

IDENTIFICATION AND CHARACTERISATION OF MARKERS LINKED TO THE LEAF RUST RESISTANCE GENE *LR37*



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PREFACE

The results represented in this thesis follow from the study, which was carried out at the Forestry and Agricultural Biotechnology Institute and Department of Genetics at the University of Pretoria under the supervision of Prof A-M. Oberholster and the co-supervision of Dr. F.J. Kloppers.

The results represented here are original and have not been submitted in any form to another University.

Gonhie

Christiaan Troskie



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LIST OF ABBREVIATIONS

AFLP	Amplified fragment length polymorphism
2AS	Chromosome 2A, short arm
BSA	Bulk segregant analysis
c	Recombination fraction
сп	Chlorotic/ Necrotic
cM	Centimorgan
D	Genetic distance between any two individuals
Dn	Russian wheat aphid resistance gene
DNA	Deoxyribonucleic acid
dNTP	Deoxyribonucleotides
DTT	Dithioerylthritol
EDTA	Ethylenediaminetetraacetic acid
et ul.	<i>Et alii</i> (and others)
F	Index of genetic similarity
IPTG	Isopropylthio-B-D-galactoside
kb	Kilobases
LB	Luria Bertani
Lr	Leaf rust resistance gene
M	Molar
m/v	Mass/volume
MAS	Marker-assisted selection
МЬ	Megabases
ng	Nanogram
NILS	Near isogenic lines
PCR	Polymerase chain reaction
PEG	Polyethylene glycol
Pm	Powdery mildew resistance gene
pM	Picomolar
PPi	Difosfate
RAPD	Random amplified polymorphic DNA
RFLP	Restriction fragment length polymorphism
Rpm	Revolutions per minute
s	Similarity coefficient
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SCAR	Sequence characterized amplified region
SDS	Sodium dodecyl sulphate
Sr	Stem rust resistance gene
SSRs	Small sequence repeats
STS	Sequence-tagged site
Taq	Termus aquaticus
TBE	Tris-borate/EDTA
TEN	Tris-EDTA-soduim chloride
Tris	Tris (hydroxymethyl)- aminomethane
U	One unit of enzyme
UHQ	Distilled and UV treated water
UPGMA	Unweighted pair-group mean arithmetic
UV	Ultraviolet
v	Volts
X	Distance of marker to the gene
X-Gal	5-bromo-4-chloro-3-indolyl-B-D-galactoside
Yr	Yellow rust resistance gene
μg	Microgram
μL	Microliter