



# WGIN Programme: TILLING in hexaploid wheat

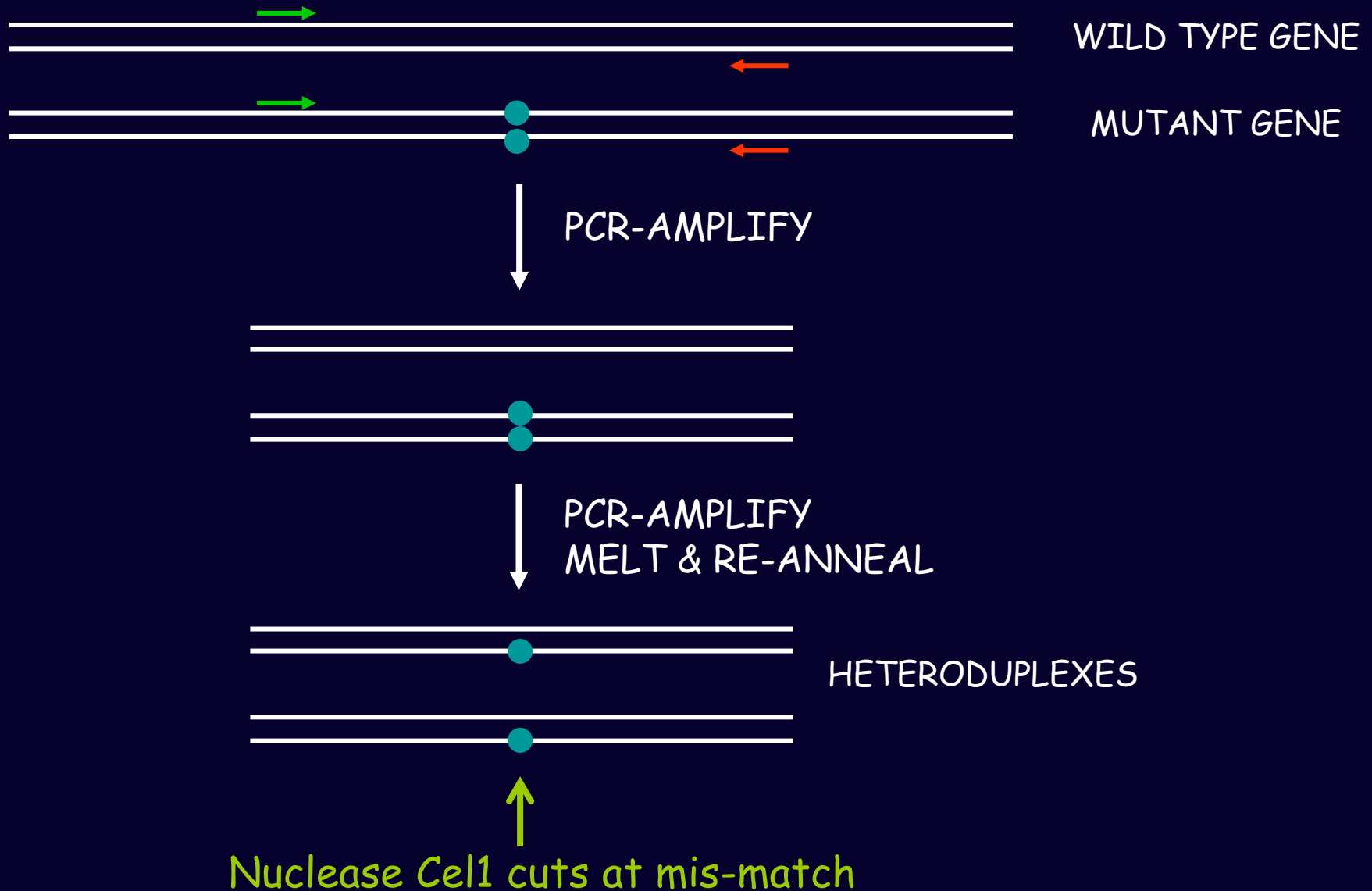
Year 1 Report

# What is TILLING?

- ❖ Targeting Induced Local Lesions in Genomes
- ❖ PCR-based reverse genetics screen for polymorphisms in genes controlling key traits
- ❖ Non-GM method of targeted gene modification in mutagenised populations
- ❖ Identifies natural variation in genes of elite varieties, historic lines and landraces.

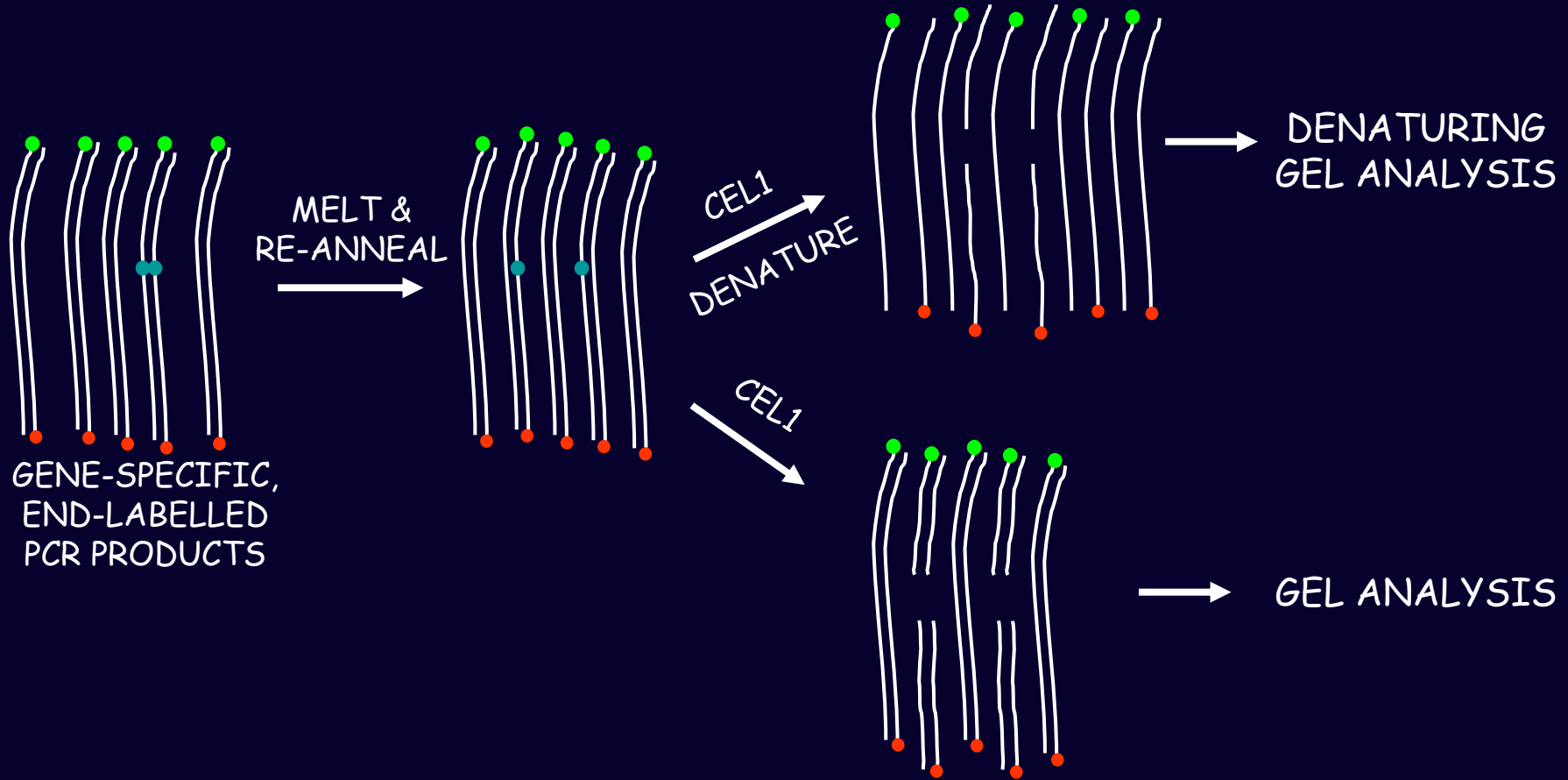
# TILLING

(Targeting Induced Local Lesions in Genomes)

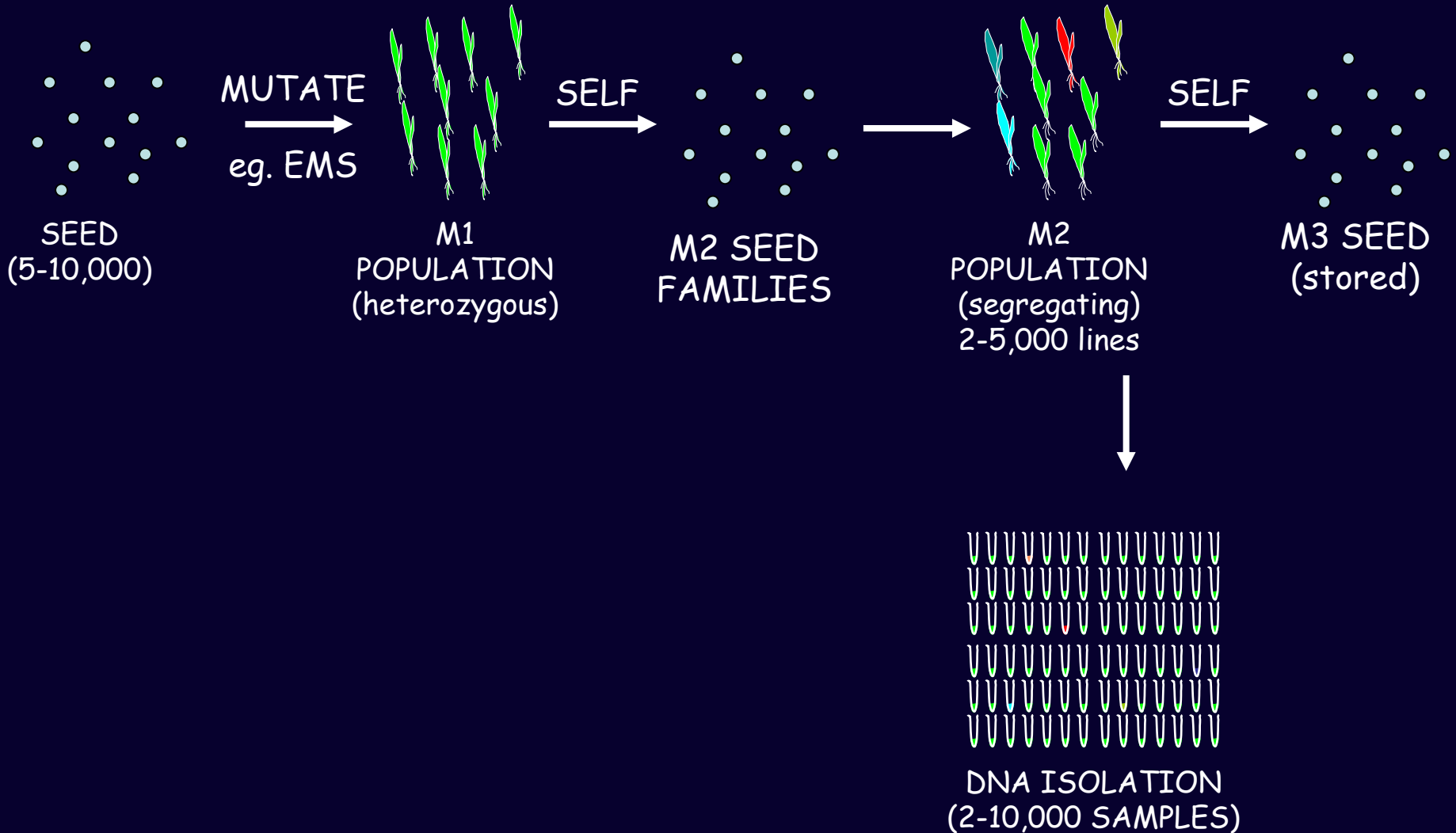


# TILLING

(Targeting Induced Local Lesions in Genomes)



# PRODUCTION OF MUTAGENIZED POPULATION



# HIGH-THROUGHPUT TILLING

eg. 3,000 M<sub>2</sub> LINES



3,000 DNA SAMPLES



375 @ 8-FOLD POOLS



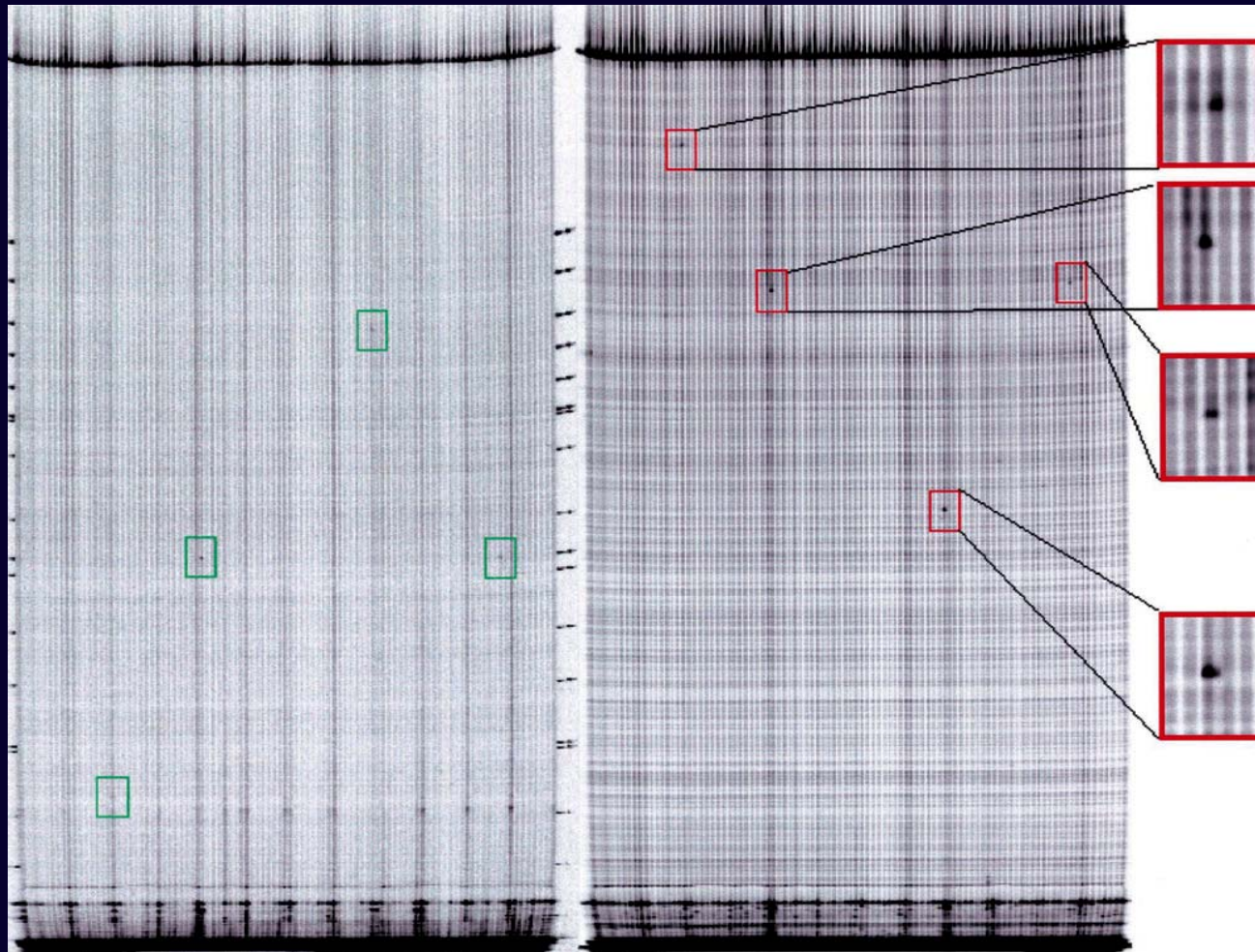
GENE-SPECIFIC PCR,  
DENATURATION,  
CEL1-CLEAVAGE



4 @ 96-LANE GEL RUNS

# TILLING EXAMPLE

## ARABIDOPSIS DNA, 8-FOLD POOLING



LEFT PRIMER

RIGHT PRIMER

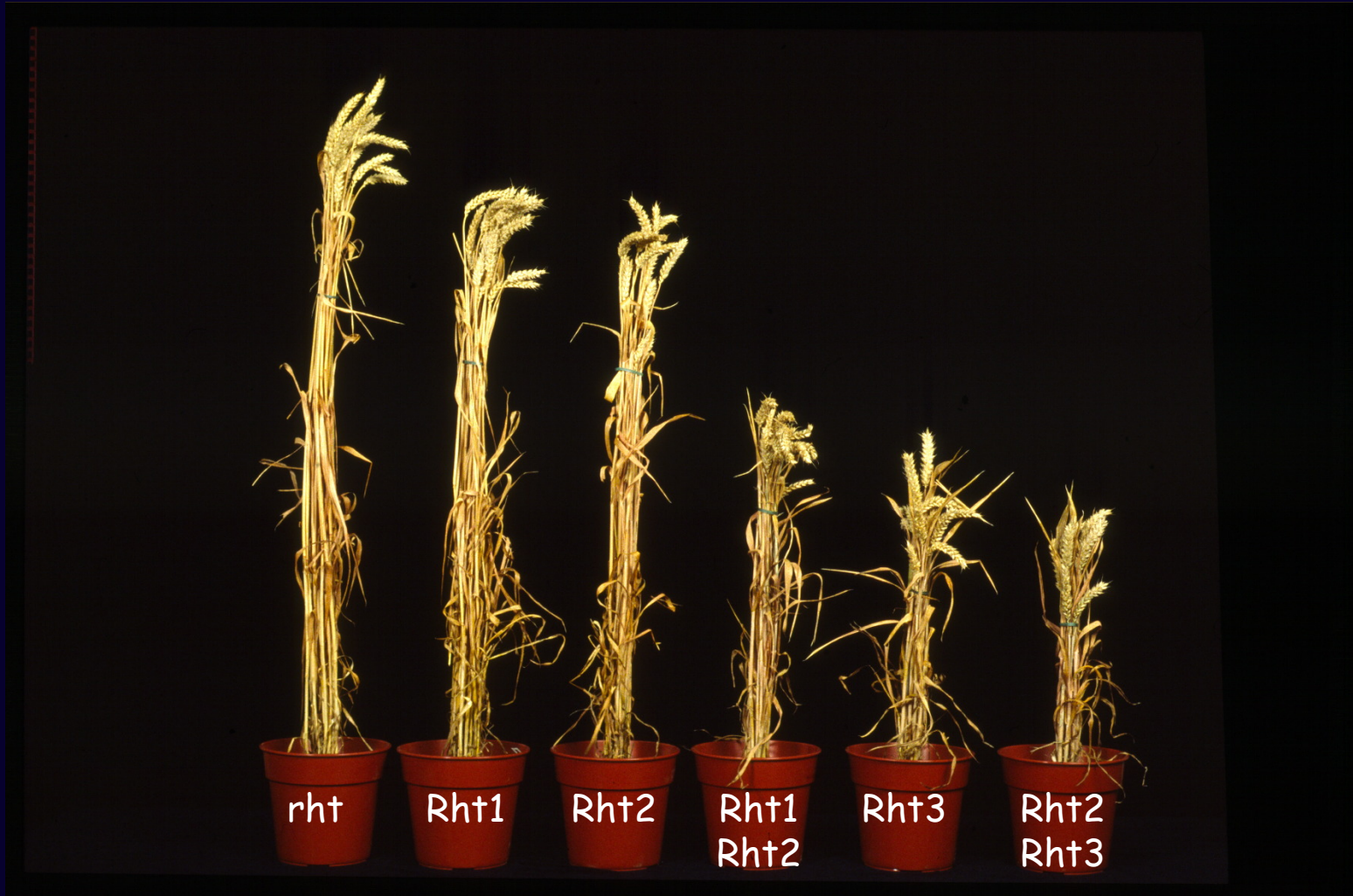
# TILLING FOR WHEAT IMPROVEMENT

- ❖ Non-GM approach
- ❖ Can screen mutagenized lines or natural variation
- ❖ Provides a series of alleles of varying strength
- ❖ Generates perfect genetic markers for breeding
- ❖ Possible to target complex multigenic traits
- ❖ Targeting individual homoeologues of wheat uncovers traits not detectable by normal screens
- ❖ But - do need specific genes as targets - dependent on good basic knowledge of gene function
- ❖ And - need to develop homoeologue-specific PCR amplification at each target locus (rate-limiting)
- ❖ However - high ploidy level of wheat allows high levels of mutagenesis



# INITIAL WGIN TILLING TARGETS

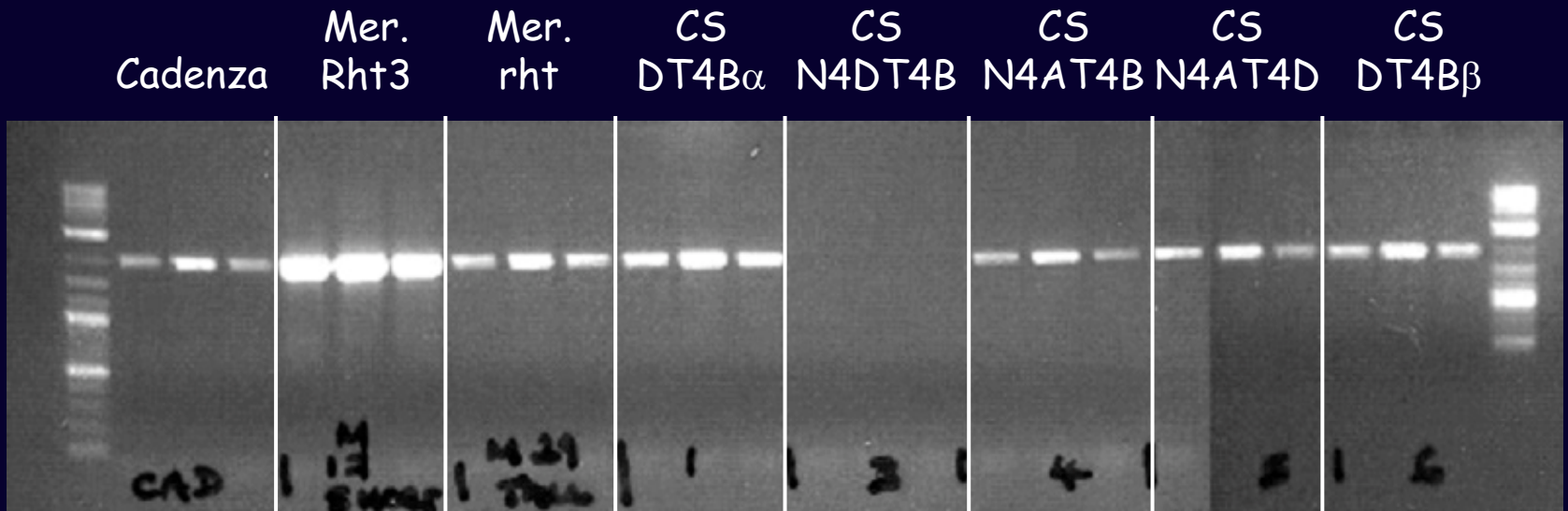
## NEW ALLELES OF RHT AS PROOF-OF-CONCEPT



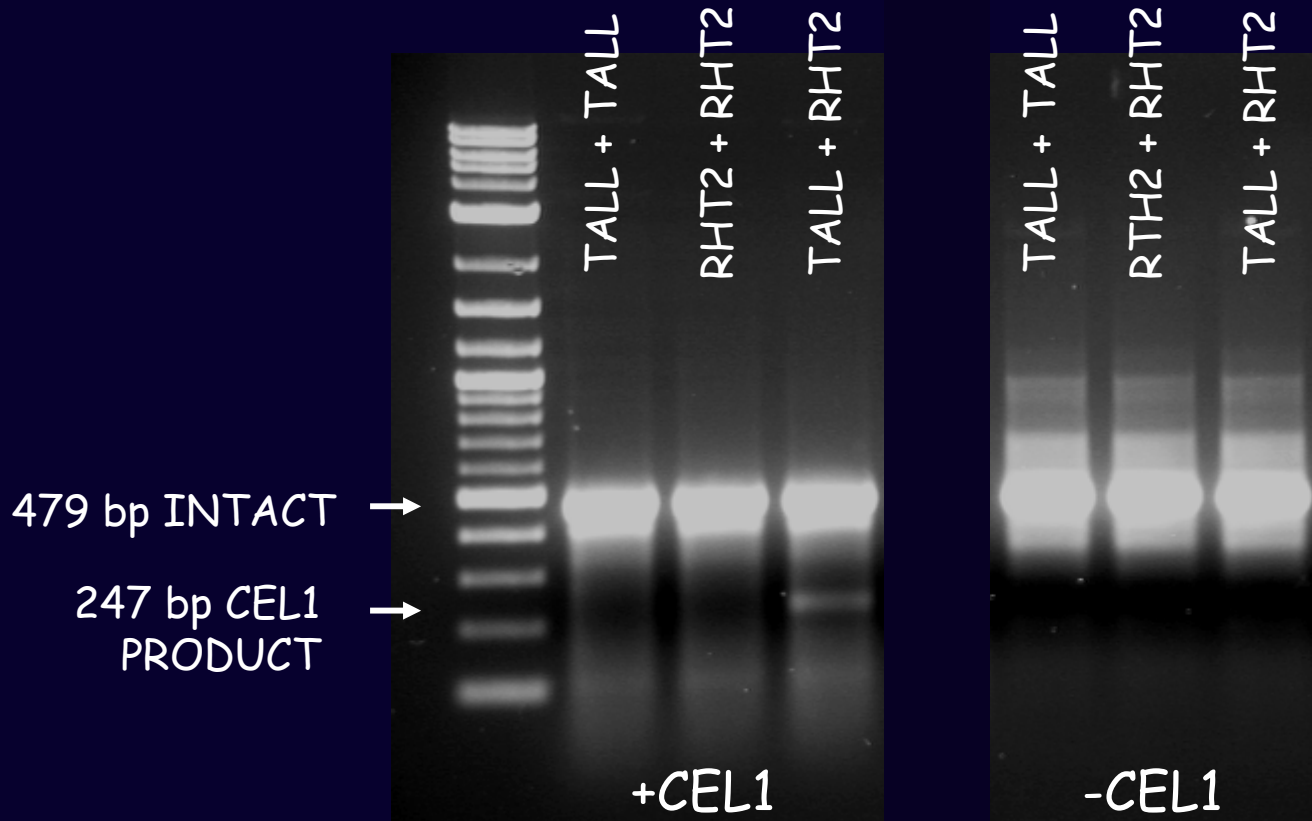
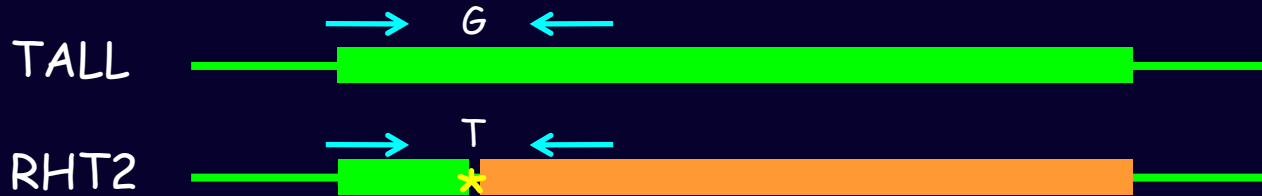
# PCR TILLING IN RHT

1. Designed and tested homeologue-specific primers for Rht-B1 (Rht1, Rht3) and Rht-D1 (Rht2).

Eg. Rht-D1-specific primers



# TESTING PCR TILLING WITH RHT



# TILLING POPULATIONS FOR WGIN

1. ~10,000 Cadenza seeds treated with 0.3%, 0.6% & 0.9% ethylmethane sulphonate for 16h, field-sown 2004.
2. Ears from ~3000 M1 individuals harvested Sept 2004. Now replanting for M2 production.
3. ~2x 4000 Cadenza seeds treated with 0.6% and 0.9% for 16h, sown under glass as for SSD.
4. Harvested leaf material for DNA from 7,000 Paragon M<sub>2</sub>, produced by Rob Koebner, JIC.



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# Cadenza M1 #2



# ESTABLISHMENT OF A TILLING FACILITY

## SUMMARY OF PROGRESS

- ❖ Visited TILLING labs at SCRI Dundee, U. Washington and JIC Norwich for advice & training
- ❖ Set up EMS-treated populations of hexaploid wheat (Cadenza) and collected from JIC M2 lines (Paragon)
- ❖ Imported high-throughput DNA extraction method from RAGT (Peter Jack)
- ❖ Established homoeologue-specific PCR for primary targets
- ❖ Purified Cel1 from celery
- ❖ Shown Cel1 cleavage of heteroduplex PCR products
- ❖ Won BBSRC grant for TILLING equipment:
  - ❖ Liquid handling robot
  - ❖ Licor sequence analysis system



# RRes WGIN Programme Team

Kim Hammond-Kosack (PPI) - Cereal pathogenesis & diploid TILLING

Andy Phillips (CPI) - Gibberellin signalling and hexaploid TILLING

Hai-Chun Jing (PPI) - Pathogenesis and diploid TILLING

Katie Tearall (CPI) - Hexaploid Tilling

Sam Irving - WGIN administration