### Introduction

It defines the objectives and the importance of the research. It focus on the the application of Next Generation Sequencing to molecular biology, wheat genetics and ultimately to breeding programs. It also mentions the current status of the wheat reference genome and other resources (genetic maps, markers) the need of tools to query them effectively.

### Literature review

It describes the current status of the wheat genome, genetics and other resources.

### 2.1 Wheat Breeding

An overview of how breeding is carried on currently, the different sources of genetic diversity and the relevance of fixing agriculturally important traits.

### 2.2 Wheat Genetics

The section describes alleles an the concept of gene, both as a locus in the genome (Quantitative Trait Locus, QTL) and an specific transcript (central dogma of molecular biology). Finally, it discuses traditional Mendelian inheritance and the effect of polyploidy.

### 2.3 Wheat Genomics

A description of the current status of the wheat genome (Mayer et al. (2014), Chapman et al. (2015)), the different available assemblies and and approaches to sort the scaffolds (Genome Zipper, the various genetic maps).

### 2.4 Sequencing

The importance of the selection of the library preparation and the sequencing platforms available. A brief summary of RNA-Seq, Exome capture, Whole Genome Shotgun, etc. and on which cases are more suitable for different experiments. Mention the new technologies developed during the years of the PhD (Ren-Seq, PacBio?)

### 2.5 Sequence analysis

This section discusses the criteria to decide analysis done after sequencing, when to do realignments or *de novo* assemblies, how to do SNP calling in diploid and polyploid organisims and the bulk frequency ratios.

### 2.6 Wheat online resources

A compilation of the currently available resource for whet genetics and genomics. MAS wheat, CeralsDB, Ensembl, etc.

### Genetic mapping of Yr15

This section describes in detail than the paper of Ramirez-Gonzalez et al. (2014)

### 3.1 (Introduction) Yr15

Breeding importance of Yr15 and original source (an introgression of T. diccocoides).

### 3.2 Segregating population and resistance essays

A description of the starting material and how the population was generated.

### 3.3 Sequencing and mapping

RNA-Seq and the decision to call SNPs on gene models rather than the whole reference. Details of the mapping against the Wheat UniGenes Pontius et al. (2002) and the UCW. Krasileva et al. (2013) gene models.

### 3.4 SNP Calling

. Ruby implementation of the methodology described by Trick et al. (2012).

### 3.5 Bulk Frequency Ratios

Results of the simple SNP calls from the progenitors and how the score of the Bulk Frequency Ratios(BFR) improve the location of the SNPs.

### 3.6 In silico mapping

Mapping of the gene models to the IWGSC CSS Mayer et al. (2014) reference and the location of the SNPs using the genetic map from Wang et al. (2014).

### 3.7 Assay selection

. The selection criteria to decide which SNPs where selected to produce the genetic map: BFR>6, in the short arm of chromosome group 1 and from the Yr15 progenitor.

### 3.8 Genetic map

The three versions of the genetic map: With a subset of the F<sub>2</sub> population

### 3.9 Assembly of the transcriptome

A comparison between the known unigenes and the transcript from the progenitors. Since Yr15 comes from an introgression with  $T.\ diccocoides$ , some novel transcripts can be extracted. Analysis of the gels from Mitaly?

### 3.10 Conclusions

Remarks on how this techinque can be used to do fine-mapping and that if I were to start the project now I would use exome capture or Ren-Seq.

## PolyMarker: A fast polyploid primer design pipeline

One of the main challenges of working with polyploid species is the design of genome specific molecular markers. This is particularly true when targeting conserved homoeologue regions, where a primer could bind to a pair, or triplet, of identical sequences. For that reason, designing primers for polyploids require to include bases that are specific to the target, in addition to the physicochemical properties of the primer. The traditional methodology to find primer candidates include a blast search and a local alignment, select the primer candidates manually, and finally, validate the primers with a tool, like Primer3 (Rozen and Skaletsky, 2000). To reduce the time invested in designed primers I have developed PolyMarker (Ramirez-Gonzalez et al., 2015), a pipeline to automate the primer design for polyploid organisms.

### 4.1 Pipeline

PolyMarker is an automated pipeline that takes as input a list of SNPs and a reference file and produces a list of primer triplets for SNP genotyping. The list of SNPs is first converted to a FASTA file with ambiguity codes(Cornish-Bowden, 1985) The sequences are searched on the genomic reference using exonerate(Slater and Birney, 2005) to find the homoeologue regions to the target sequence. Then, the alignment between homoeologues is refined using MAFFT(Katoh and Standley, 2013). A list of candidate variations is produced and used as input for Primer3(Rozen and Skaletsky, 2000). Finally, the output of Primer3 is parsed to find the best primer pair that contains a the targeted SNP and a base that is specific to the target genome (Figure 4.1). The pipeline is written as a Ruby script, using parsers and wrappers from BioRuby (Goto et al., 2010) and bio-samtools (Etherington et al., 2015; Ramirez-Gonzalez et al., 2012). The software is open source and released as a biogem (Bonnal et al., 2012), bio-polyploid-tools, the source code is available in github: https://github.com/TGAC/bioruby-polyploid-tools.

The PolyMarker input consist on SNP list with: unique name for the marker, the target chromosome and the sequence for the marker. The alternative alleles are surrounded by square brackets within the sequence. PolyMarker can take a list of several markers and design them in batch (Figure 4.2a). A FASTA file is produced with all the template sequences, with the alternative alleles substituted by the IUAPC ambiguity codes (Cornish-Bowden, 1985). The flanking sequence surrounding the SNP is limited by default to 100bp to reduce the search time and avoid missing regions that diverge near the SNP, as when the variation is near an intron-exon junction.

The template sequences are searched in the reference sequence using exonerate(Slater and Birney, 2005), figure 4.2b. The alignment is run with the --model est2genome option, to

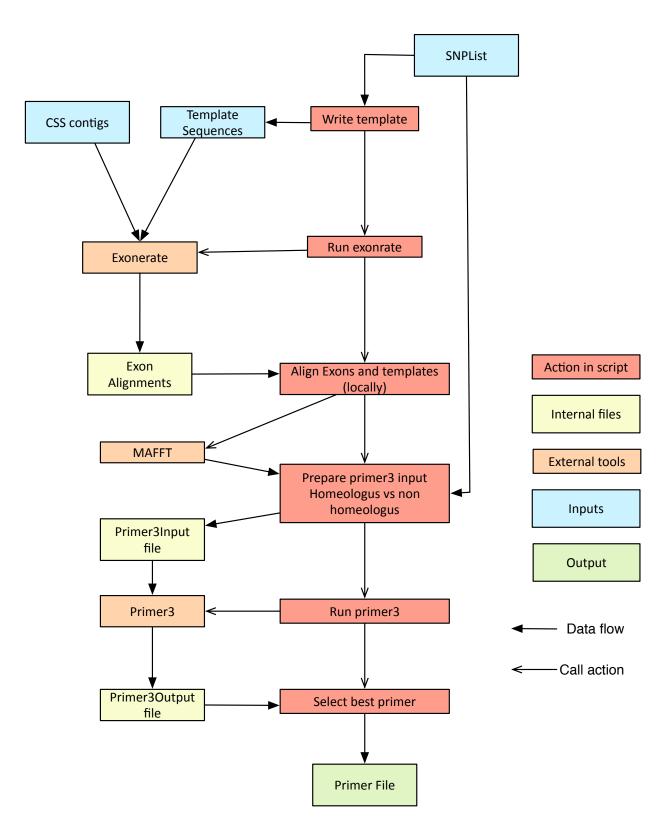


Figure 4.1: Steps and tools called by PolyMarker. The colour of the boxes represent: the step is an action inside the script(red); actions of the script(orange); temporary files(yellow); inputs(blue) and; outpus(green)

allow the search of sequences coming from transcripts, a common source of SNPs(Allen et al., 2011). The exonerate output is formatted with the --ryo (roll your own format) to get an output easy to parse. All the hits that contain the SNP are extracted from the reference with a flanking sequence that extend out of the hit, by defualt, to 100bp on each side of the SNP Figure 4.2c. The size of the flanking sequence can be set to different sizes to allow the design of different types of primers. Different homoeologues may contain small indels (Figure 4.2d). To enable a comparasion base-per-base, a local alignment with MAFFT (Katoh and Standley, 2013) is produced (Figure 4.2e. For wheat, PolyMarker uses the contigs from Mayer et al. (2014), as deposited in Ensembl. As new releases of the wheat genome are made available, different parsers to assign the chromosome to each sequence can be added with little effort to PolyMarker.

### 4.2 Primer design tools

In this section, the principles of *in silico* primer design are discussed, and why not simply selecting a genomic variation is enough (thermal stability, primers folding on themselves)

### 4.3 Primer selection algorithms

Different algorithms to select the best primer:

### 4.3.1 KASP markers

For KASP markers, the product should be as short as possible with the mutation in the first three bases.

### 4.4 Designed markers

Details of the generated primers for the 80k iSelect chip and the 820k axiom chip. This section also include counts on how many are genome specific, semi-specific and non specific. Also an analysis of how many are repeated or map to more than one chromosome perfectly.

### 4.4.1 Regular markers

PolyMarker was designed for KASP assays, but it was later extended to produce regular primers, where both primers start with a genome-specific base. This simplifies the design of primers for regular PCR and capillary sequencing.

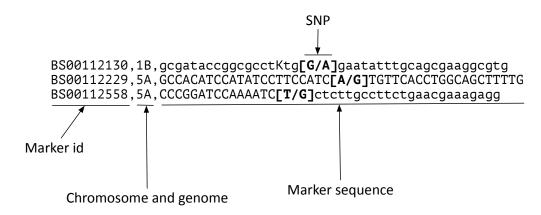
### 4.4.2 Deletion algorithms

Algorithm to produce KASP for deletions in polyploids.

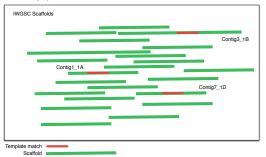
### 4.5 Conclusions

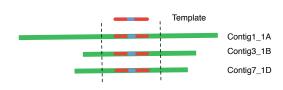
Remarks on the importance of getting the primers right, and the time saved by automating the primer selection. Also mention other primer design tools that have been inspired by polymarker: Ma et al. (2015), Wang et al. (2016)

PolyMarker has been used successfully to design genome-specific primers in several projects.

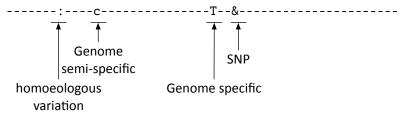


(a) PolyMarker input. The alternative alleles are sorrounded by brackets.





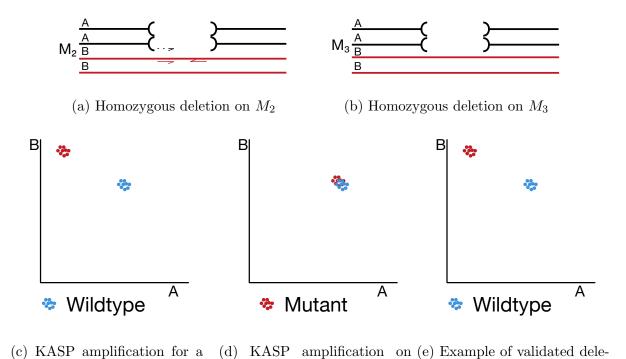
- (b) Global search of templates in the reference contigs.
- SNP-1 A cgcatttGcgcgYgcgataccggcgctKtgGgaatatttgcagcgaaggcgtg
  SNP-1 B cgcatttAcgcgYgcgataccggcgctKtgAgaatatttgcagcgaaggcgtg
  TWGSC-1A cgcatttgcgcgcggataccggcgctgtgggaatatttgcagcgaaggcgtg
  TWGSC-1B cgcatttacgcgcgcgataccggcgctttgggaatatttgcaaggaggtg
  TWGSC-1D catttgcgcgTgcgataccggcgctttggggaatatttgcaaggaggtg
- (c) Selected regions around the SNP on every chromosome.
- SNP-1 A cgcatttGcgcgYgcgataccggcgctKtgGgaatatttgcagcgaaggcgtg
  SNP-1 B cgcatttAcgcgYgcgataccggcgctKtgAgaatatttgcagcgaaggcgtg
  TWGSC-1A cgcatttGcgcgcgcgataccggcgcttGtgGgaatatttgcagcgaaggcgtg
  TWGSC-1B cccatttAcgcgcgcgataccggcgctTtpGgaatatttgc---paaggcgtg
  TWGSC-1D c--atttGcgcgTgcgataccggcgctCtgGgaatatttgcagcgaaggcgtg
- (d) Sequence of found regions around the SNP.
- (e) Local alignment on regions around the SNP detects indels.
- SNP-1 A cgcattt**G**cgcgYgcgataccggcgcctKtg**G**gaatatttgcagcgaaggcgtg
  SNP-1 B cgcattt**A**cgcgYgcgataccggcgcctKtg**A**gaatatttgcagcgaaggcgtg
  IWGSC-1A cgcatttGcgcgcgcgataccggcgcctGtgGgaatatttgcagcgaaggcgtg
  IWGSC-1B cgcatttAcgcgcg<mark>cgataccggcgcctT</mark>tgGgaatatttgc---gaaggcgtg
  IWGSC-1D c--atttGcgcgTgcgataccggcgcctGtgGgaatatttgcagcgaaggcgtg



(f) Alignment with mask and primer candidates.

Figure 4.2: Alignments done by PolyMarker.

4.5. CONCLUSIONS



(c) KASP amplification for a (d) KASP amplification on (e) Example of validated delereal deletion false positive tion on mutant poulation

Figure 4.3: PolyMarker used to find primers to detect long deletions in tetraploid wheat.

### Gene expression (expVIP)

### 5.1 Expression experiments (Introduction)

Describe the list of previously published expression experiments and how they can potentially be used as a framework for new experiments.

### 5.2 Database design

Description of how the database was designed and the flexibility given by having the factors and units as variables

### 5.3 Analysis pipeline

Implementation of the pipeline, from running kallisto to load the data in the database

### 5.4 Graphical interface

How the expression can be displayed filtered, and sorted

### 5.5 Conclusions

The use of previously published studies is a valuable resource. Also, mention that despite the fact that there are several expression/gene browsers, none of them allow comparisons between species and don't consider polyploids.

### Conclusions and final remarks

This section wraps up by showing the relationship and importance of a comprehensive approach to data analysis, from the field, genetics, molecular biology and genomics. I will also remark how the technology and the resources have changed in the last 4 years. As at the references used at beginning where superseded during the PhD.

## Appendix A<br/> PolyMarker validation

## A.1 Validation of mutations on $M_4$ on Kronos

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acAaggactgcttcagaGaC	gcttccactGggtcctgT	gcttccactGggtcctgC	Het	Het	1 H	2 0	1515	Kronos3085	IWGSC_CSS_1BL_scaff_3897513
gtgattttgccaggggagA	ccacaccttgagcctcgT	ccacaccttgagcctcgC	Het	Het	A	Ω	7449	Kronos3085	IWGSC_CSS_1AS_scaff_3284790
Common Primer	Primer 2 (mutant)	Primer 1 (Kronos)	Called on $M_4$	Predicted	Mut	WT	Pos	Line	IWGSC contig

# A.2 Validation of mutations on $M_4$ on Cadenza

ω		6019 C 2418 C 11277 G 15710 G			het	het	caggat Agt GegactetcaaaG	caggat AgtGggactgtcaaaA	ggagacGGctGtggacatT
L_scaff_6955403 L_scaff_710846 S_scaff_495135 S_scaff_495646 S_scaff_495646 S_scaff_41715962 L_scaff_7108426 S_scaff_1175962 S_scaff_7270737 S_scaff_7270737 S_scaff_7270737 L_scaff_7270737 L_scaff_727099 L_scaff_727094 L_scaff_72709409							COMMENT OF THE COLUMN TO SHARE THE COLUMN TO S		0
ω				E	het*	hom	tcagCggattgtcgggatG	tcagCggattgtcgggatA	tgtcCatgaaTcttgtccacG
ω					hom	hom	tgggatccatgcctacactG	tgggatccatgcctacactA	${\tt gatggtGgatttgccgctA}$
ω					hom	hom	ctggccctgcgctaC	ctggccctgcgctgctaT	gtggaaGttcagaaggaccaG
φ					het*	hom .	gcaggttgacttcccggaG	gcaggttgacttcccggaA	tGaggtacgaGcTaaagAaagC
ω 00					hom	hom	cagctgtggTatctcaactgG	cagctgtggTatctcaactgA	CcCtGaaACACcGtttggaI
φ		2113			nom bot	nom bet	gcgacGaacctcgagatctG	gcgacGaacctcgagatctA A+aggagattgatagagtgaA	ga I ggcaAtcgt/gtgcA
∞					het*	hom	cctgacatcattgttcacgatC	cctgacatcattgttcacgatT	cactccgagetetccatgaT
· · · · · · · · · · · · · · · · · · ·					hom		attcCTgtgttgtggCaaatgaG	attcCTgtgttgtggCaaatgaA	taaGcacaaAccetccagetgG
<sub>∞</sub>					hom	hom	ccacagtgagactcctattgaCG	ccacagtgagactcctattgaCA	atgtctgattcGtcGtagtcC
∞					het	het	catccgccGtttcctcC	catcccgccGtttcctcT	gctcgccgatgaagagcT
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<sub>∞</sub>					hom	hom	aCagatacaAttgtcatgcaggC	aCagatacaAttgtcatgcaggT	acctgggTTgtccaatacttC
	_				het	4	cgtggcCgaatCtcGacG	cgtggcCgaatCtcGacA	ttcttgtgggagccgggC
					nom L-t	hom L-t	gtcacgaaCccgctcagG	gtcacgaaCccgctcagA	aggaaagagaaaagaGcG
		5004			net hot	net bet	actotogicaagaactgatacaG	acteregreagaactgatacaA	gcaGagaatgttcttgcaacı
					hom	hom	SS'SSSS archess as Street Control	SS SSSS COSSSS CONTRACTOR A	SCSCSSCCSCSCSCC
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		2125 (			hom	het	gcaettttatcctcaetaetctteG	geagttttatecteagtagtettgA	ttctgagaaTgtaatgtgcGatG
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	538				hom	het	agatgettgtCggGceaG	agatgettgtCggGccaA	gctgaagcaacgcgatcaaT
×	538				het	het	ggcagtaatgtggtgctgagC	ggcagtaatgtggtgctgagT	tTgaCttctggtttggtggcA
oo	538				het	het	gctaaagaagcttgagagaattC	gctaaagaagagcttgagagaattT	aatttetgaagagagtgttgtatG
oo oo	538				het	1	gatateteccacaggegG	gatatctcccacacgccgA	tgagccactcttgcagtttT
oo oo	538				hom	het	agcaattetttggetatcaattagC	agcaattctttggctatcaattagT	tcatctGtcttaactctactgctG
oo	538				het	het	gctCagggaggaagacaagaaG	gctCagggaggaagacaagaaA	tgctatgaagaattccgacctC
oo	538				hom		tcagcaaaa $t$ cacc $t$ gc $C$ g $C$	tcagcaaaa $t$ cacc $t$ gc $C$ g $T$	gCtgcccatcatcgtttaT
∞					het	het	tCgttgcaagcCttTtgtgC	tCgttgcaagcCttTtgtgT	agaGttaTcaagcTactgtcacA
∞	_			A	hom	hom	ctcttcAgagatgaacgcgG	$\operatorname{ctcttcAgagatgaacgcgA}$	tcGtGagatgGtggtttGTtA
∞					het*	hom	ccgaccaAttcactaaccgG	ccgaccaAttcactaaccgA	accetetttcccAgacatgaT
∞				Α!	hom	hom	aacatttgcctTaCcaaaacGC	aacatttgcctTaCcaaaacGT	acacagcaagttataatgCAAgC
∞					het	het	caacatgagacacaacaccttC	caacatgagacacaacaccttT	gtcaacgcgtgaggattgtC
∞					hom	hom	tggTlGtagacacttggcgaG	tgg'l'GtagacacttggcgaA	catggcgaccaccAcctG
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	Cadenza0580				hom	hom	CgttgaaaaGctgcaagaacttaaC	CgttgaaaaGctgcaagaacttaaT	cagttcttccTtCaGagcagataT
	Cadenza0580 1				hom		tggattttcccgcactgttC	tggattttcccgcactgttT	gtaaacaaggcatttcaagagtcA
			7	A	hom		gCtcAttcagggatTGTcCtaTatG	gCtcAttcagggatTGTcCtaTatA	${\tt tgaCagaacagttggtcatacT}$
					hom	hom	gccgtggttgatggAgaG	gccgtggttgatggAgaA	cgtccagattactgatacttgcA
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					het.	IIIOIII	cotosootoootostttoC	Scarcing Sings nation	terantaettanantenetTC
_					hom	hom	caggaGcctggcaaataaaGG	caggaGcctggcaaataaaGA	ctttcGcagtctcttagtttcG
				A	hom	hom	gcatgctaacaggcgaaaagG	gcatgctaacaggcgaaaagA	ctcatgctcctgatcttaaggtT
					hom	hom	tacgtgcatgatgtggtagtcgtaC	${ m tacgtgcatgatgtggtagtcgtaT}$	${\tt gtttgaagtgcatcagatgTaccA}$
IWGSC_CSS_6DS_scaff_1880206 Cade				Α.	het	het	ctgCgaaggctccacaaG	ctgCgaaggctccacaaA	ggatgagaagtttgcattgctC
					het	.	ccatgtgtttccaatgttagagC	ccatgtgtttccaatgttagag.I	tgccctagctggtatgcT
					hom	hom	agtaagCGtgacagcaatggG	agtaagCGtgacagcaatggA	AtgtctTtgGtggaagtacatcA
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			י רז	_	het	het	tggAagcAcaaggggccC	tggAagcAcaaggggccT	GccgccgatggagactcG
			7	_	het	het	gtcccttgcacacagctttG	gtcccttgcacacagctttA	cctgctggactacaacttcaaT
					het	het	caagaatgATgctgatgttggaG	caagaatgATgctgatgttggaA	acatgctgaatcgccgaatC
IWGSC_CSS_3AS_scaff_3371333 Cade:		538	ر ان د	 ∢ E	het ket	het bet	gggaaaCgAgAcgagcgG	gggaaaCgAgAcgagcgA	cegtgeetteeteaceeT
	Cadenza0364				net hot	net bet	arccccacggcacagag G	atccccacggcacagagA	aAttggccttggtgattc
		2242 (			het	-	efttcTeTctctttcactG	esttcTeTectcttccactA	tetetteaaccecaaecA
								)	

IWGSC contig	Line	Pos	$_{ m TW}$	Mut	Predicted	Called on ${\cal M}_4$	Primer 1 (Cadenza)	Primer 2 (mutant)	Common Primer
	Cadenza0364	3198	С	H	het	het	${\it gagtcaTtaagttggtaagattggC}$	gagtcaTtaagttggtaagattggT	GCaGaTaaCaacaggatcacG
WGSC_CSS_3AL_scaff_4447942	Cadenza0364	$\frac{11917}{2547}$	ე ი	∃≯	het	het	gtcataaagattgctcctgtgaaG	gtcataaagattgctcctgtgaaA	ctcGgatgtgggaggaagA
WGSC_CSS_3AS_scaff_2648747	Cadenza0364	2688	Q.	Α	het	het	tggAagcAcaaggggccC	m tggAagcAcaaggggccT	GccgccgatggagactcG
IWGSC_CSS_3AS_scaff_3304956	Cadenza0364	1017	ດດ	] ⊳	het	het	gtcccttgcacacagctttG	gtcccttgcacacagctttA	cctgctggactacaacttcaaT
$WGSC\_CSS\_3AS\_scaff\_3321091$ $WGSC\_CSS\_3AS\_scaff\_3371333$	Cadenza0364 Cadenza0364	53 8 53 8 53 8	IJ C	₽⊣	het het	het het	caagaatgATgctgatgttggaG gggaaaCgAgAcgagcgG	caagaatg $\operatorname{ATgctgatgttggaA}$	acatgctgaatcgccgaatC ccgtgccttcctcacccT
IWGSC_CSS_3AS_scaff_3371815	Cadenza0364	1061	Q	H;	het	het	atccccacggcacagagG	atccccacggcacagagA	aAttggcccttggtgattcC
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IWGSC_CSS_3B_scaff_10343586	Cadenza0364	2242 2433	J G	∃≯	het	r	ggttcTgTcctctctccactG cateeCgaceetGtcctG	ggttcTgTcctctcttccactA	tgtgttgaacccgcaagcA aAccctcatTTts~CTACTtCT
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WGSC_CSS_6AS_scaff_4378990	Cadenza0281	6748	īΩ	Η	hom	hom	cccaggttctgcttcttttcC	cccaggttctgcttcttttcT	caagtatcaagaaaatgaagggTgT
IWGSC_CSS_6BL_scaff_4488310	Cadenza0281	3808	ט מ	⊳ ⊢	hom	hom	aCtactcaaatggcttGgtgtaG	aCtactcaaatggcttGgtgtaA ettctcttetaetaecaeccA	tcagtccaacatgTcaagagatT
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WGSC_CSS_3B_scaff_10626860	Cadenza0148	7847	<u>۾</u> (	<b>≯</b> +	het	het	gcagctctgggaaggagG	gcagctctgggaaggagA	gttaatgtacCTcctagcctcG
WGSC_CSS_3DL_scaff_6915683	Cadenza0148	6904	Ω	H	het	het	cgtcaaCctgtgggcaattG	cgtcaaCctgtgggcaattA	tcatgctcataatgTcatagggT
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WGSC_CSS_4DL_scaff_14455742	Cadenza0148	1946	Q	H	$_{ m hom}$	hom	gCctgagggagatcgcgC	gCctgagggagatcgcgT	aaccgGtAaCTGtGgGcA
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WGSC_CSS_5BS_scaff_1646558	Cadenza0148	2916	Ω	H	het	het	gccGtacactcacctAtcctttG	gccGtacactcacctAtcctttA	gcaaTgtccacttAtcatcccT
WGSC_CSS_1AL_scaff_3883106	Cadenza0110	27536	ם מ	Þ	het het*	het	accttccatcactggctgG	accttccatcactggctgA	gtgaagaacaacaggttgaagC
WGSC_CSS_1DL_scaff_2266648	Cadenza0110	6156	Q.	A	het	het	actgcgtggttatgggacC	actgcgtggttatgggacT	ccccatcactgaacacaacA
WGSC_CSS_1DS_scaff_1889435	Cadenza0110	4636 4636	D C	⊳⊢	hom	hom het	aaccatgaattactcggacagG	gatecatgaattacteggaeagA	gccctgaagaattgtatcaaaacaG ToctotTooatatocaottacT
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IWGSC_CSS_3AL_scaff_4384278	Cadenza0110	1276	Q	H	het	het	agcTgaactgccccTgtaG	agcTgaactgcccCTgtaA	agggacctCgGtggatgaA
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WGSC_CSS_6AS_scaff_4429974	Cadenza2103	3867	Ω	A	hom	hom	GagatgaAtttattgagcatgtggC	GagatgaAtttattgagcatgtggT	ggttccggctgcataagT
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	Cadenza0277	10124	Q (	Η;	het	het	tgacaggggacgctatacaG	tgacaggggacgctatacaA	gtctaaCTtACattAcccatcagC
	Cadenza0277	663	Ω	A	hom	hom	actgcactcatacaatActtCtgC	actgcactcatacaatActtCtgT	tcCacctggacagcaagtG
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WGSC_CSS_5AL_scaff_2736916	Cadenza0277	4296	Ω Ω	⊳	het	het	aagaactATgAaaGtaacacacgaC	aagaactATgAaaGtaacacacgaT	ttcGcTttTaagGcAttCtcG
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WGSC_CSS_2AS_scaff_2306489 WGSC_CSS_2BL_scaff_7984123 WGSC_CSS_2DL_scaff_9907477 WGSC_CSS_2DL_scaff_930886 fWGSC_CSS_2DS_scaff_5330886 fWGSC_CSS_3AL_scaff_4449951	Cadenza0097	7819	Q	H	het	het	agaggggtgctatccatAttgG	agagggtgctatccatAttgA	orrestorressoretteC
WGSC_CSS_2AS_scaff_306489 WGSC_CSS_2BL_scaff_7984123 WGSC_CSS_2DL_scaff_9907477 WGSC_CSS_2DL_scaff_330886 WGSC_CSS_2DS_scaff_5330886 WGSC_CSS_3B_scaff_10479889 WGSC_CSS_3B_scaff_10479889	Cadenza0097	6360	G	A	hom	hom	2		ag c gar g c caa g g c r c c c
WGSC_CSS_2BL_scaff_7984123 WGSC_CSS_2BL_scaff_7984123 WGSC_CSS_2BL_scaff_5907477 WGSC_CSS_2DL_scaff_5907477 WGSC_CSS_2DS_scaff_5330886 WGSC_CSS_3B_scaff_10479889 WGSC_CSS_3B_scaff_10479889 WGSC_CSS_3B_scaff_10562262 WGSC_CSS_3AL_scaff_7040796 WGSC_CSS_3AL_scaff_7064786			2	-	*****		q	acacaacattgccaccagaA	agugauguuaagguuu CAatCgattgcttgctTctcC aggcagtggatgtatgtgaagttT
WGSC_CSS_2AL_scaff_7984123 WGSC_CSS_2BL_scaff_7984123 WGSC_CSS_2BL_scaff_5907477 WGSC_CSS_2DL_scaff_530886 WGSC_CSS_2AL_scaff_530886 WGSC_CSS_3A_Lscaff_10479889 WGSC_CSS_3B_scaff_1052262 WGSC_CSS_3B_scaff_706796 WGSC_CSS_4AL_scaff_7063488 WGSC_CSS_4AL_scaff_7091701	Cadenza0097	5050	Ω Ω	≽⊣	het	het	acacaacattgccaccagaG gcctctcacCttAatttgaagctgC catgagcatctgggaggaaaatG	acacaacattgccaccagaA gcctctcacCttAatttgaagctgT catgagcatctgggaggaaaatA	CAatCgattgcttgctTctcC caggcagtggagtatgtgaagttT aggcaagtggaAtaatgaacggaaA
IWGSC_CSS_4AL_scaff_791761 IWGSC_CSS_4AL_scaff_984123 IWGSC_CSS_2BL_scaff_9984123 IWGSC_CSS_2DL_scaff_9907477 IWGSC_CSS_2DL_scaff_10479889 IWGSC_CSS_3AL_scaff_10479889 IWGSC_CSS_3B_scaff_10479889 IWGSC_CSS_3B_scaff_10562262 IWGSC_CSS_4AL_scaff_7040796 IWGSC_CSS_4AL_scaff_7040796 IWGSC_CSS_4AL_scaff_7040796	Cadenza0097 Cadenza0097	5050 7110	2 6 6 0	> > > H	het hom	het hom	at G	acacaacattgccaccagaA gcctctcacCttAatttgaagctgT catgagcatctgggaggaaaatA aatgTAgctccccatacCgA	CAatCgattgcttgctTctcC aggcagtggagtatgtgaagttT agcaagggaAtaatgaacggaaA actgaaacTgcaatcgtTtatggA

IWGSC contig	Line	Pos	WT	Mut	Predicted	Called on $M_4$	Primer 1 (Cadenza)	Primer 2 (mutant)	Common Primer
IWGSC CSS 1AL scaff 3952258	Cadenza2092	8107	۲	E	het.		toaotagaaattoacagtotoG	teaetaeaaatteacaetete A	teccaccatteacateacaC
IWGSC_CSS_1BL_scaff_3858008	Cadenza2092	10278	Ü	4	hom	hom	ttteaecaeecaeeatceC	tttgagcaggcaggatcgT	actcacegectatatcActattC
IWGSC_CSS_1DL_scaff_2265172	Cadenza2092	9094	Ö	Н	hom	hom	tgcaTGTcatttgttcttatcagC	tgcaTGTcatttgttcttatcagT	agtgtccaacttccGttcatC
IWGSC_CSS_2AL_scaff_6435867	Cadenza2092	16201	Ü	Ą	hom	hom	tttctgTaccttaacgtcaattgaC	tttctgTaccttaacgtcaattgaT	gtgaggatgatgaggtaagacC
IWGSC_CSS_2AL_scaff_6439430	Cadenza2092	25101	Ö	Η	het		caagaaagggCagCtCagC	caagaaagggCagCtCagT	tcGttAcTctttcActggtgaA
IWGSC_CSS_2DL_scaff_9760848	Cadenza2092	4733	Ö	Η	het	het	gcaccatgggtctcaggtaC	gcaccatgggtctcaggtaT	tcagtcagtttGCTCtgTCTG
IWGSC_CSS_3AL_scaff_4407012	Cadenza2092	2785	Ö	H	hom	hom	acatatAgtgttctcatccaccatC	acatatAgtgttctcatccaccatT	acctctctcatgttaataggtttgT
IWGSC_CSS_3AS_scaff_3441108	Cadenza2092	541	Ü	Ą	het	het	GtgatgaccttgagacGgaG	$\operatorname{GtgatgaccttgagacGgaA}$	$\operatorname{aggcaTgacaaCgcgcaA}$
IWGSC_CSS_3B_scaff_10449827	Cadenza1551	4779	Ü	A	hom	hom	ggcaaggtcaagaaacGgtC	ggcaaggtcaagaaacGgtT	aCagaGtgggttagaggcaG
IWGSC_CSS_3B_scaff_10550638	Cadenza1551	3250	೮	H	het	het	ctccttcacttgttgcggC	ctccttcacttgttgcggT	gcaacAtTttgatactgcaaagG
IWGSC_CSS_3DL_scaff_6945816	Cadenza1551	589	Ö	Η	hom	hom	agcatctcacctgcaaCaataC	agcatctcacctgcaaCaataT	TgtgcccTctgaAtattttcaTG
IWGSC_CSS_3DL_scaff_6954177	Cadenza1551	3508	Ö	Η	het	het	tgtagcatcacattaactttcctG	tgtagcatcacattaactttcctA	gcttggtataaaccCttacgacA
IWGSC_CSS_4AS_scaff_5938272	Cadenza1551	19080	Ü	A	hom	hom	agAcCccgAtcgccatgG	agAcCccgAtcgccatgA	$\operatorname{GggAgatAcaggtaaaActcTtcG}$
IWGSC_CSS_4AS_scaff_5977594	Cadenza1551	11092	Ö	H	het	het	gccttgattcggaacaacaaaC	gccttgattcggaacaacaaaT	gcgtctctcagtcctgcA
IWGSC_CSS_5AL_scaff_2671035	Cadenza1551	5859	೮	H	het	het	$\operatorname{cggtgatattTttagacttcgacgC}$	$\operatorname{cggtgatattTttagacttcgacgT}$	ggcagttcagcGacccatT
IWGSC_CSS_5BL_scaff_10889480	Cadenza1551	2530	Ü	A	hom	hom	gagcttaactcgcagatggaG	gagcttaactcgcagatggaA	tccatgCAacGccttggT
IWGSC_CSS_3B_scaff_10528396	Cadenza2088	8059	Ü	Ą	hom		cttttccgtccgtaagcaataG	cttttccgtccgtaagcaataA	gtgcactgttcaggcctgA
IWGSC_CSS_3B_scaff_10637573	Cadenza2088	16815	Ü	A	het	het	agcaagcttaccGgtctgC	agcaagcttaccGgtctgT	cgagcAactacgagcagctT
IWGSC_CSS_4AL_scaff_7086469	Cadenza2088	2699	Ü	A	het	het	gccgtctacttcaacgcG	gccgtctacttcaacgcA	ccaGaggcttgtTGcattttT
IWGSC_CSS_4AL_scaff_7126302	Cadenza2088	3627	Ü	Ą	hom	hom	gttcaaaaacaagtggctAatttgC	gttcaaaaacaagtggctAatttgT	cacaaggatatgaagcTcttctagA
IWGSC_CSS_4BL_scaff_7041808	Cadenza2088	10234	Ü	A	hom	hom	tcaatggatgagggtgcttC	tcaatggatgagggtgcttT	ccatagcagcatcagccacA
IWGSC_CSS_5AL_scaff_2794167	Cadenza2088	13162	Ü	Ą	het		agtattcaggacaagcatCttCaG	agtattcaggacaagcatCttCaA	caatgaaacctctcgaagaaGaG
IWGSC_CSS_5BL_scaff_10889232	Cadenza2088	3885	Ü	Ą	het	het	${ m cTc}$ aaccacaat ${ m gggcaA}$ at ${ m C}$	${ m cTc}$ aaccacaat ${ m gggcaAatT}$	tcc $t$ tca $t$ caatca $t$ caa $t$ t $g$ t $t$ g $G$
IWGSC_CSS_5BS_scaff_2267405	Cadenza2088	11113	Ö	H	hom	hom	ctttgatgatcctaggcctctTG	ctttgatgatcctaggcctctTA	tgatttggtCtggttAgagtttGA
IWGSC_CSS_3B_scaff_10475354	Cadenza1409	2203	Ü	Ą	hom	hom	agCgaacaagagGtcaaacG	$\operatorname{agCgaacaagagGtcaaacA}$	ctgaaacacaCtagaCAattAccG
IWGSC_CSS_3B_scaff_10674115	Cadenza1409	4555	Ö	H	het	het	gcttcagtgcatgccttcaG	gcttcagtgcatgccttcaA	cttcacacccGagataatGtattG
IWGSC_CSS_4AL_scaff_7153568	Cadenza1409	13073	Ö	Η	hom	hom	tccgaccgAtcaaccttgG	tccgaccgAtcaaccttgA	gaccggaactcctcggcC
IWGSC_CSS_4DL_scaff_14314966	Cadenza1409	2010	Ü	A	het	hom	${f gtaggtccctcctCAggG}$	${f gtaggtccctcctCAggA}$	$\operatorname{cggcgTcacaAgttgCcT}$
IWGSC_CSS_4DS_scaff_2324074	Cadenza1409	2092	Ü	A	het	het	tGcatgaaaatgtgtGcaGaG	${ m tGcatgaaaatgtgtGcaGaA}$	${\tt gggtaAgttcAaaactGaagtgaaG}$
IWGSC_CSS_5AS_scaff_1517889	Cadenza1409	3561	Ü	A	het	het	tetegacatetteeegtgtaC	tctcgacatcttcccgtgtaT	${f gtgcctggaacattgcttatttA}$
IWGSC_CSS_5AS_scaff_1523866	Cadenza1409	8054	Ü	A	hom		ggtgatctaccgccaGgaC	${f ggtgatctaccgccaGgaT}$	tcctgcagCcTctcctcA
IWGSC_CSS_5BL_scaff_10917655	Cadenza1409	19073	Ü	Ą	hom	hom	caaatgacatgcaaaagaagttgC	caaatgacatgcaaaagaagttgT	cgcttcatcactacaAaatatgtcT
IWGSC_CSS_1AL_scaff_3886649	Cadenza1599	5204	ರ	Η	het	het	tgatgccaaccacaatGcC	${f tgatgccaaccacaatGcT}$	ggactgactgaccatatttaG
IWGSC_CSS_1BL_scaff_3810267	Cadenza1599	6634	Ö	Η	hom	hom	ccCaggaaatgagcacctC	ccCaggaaatgagcacctT	cgcaggcgaagatgtgaTtG
IWGSC_CSS_1DL_scaff_2291677	Cadenza1599	12856	Ö	H	hom	hom	GgtagacaagtcgccgaG	GgtagacaagtcgccgaA	cetectecateacGCeG
IWGSC_CSS_2AL_scaff_6354492	Cadenza1599	7566	Ü	A	het	het	gGagaatgcaCAgtAacTtctgG	${\tt gGagaatgcaCAgtAacTtctgA}$	${ m ttccgaagaaccacaTccTG}$
IWGSC_CSS_2AS_scaff_5282937	Cadenza1599	9236	Ü	A	het	het	gctgtagattttatagctgctatgC	gctgtagattttatagctgctatgT	cacCagaattgttCactgatttTC
IWGSC_CSS_2BL_scaff_7952427	Cadenza1599	19249	Ü	A	hom	hom	$\operatorname{cgTccctCcctagcacgaC}$	$\operatorname{cgTccctCcctagcacgaT}$	aTcactccattagcgcgAG
IWGSC_CSS_2DL_scaff_9897981	Cadenza1599	5627	Ö	Η	het	het	cttggtgctTgattgcttactC	${ m cttggtgctTgattgcttactT}$	${ m gTttgctCtctctgatctTtgtG}$
IWGSC_CSS_3AL_scaff_4446105	Cadenza1599	1765	Ü	A	hom		aaatgettteetaCegetagtG	aaatgctttcctaCcgctagtA	${ m ttctAgaggcaatagctTatatgcT}$

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