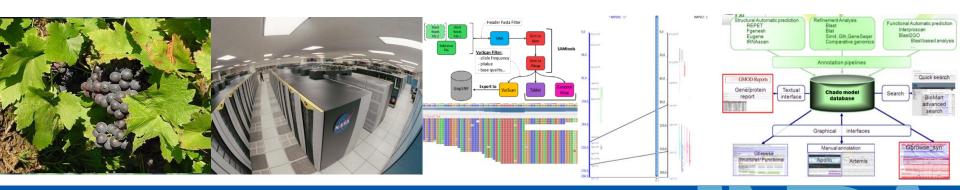




Grape@GnpIS

Overview of the GnpIS Grape Genomics & Genetics Integrative Resource

Nacer Mohellibi



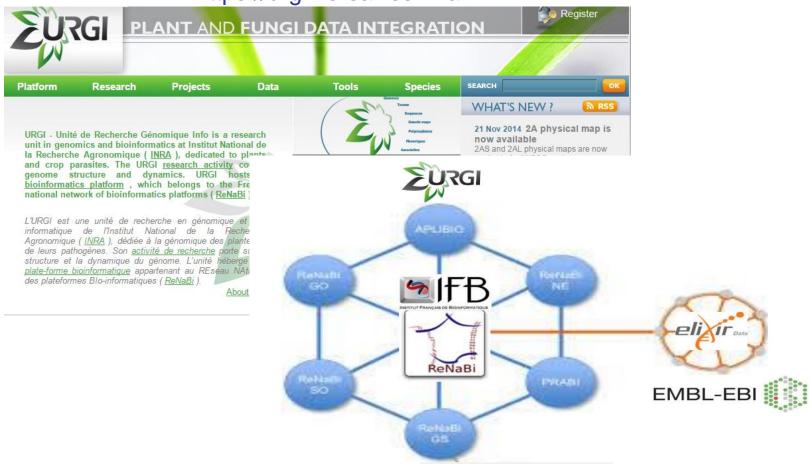
ALIMENTATION
AGRICULTURE
ENVIRONNEMENT





ABOUT URGI

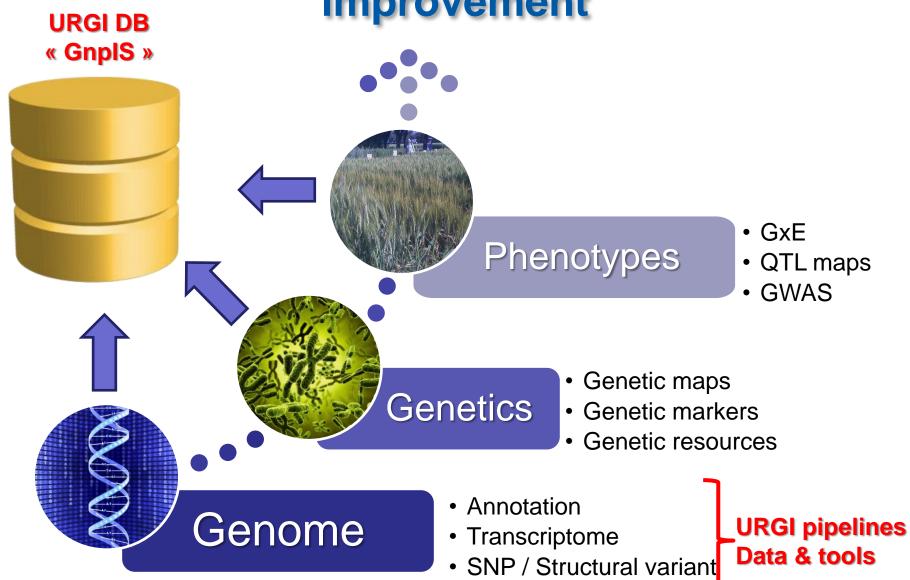
https://urgi.versailles.inra.fr







A bioinformatic plateform for crop improvement



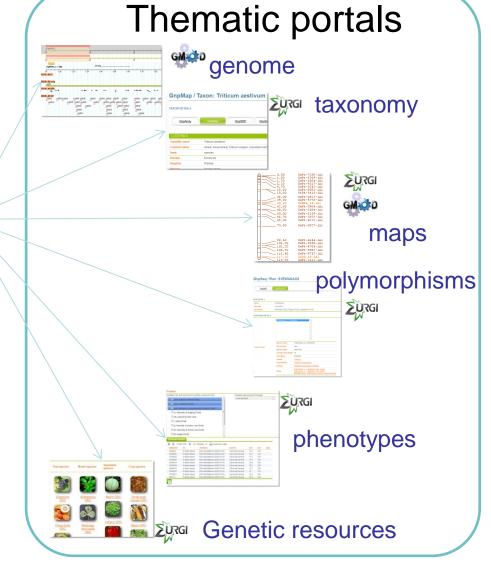
INRA SCIENCE & IMPAC

Nacer Mohellibi



GnpIS Overview

Transversal portal **EU**RGI Global Genome annotation data. GnpGenome. search Taxons **QUICK SEARCH** Sequences NGS projects description. GnpSeq. You can find the indexed data list here. Genetic maps Examples: WI, WIF52, gene, arabidopsis. Genetic maps and OTLs, GnpMap, Polymorphisms Molecular polymorphism, GnpSNP, Phenotypes ADVANCED TOOLS Phenotypic and environmental experiments. Ephesis. GALAXY Genetic resources Plant genetic ressources data, Siregal Expression data. GnpArray. **EU**RGI Grape<u>Mine</u> Genetic maps (markers, Otls), Poly enetic maps OTHER SPECIES Genetic maps OTHER SPECIES . Filters [None selected] Galaxy **Biomart**







Transversal portal - quicksearch

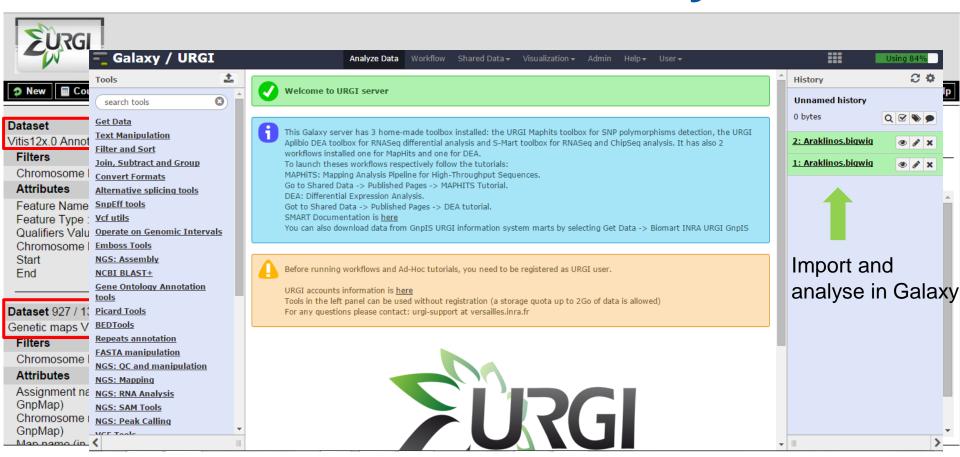


https://urgi.versailles.inra.fr/gnpis/





Transversal portal Biomart & Galaxy



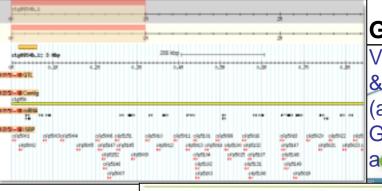


Integration of Genomic and Genetics data for Vitis markers in chr. 18





Thematic portals

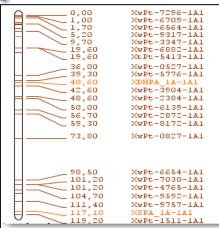


Genome Vitis 8X

& 12X

(annotations Genoscope

and CRIBI



GnpSeg / Run :61EW4AAXX

Sequencing Metadata & Files

Resequencing runs: Genetic Maps & Markers 27 Vitis genetic maps Genetic maps and QTL

2 public / 13 private

Genetic Resources 10,459 Vitis

accessions

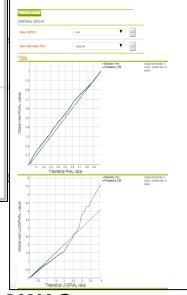
Genotyping form (4) 4 1-10 of 560 (b) (b) (J) Display 10 GrapeReSeq_Illumina_20K_experiment GrapeReSeq Illum GrapeReSeq_Illumina_20K_experiment GrapeReSeq_Illu GrapeReSeq_III GrapeReSeq Illi GrapeReSeq Illumina 20K experiment GrapeReSeq III GrapeReSeq Illumina 20K experiment ALCA? ON ALARLIE ALLAREN SCIENCE & IMPACT

Polymorphisms The Internality of well-one most (Triad) The intensity of boose root (Trial) STREET, CARRIED STREET, CO., C.P. Co. CPW0000 It fulfile behand This letter (Stations 2006 CF-SA OCHROST.

Phenotyping 1trial

Polymorphism (genotypes of >1100 accessions for > 14000

Nacer Mohellibi markers)



GWAS (in progress)



1 Workflow / 2 use cases

Annotation

Variations

Accession

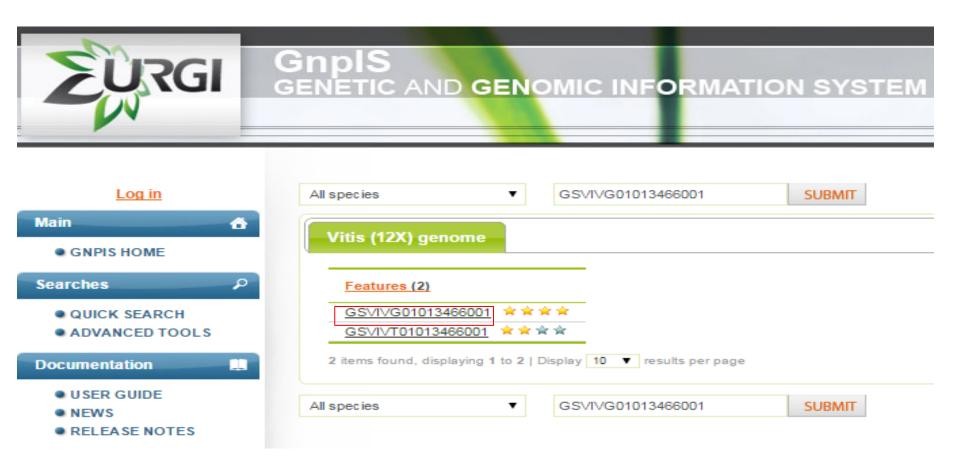
Phenotype







Use case 1: Zoom on a gene in Flb Locus

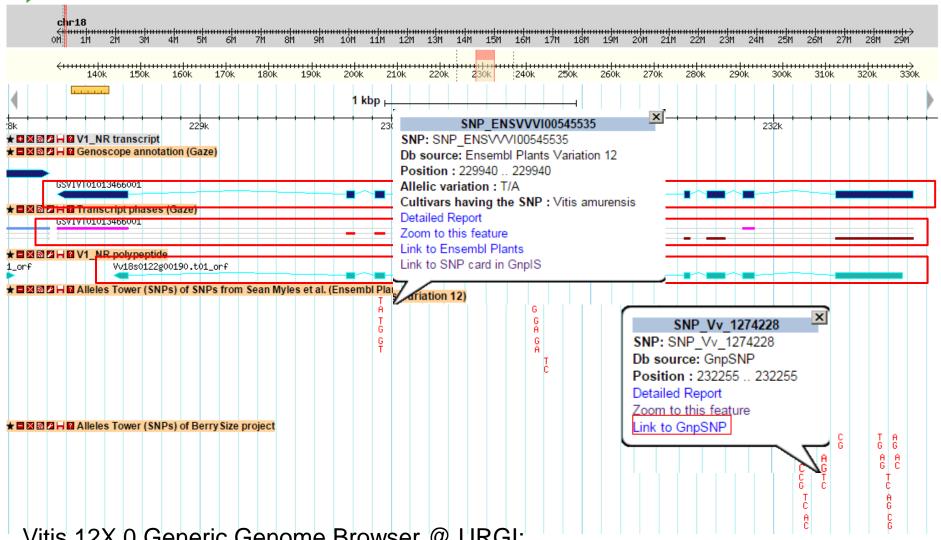


GnpIS QuickSearch: https://urgi.versailles.inra.fr/gnpis/





Vitis 12X Genome Browser



Vitis 12X.0 Generic Genome Browser @ URGI:

https://urgi.versailles.inra.fr/gb2/gbrowse/vitis_12x_pub





SNP details

Polymorphic locus card

DETAILS					
Name :		Vv_1274238			
Ref. Sequences :		Major allele of variants in VVC29	72A-bat ▼	٩	
Position on ref. seq. :		512			
Source :		GnpSNP			
COMPLEMENTS					
Туре:		SNP			
Sequence variation :		C/G 🕖			
Linked with variations :		VVC2972A_512_45 View list)	•	٩	
Linked with lines :		294Mtp1 View list	7	٩	
MAPPING INFORMATIONS	(NGS)				
Reference Genome :		Vitis vinifera 12X			
Taxon (mapping) :		Vitis vinifera L.			
Chromosome :		chr18			
Mapping position :		176646			
EXTERNAL REFERENCES					
Database		Reference name	Ref	erence value	
Vitis vinifera 12x Genome B	Browser	name	SNI	P VV 1274238	
SEQUENCES					
5' flanker on ref.seq. :	TTTTTCTTTTCTTTC	CATCACTCTTCCTCCTCAGAGGCGAGGCTGTCACCAI CTTGTTCCTCCTCTGGATTGAAAAACCTCTTCTCI CTAATCCATTCAATTCAATTCCCTAACTTCATCGCTI	AGTCCATCTGAA	ATTCCACCTCCCTAGACT	A
3' flanker on ref.seq. :	>VV_1274238-3' TTATTAATTTCATAA	ATTAATTTGTGCTATTTTTCTTTTTTCAAAAAAAAA	4		M
Genomic context on ref. seq. :	TTTTTCTTTTCTTTC	nomic_context :ATCACTCTTCCTCCTAGAGGCGAGGCTGTCACCAG :TTTGTTCCTCCTGGATTGAAAAACCTCTTCTC :TAATCCATTCATATTCATAATTAACTTCATCCTA !TTATTAATTTCATAATTAATTTGTGCTATTTTTCT!	AGTCCATCTGAA ACCGAATTCCCT	ATTCCACCTCCCTAGACT AAATTTTGCCTTGGGTCT	M





Sequence variations of the locus

Sequence variation results

RESULTS



?								
1234	31 Items found, displaying 1 t	to 10 Displa	ay 🔻 r	esults per page				
#	Name	Туре	Project	Experiment	Line	Ref. sequence	Position on ref. seq.	Gene/marker name
1	VVC2972A 512 10	SNP	BerrySize	VVC2972A- batch1	9006Mtp1	Major allele of variants in VVC2972A- batch1	512	VVC2972A
2	VVC2972A 512 11	SNP	<u>BerrySize</u>	VVC2972A- batch1	8056Mtp1	Major allele of variants in VVC2972A- batch1	512	VVC2972A
3	VVC2972A 512 12	SNP	BerrySize	VVC2972A- batch1	6541Mtp4	Major allele of variants in VVC2972A- batch1	512	VVC2972A
4	VVC2972A 512 16	SNP	<u>BerrySize</u>	VVC2972A- batch1	8010Mtp1	Major allele of variants in VVC2972A- batch1	512	VVC2972A
5	VVC2972A 512 18	SNP	BerrySize	VVC2972A- batch1	8003Mtp1	Major allele of variants in VVC2972A- batch1	512	VVC2972A





Resequencing batch genotypes summary

Genotype table

RESULTS



1234 ▶ № | 36 Items found, displaying 1 to 10 | Display | 10 ▼ results per page

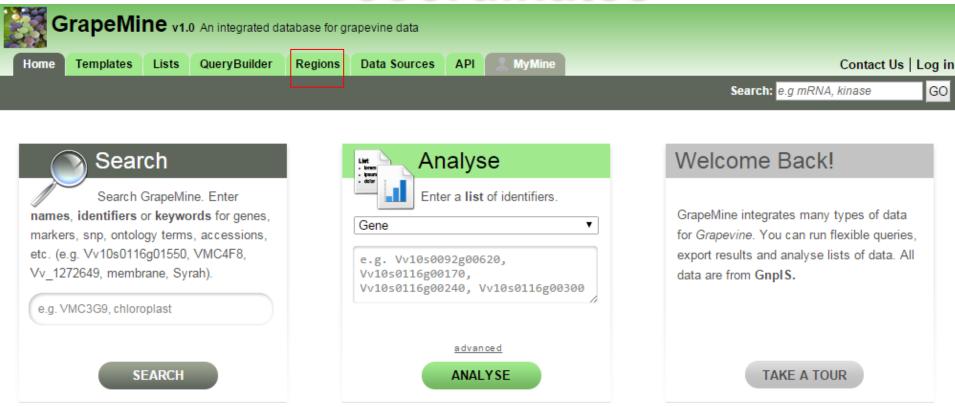
The experiment WC2972A-batch1 contains 32 marker(s)

Positions			4	15	17	21	23	46	51	53	84	99	176	283	303	321	340	346	360	391	396	427	482	487
Ref. Sequence	Major allele of variants in VVC2972A- batch1		A	ы	ю	A	DEL	e e	e e	T	ы	ю	A	C	G	T	T	DEL	IN	C	U	e e	G	T
Genotype Taxon	9006Mtp1	Vitis riparia cv. Riparia Gioire de Montpellier	Α	O I	01	Ol	<u></u>	GI	Øl.	Ī	G)	<u>0</u> 1	Α	CI	CI	Ī	CI	- 11	ØI	CI	CI	Δ	CI	I
Genotype Taxon	0Mtp567	<u>Vitis vinifera</u> <u>L.</u>	Α	G	G	Δ	TTCTGGTATTGGCGTTGGC	G	G	<u>C/T</u>	G	G	<u>A/C</u>	CI	CI	Ī	Ī	Ī	1	<u>C</u>	CI	GI	CI	I
Genotype Taxon	0Mtp636	<u>Vitis vinifera</u> <u>L.</u>	Α	G	G	Α	TTCTGGTATTGGCGTTGGC	G	G	<u>C/T</u>	G	G	A/C	<u>C</u>	<u>C</u>	Ī	Ī	Ī	1	<u>C/T</u>	<u>C</u>	G	<u>C</u>	I
Genotype Taxon	1498Mtp1	Vitis vinifera subsp vinifera cv. Espadeiro tinto	Α	Ø!	10	Α	TTCTGGTATTGGCGTTGGC	GI	Ol.	Ī	GI.	01	A/C	CI	O	I	I	I	1	CI	CI	(O)	O	I
Genotype Taxon	225Mtp2	Vitis vinifera subsp vinifera cv. Cesar	Α	OI.	01	A	TTCTGGTATTGGCGTTGGC	<u>0</u> 1	<u>0</u> 1	I	<u>0</u> 1	01	Α	Ю	Ol	Ī	I	I	ī.	Ю	Ю	lo	Ol	I
Genotype Taxon	0Mtp1323	<u>Vitis vinifera</u> <u>L.</u>	Α	G	G	Α	TTCTGGTATTGGCGTTGGC	G	G	<u>c</u>	G	G	<u>c</u>	<u>c</u>	<u>c</u>	Ī	Ī	Ī	1	<u>c</u>	<u>c</u>	G	<u>c</u>	I
Genotype Taxon	8500Mtp23	<u>Vitis vinifera</u> <u>L.</u>	Δ	g	g	Α	TTCTGGTATTGGCGTTGGC	G	g	Ī	G	g	Α	<u>C</u>	<u>C</u>	Ī	<u>C/T</u>	Ī	n/a	n/a	n/a	n/a	n/a	n/a





Use case 2: Enter by genome coordinates



GrapeMine: http://urgi.versailles.inra.fr/GrapeMine/begin.do

More details on Intermine functionnalities Wheat3BMine Poster #654: Letellier T. et al. and computer demo: January 13th 11:00 am



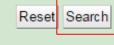
GrapeMine: Region form

Search for features within Genomic Regions

Search for features that overlap a list of genome coordinates you enter or upload, e.g. chr1:201000000
--

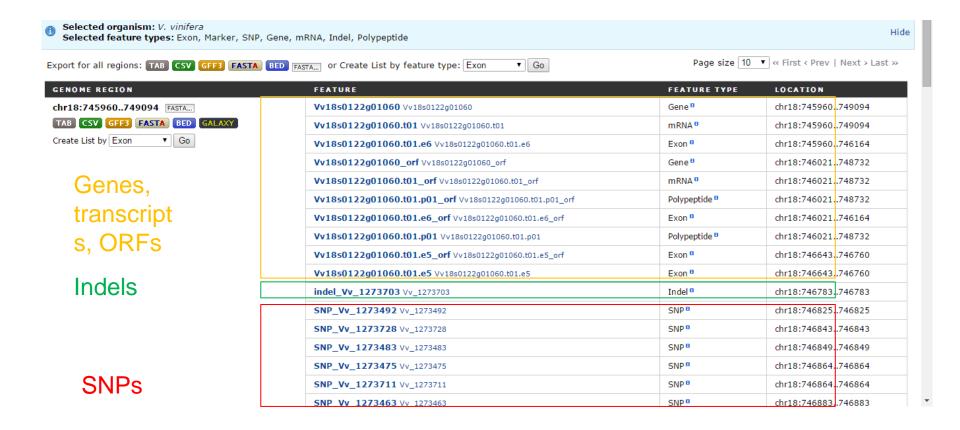
_				
Gon	$\alpha m \alpha$	coordina	toe ha	Mn.
oen	ome	COOI UIII a	ces ne	210

	Select Organism: V. vinifera ▼ genome ✓ Select Feature Types:	e build: not availa	ble	
	✓ Exon ^B ✓ Gene ^B ✓ Indel ^B	✓ Marker ⁰ ✓ mRNA ⁰ ✓ Polypeptide ⁰		 SNP□
3.	Type/Paste in genomic regions in [®] base (click to see an example)▼	coordinate ⁸ O i	interbase coordin	ate ⁰
	chr18:745,960749,094		/2	
	or Upload genomic regions from a .txt file Choisissez un fichier Aucun fichier choi			
4.	Extend your regions at both sides:			
	0 1k	10k 	100k 	1M 10M













Zoom on a SNP

SNP: SNP_V	v_1273492 V. vinifera	7			
DB identifier Allele Taxon	Vv_1273492 A/G Vitis vinifera L.	Organism . Name Gnpsnpid	Vitis vinifera 1273492		
Genome feature					Lists
Region:	SNP	Length:		1	This SNP isn't in any lists. U
Location:	chr18:746825-746825 f	orward strand			a list.
					External Links
Overlapping Features Genome features that over Genes: 2, mRNAs: 2, Polyp	erlap coordinates of this SNP				GRrowse URGI VITIS12X GNP-SNP URGI VITIS12X
		Show all in a table »			
1 Organism					
Name		Tax	on Id		
Vitis vinifera		2976	60		



1 Data Sets

Name markersvitis12X URL



List of the species concerned by this marker

Genotype results

AVAILABLE GENOTYPES



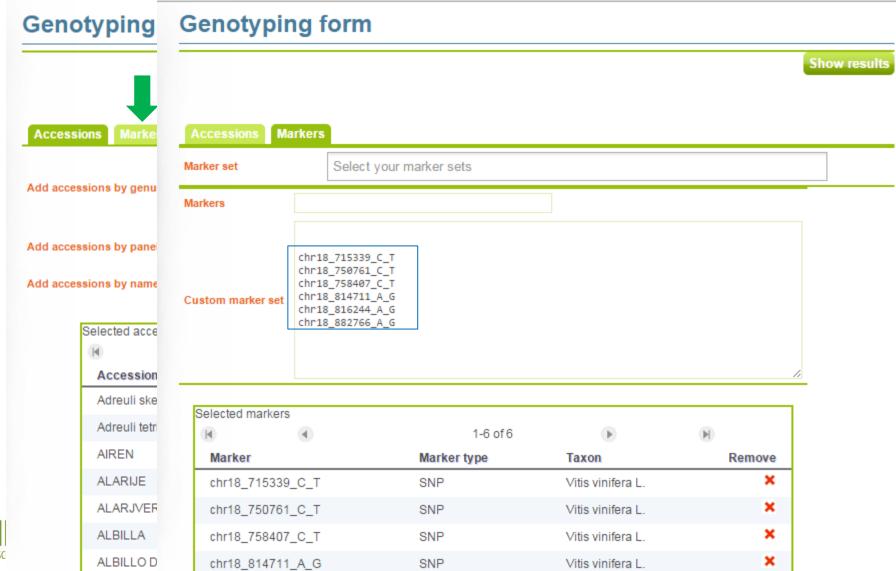
#	Genotype name	Taxon	
1	2082Mtp1	√itis vinifera subsp vinifera cv. Mehdik	
2	9006Mtp1	Vitis riparia cv. Riparia Gloire de Montpellier	
3	0Mtp567	<u>Vitis vinifera L.</u>	
4	0Mtp636	<u>Vitis vinifera L.</u>	
5	8011Mtp12	<u>Vitis berlandieri Planch.</u>	
6	225Mtp2	<u>Vitis vinifera subsp vinifera cv. César</u>	
7	0Mtp1323	<u>Vitis vinifera L.</u>	
8	8003Mtp1	<u>Vitis amurensis Ruprecht</u>	
9	324Mtp1	<u>Vitis vinifera subsp vinifera cv. Cabernet franc</u>	
10	1186Mtp1	Vitis vinifera subsp vinifera cv. Chirai obak	





Diversity in V. vinifera in the chr18: 700K-800k region (1)

GnpIS genotyping form: https://urgi.versailles.inra.fr/GnpSNP/snp/genotyping/form.do





EURGI

Diversity in V. vinifera in the chr18: Genotyping form 700K-800k region (2)

Back to form

	A	Α	В	С	D	E	F	G	Н	- 1
ENOTYPING RESULTS	1 markerNa	ne	AccessionName	Genotype type	Genotype	Quality type	Quality	Experiment	Genotyping a	irray
	2 chr18_814	711_A_G	Wildbacherfruehblau	genotype	AA			GrapeReSeq_Illumina_20K_experiment	GrapeReSeq_	_Illumina_
1-10 of 560	3 chr18_814	711_A_G	LAIREN	genotype	AA			GrapeReSeq_Illumina_20K_experiment	GrapeReSeq	_Illumina_
	4 chr18_814	711_A_G	SENA	genotype	AA			GrapeReSeq_Illumina_20K_experiment	GrapeReSeq	_Illumina_
<u>Marker</u>	5 chr18_814	711_A_G	TROUSSEAU NOIR	genotype	AG			GrapeReSeq_Illumina_20K_experiment	GrapeReSeq	_Illumina_
chr18 814711 A G	6 chr18_814	711_A_G	LADO	genotype	AA			GrapeReSeq_Illumina_20K_experiment	GrapeReSeq_	_Illumina_
	7 chr18_814	711_A_G	ALVARELHAO	genotype	AA			GrapeReSeq_Illumina_20K_experiment	GrapeReSeq_	_Illumina_
chr18 814711 A G	8 chr18_814	711_A_G	Tauberschwarz	genotype	AA			GrapeReSeq_Illumina_20K_experiment	GrapeReSeq_	_Illumina_
chr18 814711 A G	9 chr18_814	711_A_G	Oeilladenoire	genotype	AA			GrapeReSeq_Illumina_20K_experiment	GrapeReSeq_	_Illumina_
chr18 814711 A G	10 chr18_814	711_A_G	TOURIGAFRANCA	genotype	AA			GrapeReSeq_Illumina_20K_experiment	GrapeReSeq_	_Illumina_
chr18 814711 A G	11 chr18_814	711_A_G	Javorweiss	genotype	AA			GrapeReSeq_Illumina_20K_experiment	GrapeReSeq_	_Illumina_
	12 chr18_814	711_A_G	TrincadeiradasPratas	genotype	AA			GrapeReSeq_Illumina_20K_experiment	GrapeReSeq_	_Illumina_
chr18 814711 A G	13 chr18_814	711_A_G	DERECHERO DE MUNIES	genotype	AA			GrapeReSeq_Illumina_20K_experiment	GrapeReSeq_	_Illumina_
chr18 814711 A G	14 chr18_814	711_A_G	MIGUELDEARCO	genotype	AA			GrapeReSeq_Illumina_20K_experiment	GrapeReSeq_	Illumina_
chr18 814711 A G	15 chr18_814	711_A_G	Heunischrot	genotype	AA			GrapeReSeq_Illumina_20K_experiment	GrapeReSeq_	Illumina
chr18 814711 A G	16 chr18_814	711_A_G	TARRAGONI	genotype	AA			GrapeReSeq_Illumina_20K_experiment	GrapeReSeq_	Illumina
	17 chr18_814	711_A_G	E32 07	genotype	AA			GrapeReSeq_Illumina_20K_experiment	GrapeReSeq_	Illumina
<u>chr18 814711 A G</u>	18 chr18_814	711_A_G	FORCALLAT TINTA	genotype	AA			GrapeReSeq_Illumina_20K_experiment	GrapeReSeq_	Illumina
	19 chr18_814	711_A_G	Savagninblanc	genotype	AA			GrapeReSeq_Illumina_20K_experiment	GrapeReSeq_	Illumina
	20 chr18_814	711_A_G	VERDIL	genotype	AA			GrapeReSeq_Illumina_20K_experiment	GrapeReSeq	Illumina
V	21 chr18_814	711_A_G	CONCA D???ORO	genotype	AA			GrapeReSeq_Illumina_20K_experiment	GrapeReSeq_	Illumina
	22 chr18_814	711_A_G	Limnio	genotype	AA			GrapeReSeq_Illumina_20K_experiment	GrapeReSeq	Illumina
	23 chr18_814	711_A_G	ZALEMA	genotype	AA			GrapeReSeq_Illumina_20K_experiment	GrapeReSeq	Illumina
	24 chr18_814	711_A_G	BERMEJUELA	genotype	AA			GrapeReSeq_Illumina_20K_experiment	GrapeReSeq	Illumina
ENOTYPING MATRIX	25 chr18_814	711_A_G	Bakatorroz	genotype	AA			GrapeReSeq_Illumina_20K_experiment	GrapeReSeq	_ _Illumina_
100 of 6	H 4 → H ger	otyping2928	940581837811622					14		

Marker \ Accession	ALBILLA	Adreuli skelkana	ALARIJE	ALCA? ON	AIREN	ALLAREN	ALEDO	Adreuli tetri	ALMERIA NERA
chr18_882766_A_G	GG	AA	AA	AA	AG				
chr18_758407_C_T	CC				CC	CC	CC	CC	CC
chr18_816244_A_G			AA			AA		AG	AA
chr18_750761_C_T	TC	TC		CC		CC	CC		CC
chr18_715339_C_T	TC				CC	CC	TT	CC	
chr18_814711_A_G	AA	AA	AA	AA		AA			





Prospects

Jbrowse based on the Vitis12X.2 assembly

Learn more about this assembly:

https://urgi.versailles.inra.fr/Species/Vitis/Data-Sequences/Genome-sequences

Add GWAS Vitis data (INRA)





Available soon: Vitis GWAS

data

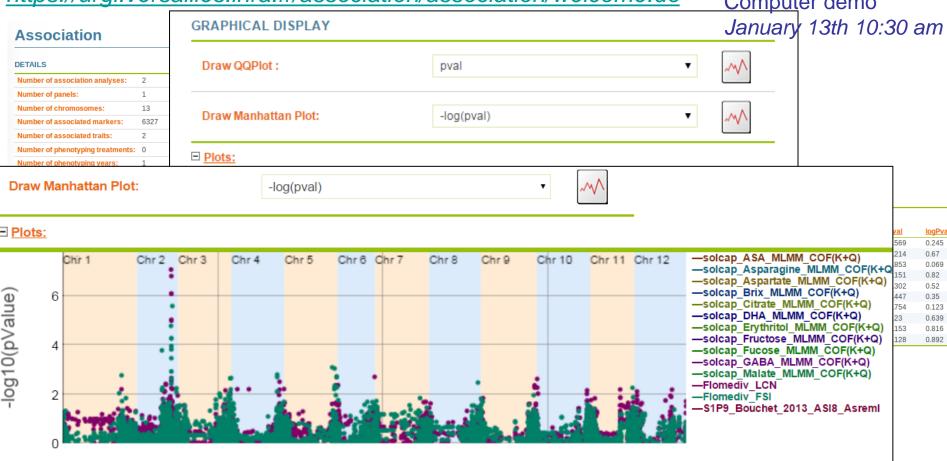
GnpIS Genetic Association form:

https://urgi.versailles.inra.fr/association/association/welcome.do

More details on GnpAsso:

Poster #1128

Computer demo



Position of markers by chromosome



0.1

Acknowledgments

URGI TEAM



- INRA PARTNERS
 INRA EPGV, INRA AGAP,
 INRA URGV...
- PROJECT PARTNERS Grapereseq, Muscares, SNPGrapeMap...





THANK YOU FOR YOUR ATTENTION!





Contact: nacer.mohellibi@versailles.inra.fr

