

Sanger sequencing results of *E. coli* strains

StreptoCAD: An open-source software toolbox automating genome engineering workflows in streptomycetes

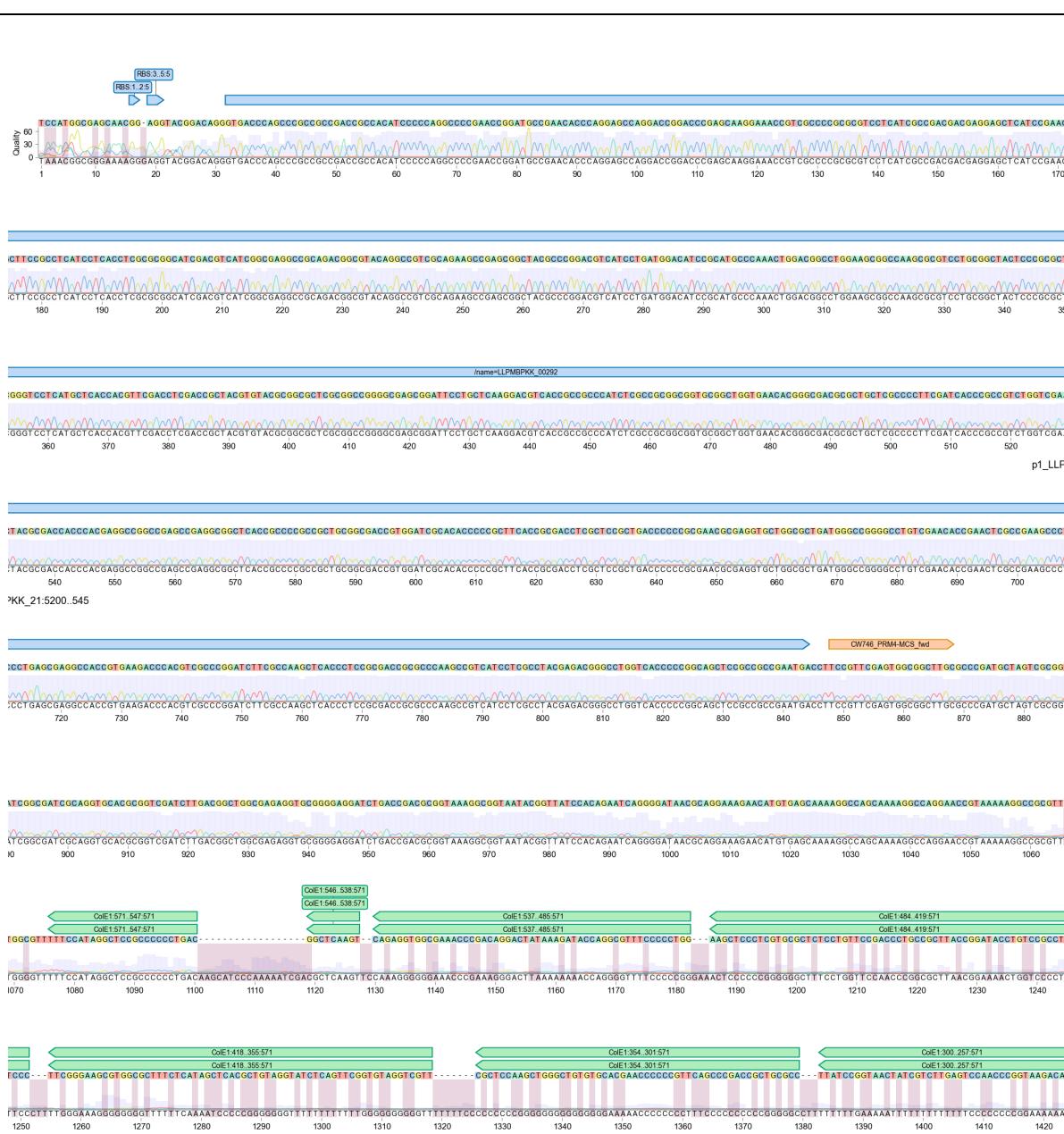
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¹The Novo Nordisk Foundation Center for Biosustainability, Technical University of Denmark, Kgs. Lyngby, Denmark

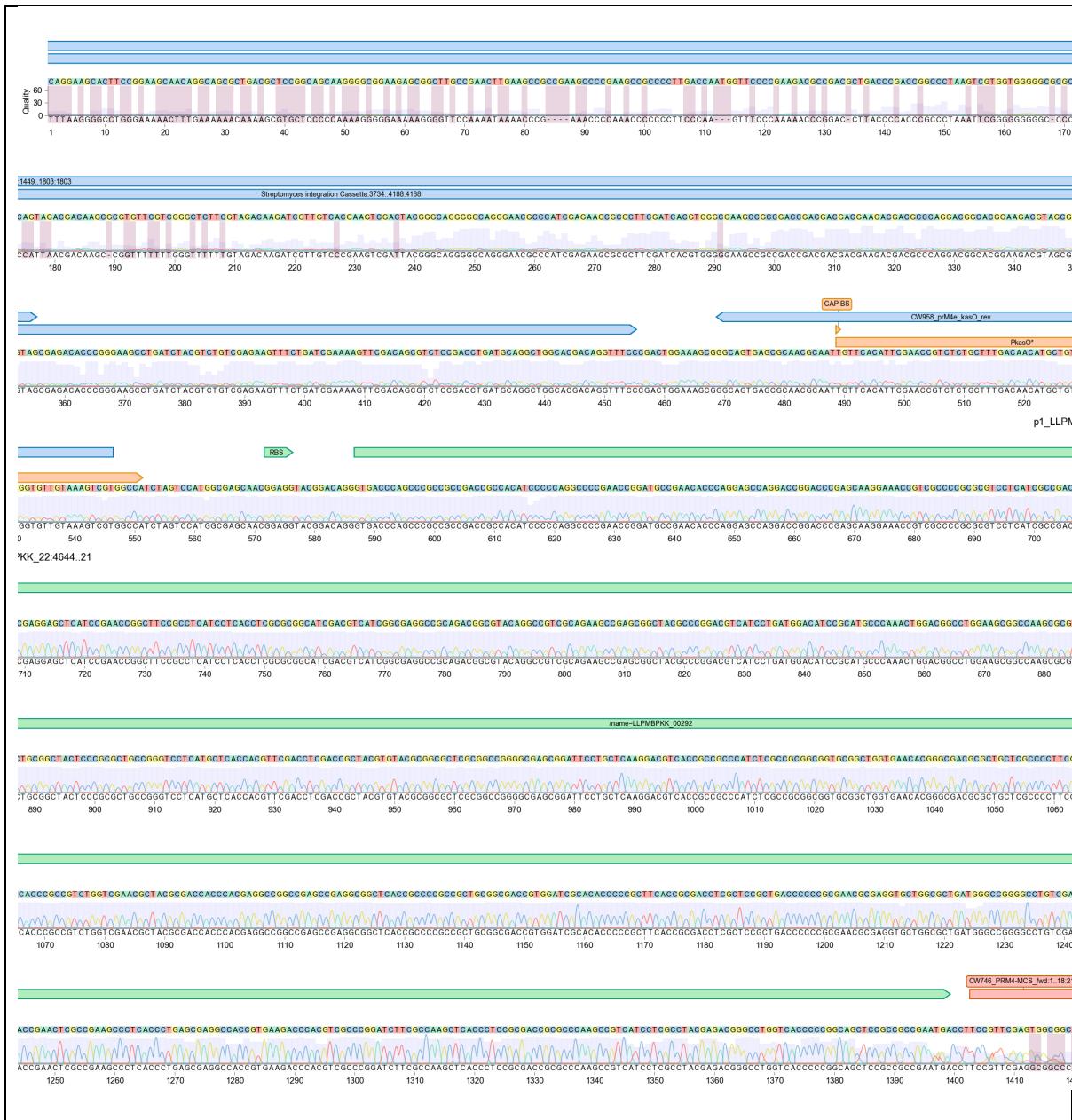
² Department of Biotechnology and Biomedicine, Technical University of Denmark, Kgs. Lyngby, Denmark

= to whom correspondence should be addressed: Tilmann Weber, Rasmus J.N. Frandsen

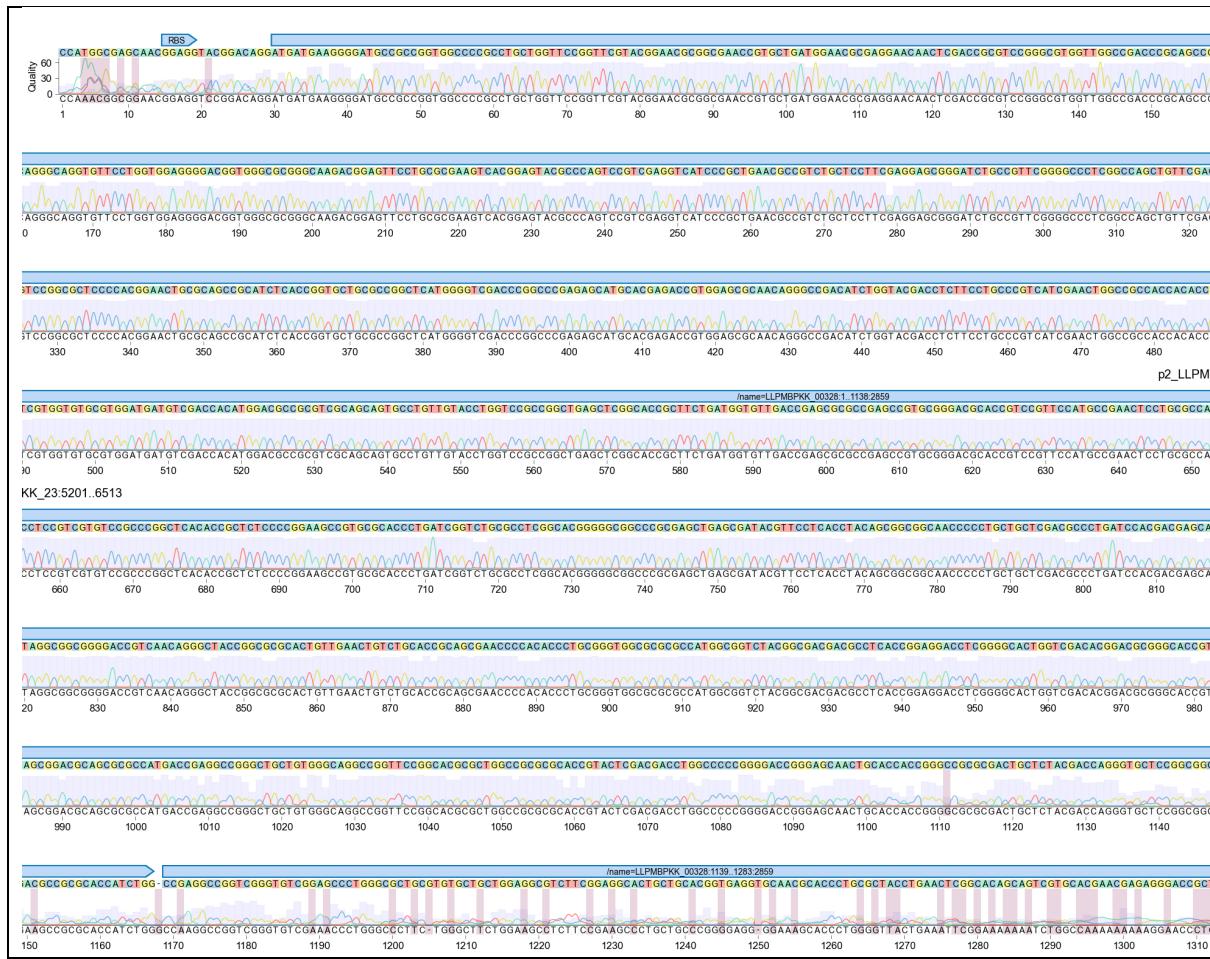
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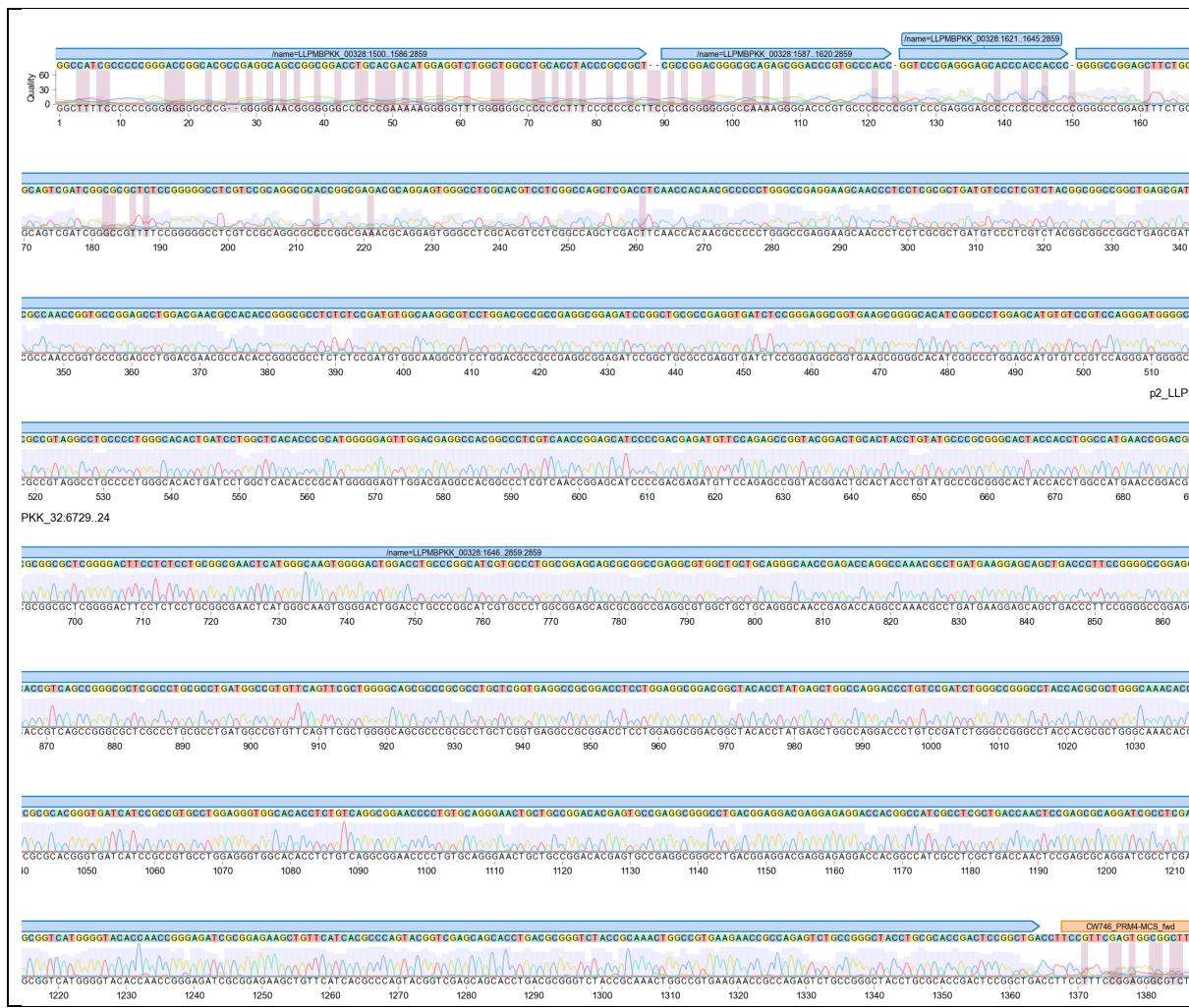
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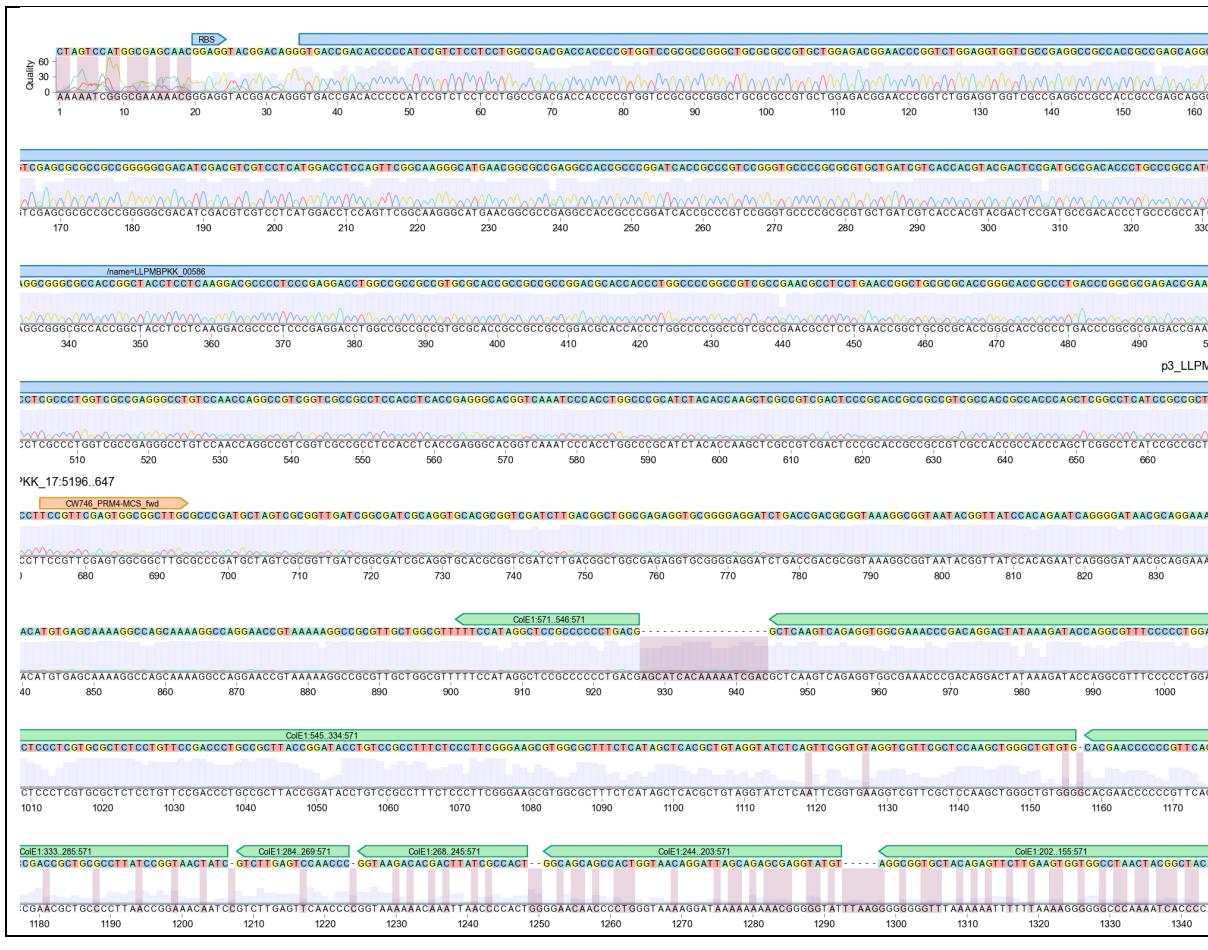
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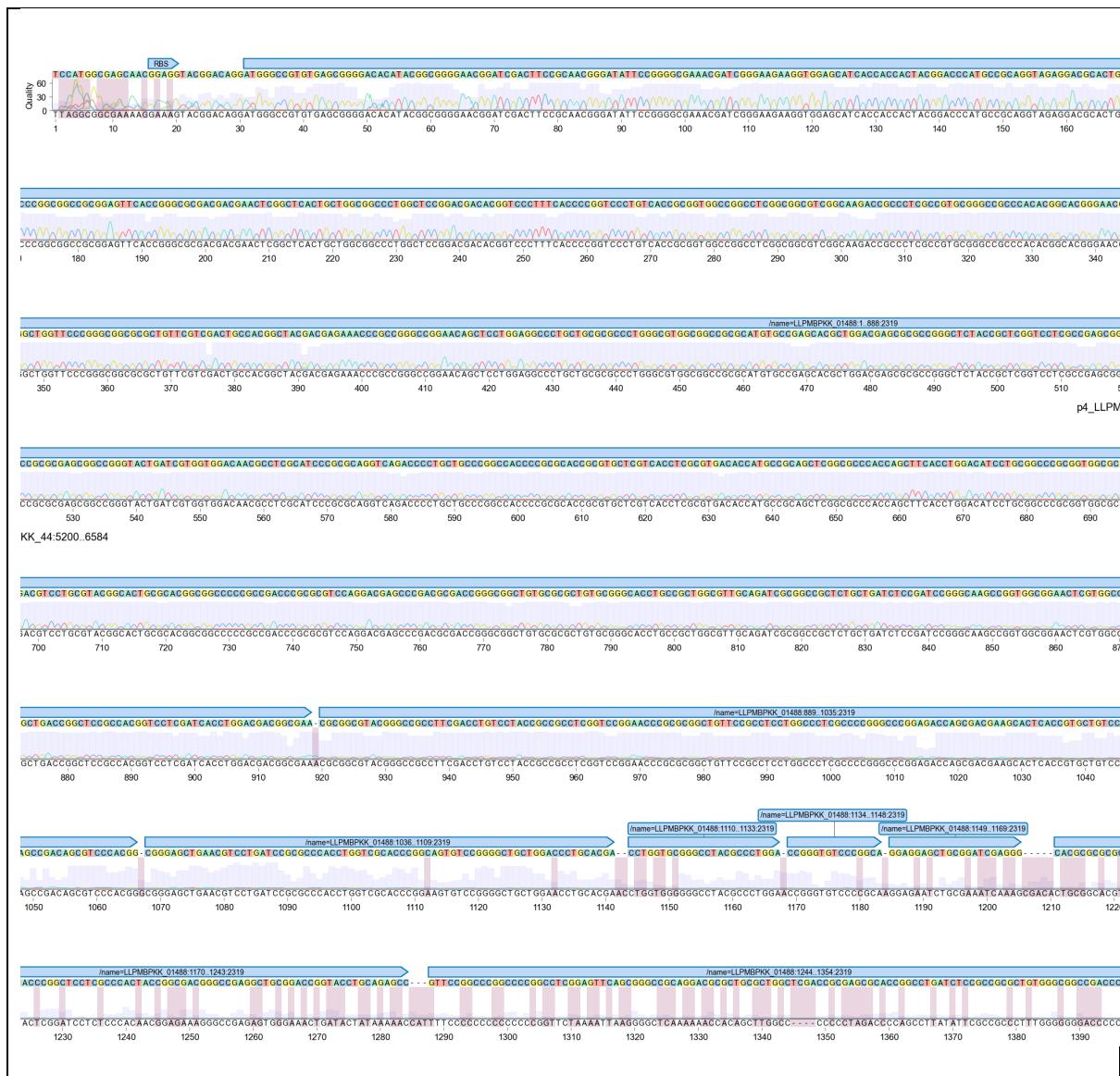
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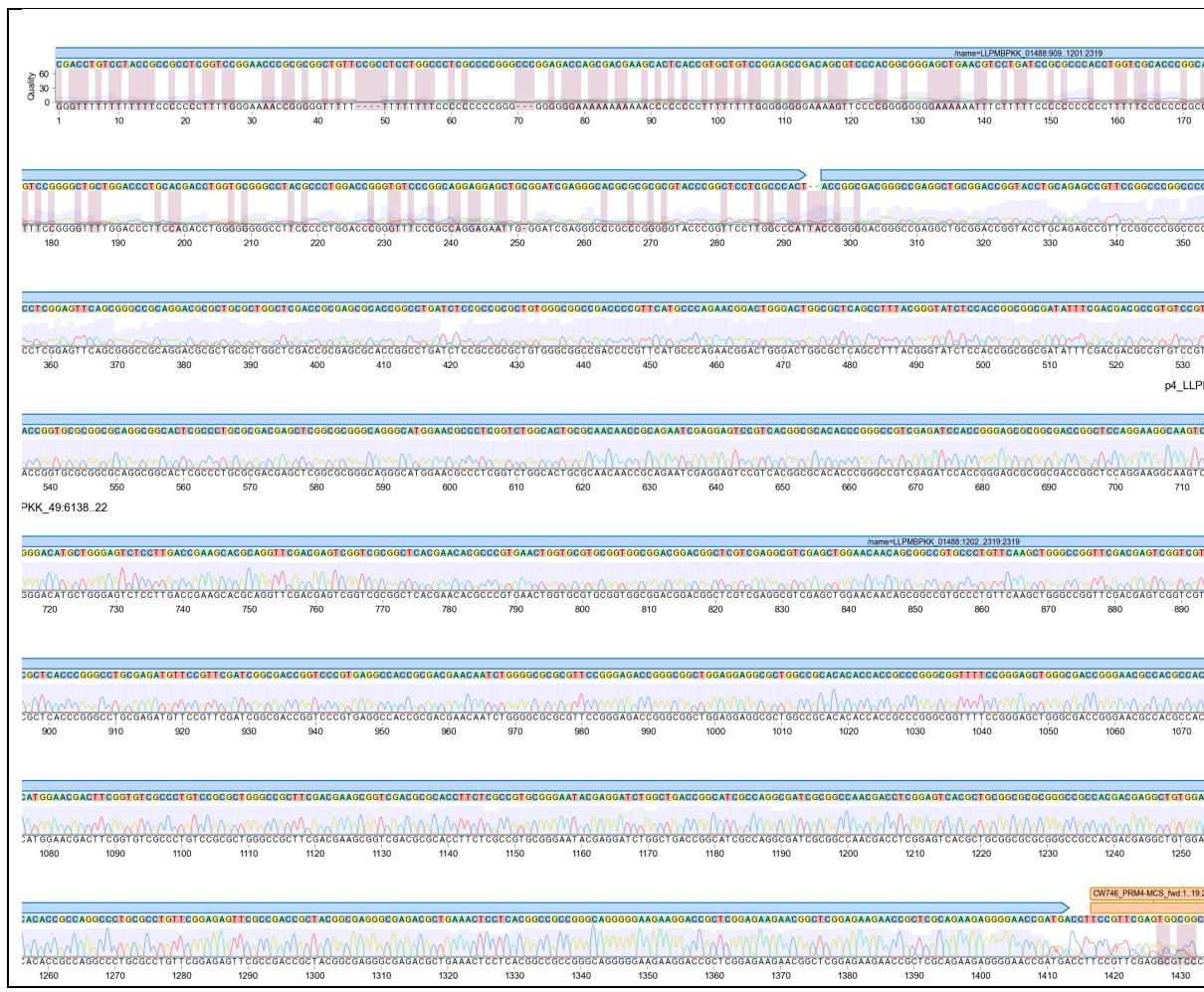
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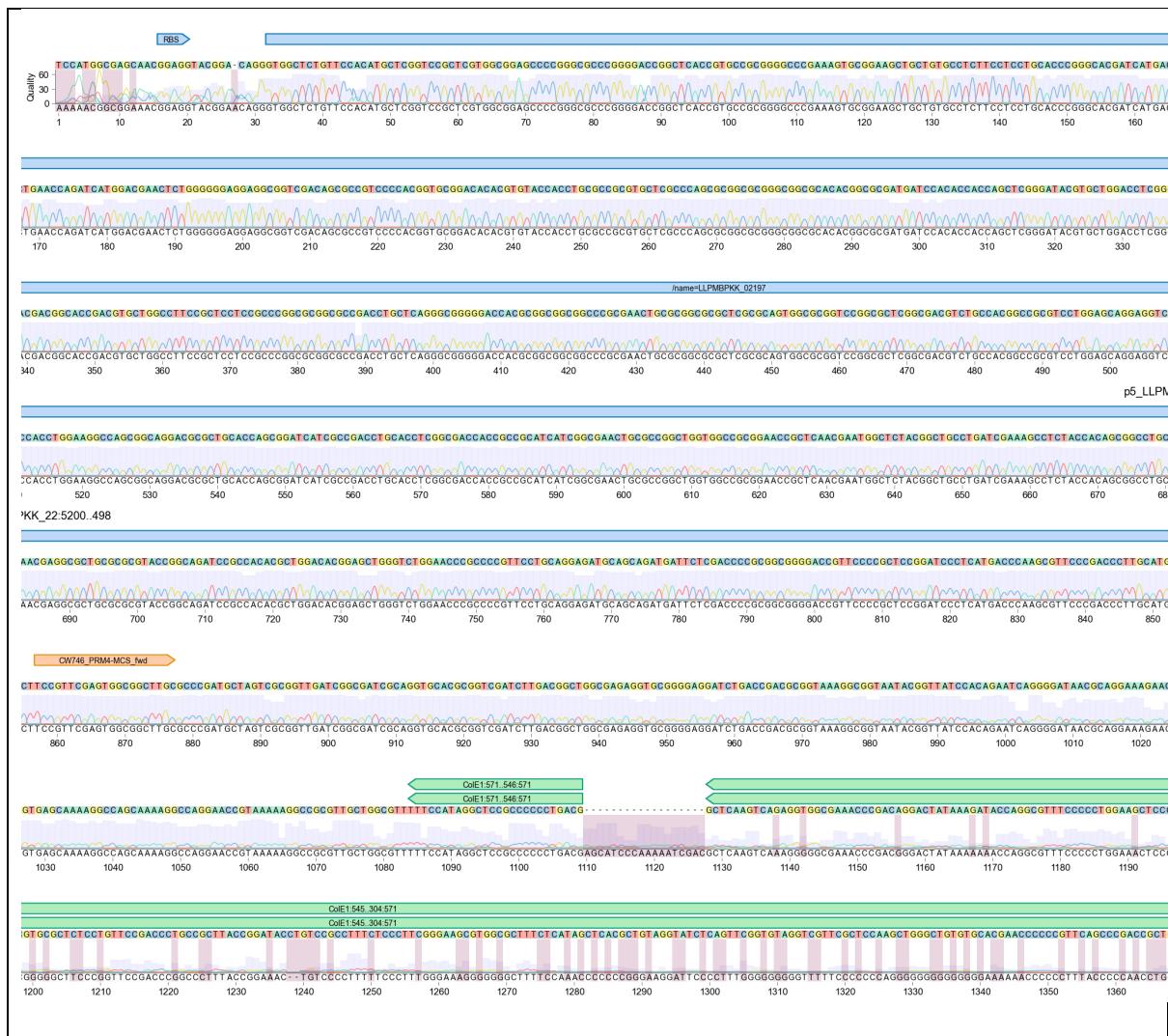
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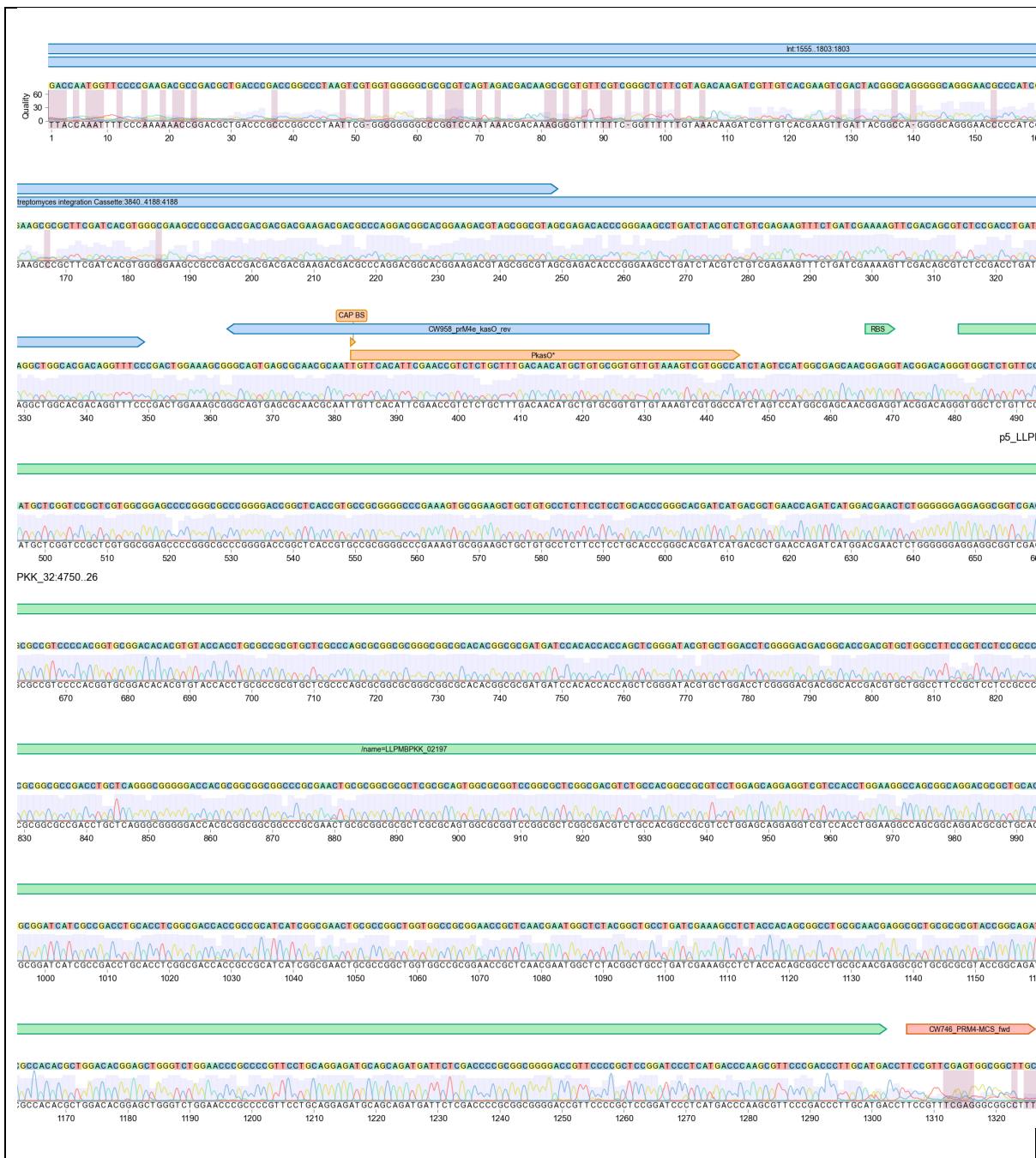
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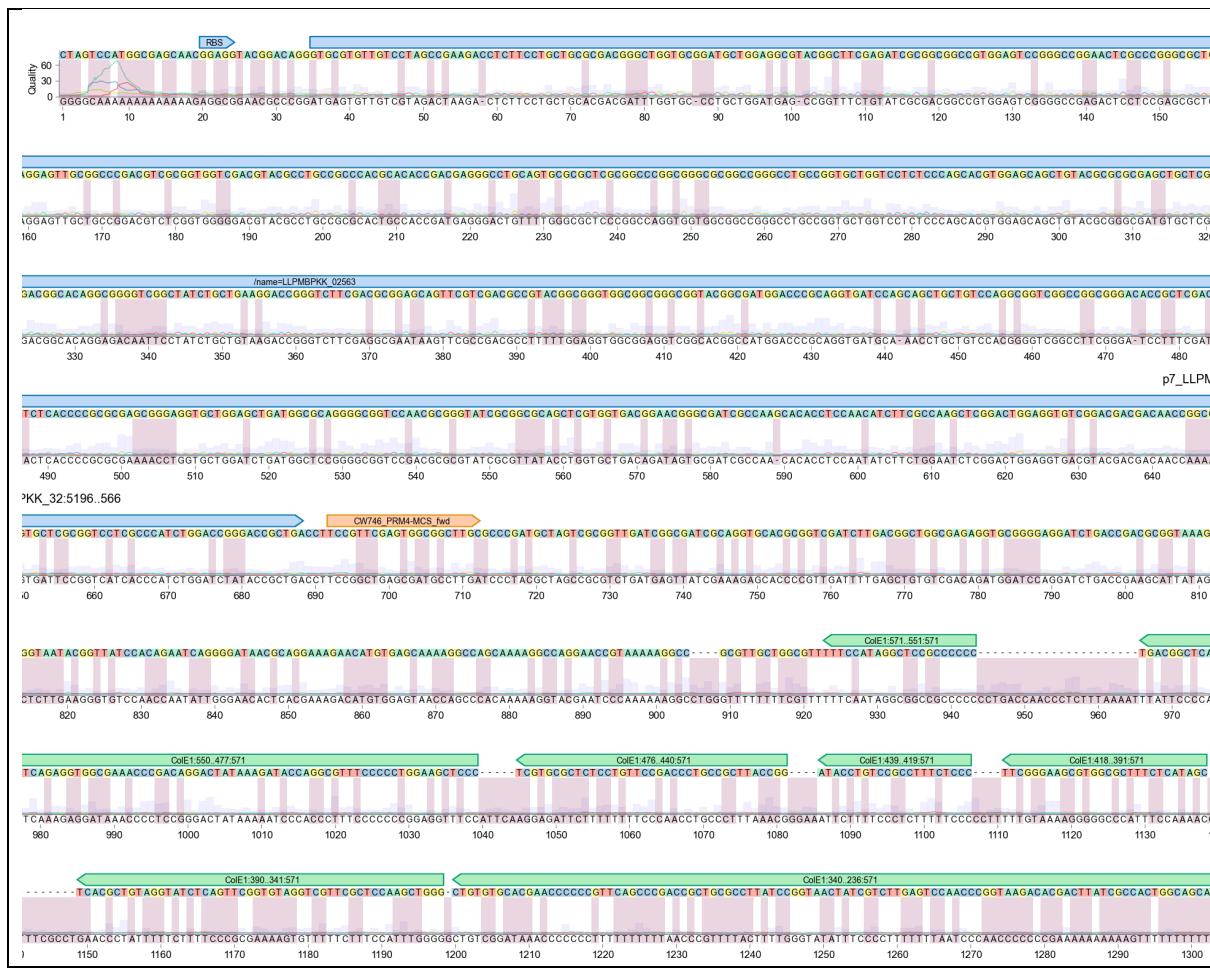
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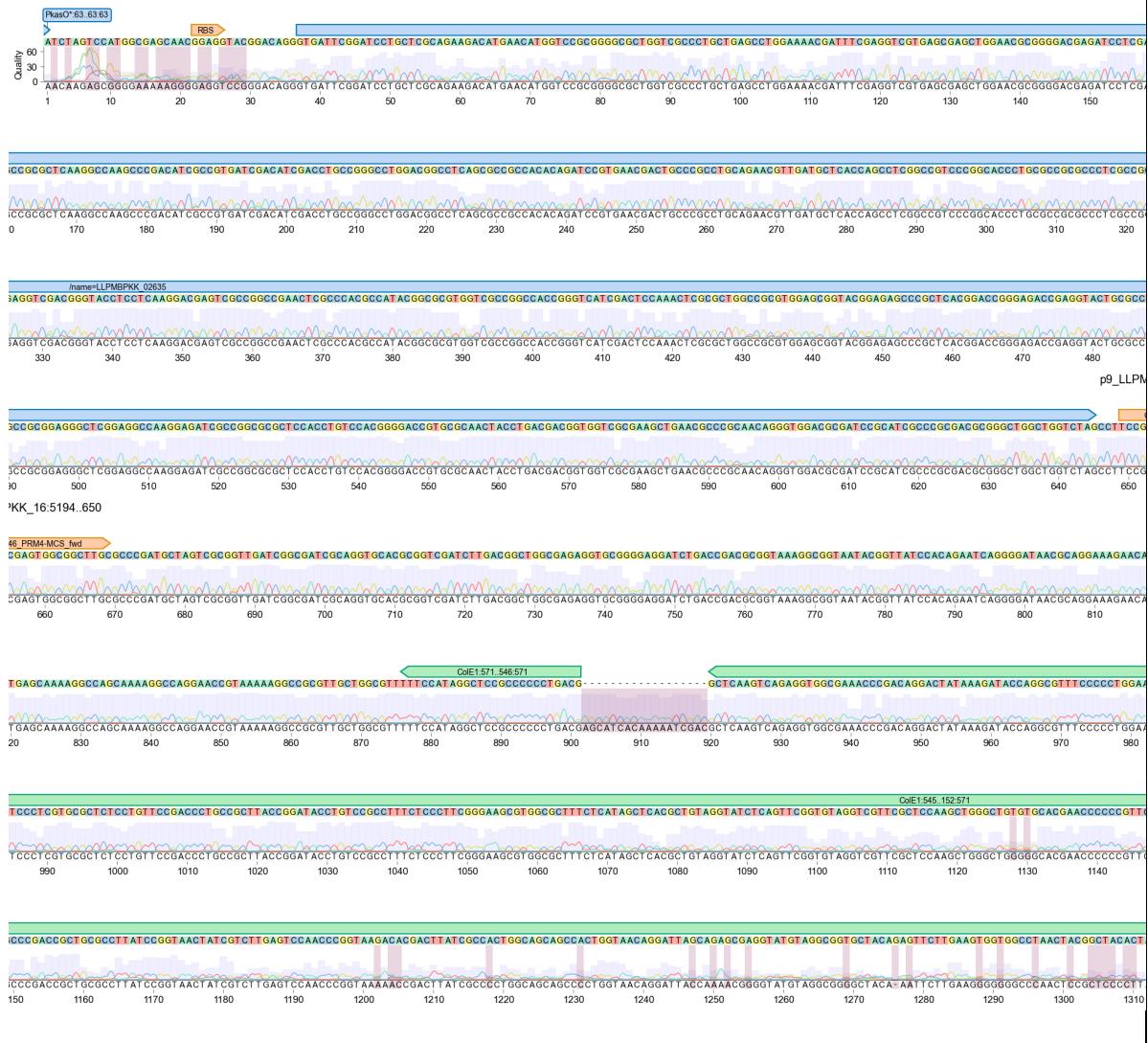
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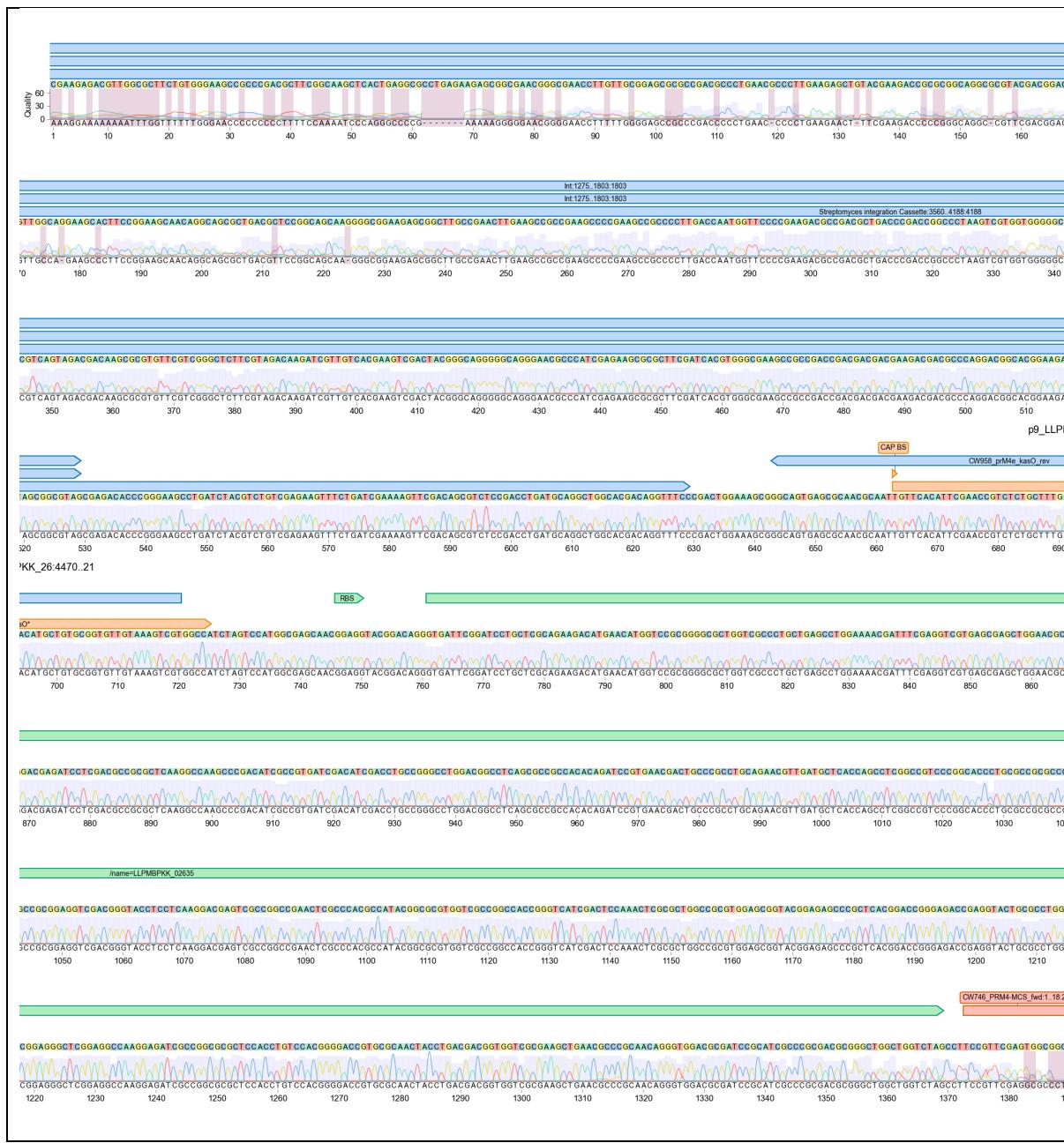
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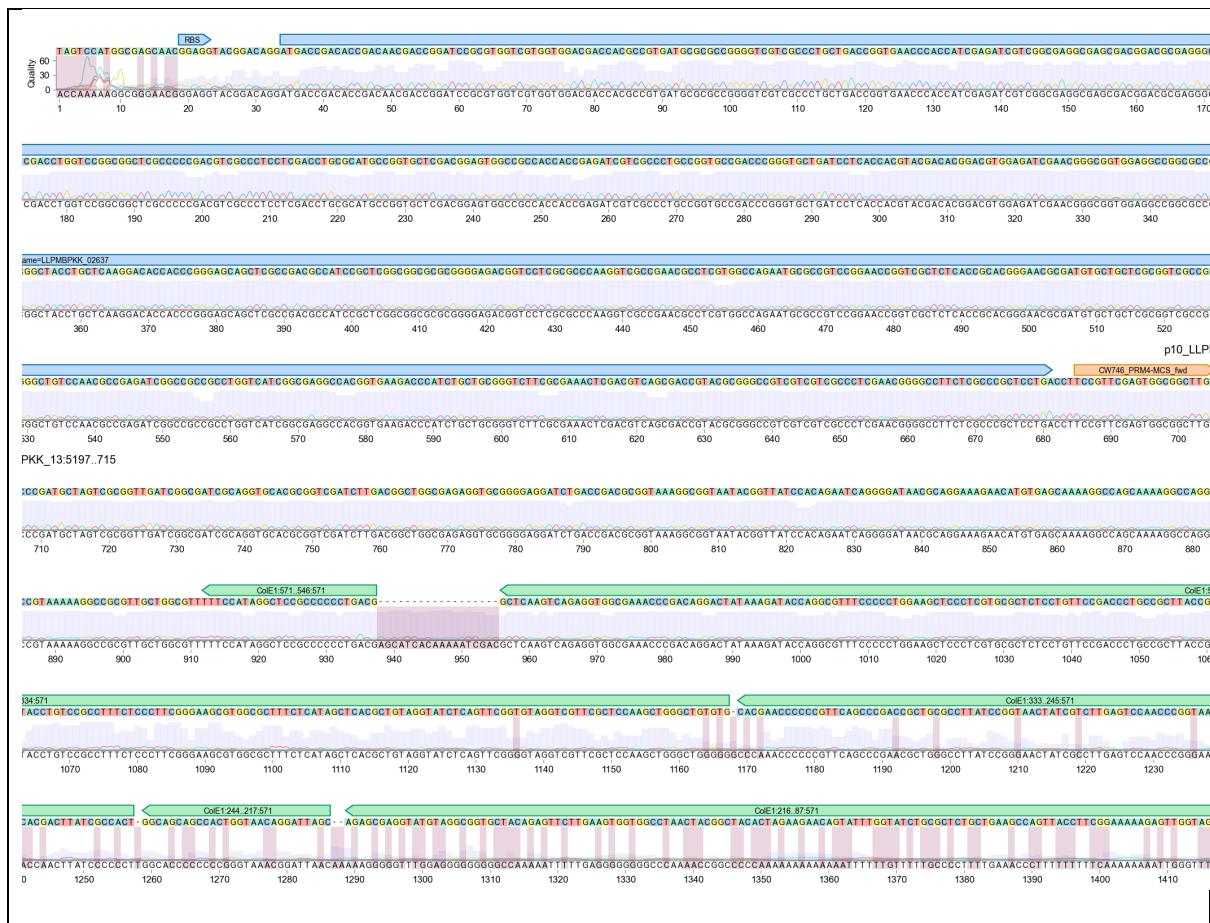
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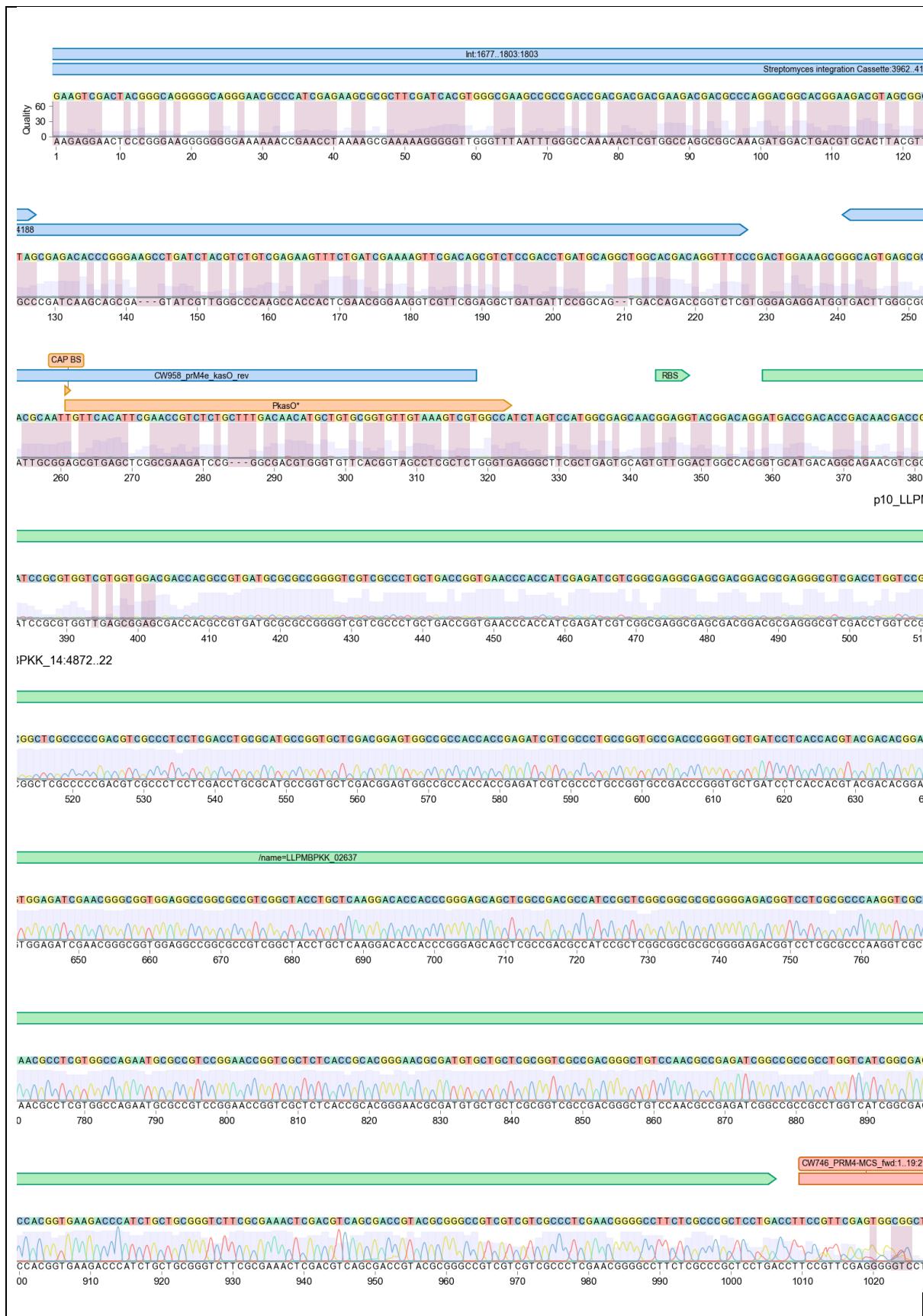
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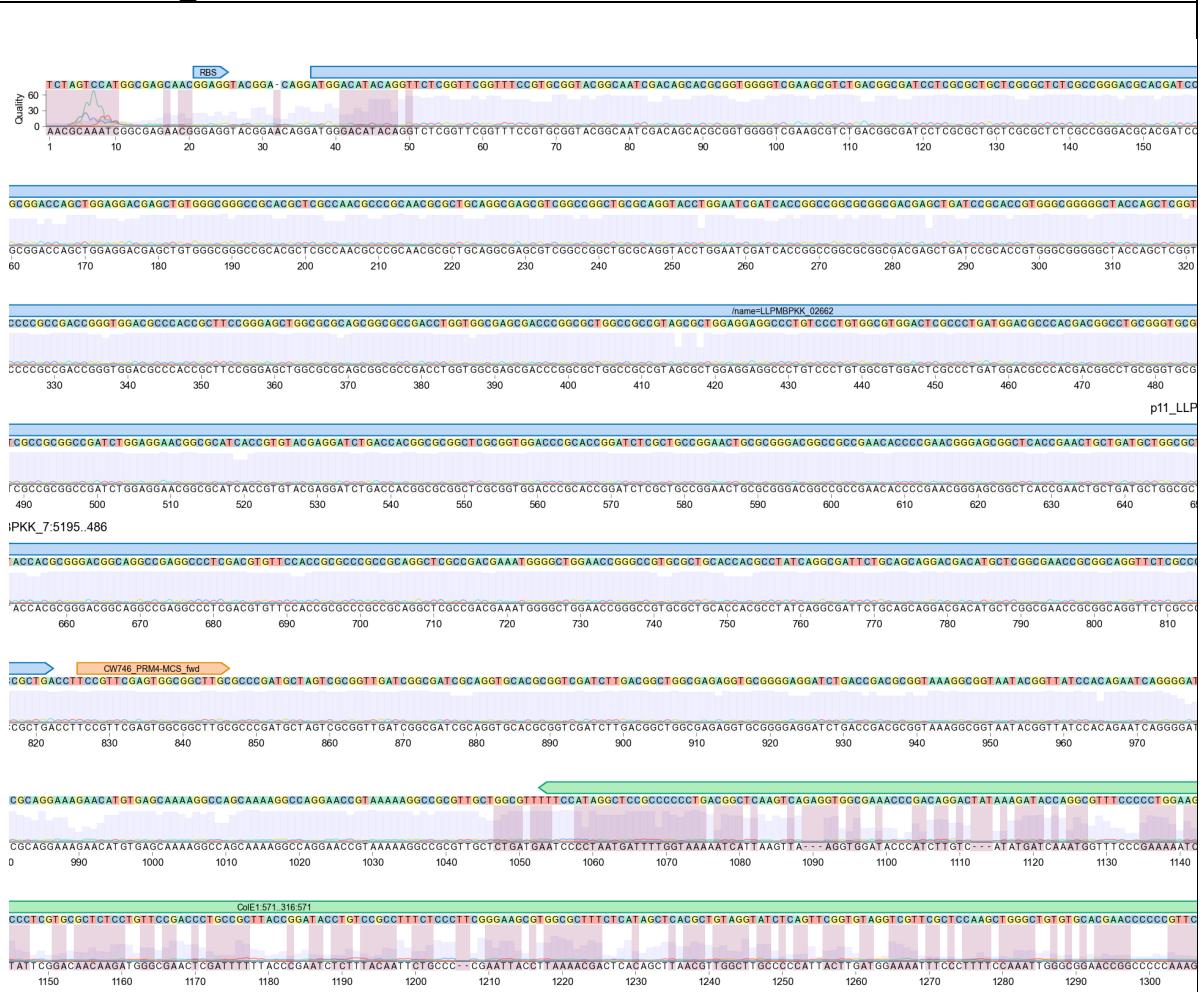
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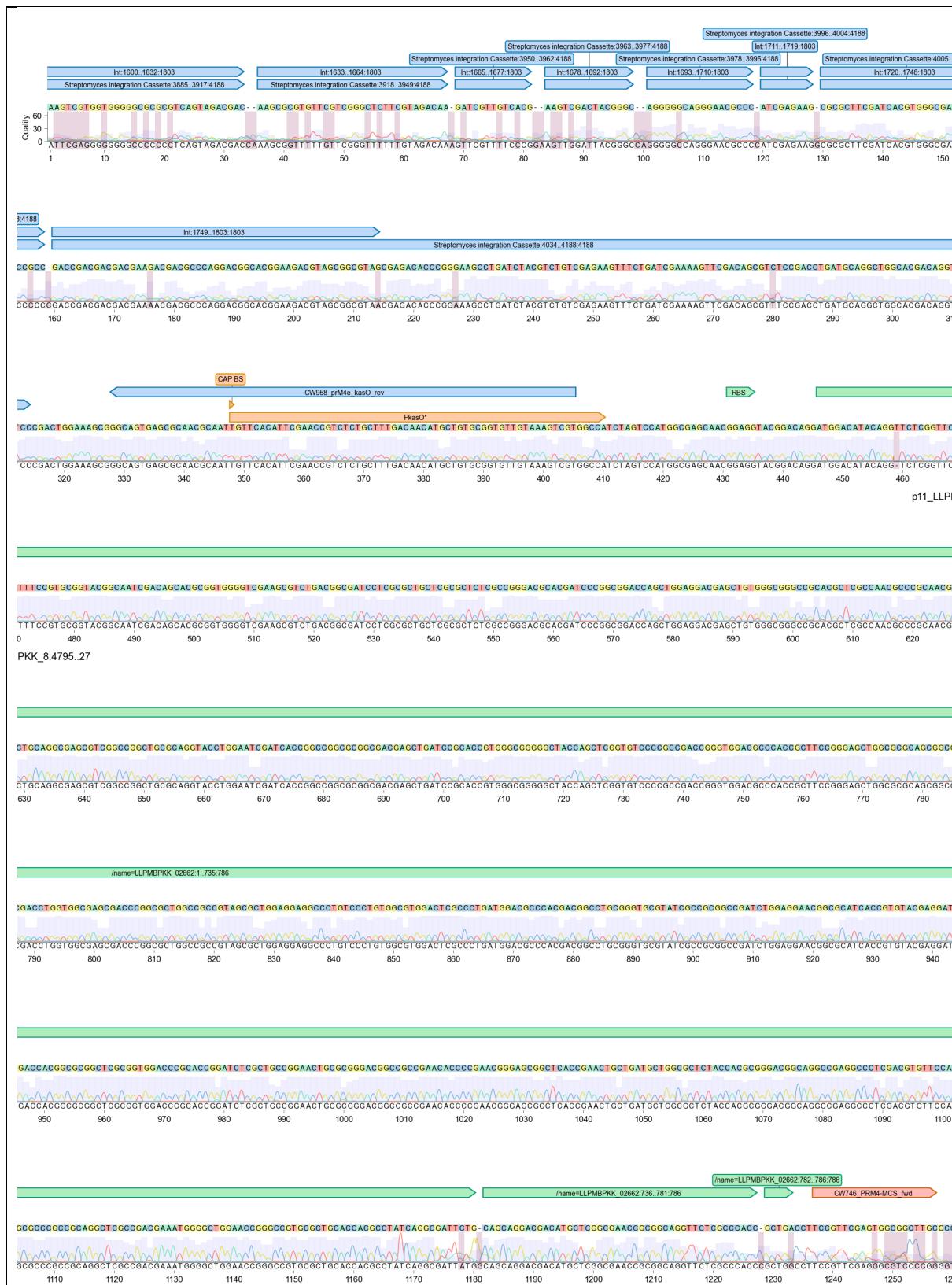
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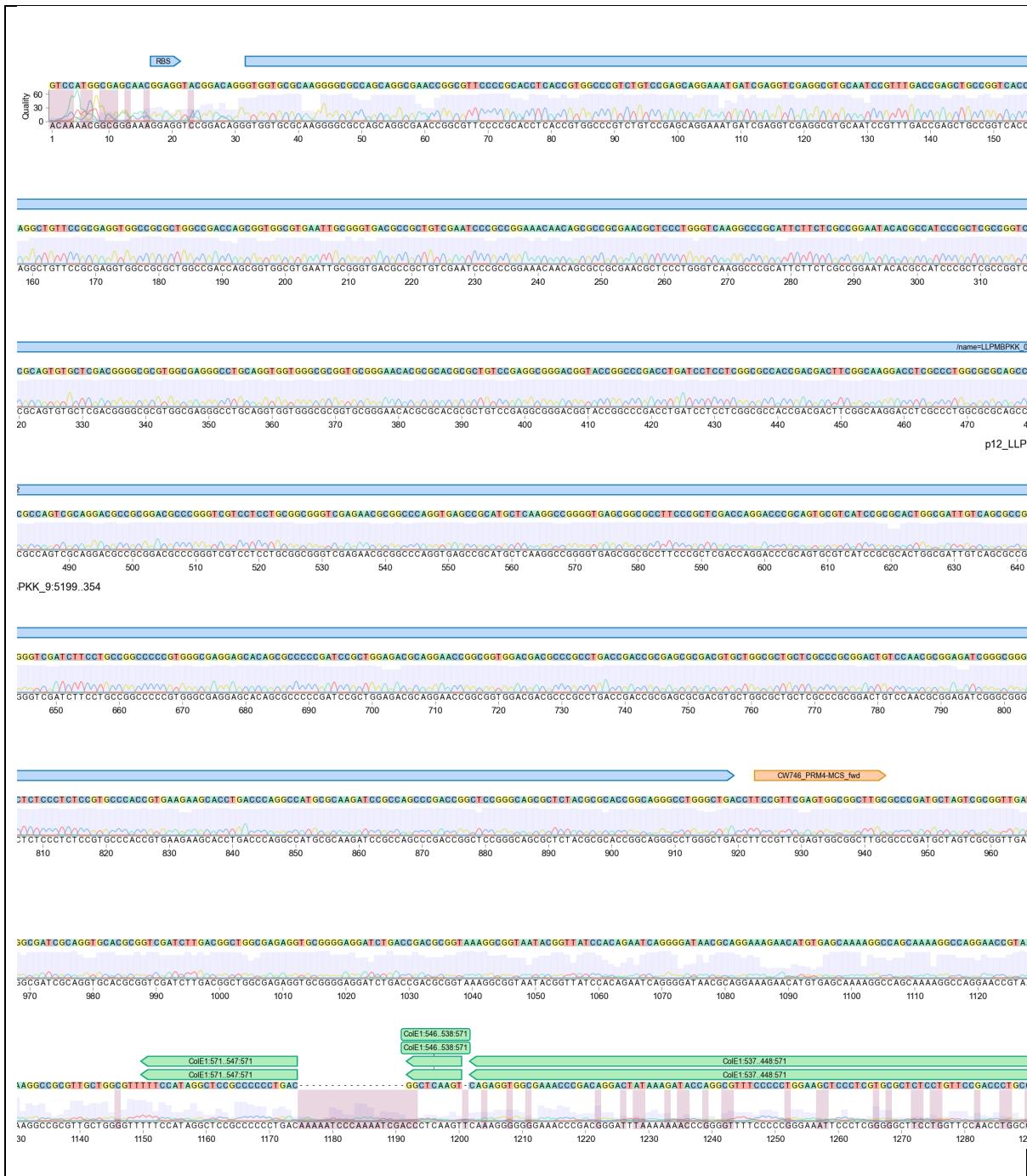
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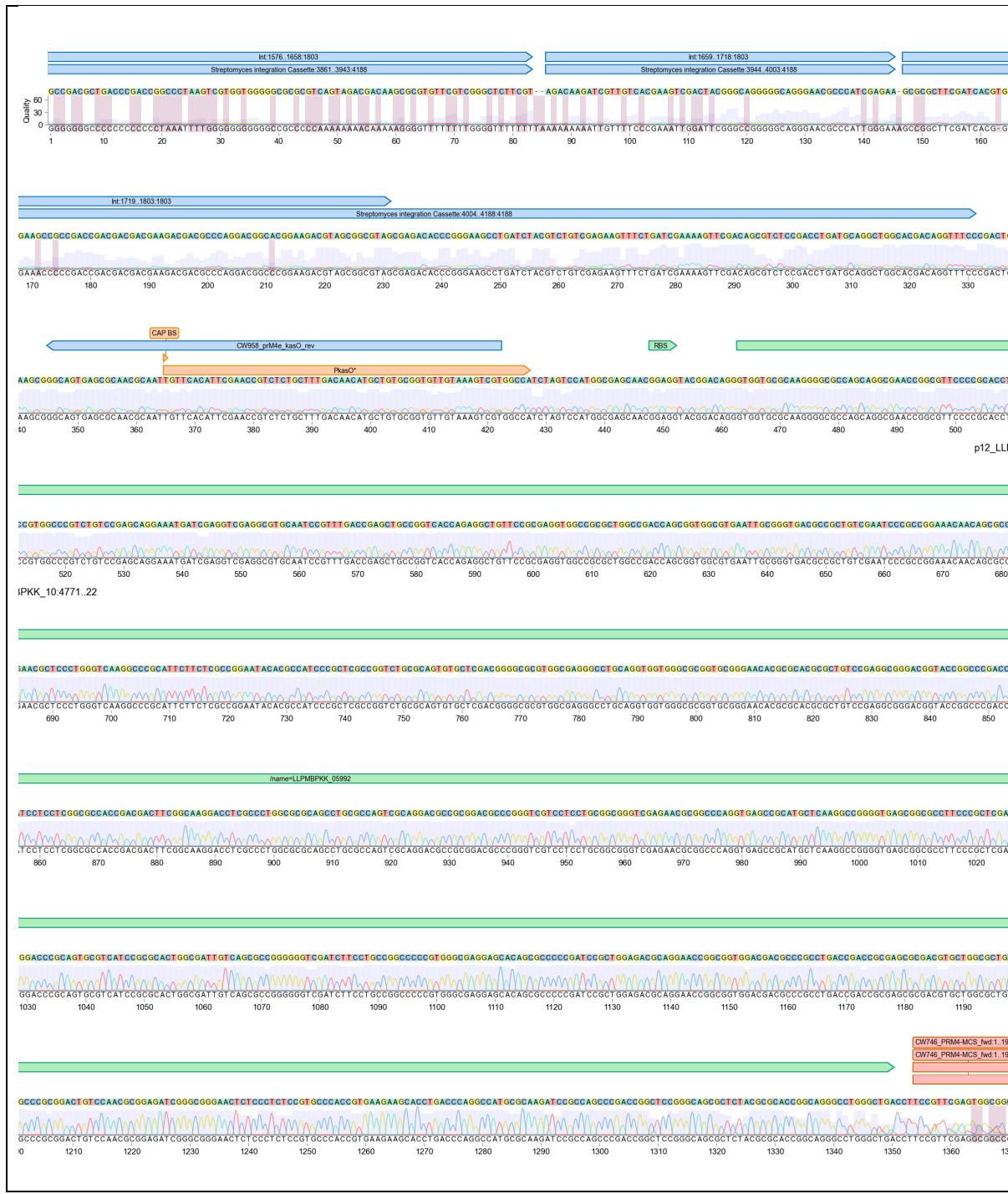


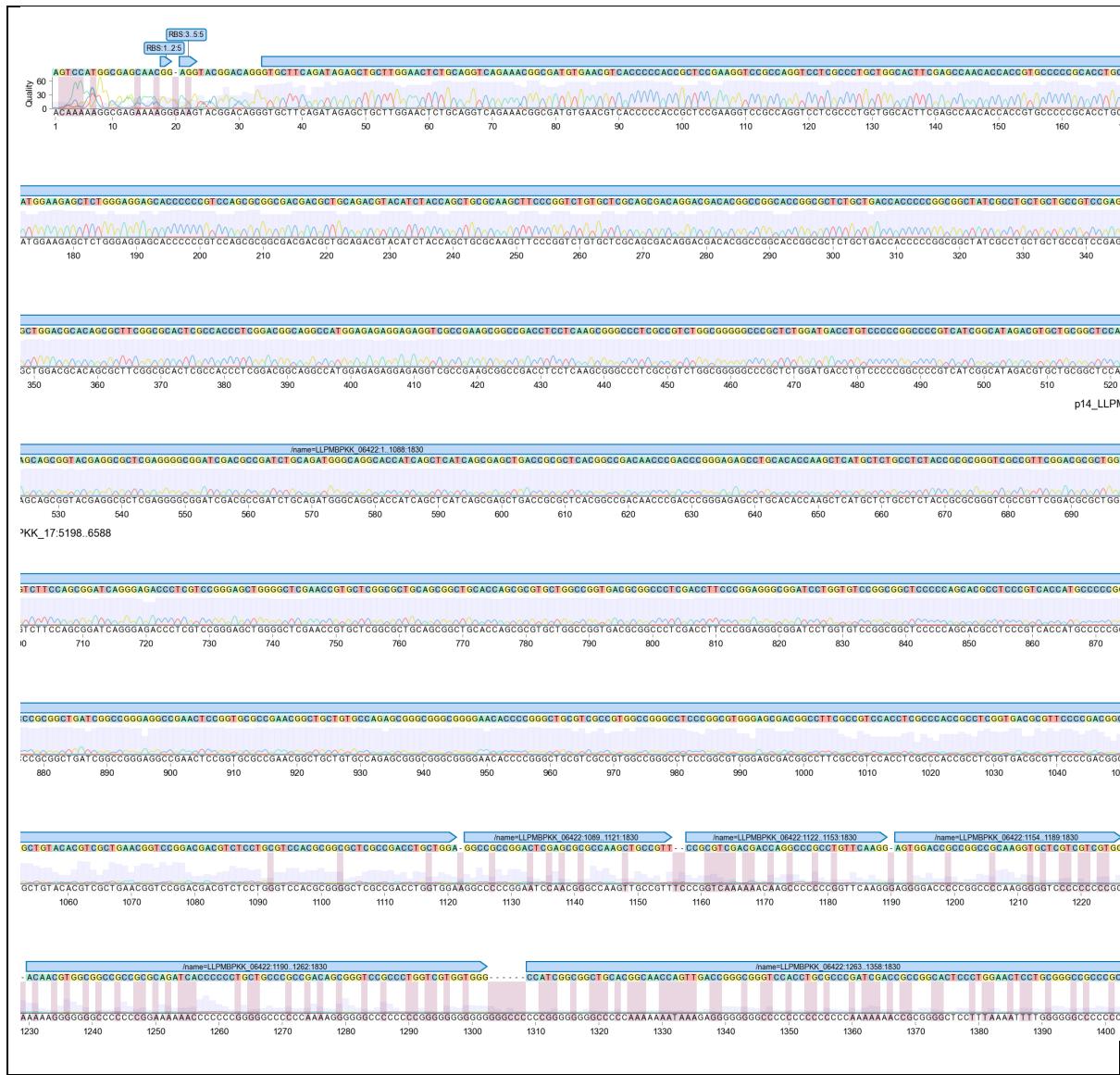
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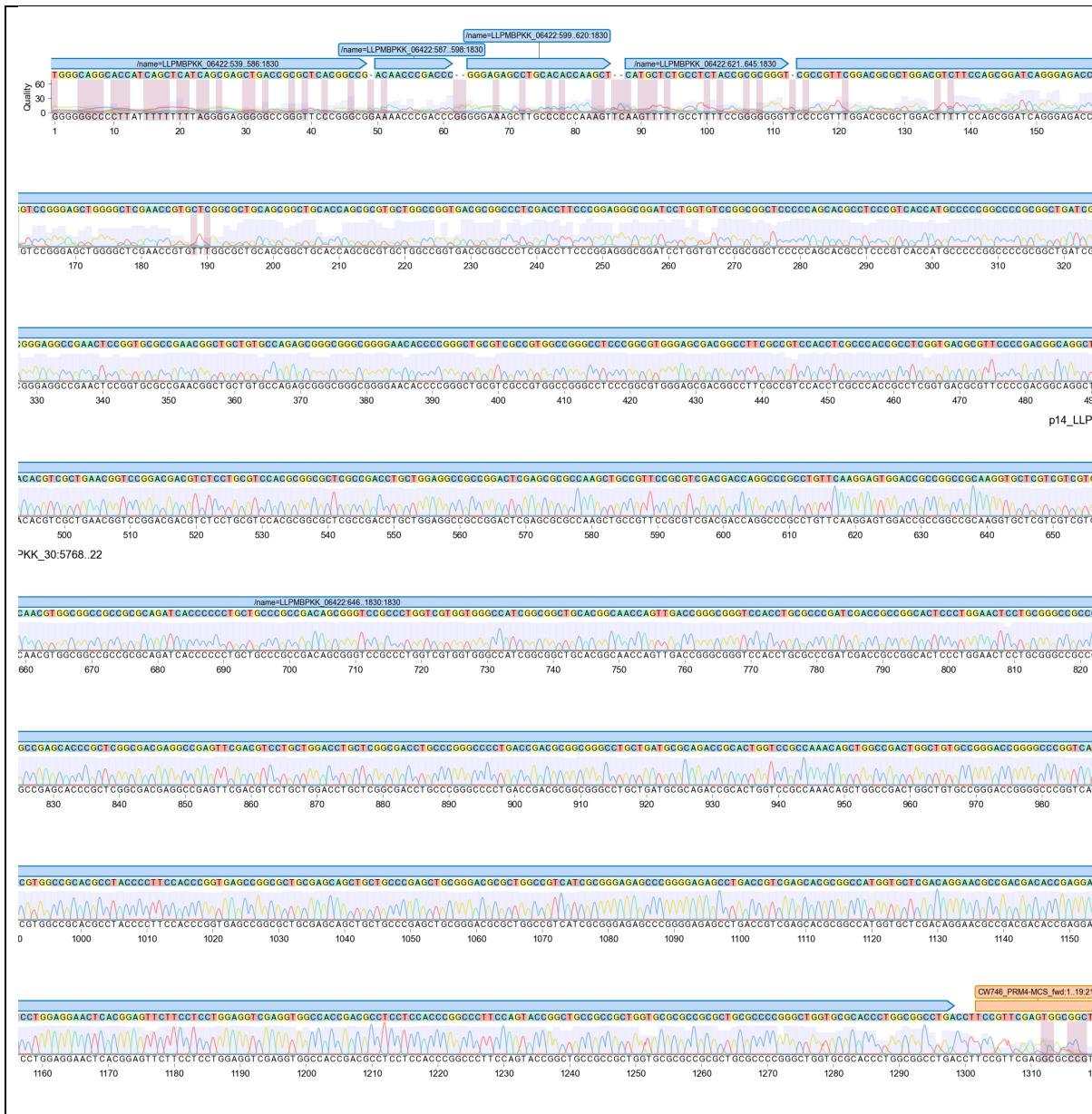
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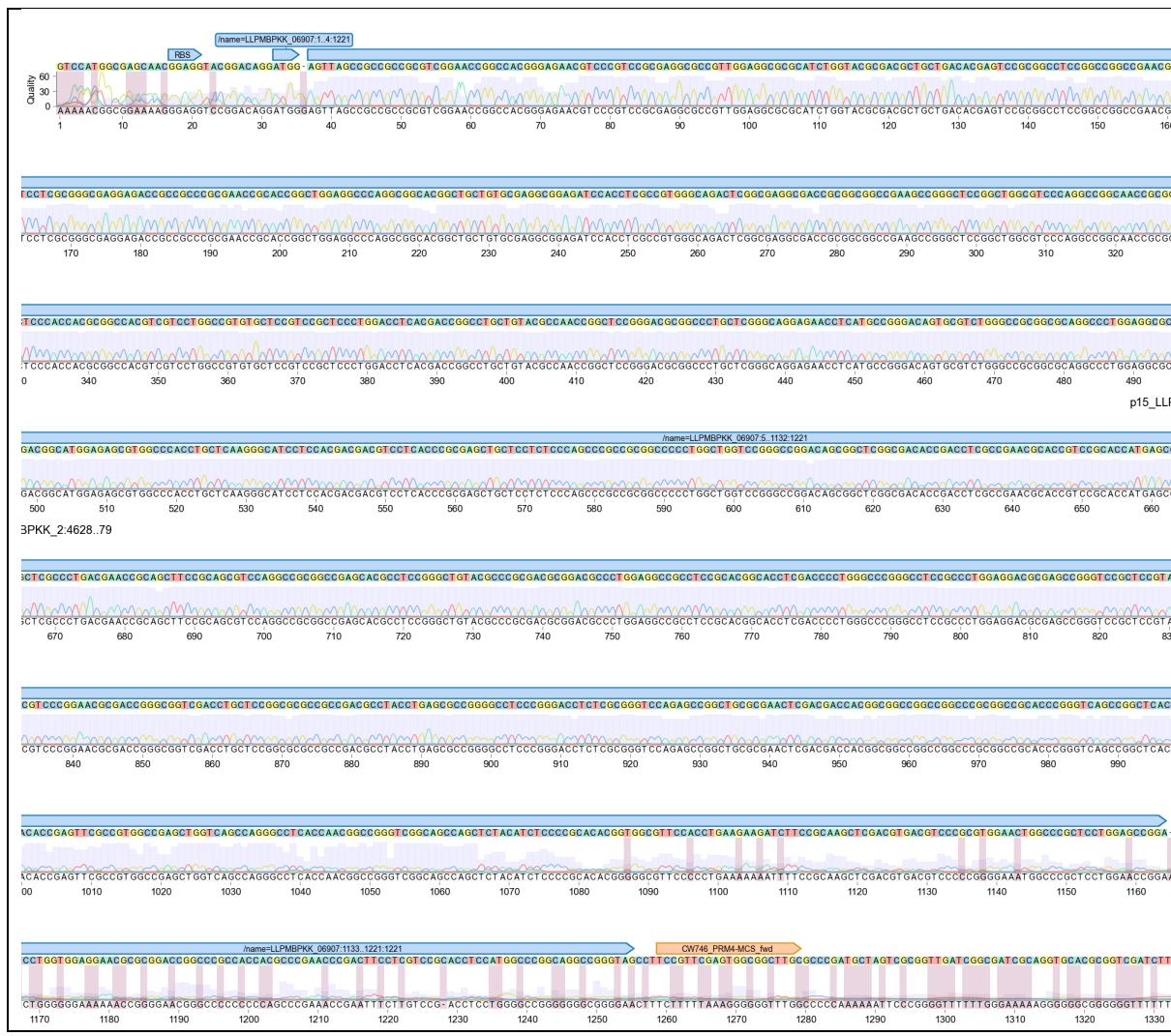




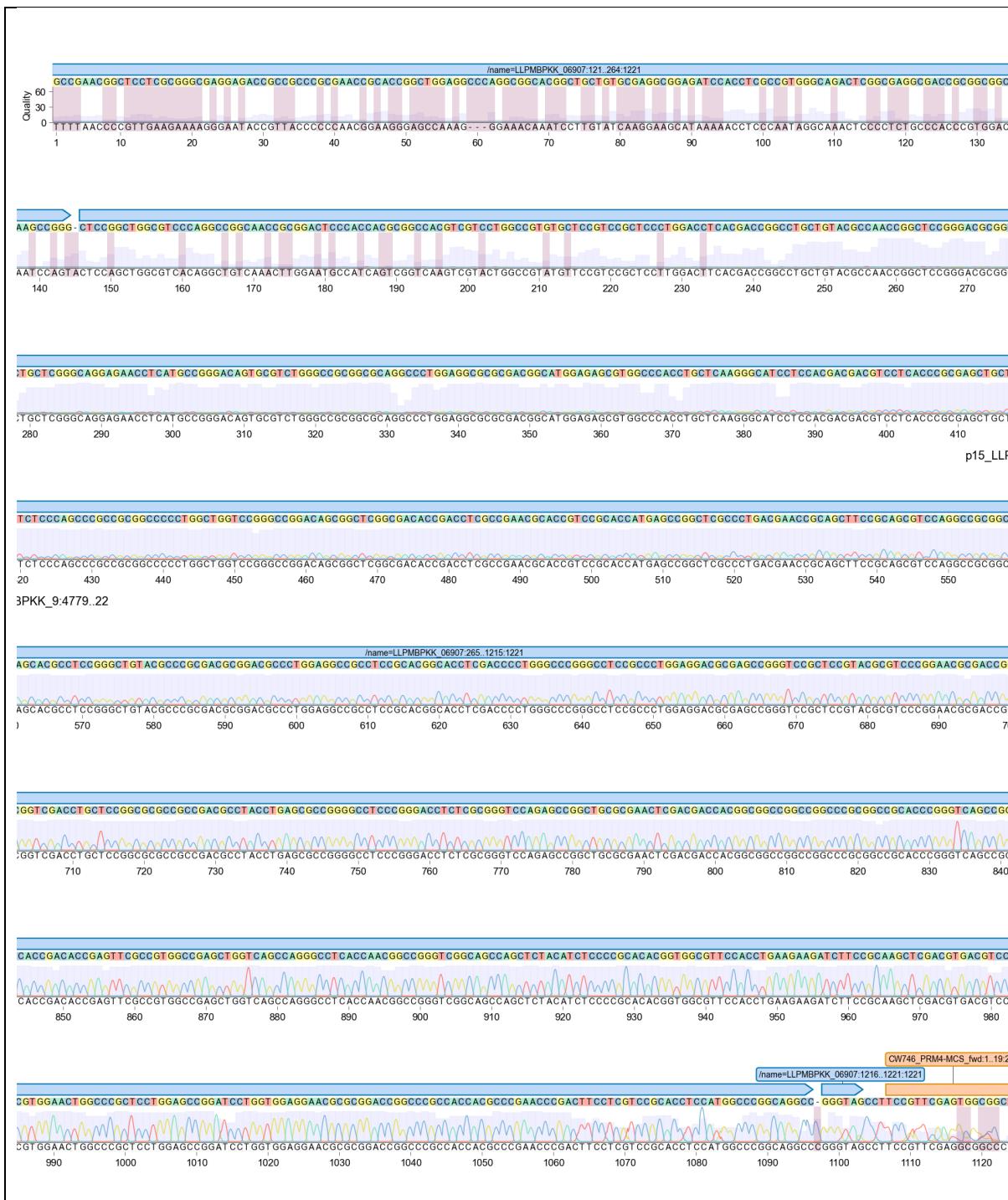
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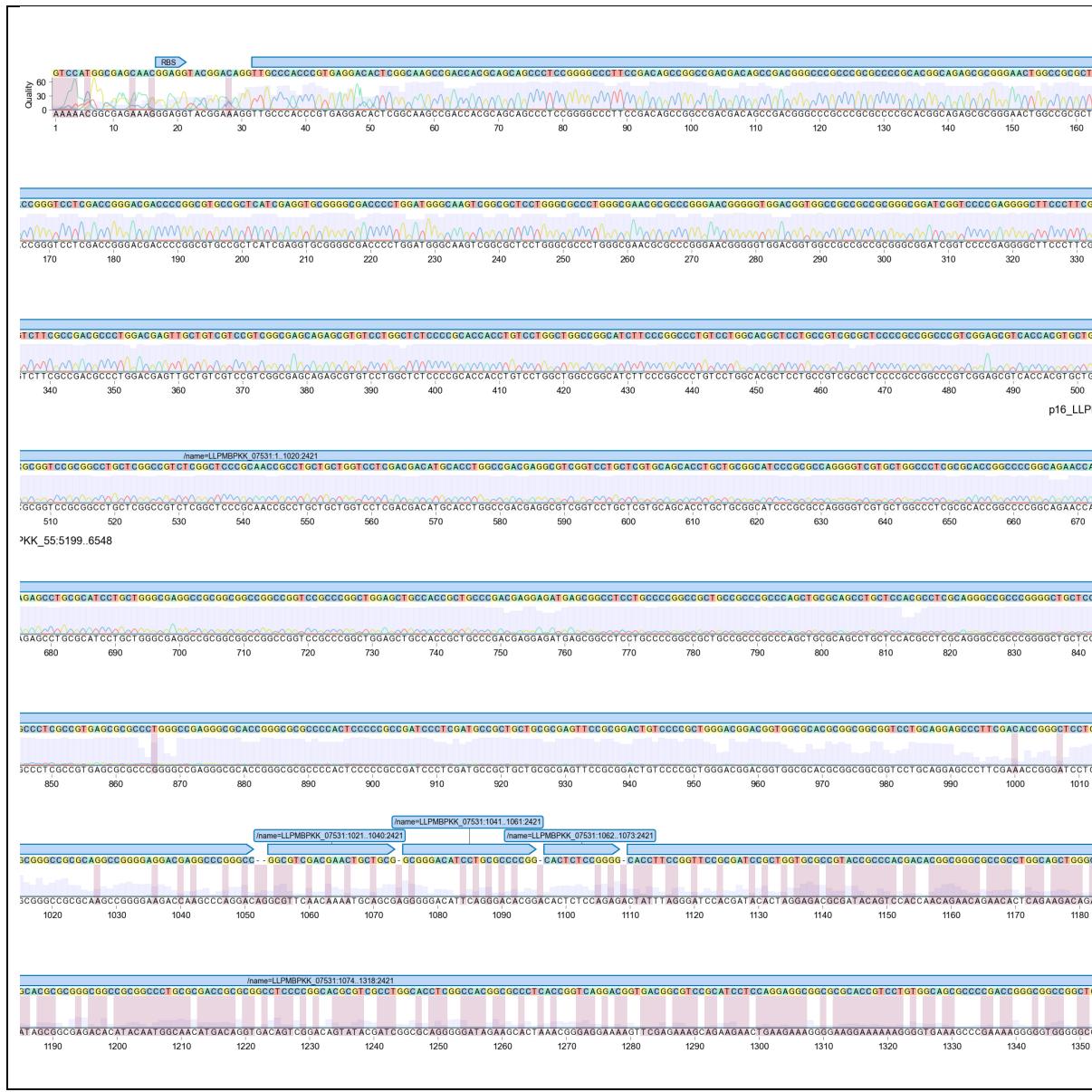
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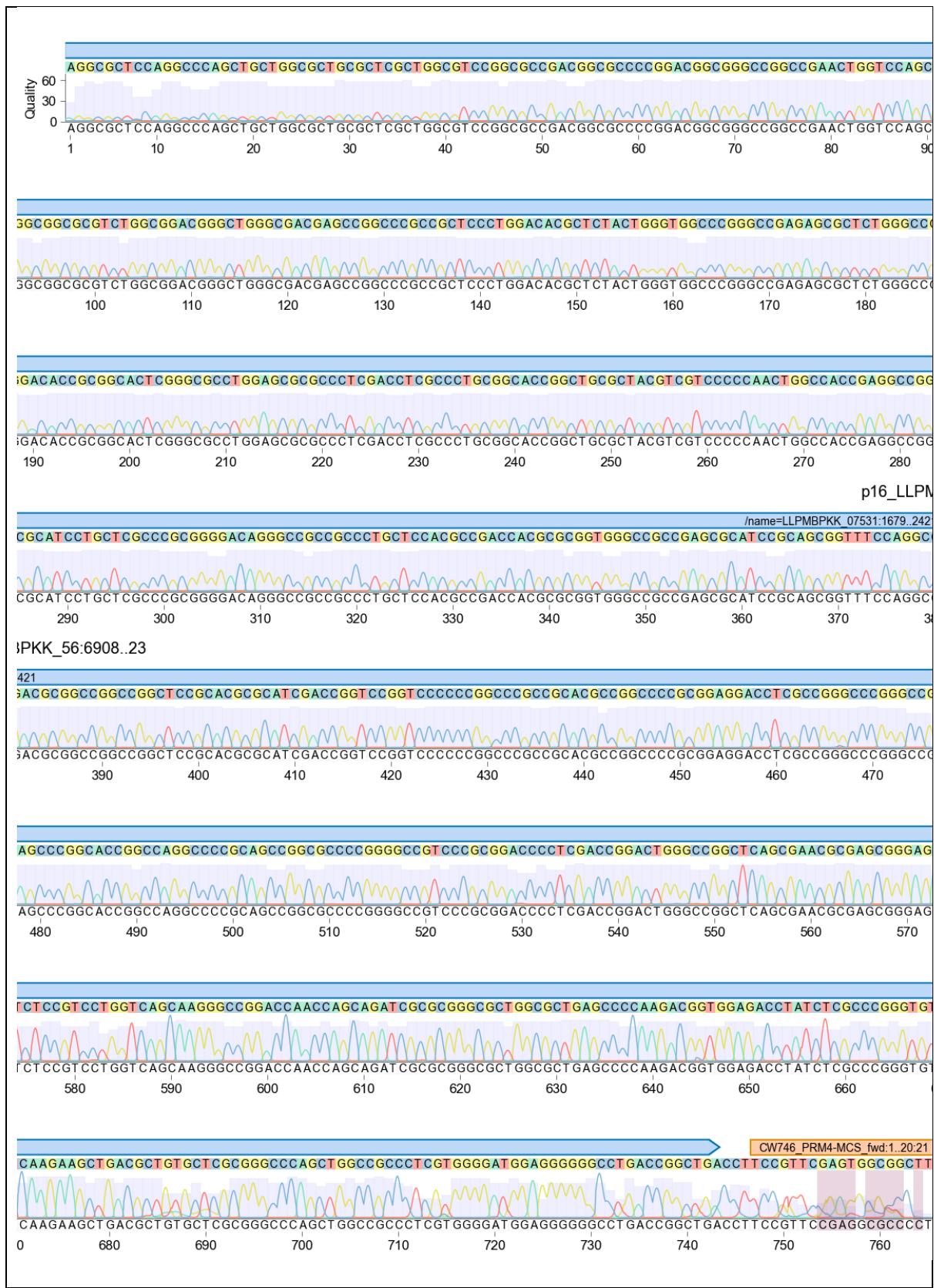
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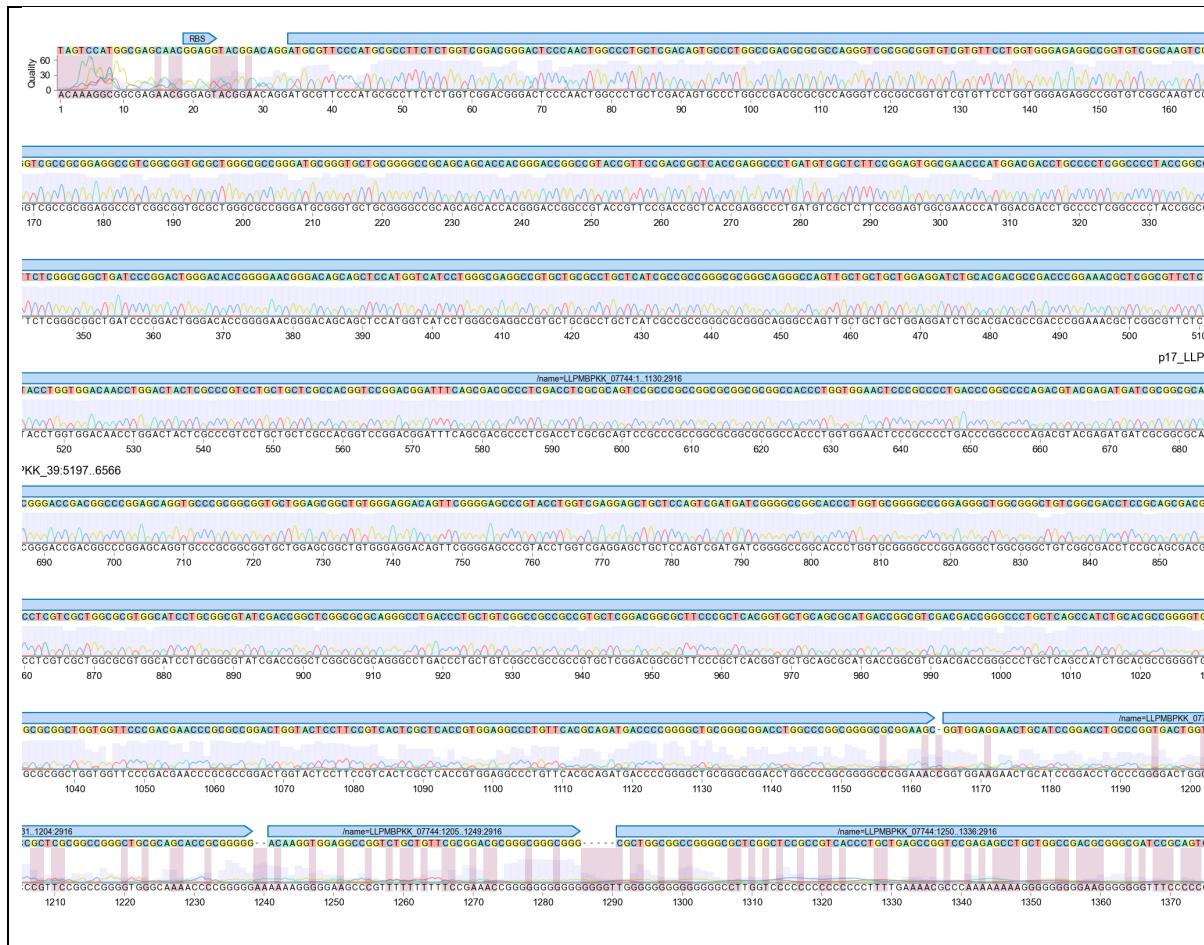
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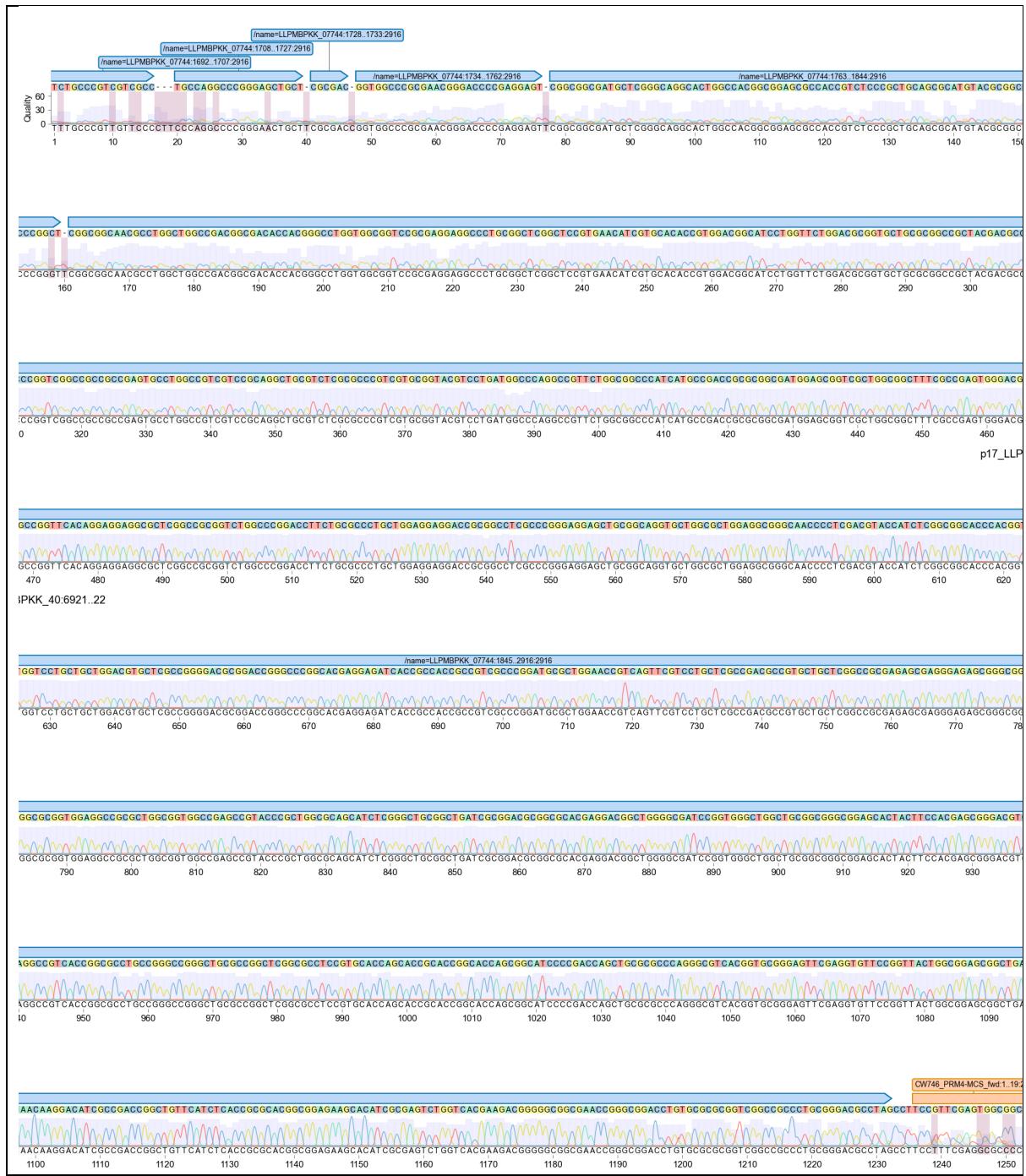
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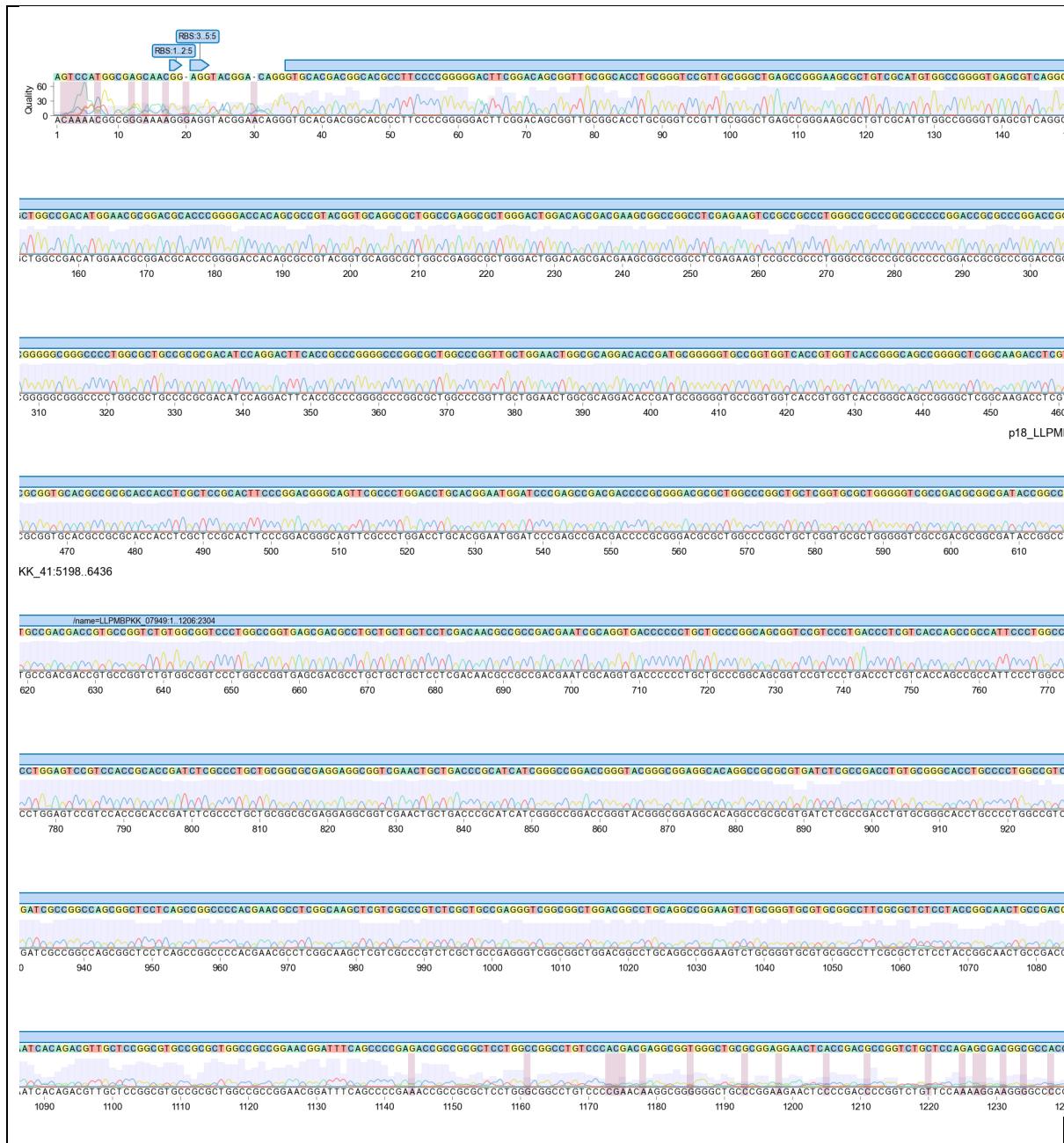
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