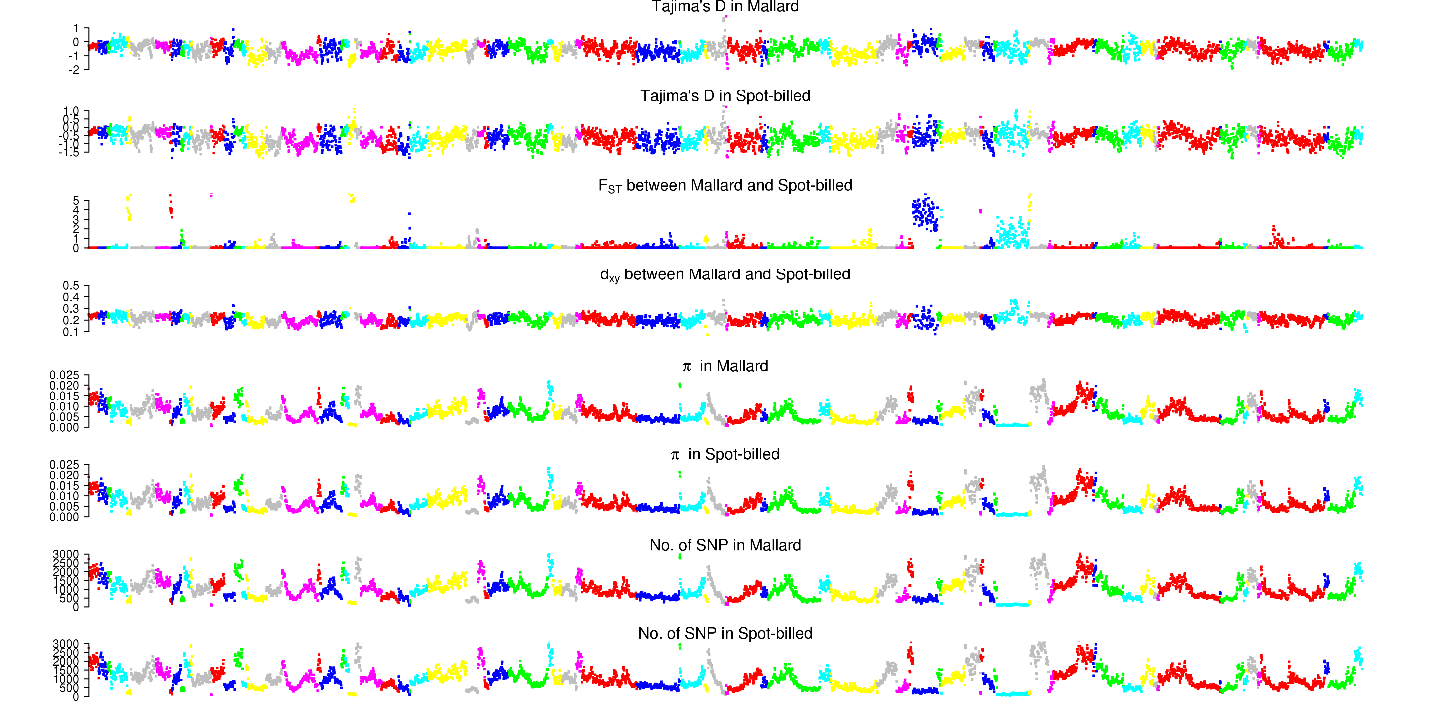
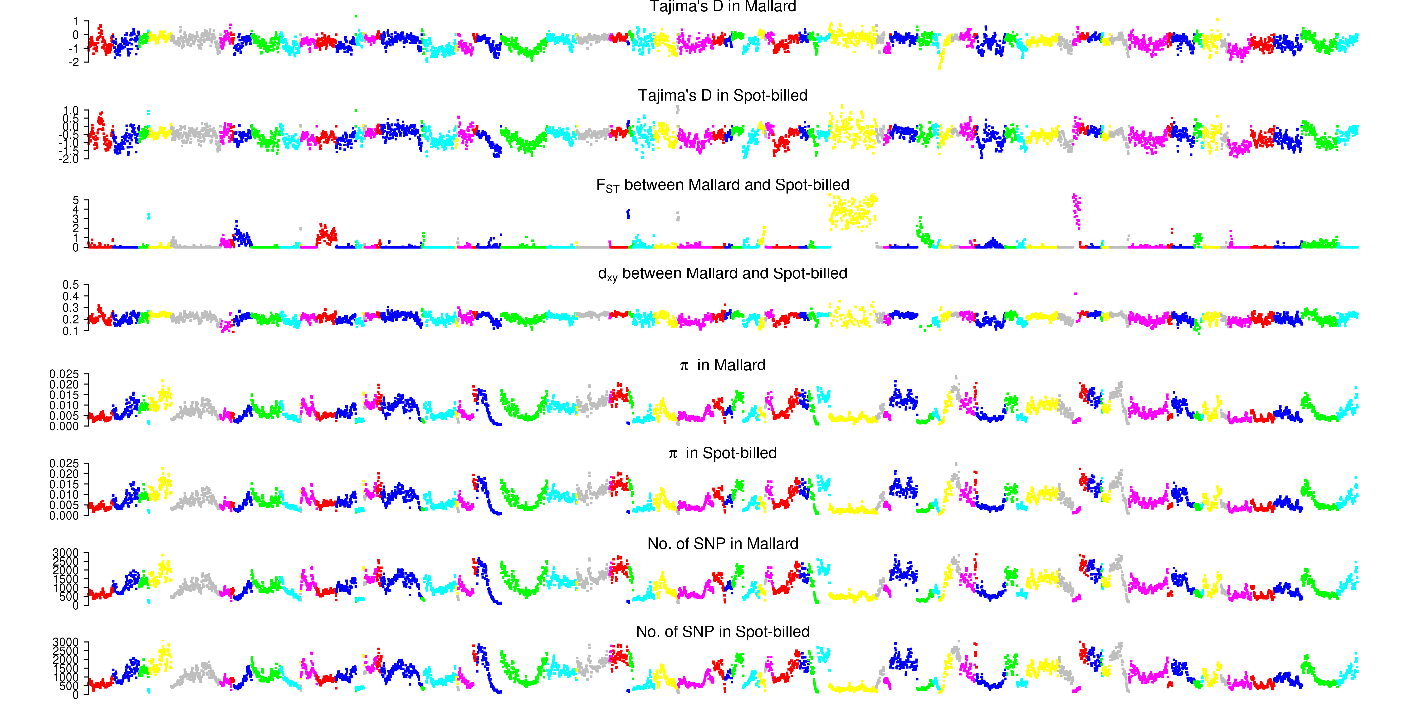
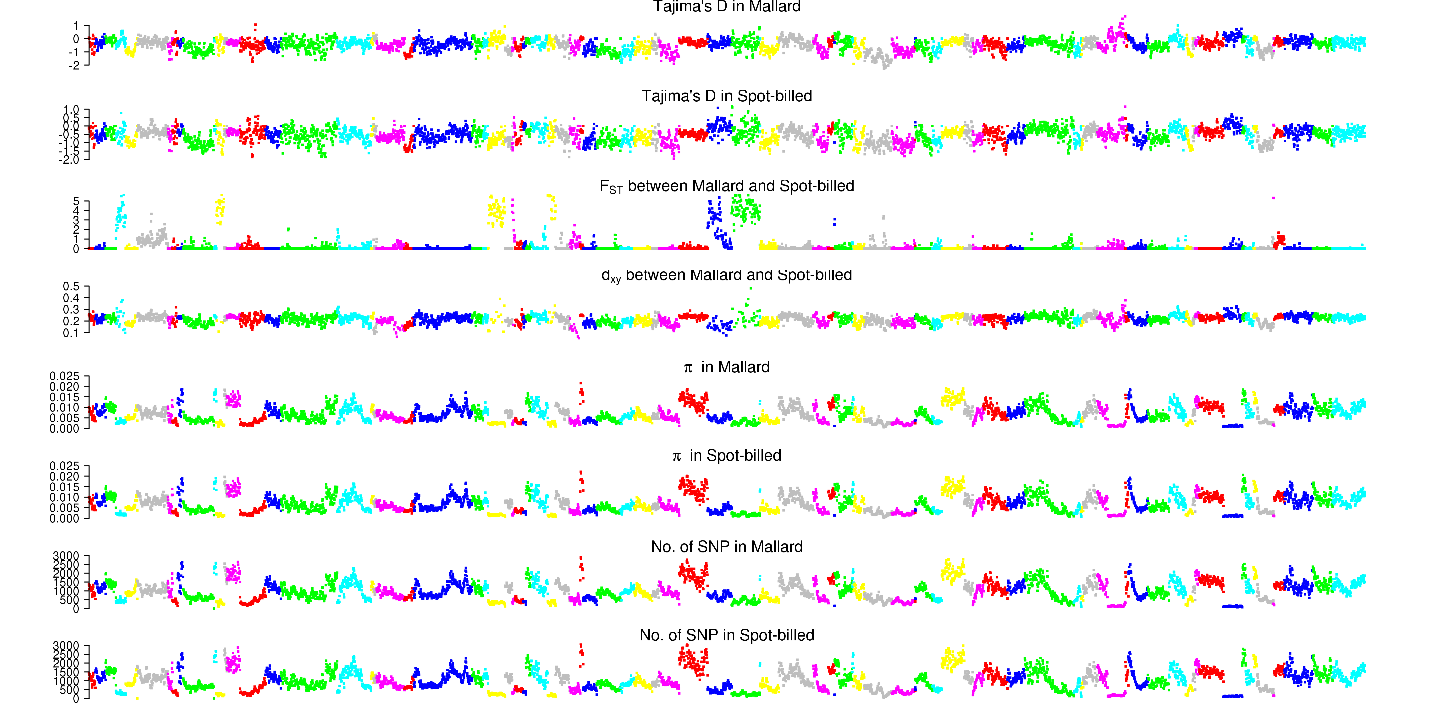
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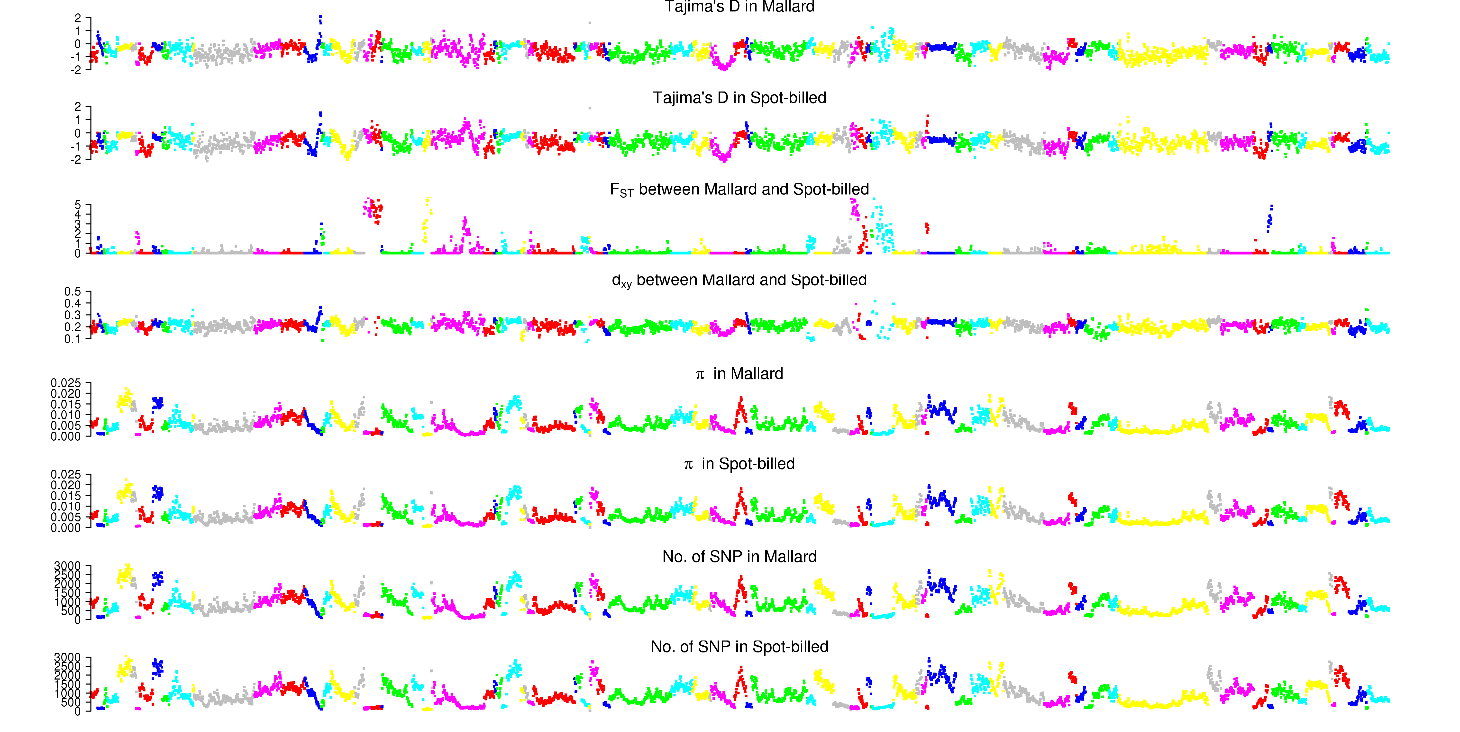
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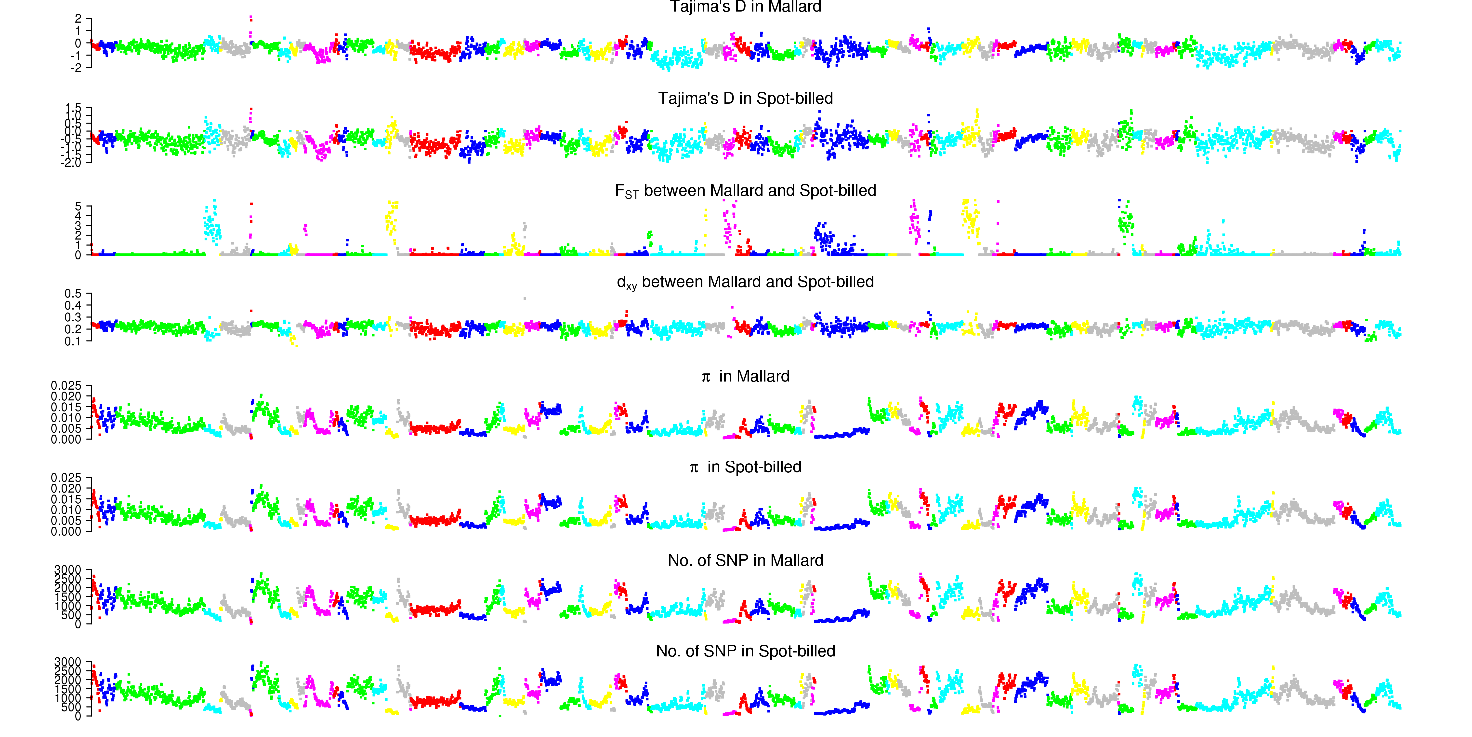
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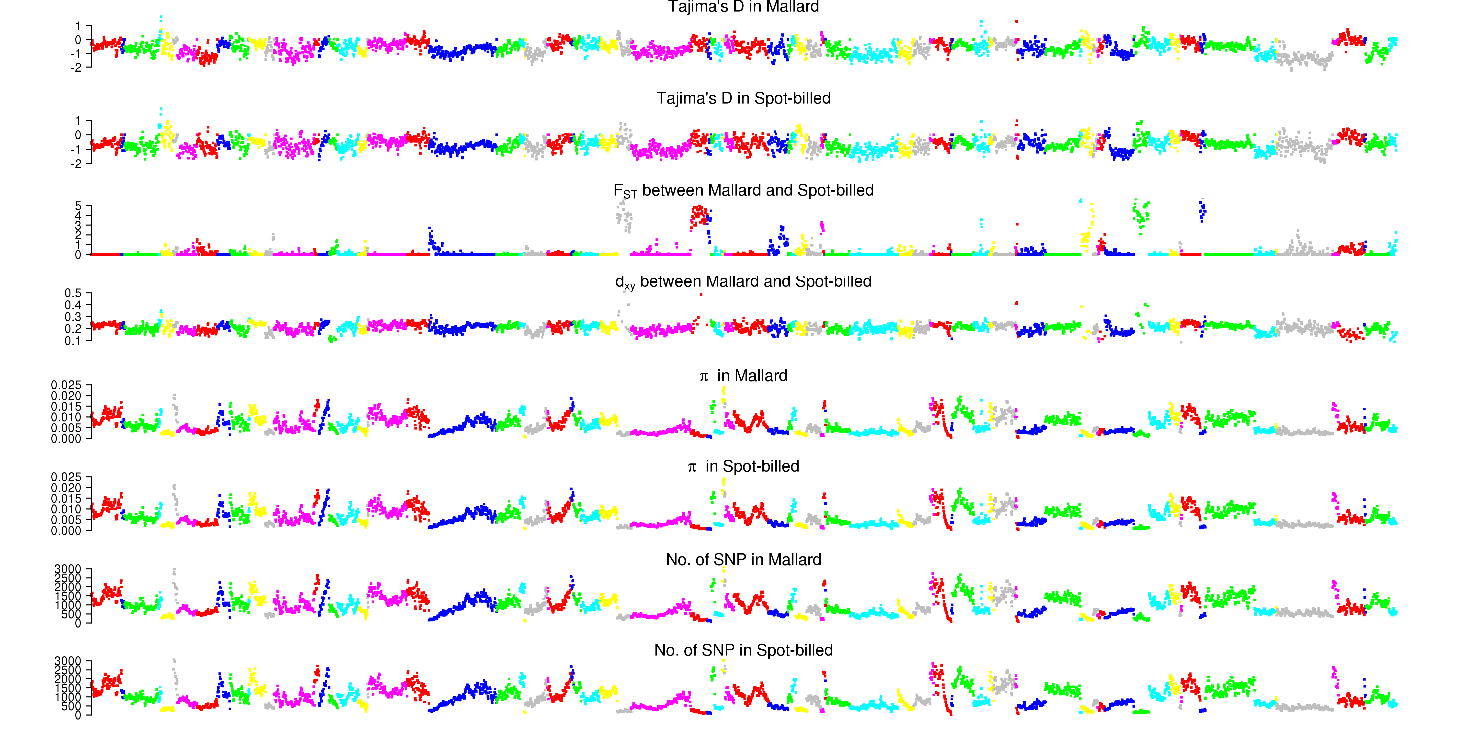
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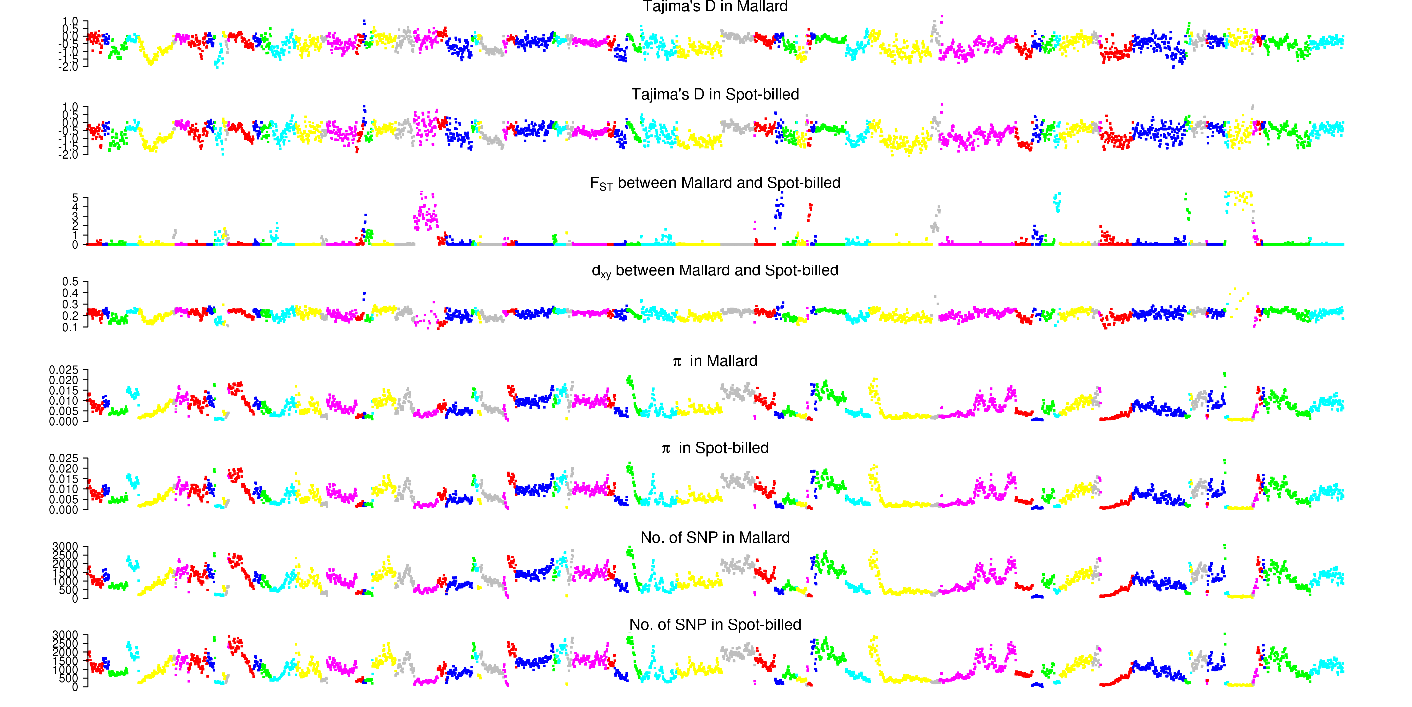
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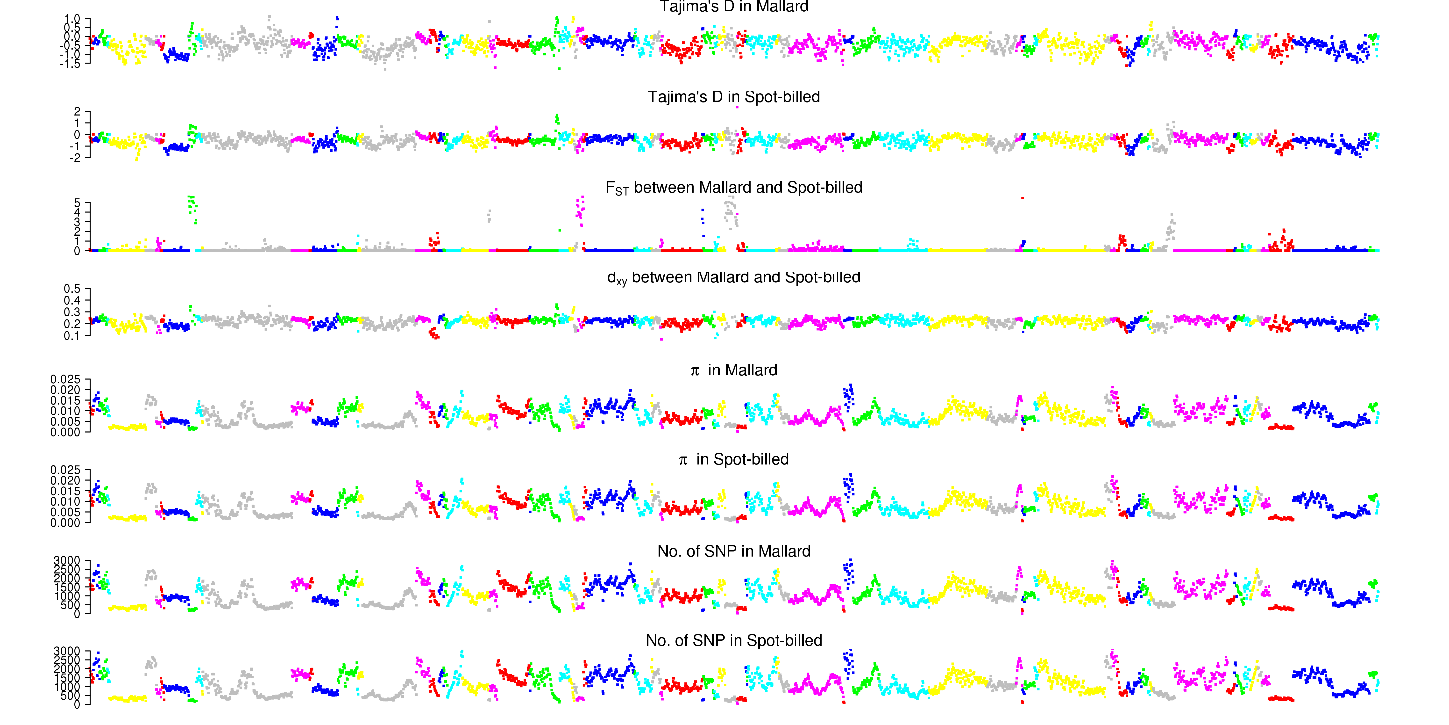
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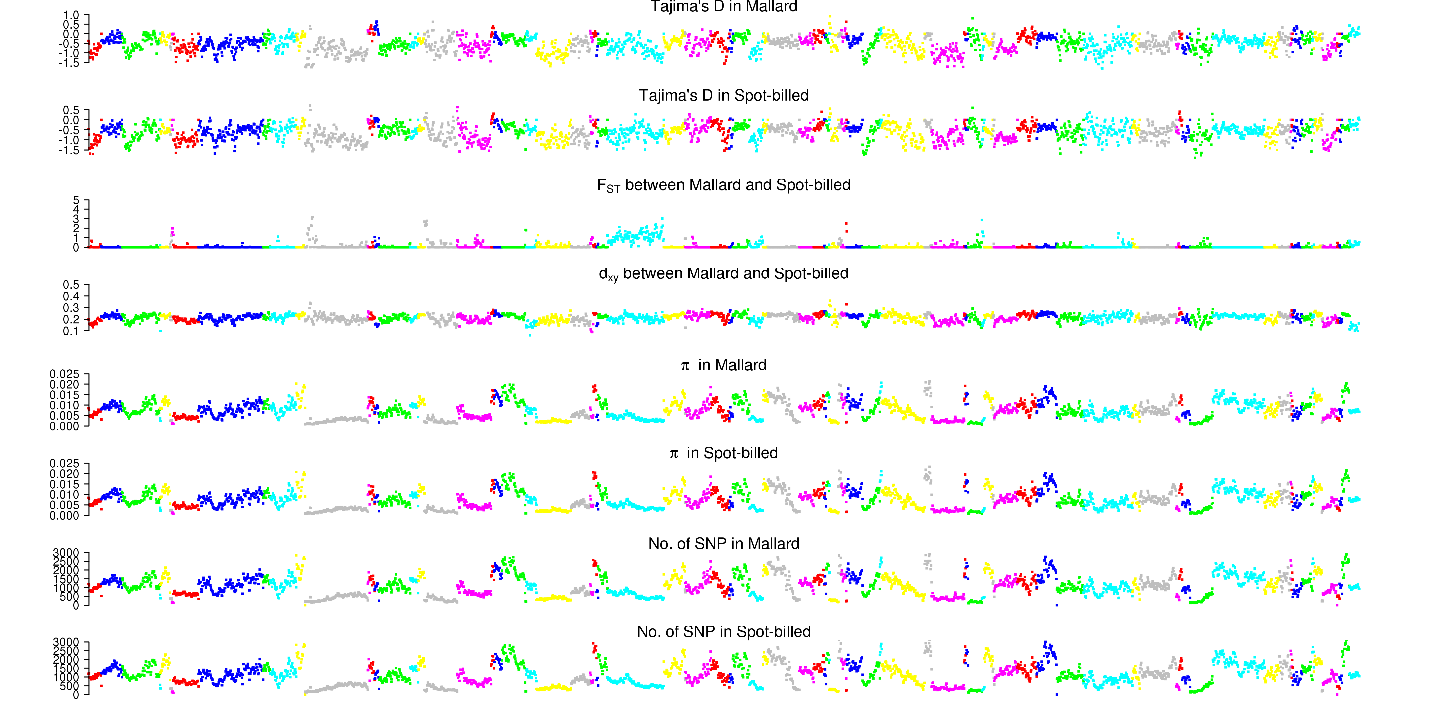
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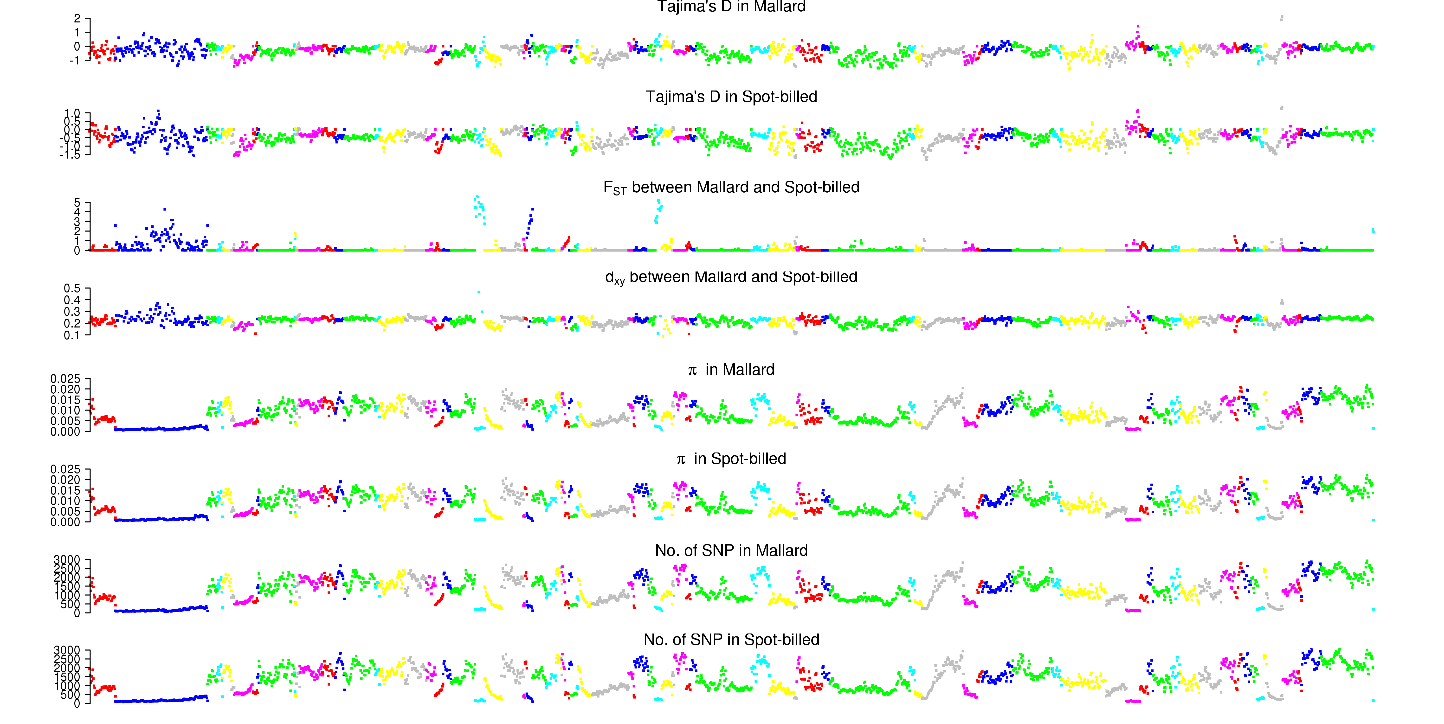
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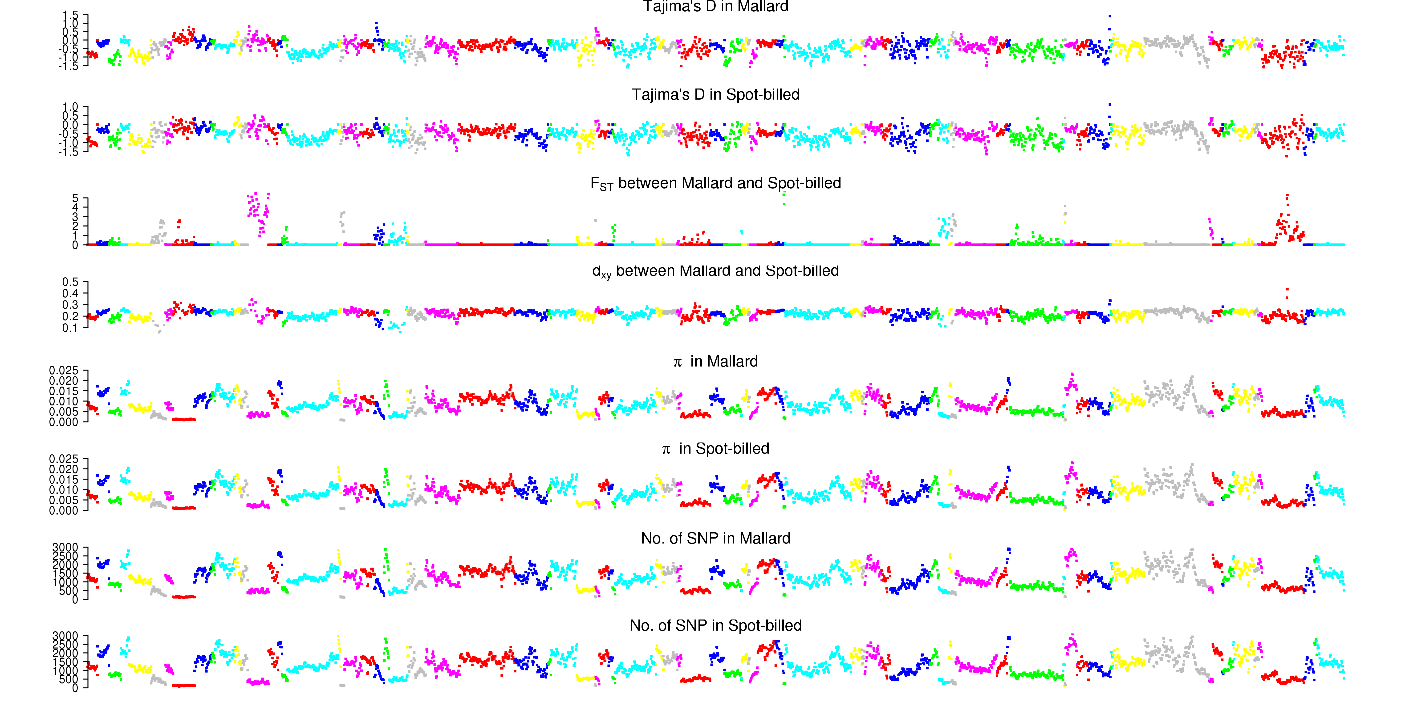
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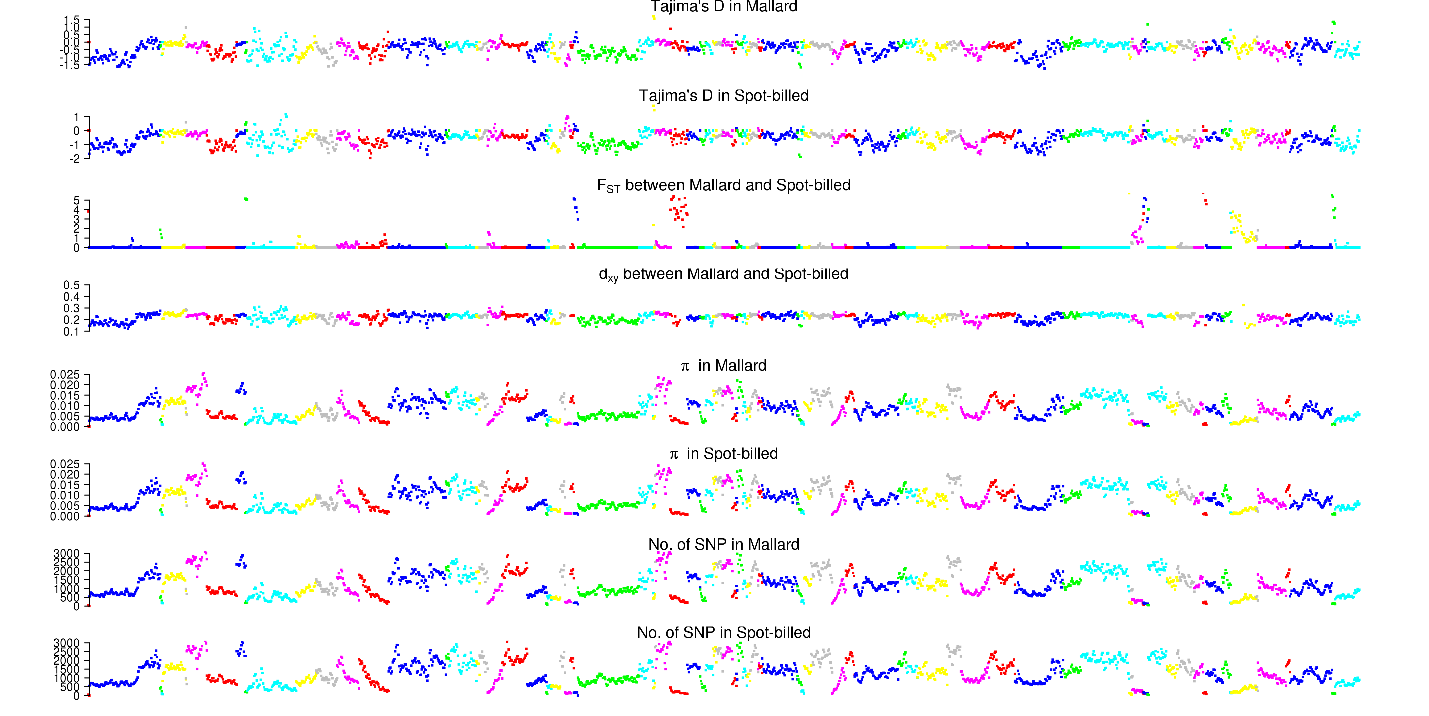
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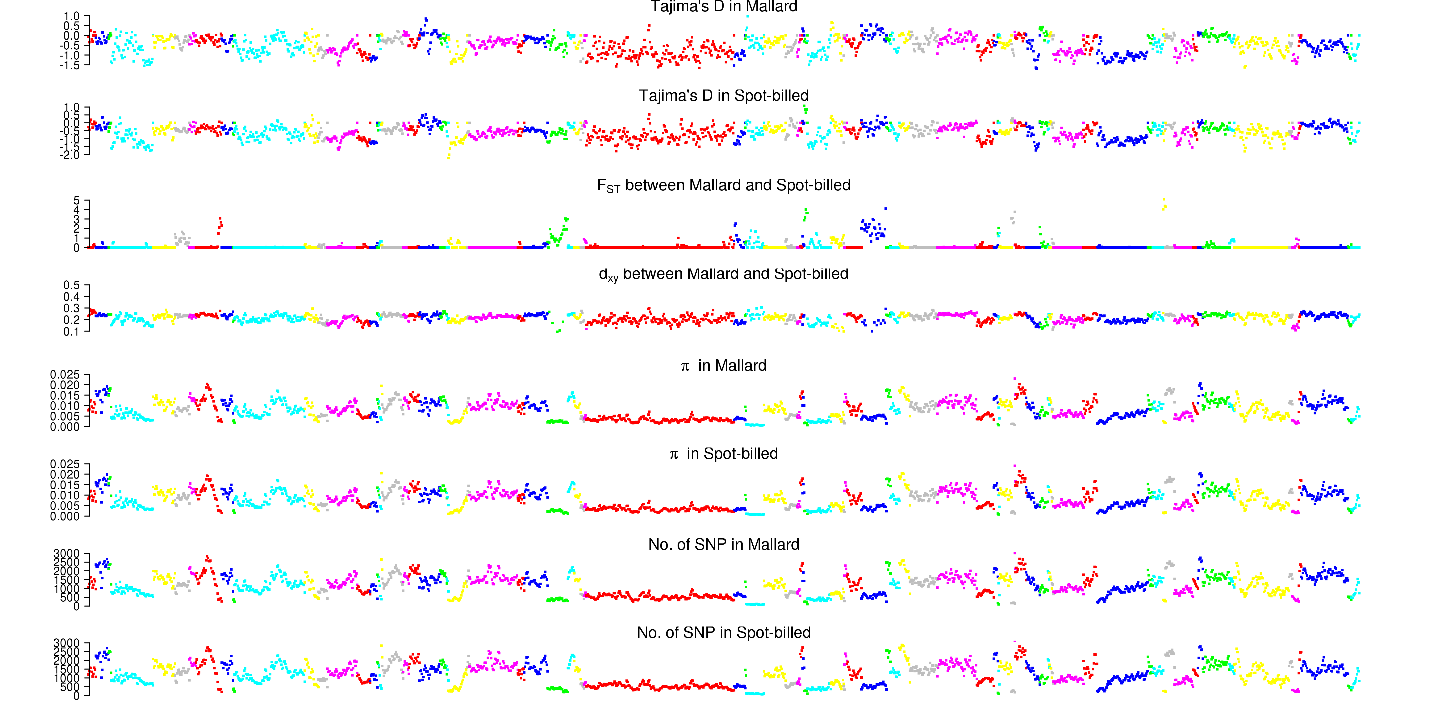
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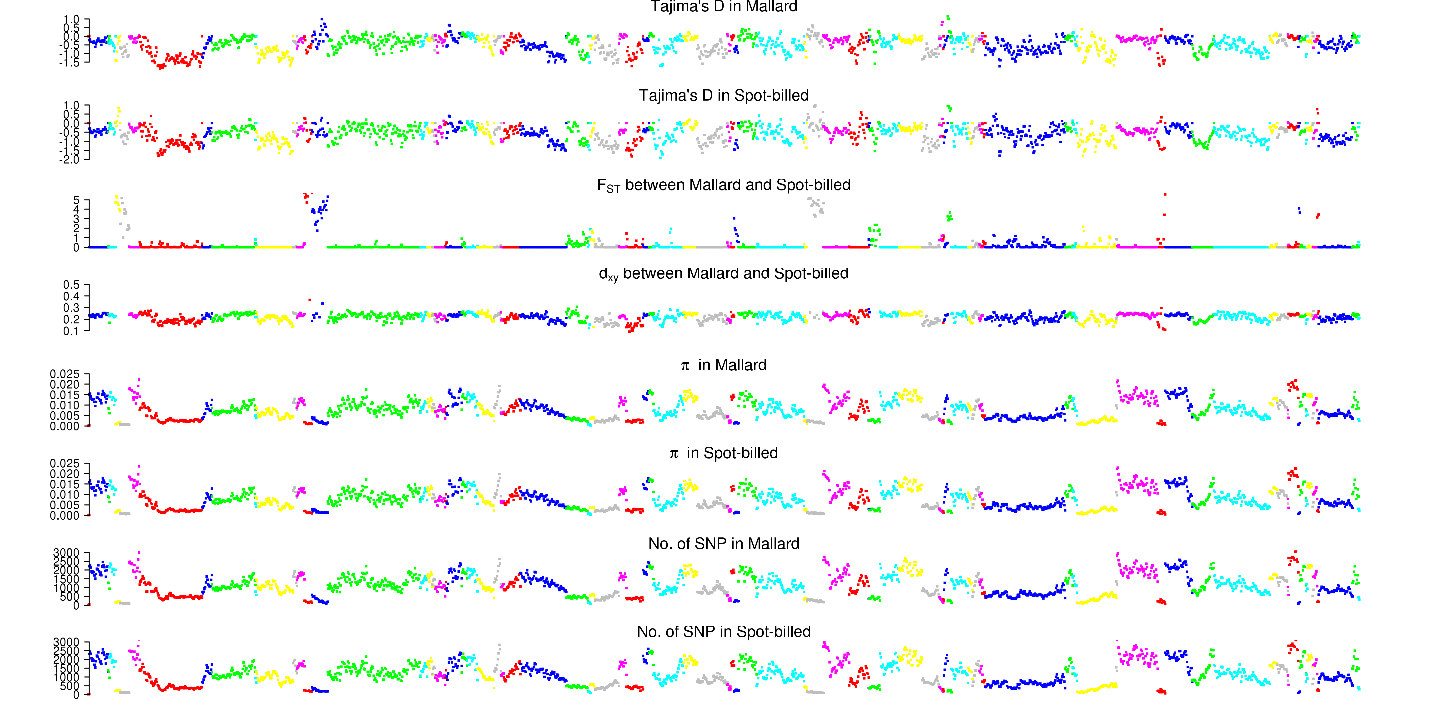
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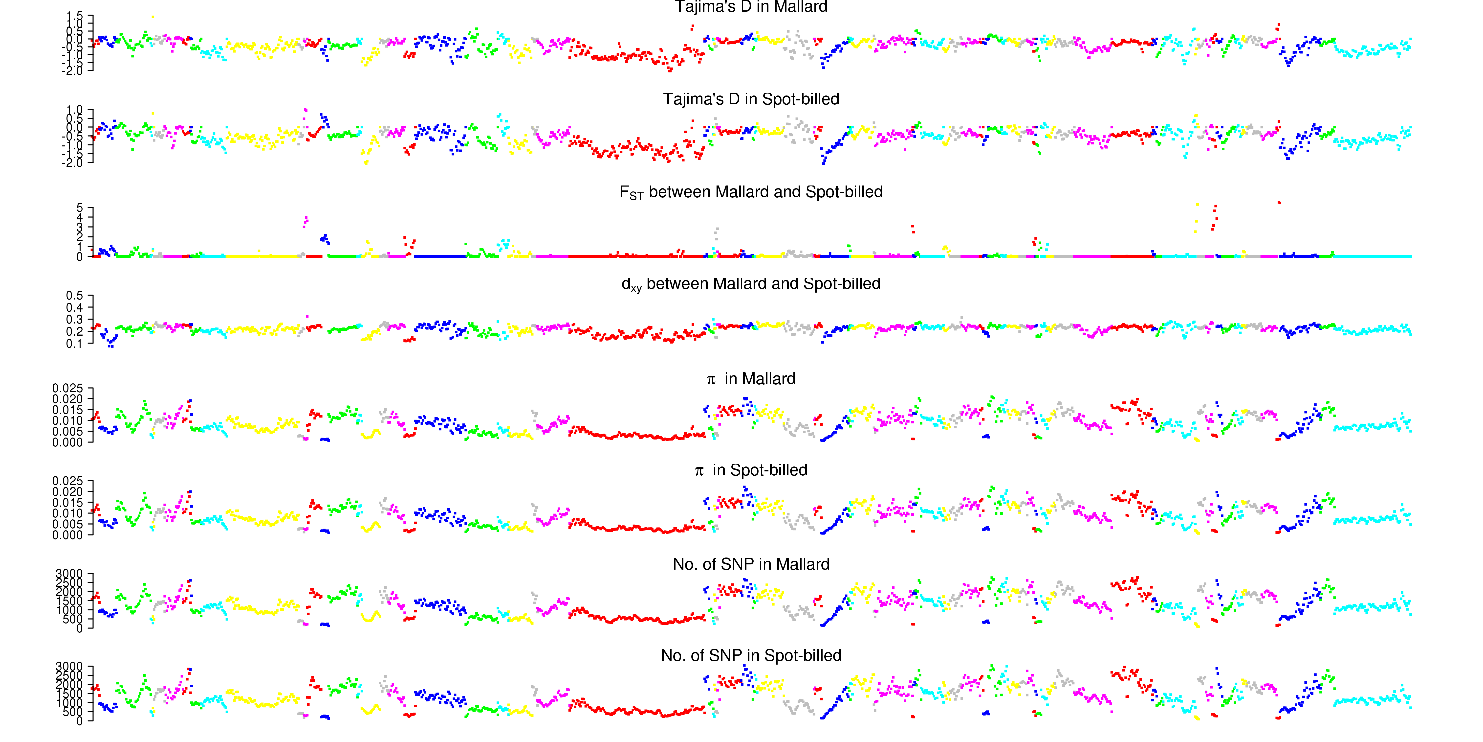
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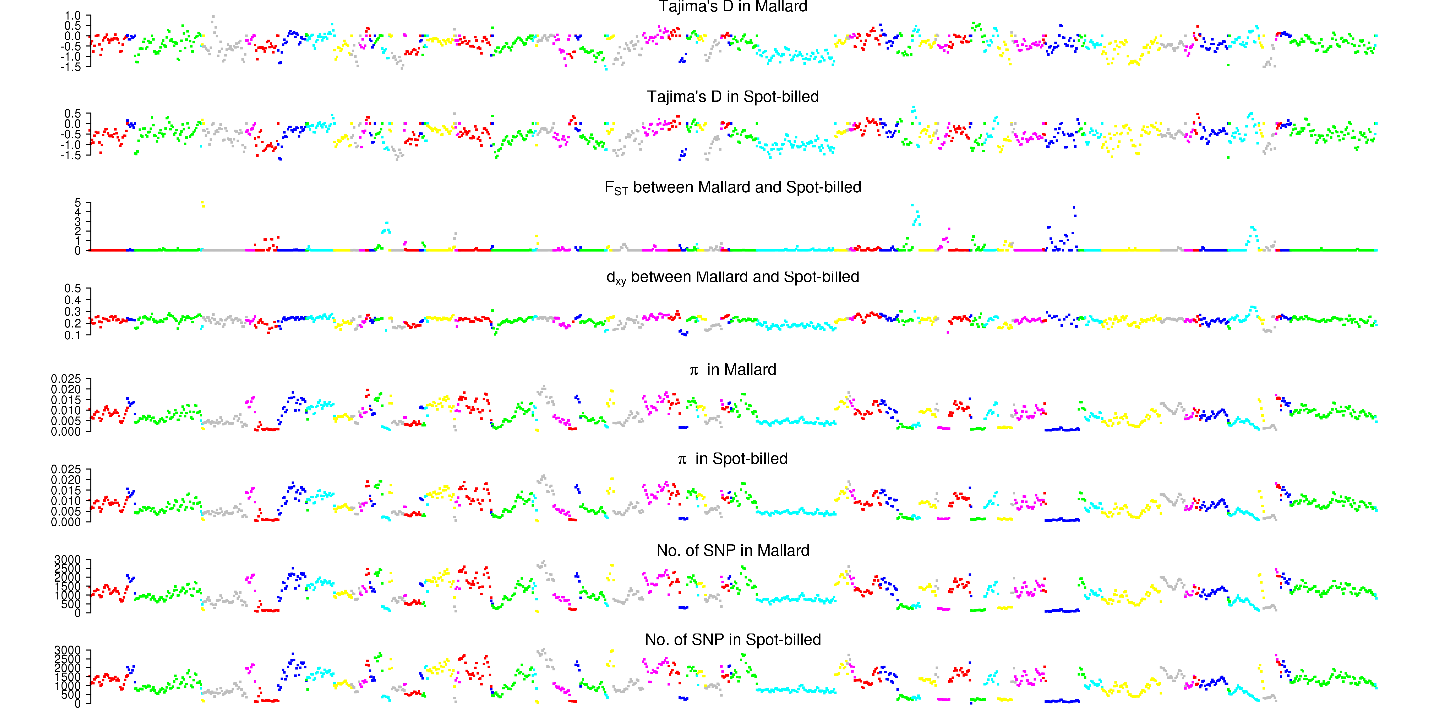


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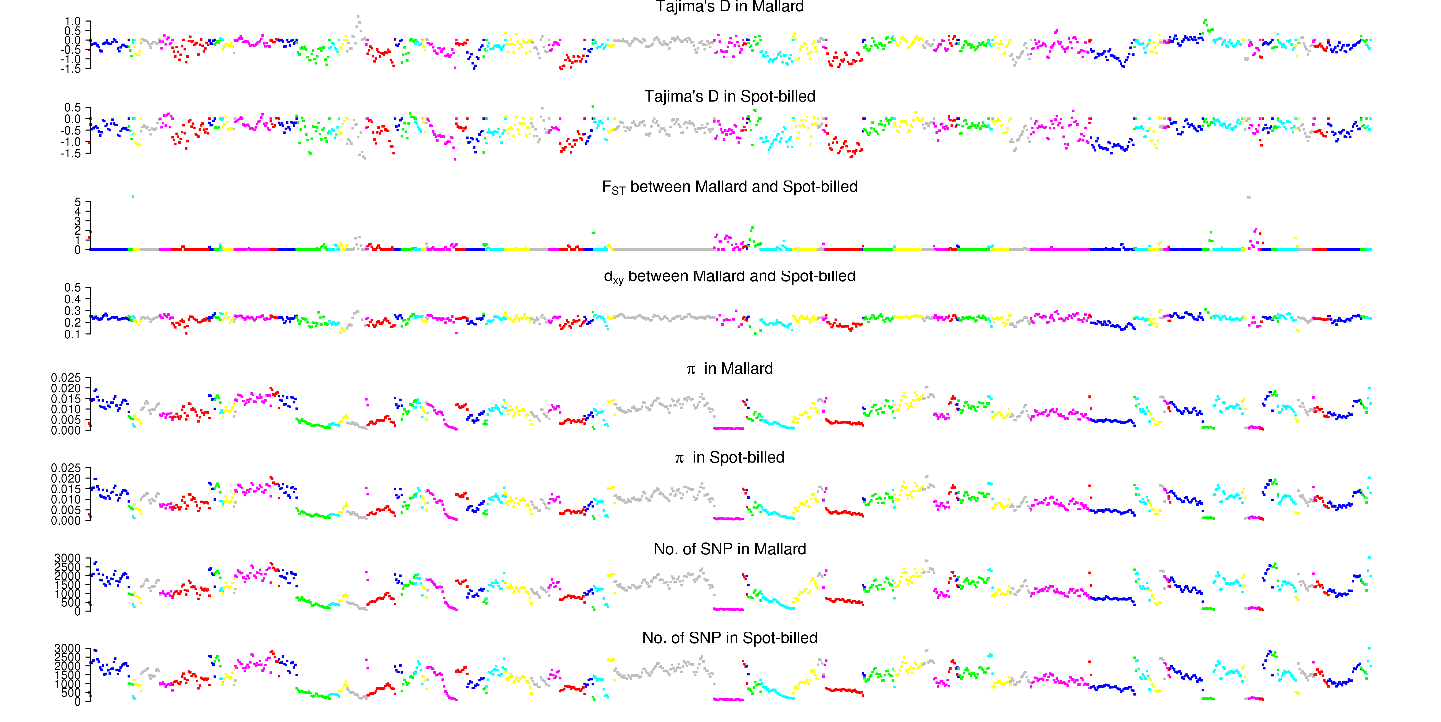


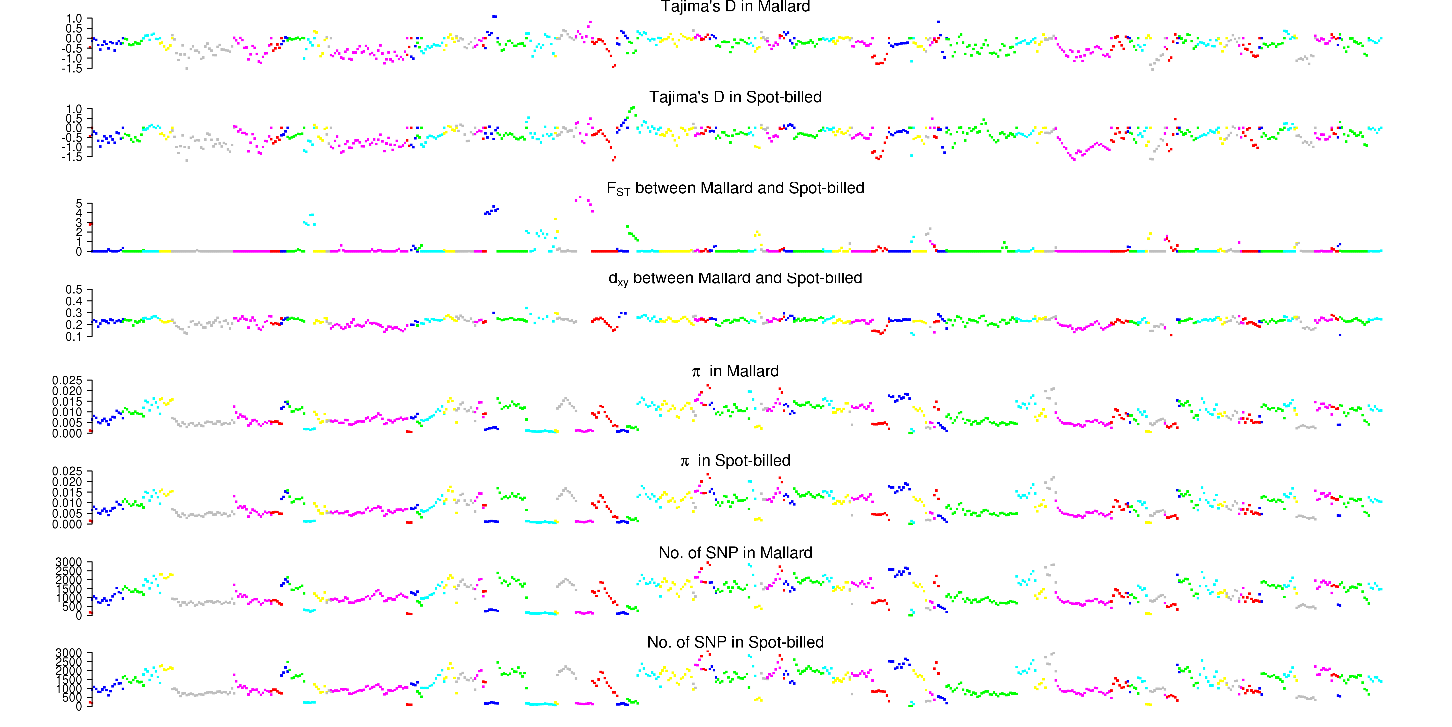
Scaffolds from KB744475.1 to KB744721.1



Scaffolds from KB744725.1 to KB745078.1

Scaffolds from KB745085.1 to KB745764.1



Scaffolds from KB745806.1 to KB747284.1

## Fig. S23. Distribution of Tajima’D, positive end of z*FST*, dxy, number of SNPs per window, nucleotide diversity (π) and zHp along scaffolds in ducks.

These numbers were counted with 40 kb window by step size of 20 kb.