

Using whole genome comparison to detect sequence similarities between plants and microbes

By

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DECLARATION

I declare that this thesis, which I hereby submit for the degree, Philosophiae Doctor (Plant Science-Plant Biotechnology) at the University of Pretoria, is my own work and has not previously been submitted by me for a degree at any other University

Horses	
4	15 December 2007
Signed	Date



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Summary

With an increasing amount of whole genome sequence data becoming available on a daily basis we have an opportunity to study the interactions and dynamics of different organisms on a whole genome level. In the past, reports of horizontal gene transfer have focused mainly on the identification of single genes that show distorted phylogenetic profiles to that of the organism it was isolated from. This study firstly did whole genome comparisons between the rice nuclear and plastid genomes to determine the level and dynamics gene transfer and insertion of the chloroplast ad mitochondrial genomes into that of the nuclear genome of rice. Secondly, it looked to identify sequence similarities between the rice genome and microbial genomes by performing whole genome comparisons between the rice genome and that of several microbial genomes. These sequences were analyzed further to identify possible instances of horizontal transfer of DNA from microbes to the rice genome. Using this approach, this study reports several fragments in the rice genome with significant sequence similarity to that of microbial DNA fragments. This study also provides evidence supporting horizontal transfer of several of these fragments. This study provides valuable information regarding intra- as well as inter-genome DNA transfer dynamics.



Thesis Composition

The following thesis comprises of five sections. 1 Introduction and literature review presents an overview regarding the origin and evolution of the eukaryotic genome as well as the origin of the chloroplast and mitochondrion. It also review previous reports regarding DNA transfer between the organelles and nucleus, as well as those regarding transfer of DNA between various microbes and plants. It also discusses the proposed models whereby transfer occurs. 2 Materials and Methods outlines the different strategies and methods and sequences used in the whole genome comparisons and the analysis of the results. 3 Results describes the results obtained from each of the different comparisons. 4 Discussion provides an overall discussion of the study and provide some future perspectives of research in the field of horizontal DNA transfer. In the Annex the tables detailing the sequence similarities found between the rice genome and the various other genomes used are presented.

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