

## ПРИЛОЖЕНИЕ

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«Алельное разнообразие генов *Vrn* и контроль типа и скорости развития у пшениц»

### Allelic variation in VRN loci of cultivated wheat species and relationship with heading date

Locus/ gene	Allele	Mutation	Accession	Heading date	References
Diploid wheat ( $2n = 2x = 14$ )					
VRN-A1	<i>Vrn-A<sup>m</sup>1</i>	3-bp del. in the promoter region	<i>T. monococcum</i> ssp. <i>aegilopoides</i> G2528		Yan et al., 2003
	<i>Vrn-1f</i>	1-bp del. in a CArG-box motif in the promoter region + 502-bp ins. in the intron 1	<i>T. monococcum</i> PI 503874		Dubcovsky et al., 2006
	<i>Vrn-1f</i>	1-bp del. in a CArG-box motif in the promoter region	<i>T. monococcum</i> TRI 28871		Shcherban et al., 2015b
	<i>Vrn-1g like</i>	34-bp del. in a CArG-box motif in the promoter region	<i>T. monococcum</i> K-18105		Golovkina et al., 2010
	<i>Vrn-1g like</i>	34-bp del. in a CArG-box motif in the promoter region	<i>T. boeoticum</i> K-40117 (w)		Golovkina et al., 2010
	<i>Vrn-1h</i>	9-bp del. in the promoter region + 493 bp ins. in intron 1	<i>T. monococcum</i> PI 306540 (w)	+	Dubcovsky et al., 2006
	<i>Vrn-1h like</i>	20 bp or 34 bp del. in the promoter region	<i>T. boeoticum</i> IG 116198, PI 427328		Golovkina et al., 2010
	<i>Vrn-A1ins</i>	0.5 kb ins. in intron 1	<i>T. monococcum</i> KM 586653		Shcherban et al., 2015b
	<i>Vrn-A1<sup>u</sup></i>		<i>T. boeoticum</i> TRI 17174, <i>T. urartu</i> TRI 17118		Shcherban et al., 2015b
<i>WAP1</i>	<i>mvp-1</i> mutant	19 kb del., includes promoter and coding region	<i>T. monococcum</i> KU-104-2	+	Shitsukawa et al., 2007
VRN-D	<i>Vrn-D<sup>1</sup>1</i>	5437 del. in intron 1	<i>Ae. tauschii</i> Coss. KU-20-6		Takumi et al., 2011
	New allele	3273 bp del. in the intron 1	<i>Ae. tauschii</i> K-992	+	Chepurinov et al., 2023
Tetraploid wheat ( $2n = 4x = 28$ )					
VRN-1 VRN-A1		7222-bp del. in intron 1	<i>T. durum</i> cv. Langdon		Fu et al., 2005
VRN-A1	<i>Vrn-A1b</i>	20-bp del. + 1-bp del. in the promoter region (other than $2n = 42$ )	<i>T. durum</i> US:ST 36		Yan et al., 2004a
	<i>Vrn-A1b like</i>	20-bp del. in the promoter region	<i>T. turanicum</i>		Golovkina et al., 2010
	<i>Vrn-A1b</i>	19-bp del. in the promoter region	<i>T. aethiopicum</i> IG 46273, IG 46306, IG 115810		Shcherban et al., 2015b
	<i>Vrn-A1b.1 - Vrn-A1b.5</i>	20-bp del. + 1-bp del. in the promoter region + polymorphism of the A-tract within the VRN-box in the promoter region	<i>T. turgidum</i> PI 264954, PI 223173 (w), <i>T. spelta</i> PI 348700, <i>T. macha</i> PI 428178 (i)		Muterko et al., 2016
	<i>Vrn-A1d</i>	32-bp del. in the promoter region	<i>T. dicoccoides</i> FA 15, Tabigha 15, Amrim 34, Iraq 8736		Yan et al., 2004b
	<i>Vrn-A1e</i>	54-bp del. in the promoter region	<i>T. dicoccum</i> ST 27 (=Vernal)		Yan et al., 2004b
	<i>Vrn-A1e like</i>	54-bp del. in the promoter region	<i>T. durum</i> (spring line of cv. Kristall ( <i>Vrn-D4</i> ), Odessa)		Golovkina et al., 2010
	<i>Vrn-A1f</i>	50-bp del. in the promoter region	<i>T. turanicum</i> K-15993		Golovkina et al., 2010
	<i>Vrn-A1f like</i>	short deletions in promoter region + MITE element ins. in the intron 1 + wide deletion in the intron 1	Line 8.1 ( <i>T. militinae</i> $2n = 28$ ) × cv. Tähti ( $2n = 42$ )	+	Ivaničová et al., 2016
	<i>VRN-A1f-del</i>	50-bp del. in the promoter region + 2.7- kb deletion in a central part of intron 1,	<i>T. araraticum</i> PI 427403 (w)		Shcherban et al., 2016
	<i>Vrn-A1f-ins</i>	50-bp del. in the promoter region + 0.4 kb ins. of MITE element in the intron 1	<i>T. timopheevii</i>		Shcherban et al., 2016
	<i>Vrn-A1f-del/ins</i>	50-bp del. in the promoter region + 0.4 kb ins. of a MITE element + 2.7-kb del. in the intron 1	<i>T. timopheevii</i> K-29551, K-29558, PI 119442		Shcherban et al., 2016

## Продолжение приложения

Locus/ gene	Allele	Mutation	Accession	Heading date	References
	<i>Vrn-1g</i>	34-bp del. in a CArG-box motif in the promoter region	<i>T. dicoccum</i> PI 326317		Dubcovsky et al., 2006
	<i>Vrn-A1g</i>	34-bp del. in a CArG-box in the promoter region	Tetraploid line from PI 428276		Golovnina et al., 2010
	<i>Vrn-A1i</i>	Mutation of A to C (A3 instead of A5) within the VRN box	<i>T. turgidum</i> <i>T. durum</i>	+	Muterko et al., 2016
	<i>Vrn-A1k</i>	42-bp ins. in the promoter region	<i>T. dicoccum</i>		Muterko, Salina, 2017
VRN-B1	<i>Vrn-B1a</i>	127-bp ins. in the promoter region	<i>T. turanicum</i> Jakubz. K-11597		Golovnina et al., 2010
	<i>Vrn-B1c</i>	5463-bp ins. of the retrotransposon. _VRN element within the 5'UTR of <i>Vrn-B1</i> in a position near the CArG-box and VRN-box regulatory sites of VRN-A1 in di- and hexaploid wheat	<i>T. turgidum</i> subsp. <i>carthlicum</i> , PI 94749 <i>T. turgidum</i> subsp. <i>dicoccum</i>		Chu et al., 2011
	<i>Vrn-B1s</i>	2, 3 and 7 bp short deletions in the promoter region	<i>T. dicoccum</i> UA 0300212, <i>T. dicoccoides</i> PI 256029, <i>T. turgidum</i> PI 208912, <i>T. durum</i> PI 74830	+	Muterko et al., 2016
VRN-G1	<i>Vrn-G1a</i>	196-bp ins. in the promoter region	<i>T. timopheevii</i> K-38555		Golovnina et al., 2010
	<i>Vrn-G1a</i>	0.2-kb ins. MITE transposable elements in the promoter	<i>T. araraticum</i> , PI 654340 (w), <i>T. timopheevii</i> PI 119442		Shcherban et al., 2016
VRN-3	<i>Vrn-A3</i>	1(C) SNP +7 and 25 bp indel in the promoter region	<i>T. dicoccum</i> (TN 26)	+	Nishimura et al., 2018
VRN-A3 chr 7AS**	<i>Vrn-A3a-h1</i>	7-bp ins. in the promoter region of VRN-A3 locus	<i>T. dicoccum</i> (TN 26)	+	Nishimura et al., 2021
	<i>Vrn-A3</i>	1(T) SNP + 7 and 25 bp indel in the promoter region	<i>T. dicoccum</i> (TN 28)		Nishimura et al., 2018
	<i>Vrn-A3a-h2</i>	7-bp ins. that included a <i>cis</i> -element GATA box sequence at the promoter region of VRN-A3 locus	<i>T. dicoccum</i> (TN 28)		Nishimura et al., 2018
Hexaploid wheat ( $2n = 6x = 42$ )					
VRN-A1	<i>vrn-A1</i>	Native	i: Triple Dirk C		Yan et al., 2004b
	<i>Vrn-A1a</i>	222-bp or 131-bp insertion in the promoter region	i: Triple Dirk D	+	Yan et al., 2004b
		222-bp insertion in the promoter region	cv. Jupateko		Golovnina et al., 2010
	<i>Vrn-A1a.1</i>	222-bp insertion in the promoter region	<i>T. compactum</i>		Muterko et al., 2016
	<i>Vrn-A1a.2</i>	16 bp del. and 4 single nucleotide deletions within the MITE insertion when compared to <i>Vrn-A1a.1</i>	<i>T. compactum</i> PI 352302		Muterko et al., 2016
	<i>Vrn-A1b</i>	20-bp deletions in the promoter region	cv. Marquis (PI 94548), (IL 12 and IL 425), (GR 46), (IL 63 and IL 66)		Yan et al., 2004b
	<i>Vrn-A1b like</i>	(8 unique SNPs + 5 unique 1-bp indels + 5504-bp del.) in intron 1	IL 369 (Afghanistan)		Fu et al., 2005
	<i>Vrn-A1b like</i>	20-bp deletions in the promoter region	s: Saratovskaya 29/ Vietnamskaya 5R(5A)		Golovnina et al., 2010
	<i>Vrn-A1b</i>	20-bp deletions in the promoter region +177 bp ins. in intron 1	cv. Pyrothrix 28, Rescue, VL-30		Strejčková et al., 2021, 2023
	<i>Vrn-A1c like</i> <i>vrn-A1</i>	Three diagnostic SNP nucleotide polymorphisms within an 810-bp region between intron 4 and exon 8 of the <i>VRN-A1</i> gene	IL 369 (Afghanistan), IL 162 (Egypt)		Yan et al., 2004b

## Окончание приложения

Locus/ gene	Allele	Mutation	Accession	Heading date	References
	<i>Vrn-A1f-like</i>	8-bp + 50-bp del. in the promoter region	nulli5B-tetra5D Chinese Spring		Golovnina et al., 2010
	<i>Vrn-Ai2</i>	1807 and 7804-bp del. in intron 1	NILs cv. Sunstate	+	Steinfert et al., 2017
VRN-B1	<i>VrnB1a</i>	6850-bp del. in intron 1	i: Triple Dirk B		Fu et al., 2005
	<i>VrnB1b</i>	6850-bp del. in intron 1+36-bp del. in intron 1	Alpowa	+	Santra et al., 2009
	<i>VrnB1c</i>	6850-bp del. in intron 1+ 0.8-kb del. and 0.4-kb duplication in intron 1	NILs Saratovskaya 29 & Diamant II	+	Shcherban et al., 2012a
	<i>VrnB1c</i>	800-bp del. in intron 1 + duplications/translocatios around 450-bp	cvs Granny, Kaerntner Frueher, Linda, Paragon, Quarna, Septima, Sirael		Milec et al., 2012
	<i>VrnB1d</i>	(6850-bp del. +187-bp del.+ SNP (T to C), + mutation 4 bp (TTTT to ACAA)) in intron 1	cv. Hongchunmai	+	Zhang B. et al., 2018
	<i>VrnB1f</i>	836-bp ins. in intron 1	cvs Anza, Barta and Marquis	+	Strejčková et al., 2021
VRN-D1	<i>VrnD1a</i>	4235-bp del. in intron 1	i: Triple Dirk E		Fu et al., 2005
	<i>Vrn-D1b</i>	4235-bp del. in the intron 1+SNP in the CArG-box at the promoter region	<i>T. aestivum</i>	+	Zhang J. et al., 2012
	<i>VrnD1c</i>	174-bp insertion in 5'-UTR	cv. Yunong 876	+	Zhang X. et al., 2015
	<i>Vrn-D1s</i>	844-bp ins. in intron 1	<i>T. spelta</i> L., <i>T. compactum</i> Host		Muterko et al., 2015
	<i>Vrn-D1x</i>	163-bp ins. in intron 1	Mex. 3, Mex. 17	(w)	Makhoul et al., 2022
VRND1	<i>vrn-D1r</i>	17-bp del. in intron 1	cv. Robigus	(w)	Makhoul et al., 2022
VRN-2*					Kippes et al., 2016
VRN-3 VRN-B3 <i>TaFT</i> chr 7BS		5,295-bp (retrotransposon) ins. in the promoter region +6SNP+3SNP in intron 1	CS/Hope 7B		Yan et al., 2006
VRN-B3	<i>Vrn-B3a</i>	5300-bp ins. in the promoter region	cv. Hemai 26	<i>Vrn-A1</i> -null allele	Chen F. et al., 2013
	<i>Vrn-B3b</i>	5' UTR 890-bp ins. in the promoter region	cv. Chadianhong	+	Chen F. et al., 2013
	<i>Vrn-B3c</i>	(5300-bp ins. + 20-bp del. +4-bp del.) in the promoter region	cv. Ji		Chen F. et al., 2013
	<i>Vrn-B3d</i>	1617 bp ins. (retrotransposon) in the promoter region	line Velut	+	Berezhnaya et al., 2021
	<i>Vrn-B3e</i>	160 bp ins. in the promoter region	Altaiskaya, Mana2, Mariya, Obskaya 14, Omskaya 23, Novosibirskaya 29, 67, 89	+	Berezhnaya et al., 2021
VRN-A3		SNP (T–G) within the third exon	RIL Yi5029 × Nongda4332	+	Chen Z. et al., 2020
VRN-D4		3SNP in RIP-3 region of the intron 1	<i>T. sphaerococcum</i>		Kippes et al., 2015
VRN-D4			i: Triple Dirk F		Kippes et al., 2015

Note. Bp – base pairs; del. – deletion; ins. – insertion; indel – insertion/deletion; “+” – allele is associated with heading date; (w) – winter.

\*To study the *Vrn-2* gene in hexaploid wheat, a synthetic *vrn2-null* mutant was produced by combining the non-functional *Vrn-A2* allele, which are present in most polyploid wheats, with the *Vrn-B2* deletion from tetraploid wheat and the non-functional *Vrn-D2* allele from *Ae. tauschii* (Kippes et al., 2016).

\*\*To study the *Vrn-2* gene in tetraploid wheat, line #3089 (BC3F2 521 × Kronos) with non-functional alleles *vrn-A2* and *vrn-B2* was produced (Dietfeld et al., 2009).