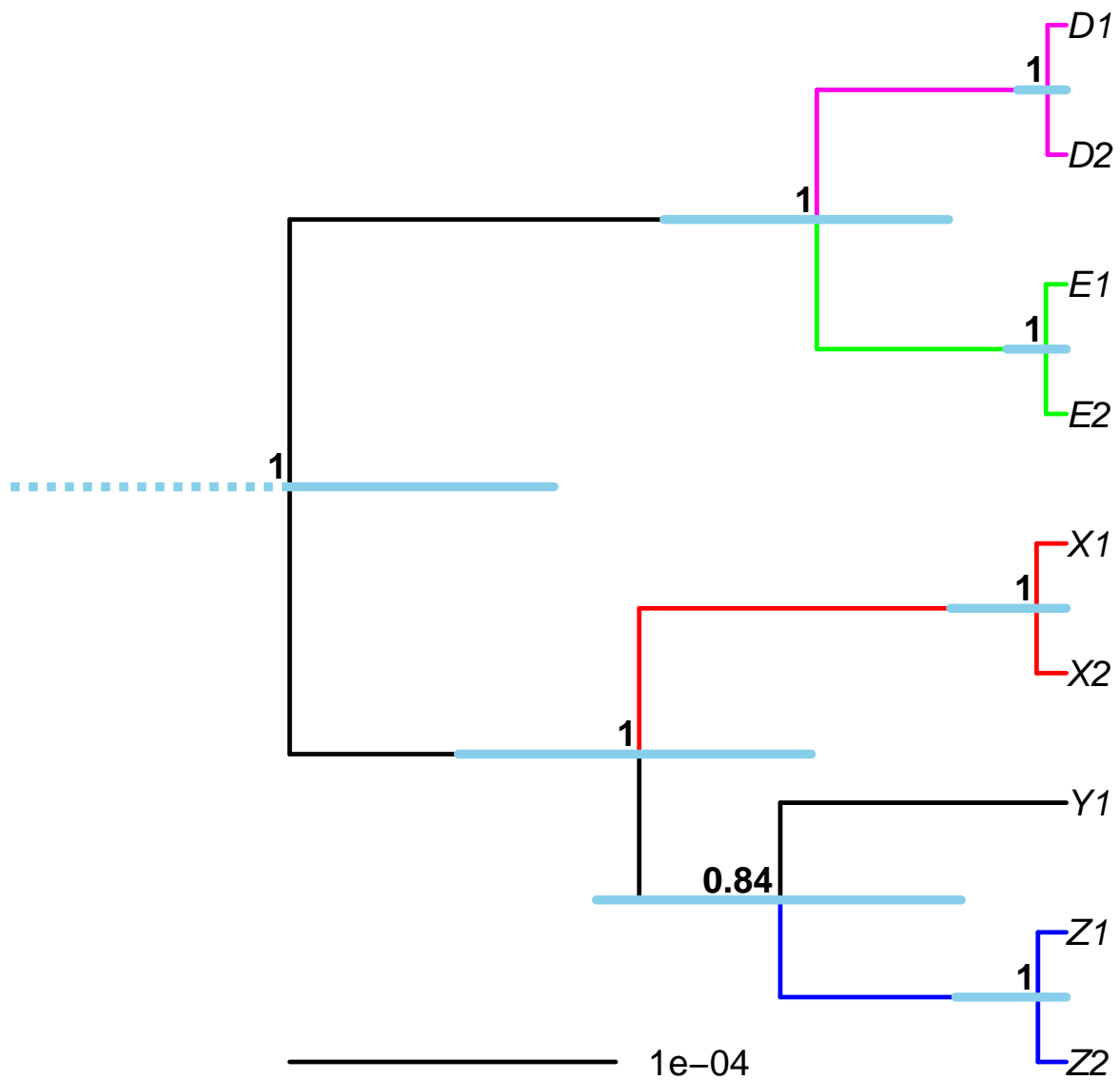
Sim-rep 8: Phased allele alignments (n=150), STACEY species tree



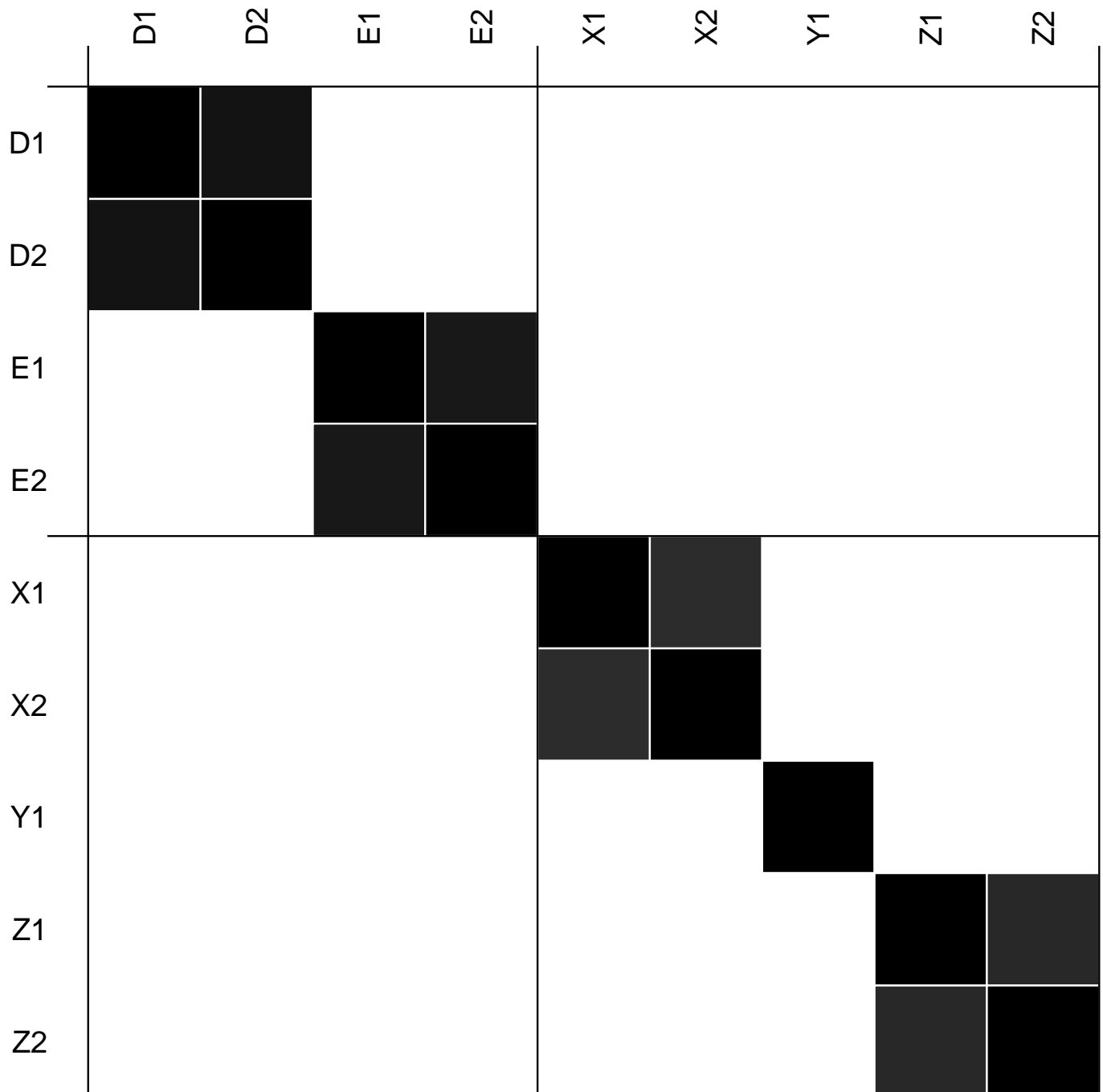
Sim-rep 8: Phased allele alignments (n=150), STACEY similarity matrix



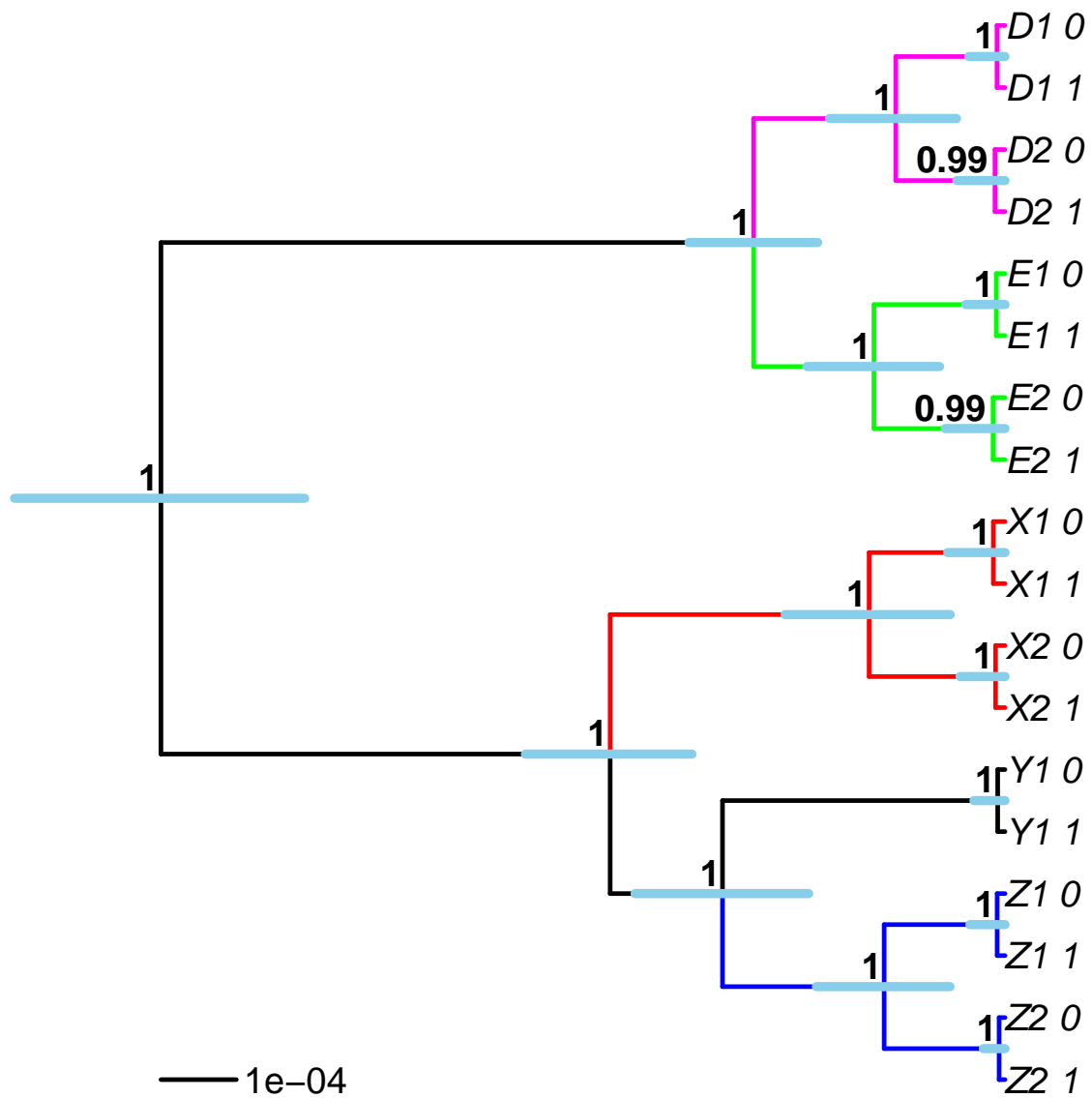
Sim-rep 8: IUPAC consensus alignments (n=150), STACEY species tree



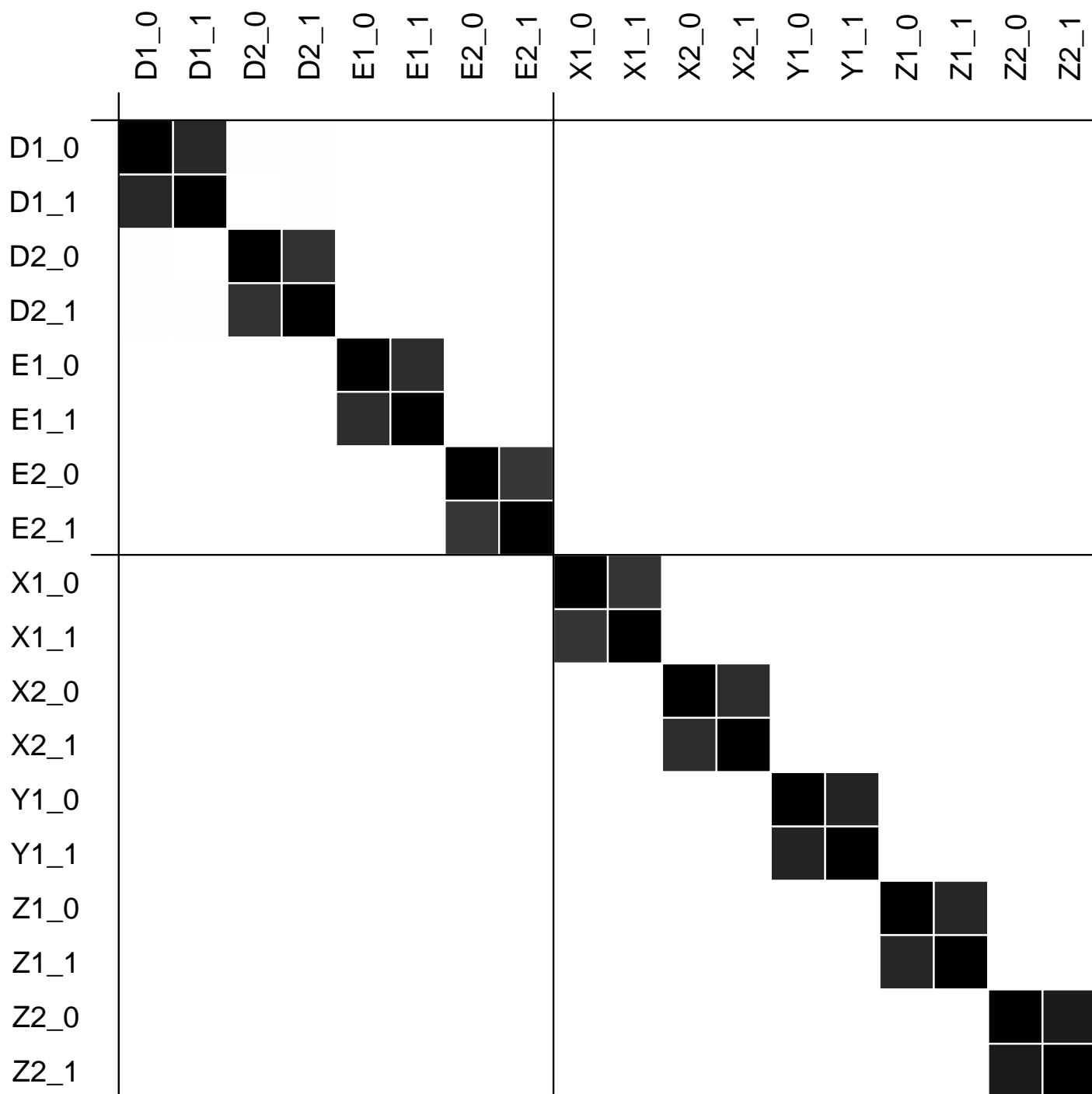
Sim-rep 8: IUPAC consensus alignments (n=150), STACEY similarity matrix



Sim-rep 8: Chimeric allele alignments (n=150), STACEY species tree

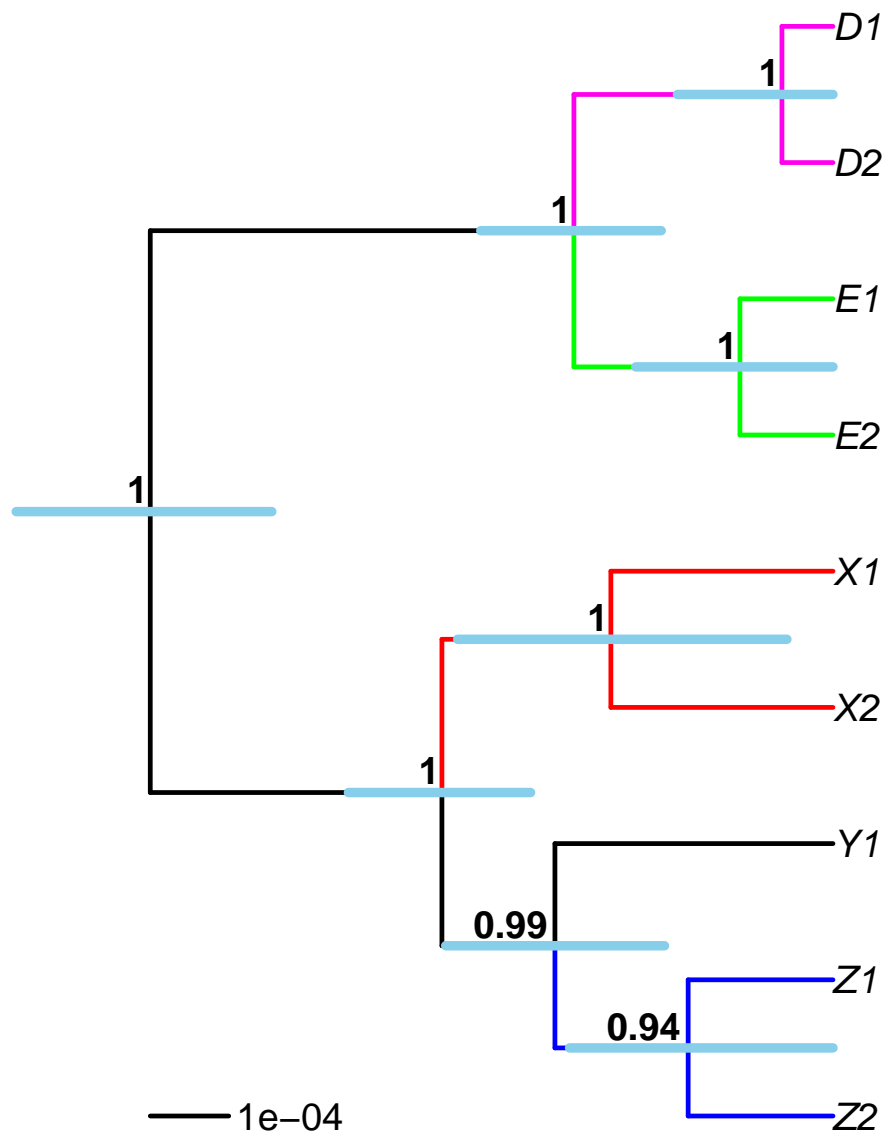


Sim-rep 8: Chimeric allele alignments (n=150), STACEY similarity matrix

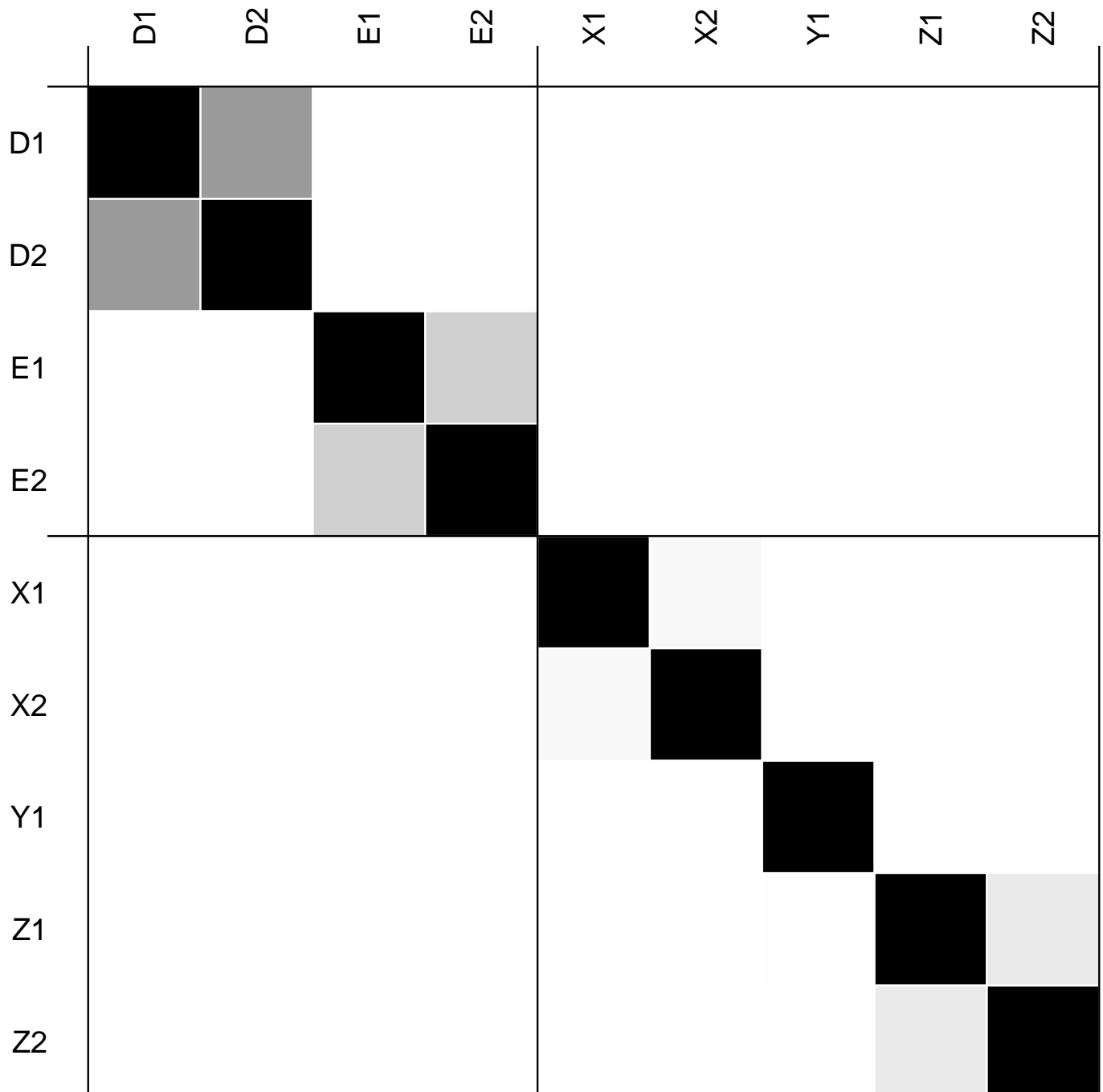




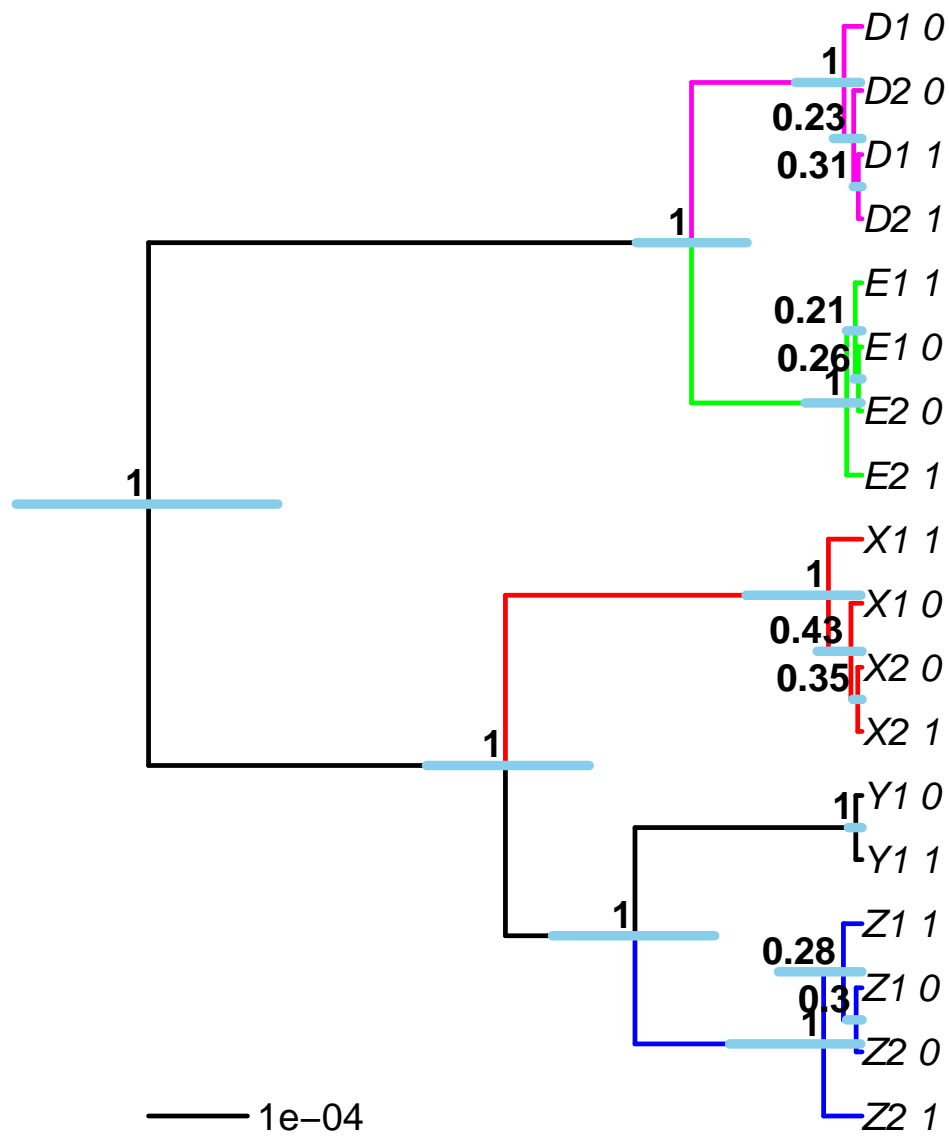
Sim-rep 9: Contig sequence alignments (n=150), STACEY species tree



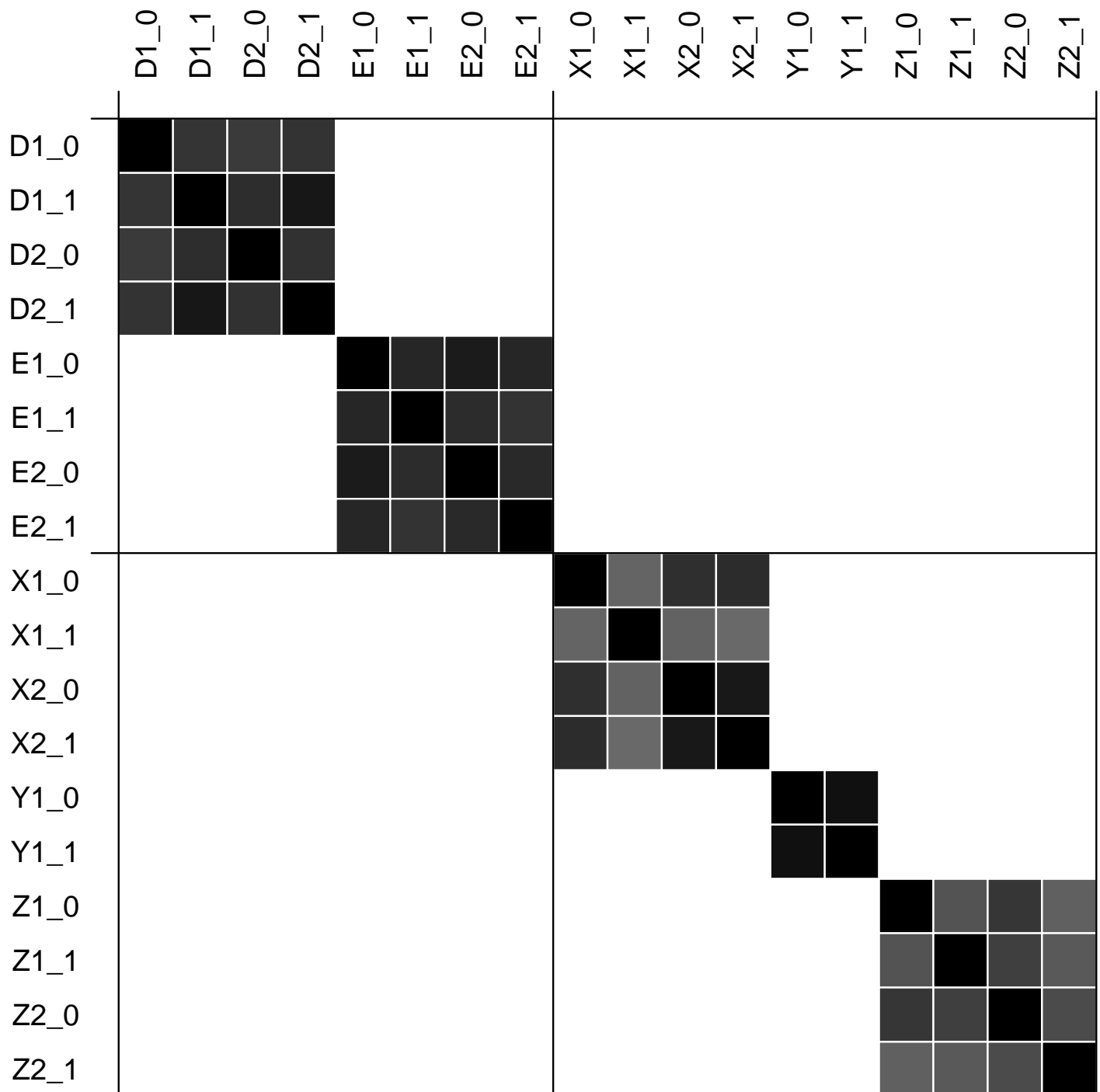
Sim-rep 9: Contig sequence alignments (n=150), STACEY similarity matrix



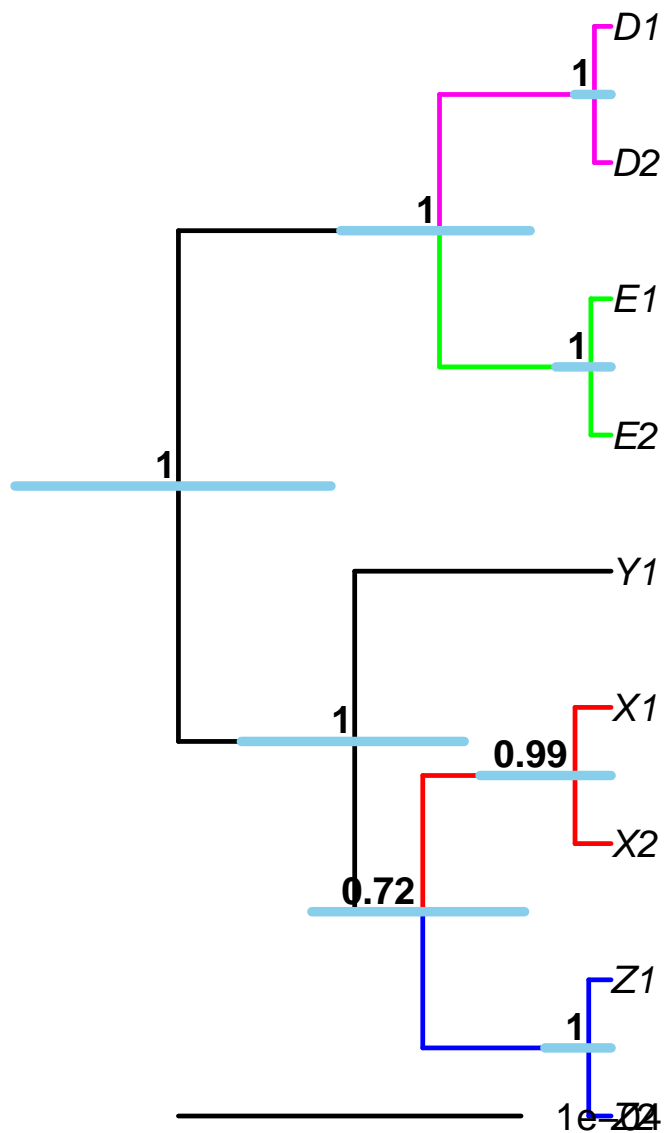
Sim-rep 9: Phased allele alignments (n=150), STACEY species tree



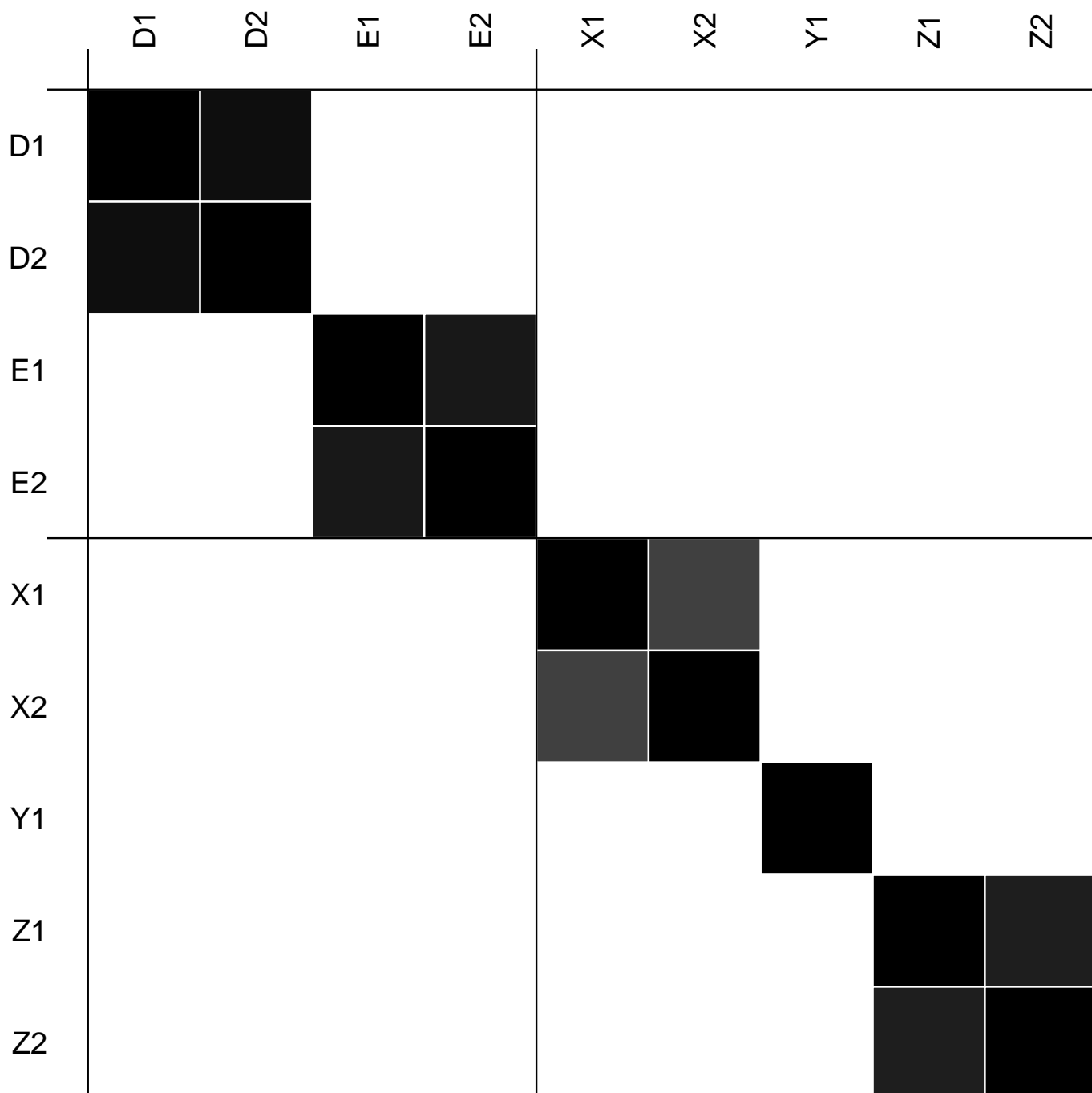
Sim-rep 9: Phased allele alignments (n=150), STACEY similarity matrix



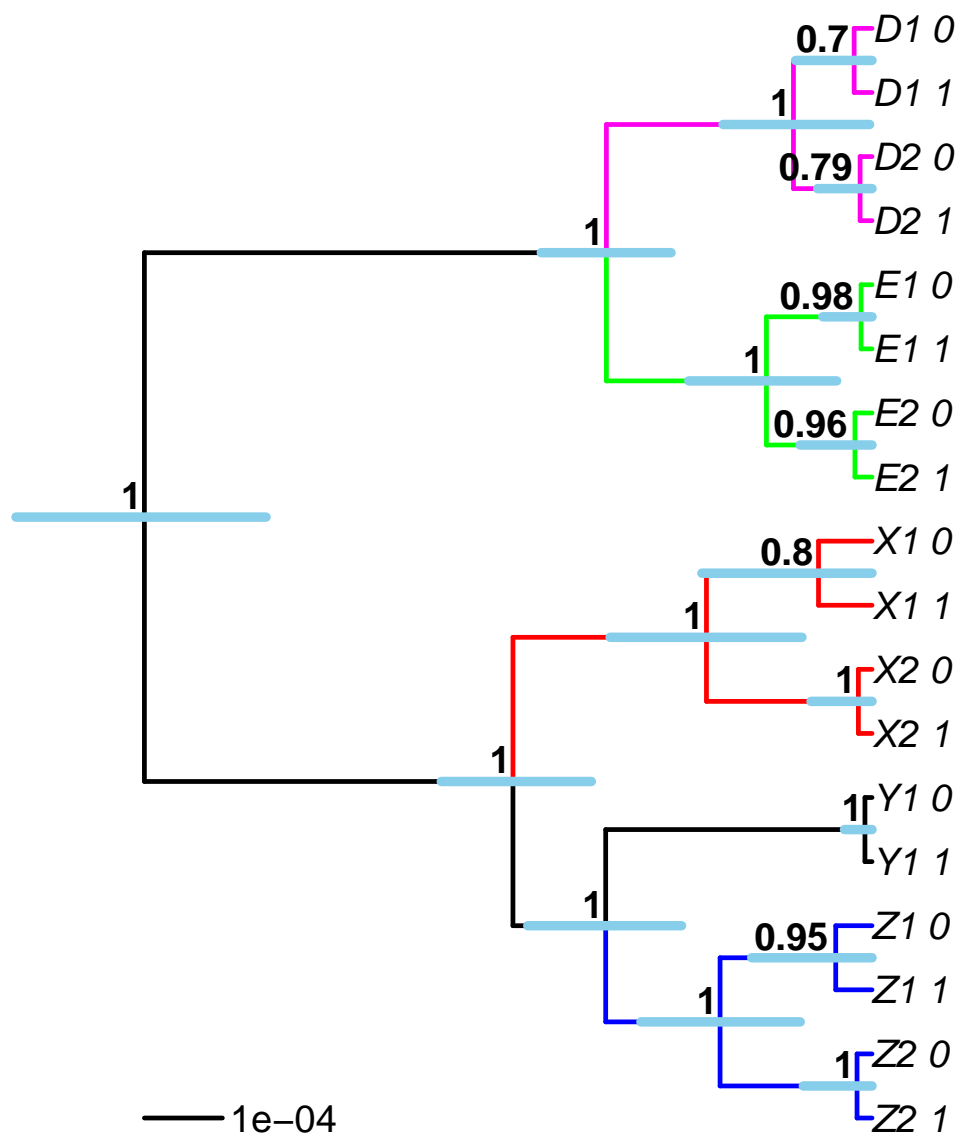
Sim-rep 9: IUPAC consensus alignments (n=150), STACEY species tree



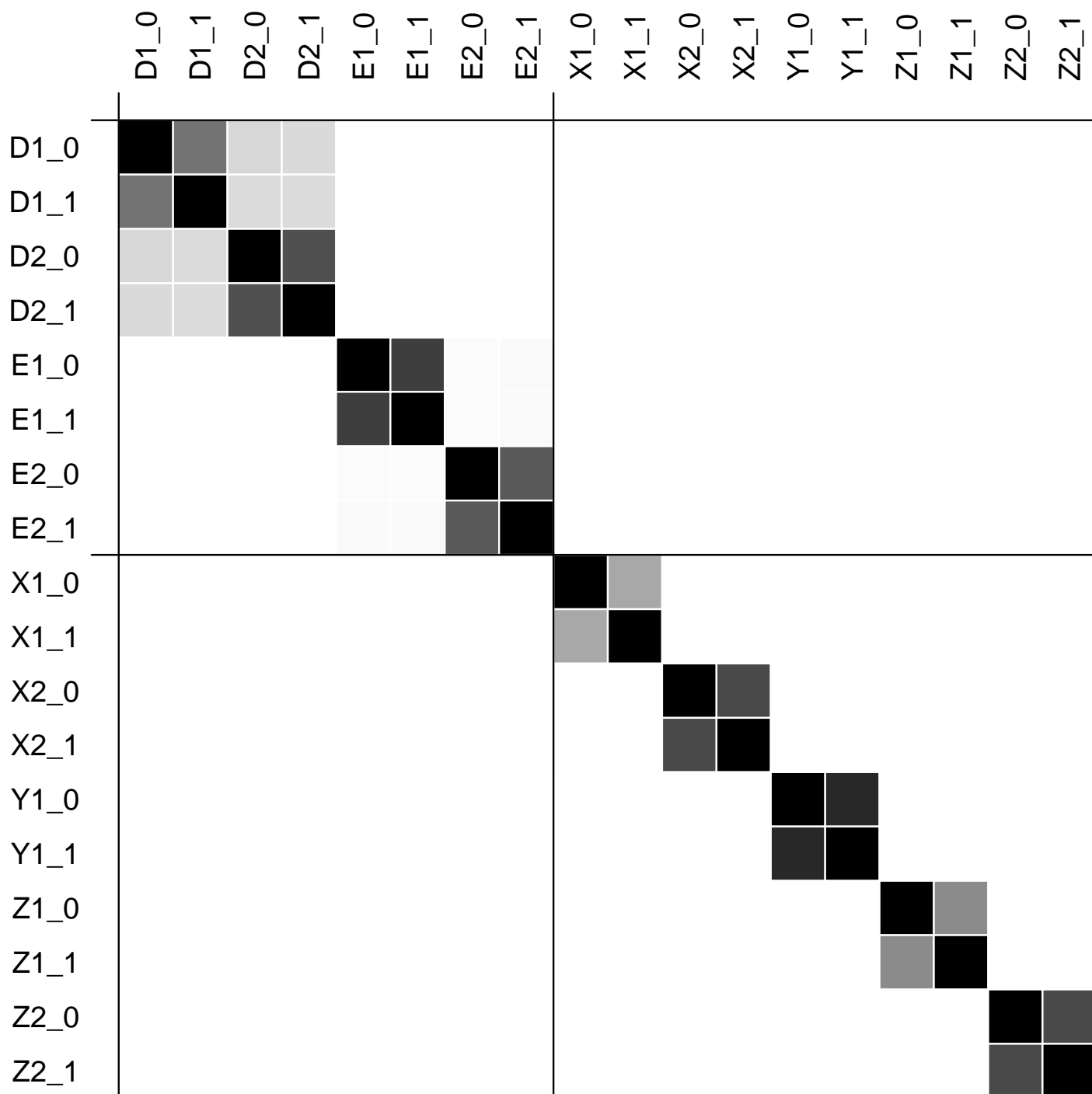
Sim-rep 9: IUPAC consensus alignments (n=150), STACEY similarity matrix



Sim-rep 9: Chimeric allele alignments (n=150), STACEY species tree

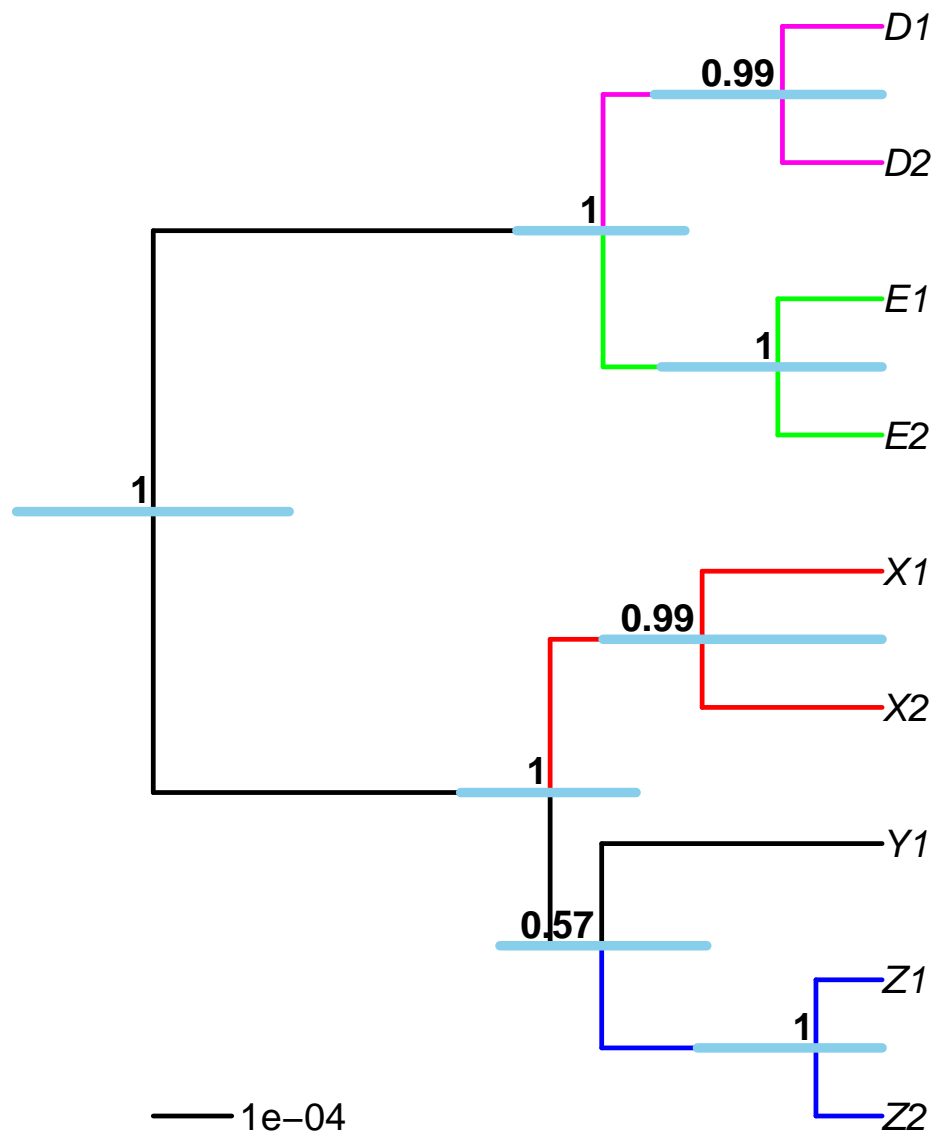


Sim-rep 9: Chimeric allele alignments (n=150), STACEY similarity matrix

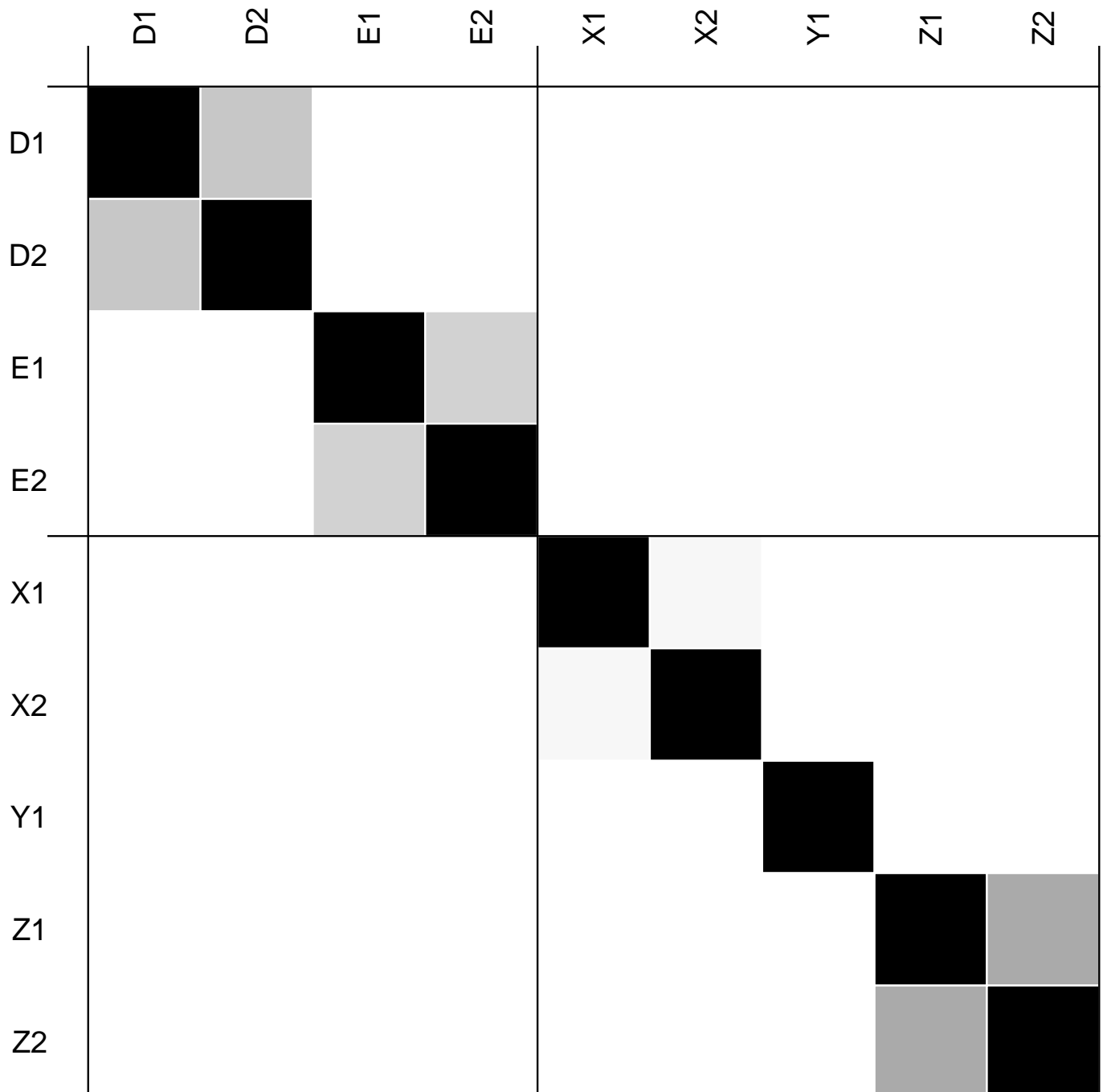




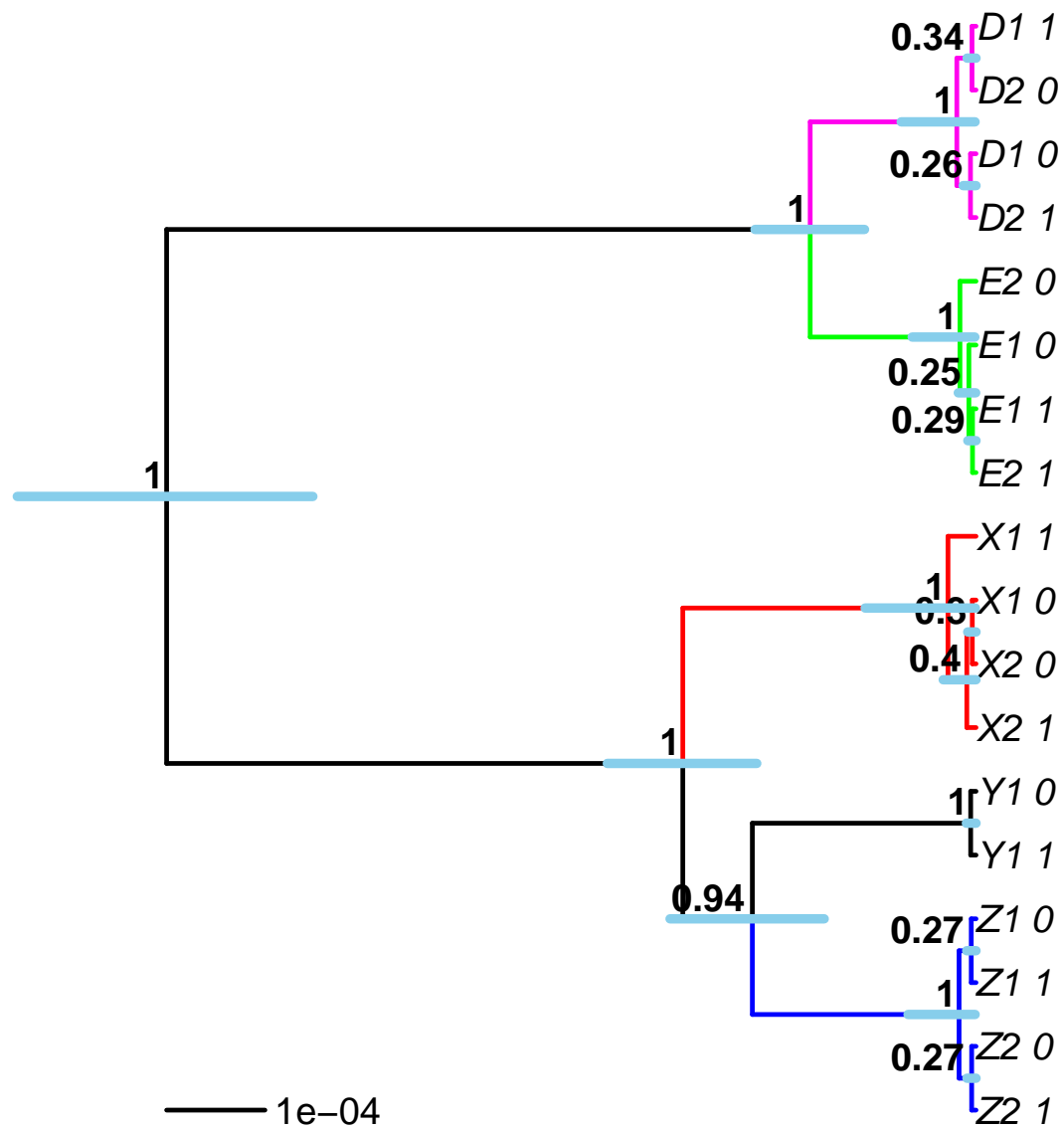
Sim-rep 10: Contig sequence alignments (n=150), STACEY species tree



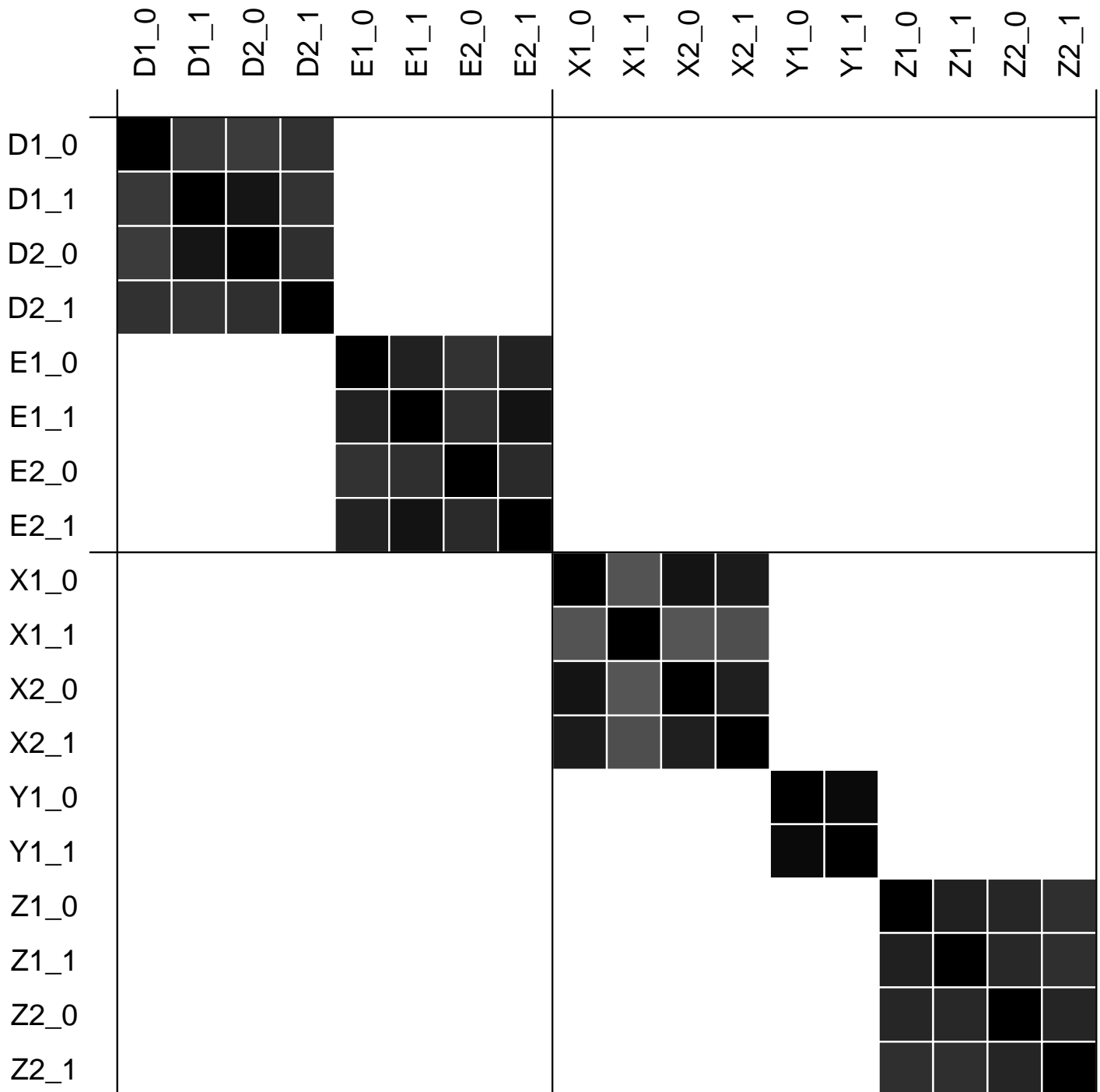
Sim-rep 10: Contig sequence alignments (n=150), STACEY similarity matrix



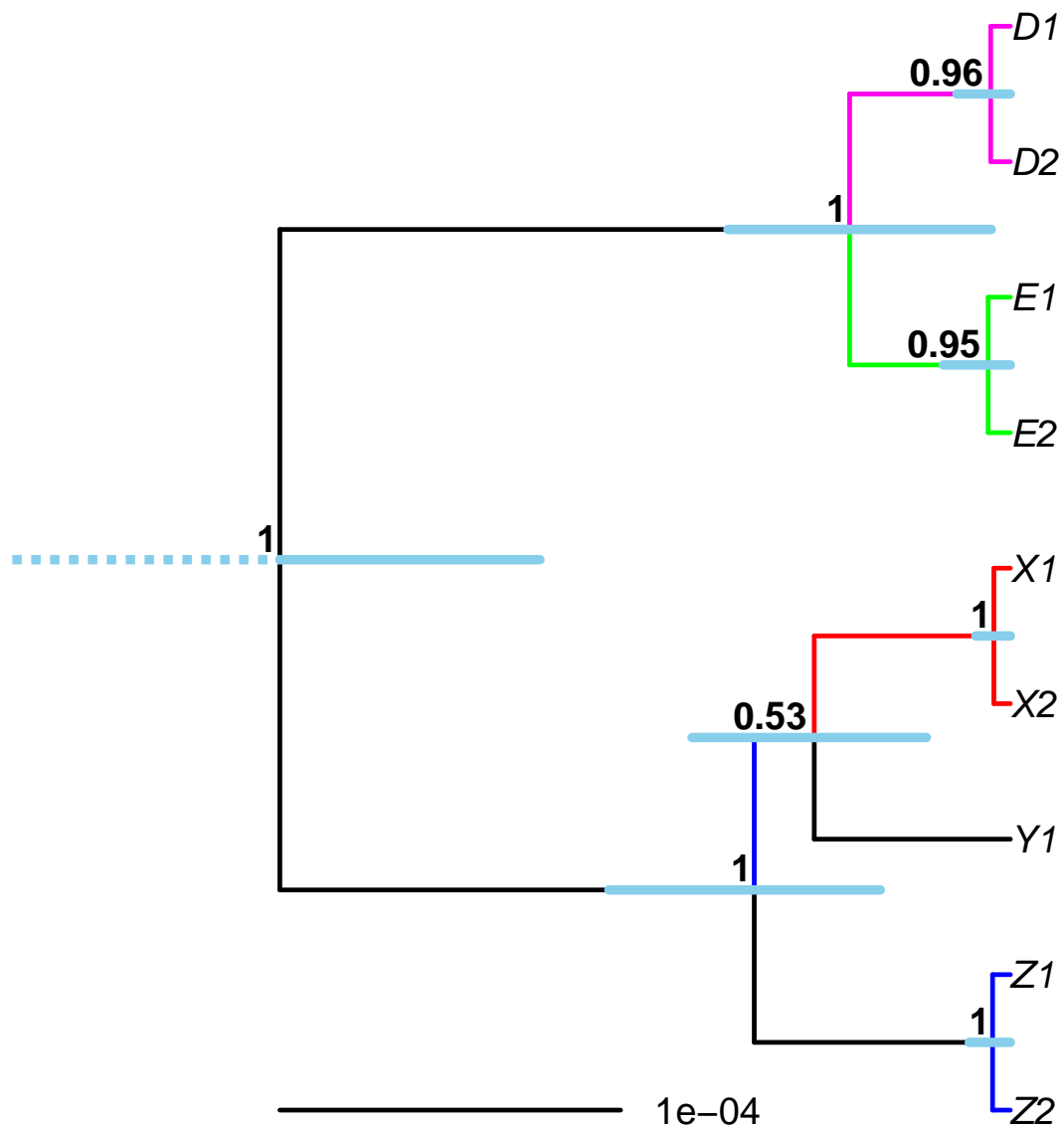
Sim-rep 10: Phased allele alignments (n=150), STACEY species tree



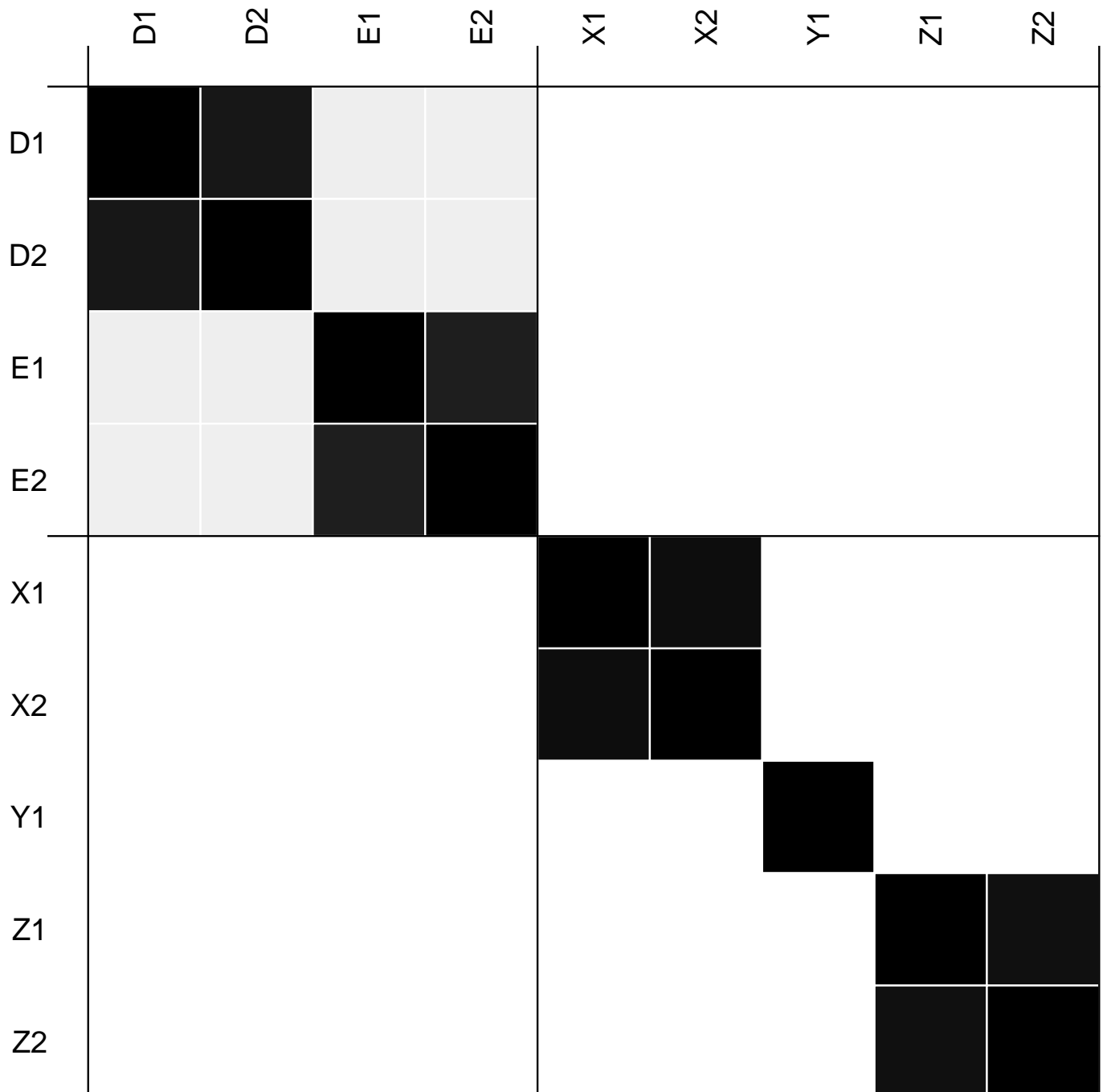
Sim-rep 10: Phased allele alignments (n=150), STACEY similarity matrix



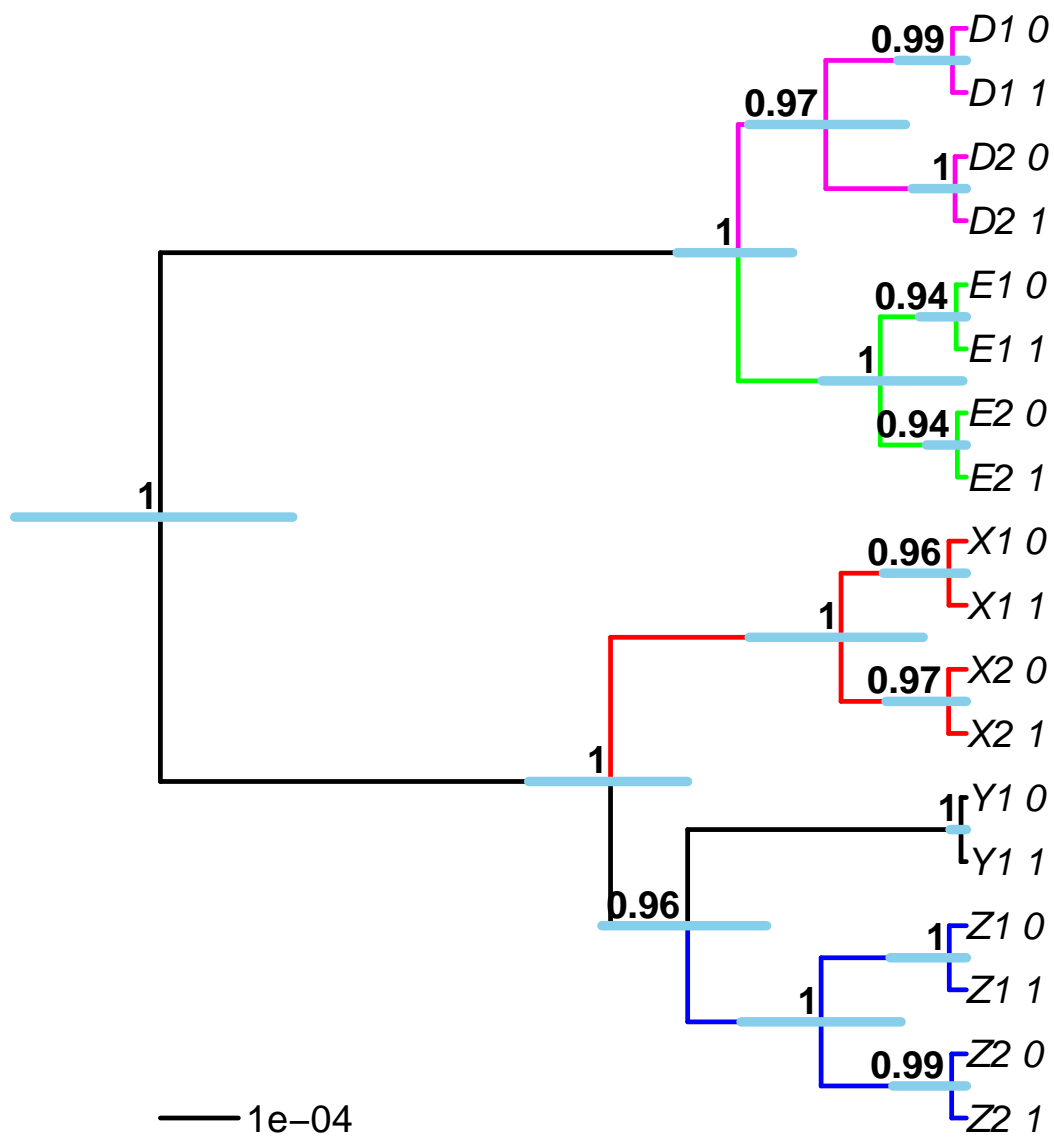
Sim-rep 10: IUPAC consensus alignments (n=150), STACEY species tree



Sim-rep 10: IUPAC consensus alignments (n=150), STACEY similarity matrix



Sim-rep 10: Chimeric allele alignments (n=150), STACEY species tree



Sim-rep 10: Chimeric allele alignments (n=150), STACEY similarity matrix

