

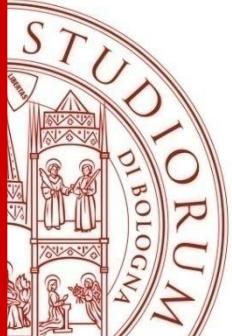
Dalla Lunasina al nsLTP2: identificazione e suo contenuto nei frumenti

University of Bologna (UNIBO)
Deptment of Agricultural Science (DIPSA)
Giovanni Dinelli



Bologna, 28.05.2016





Available literature about lunasin in wheat and other species

(after lunasin from soybean identification)



Review

Lunasin, a novel seed peptide for cancer prevention

Blanca Hernández-Ledesma¹, Chia-Chien Hsieh¹, Ben O. de Lumen*

Peptides 30 (2009) 426–430

J. Agric. Food Chem. 2007, 55, 10707–10713

Cancer-Preventive Peptide Lunasin from *Solanum nigrum* L. Inhibits Acetylation of Core Histones H3 and H4 and Phosphorylation of Retinoblastoma Protein (Rb)

JIN BOO JEONG,^{†,‡} HYUNG JIN JEONG,^{†,‡} JAE HO PARK,^{†,‡} SUN HEE LEE,[‡]
JEONG RAK LEE,[‡] HEE KYEONG LEE,[‡] GYU YOUNG CHUNG,[‡] JEONG DOO CHOI,[§]
AND BEN O. DE LUMEN*,^{¶,II}

Nutrition and Cancer, 61(5), 680–686
Copyright © 2009, Taylor & Francis Group, LLC
ISSN: 0163-5581 print / 1532-7914 online
DOI: 10.1080/01635580902850082

The Cancer Preventive Seed Peptide Lunasin From Rye Is Bioavailable and Bioactive

Hyung Jin Jeong, Jeong Rak Lee, Jin Boo Jeong, and Jae Ho Park
Young-keun Cheong
Ben O. de Lumen

Nutrition and Cancer, 62(8), 1113–1119
Copyright © 2010, Taylor & Francis Group, LLC
ISSN: 0163-5581 print / 1532-7914 online
DOI: 10.1080/01635581.2010.515529

Lunasin Is Prevalent in Barley and Is Bioavailable and Bioactive in In Vivo and In Vitro Studies

Hyung Jin Jeong^a, Jin Boo Jeong^a, Chia Chien Hsieh^b, Blanca Hernández-Ledesma^b &
Ben O. de Lumen^b

J. Agric. Food Chem. 2002, 50, 5903–5908

Barley Lunasin Suppresses ras-Induced Colony Formation and Inhibits Core Histone Acetylation in Mammalian Cells

HYUNG J. JEONG,[†] YI LAM,[‡] AND BEN O. DE LUMEN*,[§]

Journal of Cereal Science xxx (2012) 1–5

Discovery of lunasin peptide in triticale (X *Triticosecale* Wittmack)

Ilva Nakurte^a, Kristaps Klavins^b, Inga Kirhnere^c, Jana Namniece^d, Liene Adlere^d, Jaroslavs Matvejevs^a,
Arta Kronberga^c, Aina Kokare^c, Vija Strazdina^e, Linda Legzdina^c, Ruta Muceniece^{d,*}

J. Agric. Food Chem. 2008, 56, 1239–1240

Bioactive Peptides in Amaranth (*Amaranthus hypochondriacus*) Seed

C. SILVA-SÁNCHEZ,^{†,‡} A. P. BARBA DE LA ROSA,[†] M. F. LEÓN-GALVÁN,[†]
B. O. DE LUMEN,[†] A. DE LEÓN-RODRÍGUEZ,[†] AND E. GONZÁLEZ DE MEJIA*,[§]

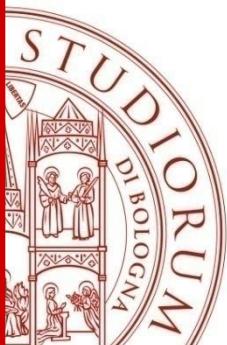
Cancer Letters 255 (2007) 42–48

The cancer preventive peptide lunasin from wheat inhibits core histone acetylation

Hyung Jin Jeong^{1,a}, Jin Boo Jeong^{1,a}, Dae Seop Kim^{1,a}, Jae Ho Park^a,
Jung Bok Lee^a, Dae-Hyuk Kweon^b, Gyu Young Chung^a, Eul Won Seo^a,
Ben O. de Lumen^{c,*}

Synthesis of the Cancer Preventive Peptide Lunasin by Lactic Acid Bacteria During Sourdough Fermentation

Carlo G. Rizzello, Luana Nionelli, Rossana Coda, and Marco Gobbetti

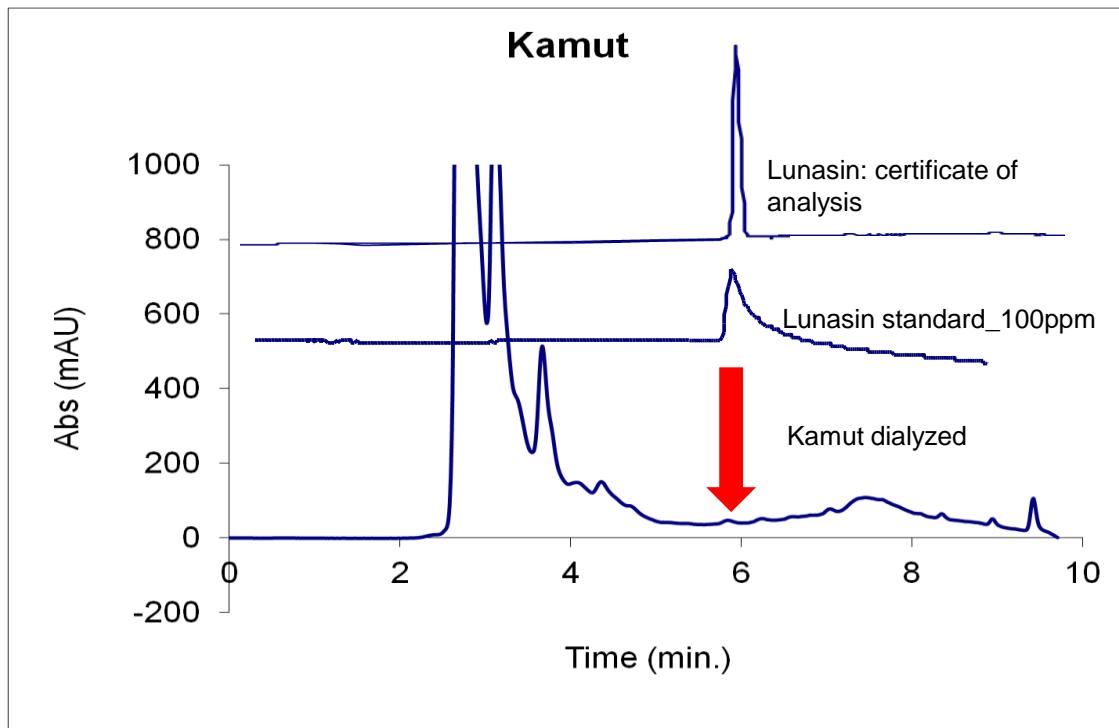


The UNIBO work: the beginning of the doubt!!



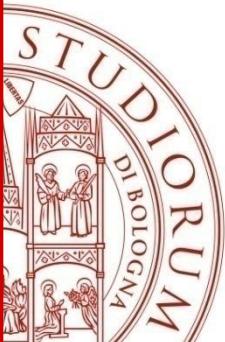
December 2012

HPLC analysis: our attempts in separation of lunasin standard and its identification in Kamut dialyzed sample.



BAD RESOLUTION

**DIFFICULT
IDENTIFICATION
OF LUNASIN IN WHEAT
SAMPLE**

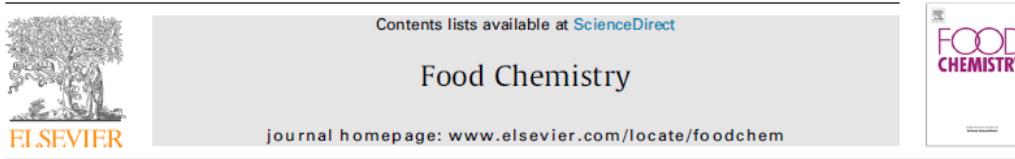


LUNASIN is not present in wheat

Jan-Dec 2013: we demonstrated the lacking of lunasin in wheat

- Lunasin gene not identified in wheat (*Dinelli et al. J. Food Chem, 2014*)

Food Chemistry 151 (2014) 520–525



Rapid Communication

Lunasin in wheat: A chemical and molecular study on its presence or absence



Giovanni Dinelli ^{a,*}, Valeria Bregola ^a, Sara Bosi ^a, Jessica Fiori ^b, Roberto Gotti ^b,
Emanuela Simonetti ^c, Caterina Trozzi ^c, Emanuela Leoncini ^d, Cecilia Prata ^b, Luca Massaccesi ^d,
Marco Malagutti ^d, Robert Quinn ^e, Silvana Hrelia ^d

^aDepartment of Agricultural Sciences, Alma Mater Studiorum University of Bologna, Viale Fanin 44, 40127 Bologna, Italy

^bDepartment of Pharmacy and Biotechnology, Alma Mater Studiorum University of Bologna, Via Belmeloro 6, 40126 Bologna, Italy

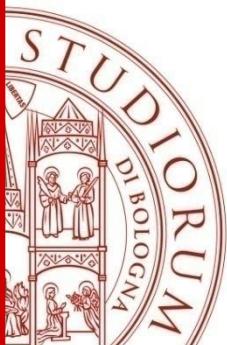
^cBioavantis, Via Rocco Scotellaro 1/A, 60035 Jesi (AN), Italy

^dDepartment for Life Quality Studies, Alma Mater Studiorum University of Bologna, Corso D'Augusto 237, 47921 Rimini, Italy

^eKamut International, 333 Kamut Lane, Big Sandy, MT 59520, USA

-NEWLY ISOLATED WHEAT PEPTIDE

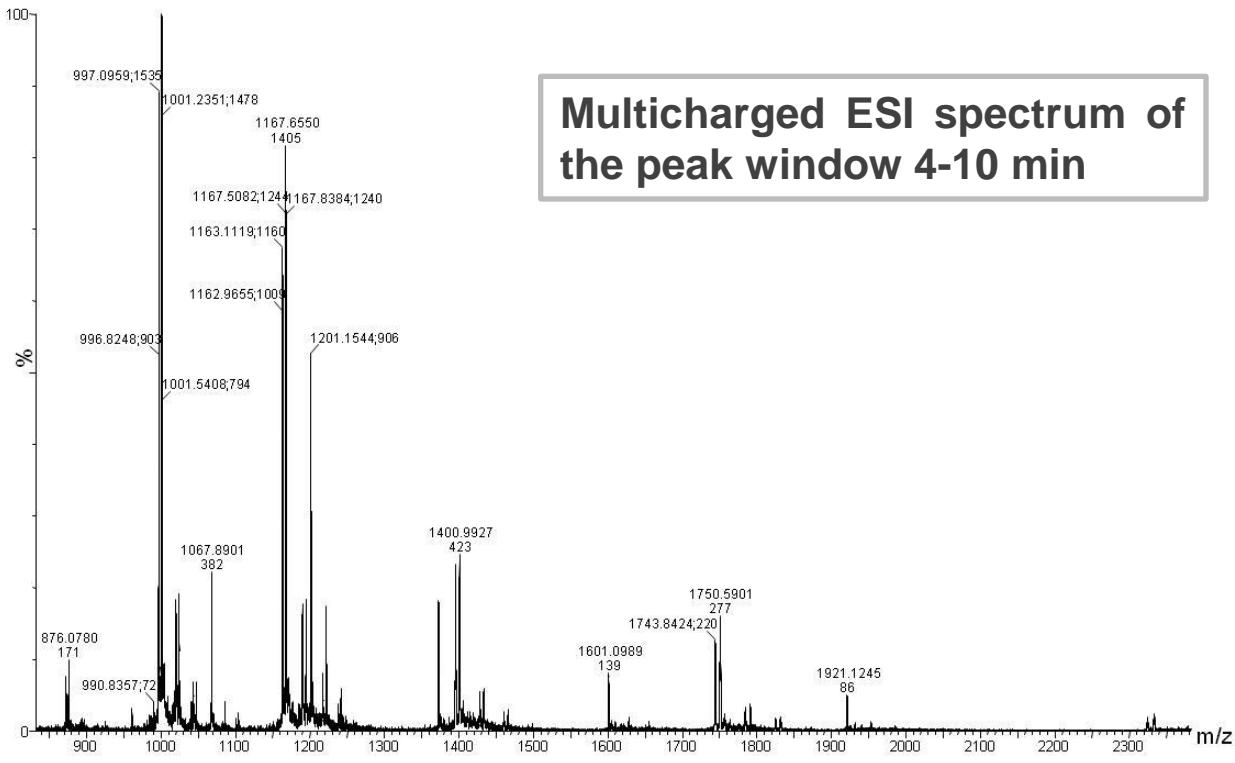
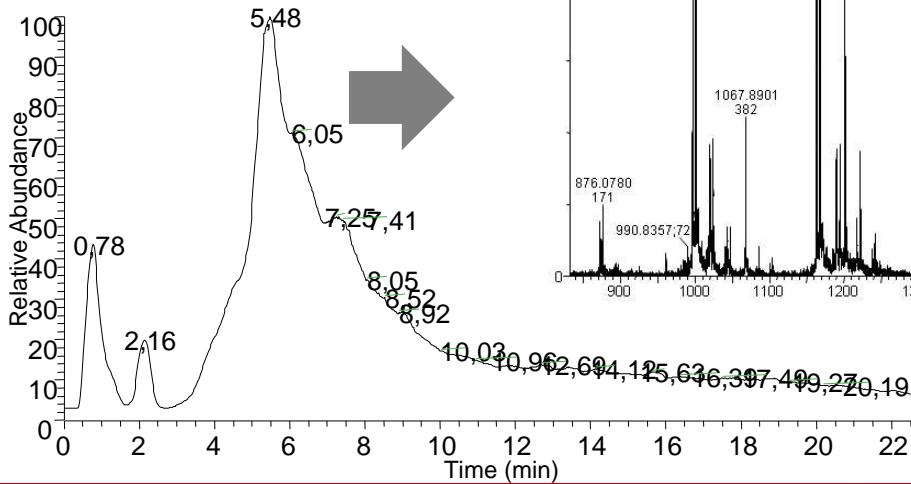
- Sequenced and identified according to the following database



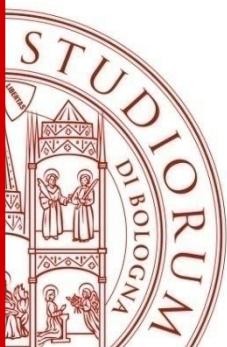
Where did the mistake occur?

February 2014: we identified a new wheat peptide

TIC MS chromatogram

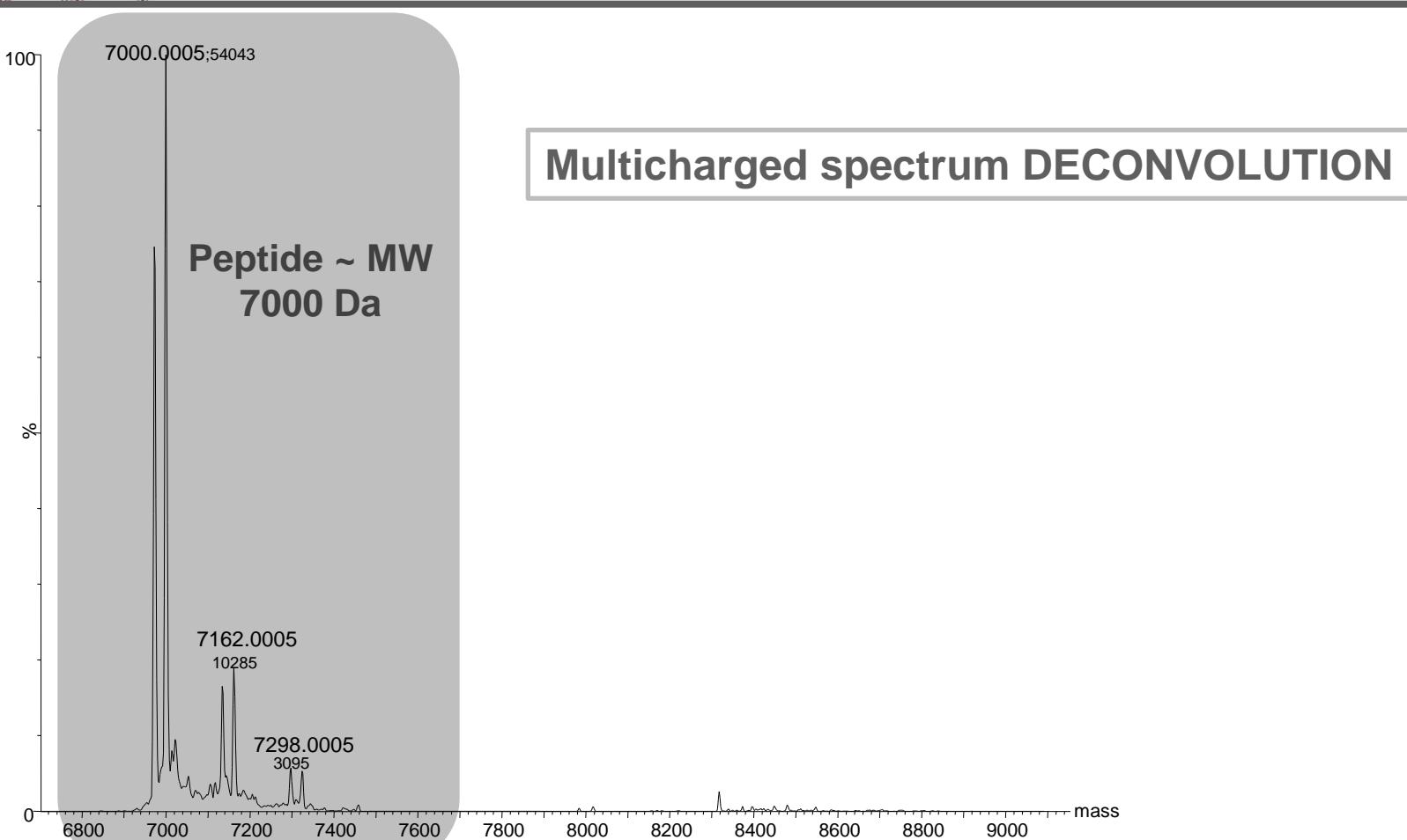


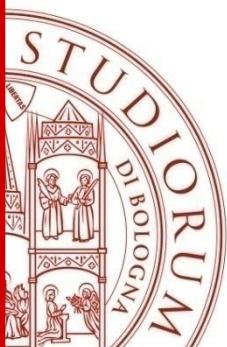
Multicharged ESI spectrum of the peak window 4-10 min



WHERE did the mistake occur?

The multicharged spectrum deconvolution
showed the presence of one main peptide





NEW WHEAT PEPTIDE IDENTIFICATION

MASCOT

Peptide MW around 7000 Da

SwissProt, NCBI

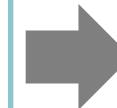
NLT2P Wheat; PM 7046

Non-specific lipid-transfer protein 2P OS=Triticum aestivum

Protein sequence coverage: 76%

1 ACQASQLAVC ASAILSGAKP SGECCGNLRA QQPCFCQYAK DPTYGQQYIRS

51 PHARDTLQSC GLAVPHC



NsLtP type 2

NCBI

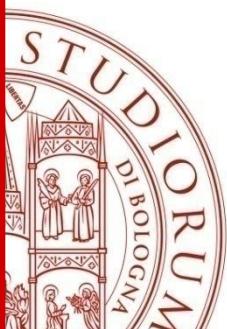
Chain A, Solution Structure Of A Liganded Type 2

Wheat Non-Specific Lipid Transfer Protein; PM 6979

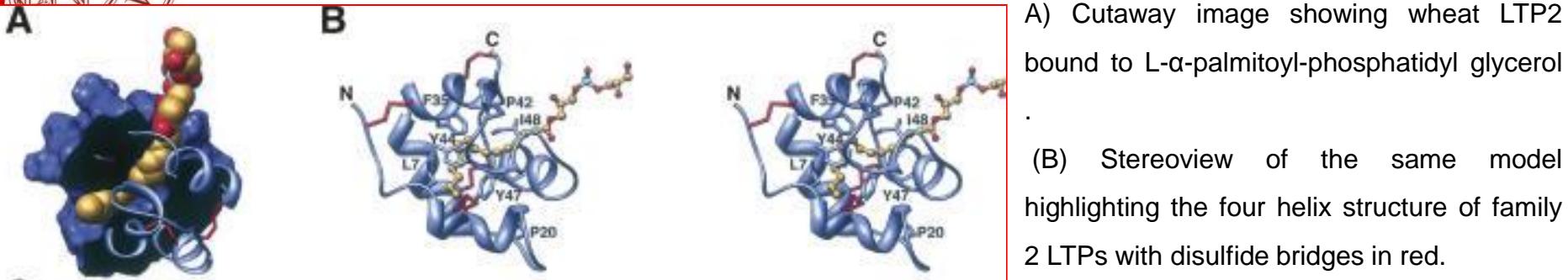
Protein sequence coverage: 92%

1 ACQASQLAVC ASAILSGAKP SGECCGNLRA QQGCFCQYAK DPTYGQQYIRS

51 PHARDTLTSC GLAVPHC

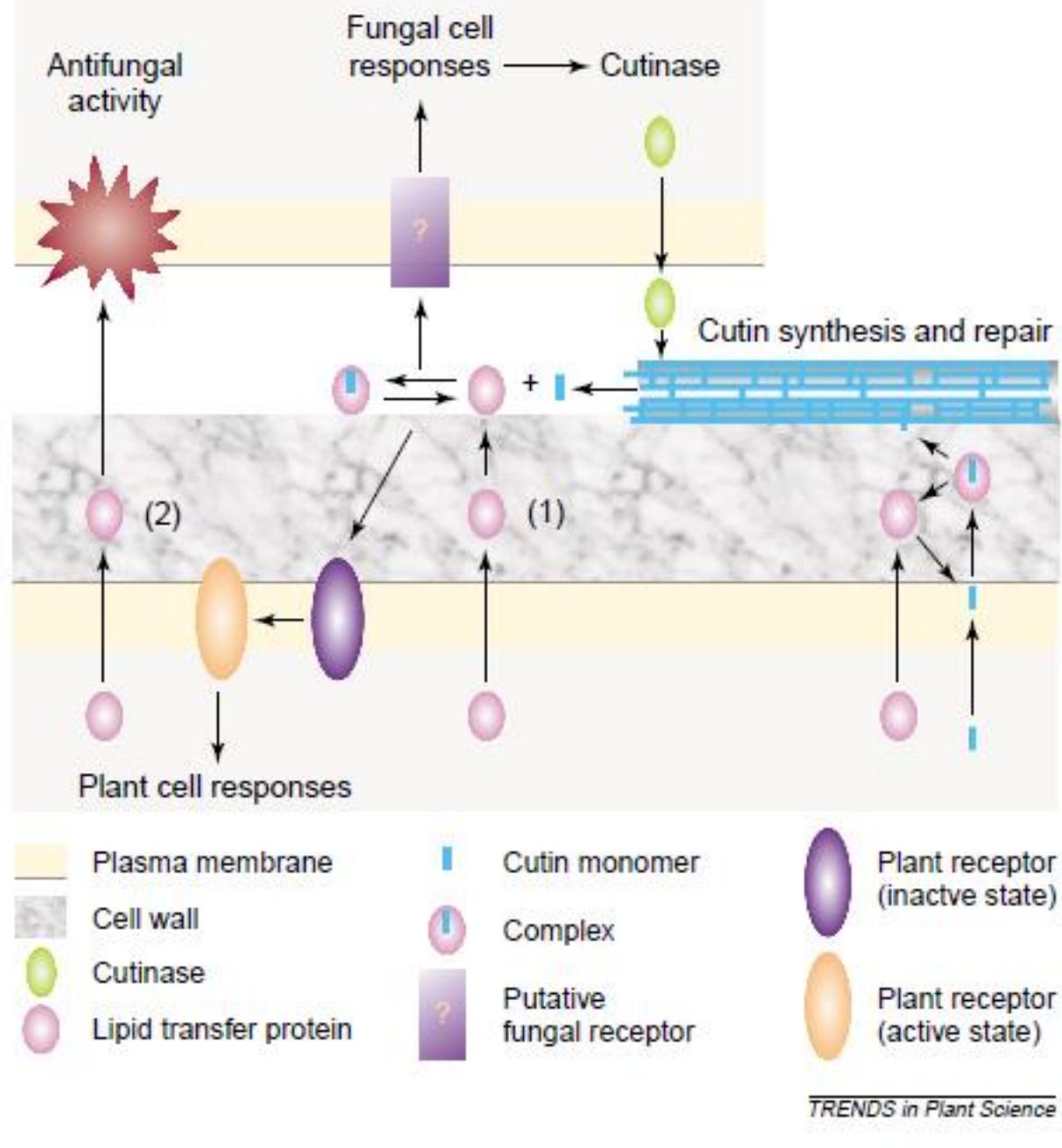


NEW WHEAT PEPTIDE IDENTIFICATION



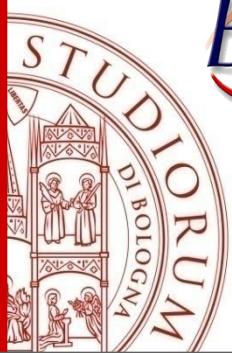
- ✓ The peptide appears to belong to the family of lipid-transfer protein 2 (LTP).
- ✓ LTP fold is characterized by a compact domain composed of 4 α -helices bound by four disulfide bonds enclosing a hydrophobic lipid binding cavity.
- ✓ A number of biological roles (antimicrobial defense, signaling, assembly of cutin) have been proposed, but conclusive evidence is generally lacking [Yeats TH, Rose JK. *Protein Sci*, 2008].
- ✓ Although both LTP1 and LTP2 proteins can be found in plant seeds, thus far, only LTP1 proteins have been characterized as allergens. In fact, the official allergen list of the International Union of Immunological Societies currently contains no entries for allergens belonging to the LTP2 subfamily [Egger M et al., *Curr Allergy Asthma Rep*, 2010]

NEW WHEAT PEPTIDE IDENTIFICATION



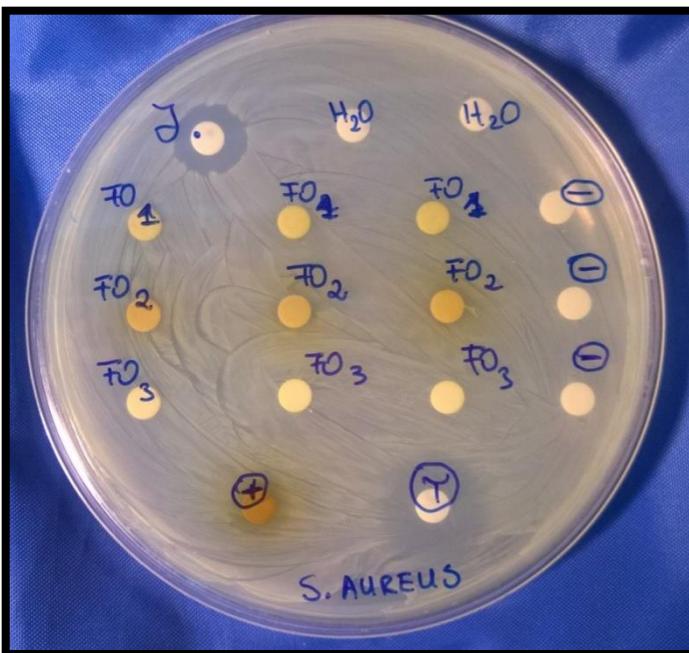
nsLTP is involved in plant defense mechanism against abiotic (ie salinity) and biotic stresses (bacteria, fungi)

Double action: direct (ie antifungal activity) and indirect (ie «SHIELD» effect)

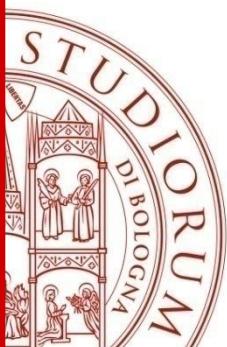


NEW WHEAT PEPTIDE IDENTIFICATION

Since the mentioned **NsLtP type 2** is reported to have antibacterial activity, a specific test carried out with *S. aureus* confirmed the chemical identification.



J = new peptide
T = Tetracycline



NEW WHEAT PEPTIDE IDENTIFICATION

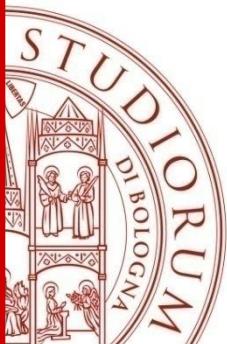
TRENDS IN MICROBIOLOGY 73 VOL. 3 NO. 2 FEBRUARY 1995

The defensive role of nonspecific lipid-transfer proteins in plants

**Francisco García-Olmedo, Antonio Molina,
Ana Segura and Manuel Moreno**

Questions

- What function(s), other than their defensive role, do nonspecific lipid-transfer proteins (nsLTPs) have in plants?
- What mechanism(s) are responsible for the toxicity of nsLTPs to pathogens?
- What mechanisms are responsible for possible microbial resistance to nsLTPs?
- Are nsLTPs essential for plants?
- Are nsLTPs a lead for drug development?



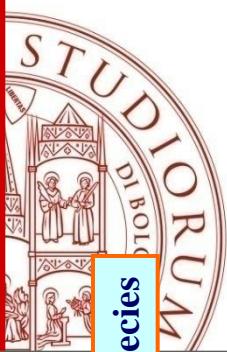
WP 3 : Extending eBASIS to study habitual intakes of bioactive compounds in the diet

Task 3.1.4.2: nsLPT2 composition of European wheats

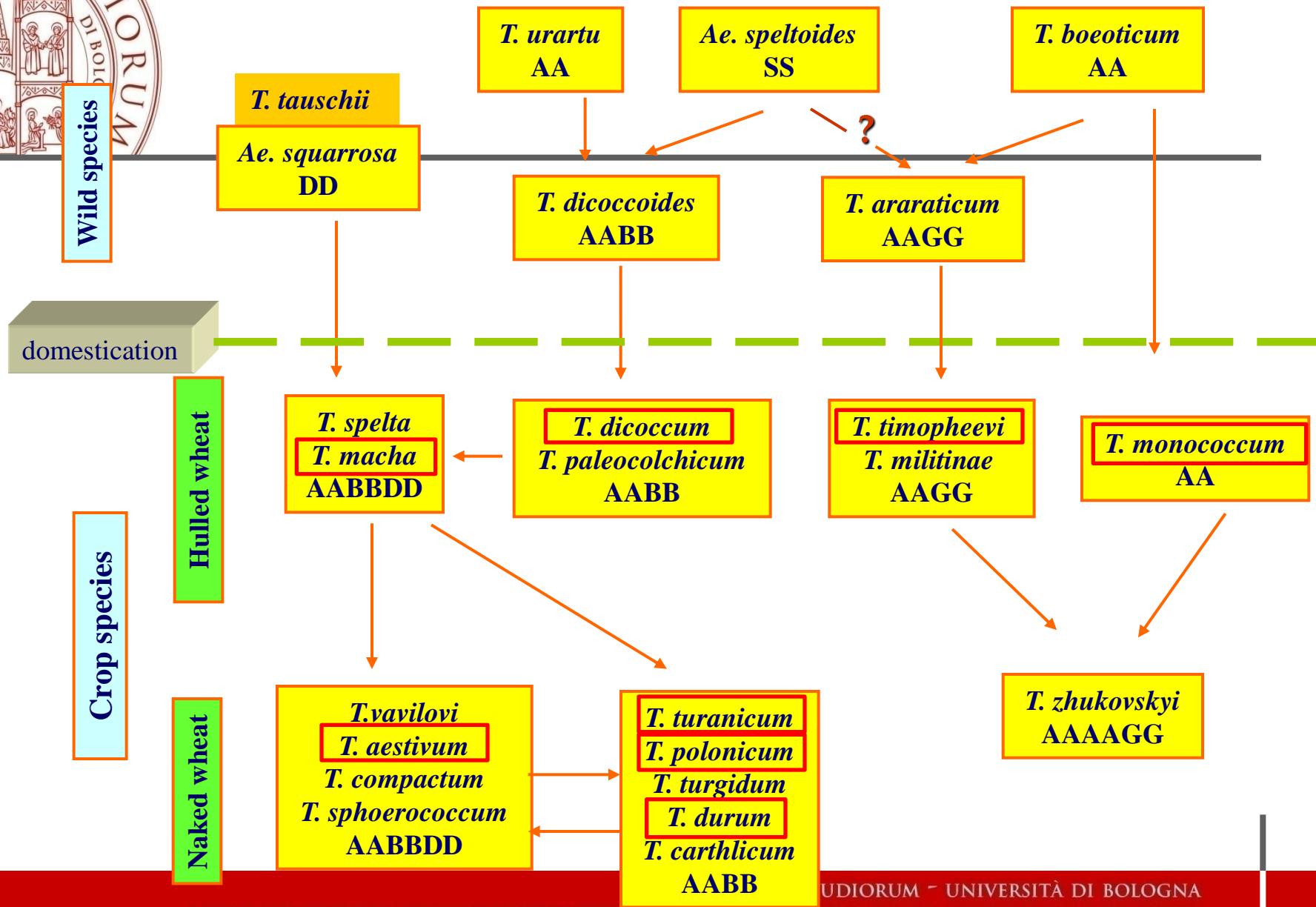
A germplasm collection of **60 accessions** of wheat genotypes held at UNIBO, and representative of the main cultivars cultivated in Europe, have been screened for their nsLTP2 content in whole grain.

- 11 durum wheat;
- 2 *Triticum turgidum* ssp. *turanicum*,
- 38 *Triticum aestivum*;
- 1 *Triticum polonicum*;
- 1 *Triticum macha*,
- 1 *Triticum timophevi*,
- 3 *Triticum monococcum*,
- 3 *Triticum dicoccum*.

The results produced have been added to the eBASIS database, using a quality assessment system identical to that used for peer reviewed publications.



WHEAT GENEALOGY



ALL HEXAPLOID SPECIES

Species	Variety	nsLTP 2 (mg/g grain)
<i>Tr. aestivum</i>	Abbondanza	0.172 ± 0.021
<i>Tr. aestivum</i>	Amarok	0.107 ± 0.006
<i>Tr. aestivum</i>	Andriolo	0.219 ± 0.017
<i>Tr. aestivum</i>	Argelato	0.095 ± 0.008
<i>Tr. aestivum</i>	Aubusson	0.118 ± 0.009
<i>Tr. aestivum</i>	Autonomia A	0.224 ± 0.034
<i>Tr. aestivum</i>	Autonomia B	0.201 ± 0.025
<i>Tr. aestivum</i>	Benco	0.232 ± 0.017
<i>Tr. aestivum</i>	Bianco nostrale	0.262 ± 0.041
<i>Tr. aestivum</i>	Bilancia	0.121 ± 0.012
<i>Tr. aestivum</i>	Blasco	0.098 ± 0.002
<i>Tr. aestivum</i>	Bolero	0.103 ± 0.005
<i>Tr. aestivum</i>	Bologna	0.114 ± 0.010
<i>Tr. aestivum</i>	Bramante	0.109 ± 0.007
<i>Tr. aestivum</i>	Canove	0.274 ± 0.028
<i>Tr. aestivum</i>	Carosello	0.399 ± 0.051
<i>Tr. aestivum</i>	Fiorello	0.214 ± 0.022
<i>Tr. aestivum</i>	Frassineto	0.175 ± 0.012
<i>Tr. aestivum</i>	Funo	0.205 ± 0.016

mean 0.208 mg/g ($0.095 \div 0.469$)

<i>Tr. aestivum</i>	Gamba di Ferro	0.396 ± 0.037
<i>Tr. aestivum</i>	Gentil Bianco	0.301 ± 0.022
<i>Tr. aestivum</i>	Gentil Rosso	0.205 ± 0.017
<i>Tr. aestivum</i>	Guà 113	0.305 ± 0.033
<i>Tr. aestivum</i>	Inallettabile	0.224 ± 0.021
<i>Tr. aestivum</i>	Marzuolo d'Aqui	0.194 ± 0.011
<i>Tr. aestivum</i>	Marzuolo Val Pusteria	0.149 ± 0.010
<i>Tr. aestivum</i>	Mentana	0.169 ± 0.013
<i>Tr. aestivum</i>	Mieti	0.106 ± 0.008
<i>Tr. aestivum</i>	Nobel	0.101 ± 0.006
<i>Tr. aestivum</i>	Palesio	0.112 ± 0.011
<i>Tr. aestivum</i>	Piave	0.255 ± 0.023
<i>Tr. aestivum</i>	Rieti	0.243 ± 0.018
<i>Tr. aestivum</i>	Serio	0.119 ± 0.009
<i>Tr. aestivum</i>	Sieve	0.469 ± 0.048
<i>Tr. aestivum</i>	Solina	0.335 ± 0.026
<i>Tr. aestivum</i>	Terricchio	0.170 ± 0.022
<i>Tr. aestivum</i>	Verna	0.162 ± 0.016
<i>Tr. aestivum</i>	Villa Gloria	0.428 ± 0.037
<i>Tr. macha</i>		0.242 ± 0.023

The present study highlighted high variability among the investigated ***Triticum aestivum*** genotypes (AABBDD); in fact the observed values ranged between 0.095 (cv. Argelato) and 0.469 mg/g whole flour (cv. Sieve).

Main source of variation: **plant habitus** and the **breeding year**

In the 11 post "green revolution" dwarf genotypes, the nsLTP-2 content was approximately halve than that found in the 27 not dwarf pre "green revolution" varieties ($0.110 \text{ mg.g}^{-1} \pm 0.008$ whole flour vs $0.247 \text{ mg.g}^{-1} \pm 0.091$ whole flour respectively).

DIPLOID SPECIES mean 0.318 mg/g (0.311 ÷ 0.325)

Species	Variety	nsLTP 2 (mg/g grain)
<i>Tr. monococcum</i>	white-type	0.319 ± 0.018
<i>Tr. monococcum</i>	black-type	0.325 ± 0.021
<i>Tr. monococcum</i>	white-type	0.311 ± 0.016

Among the eight different species analyzed, the diploid species ***Triticum monococcum* (AA genome)** was characterized by the highest mean value content of nsLTP2 ($0.318 \text{ mg.g}^{-1} \pm 0.007$ whole flour), with concentration range between 0.311 mg.g^{-1} and 0.325 mg.g^{-1} whole flour.

ALL TETRAPLOID SPECIES mean 0.181 mg/g ($0.059 \div 0.312$)

Species	Variety	nsLTP 2 (mg/g grain)
<i>Tr. turgidum</i> ssp. <i>turanicum</i>	Etrusco	0.297 ± 0.035
<i>Tr. turgidum</i> ssp. <i>turanicum</i>	KAMUT® khorasan wheat	0.304 ± 0.053
<i>Tr. turgidum</i> ssp. <i>durum</i>	Alemanno	0.220 ± 0.018
<i>Tr. turgidum</i> ssp. <i>durum</i>	Anco Marzio	0.171 ± 0.075
<i>Tr. turgidum</i> ssp. <i>durum</i>	Ciccio	0.224 ± 0.027
<i>Tr. turgidum</i> ssp. <i>durum</i>	Claudio	0.187 ± 0.062
<i>Tr. turgidum</i> ssp. <i>durum</i>	Iride	0.167 ± 0.027
<i>Tr. turgidum</i> ssp. <i>durum</i>	Levante	0.162 ± 0.023
<i>Tr. turgidum</i> ssp. <i>durum</i>	Orobel	0.180 ± 0.043
<i>Tr. turgidum</i> ssp. <i>durum</i>	Senatore Cappelli	0.205 ± 0.032
<i>Tr. turgidum</i> ssp. <i>durum</i>	Solex	0.175 ± 0.045
<i>Tr. turgidum</i> ssp. <i>durum</i>	Svevo	0.170 ± 0.051
<i>Tr. turgidum</i> ssp. <i>durum</i>	Urria 12	0.198 ± 0.017
<i>Tr. turgidum</i> ssp. <i>dicoccum</i>	white-type	0.059 ± 0.006
<i>Tr. turgidum</i> ssp. <i>dicoccum</i>	white-type	0.067 ± 0.003
<i>Tr. turgidum</i> ssp. <i>dicoccum</i>	white-type	0.063 ± 0.004
<i>Tr. turgidum</i> ssp. <i>polonicum</i>		0.312 ± 0.049
<i>Tr. timophevi</i>		0.098 ± 0.011

mean 0.300 mg/g

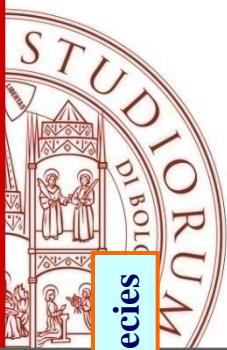
mean 0.187 mg/g

mean 0.063 mg/g

The lowest nsLTP-2 concentration was observed in the three **Emmer genotypes** (concentration range between 0.059 mg.g^{-1} and 0.063 mg.g^{-1} whole flour) and in the ***Triticum timophevi*** genotype ($0.098 \text{ mg.g}^{-1} \pm 0.011 \text{ mg.g}^{-1}$).

The ***Tr. turgidum* ssp. *durum*** showed intermediate mean value content with a limited variability among the eleven investigated varieties (0.162 mg.g^{-1} and 0.224 mg.g^{-1} whole flour in the varieties Levante and Ciccio, respectively).

The highest concentration of the peptide was observed for ***Triticum turanicum*** and ***Triticum polonicum*** with a concentration ranging between 0.297 mg.g^{-1} and 0.312 mg.g^{-1} whole flour.



WHEAT GENEALOGY

