

Exploring Indian wheat genotypes for less Celiac disease toxic epitopes

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Wheat is an important cereal having significant role in food and nutritional security of large part of human population in the world. Large numbers of end-products are made from wheat because of the presence of unique proteins which when mixed with water form visco-elastic complex called gluten. However, around 1% of world population suffers from celiac disease (CD) worldwide which varies with sex, age and location. The disease occurs in individuals with specific genetic background (with HLA-DQ2 and DQ8) which are intolerant to the gluten proteins (Gliadins and Glutenins) of not only wheat but also to the prolamins fractions of barley (hordiens), rye (secalins) and oats (avenins) (McAllister *et al.*, 2019). An autoimmune response starts when these proteins bind to the T-cells of the susceptible individuals. As a result, the villi in the small intestine get flattened and the surface area for the nutrient absorption is highly reduced and leads to malnutrition and other gastrointestinal problems. No pharmacological treatment is available to gluten-intolerant patients, and a strict, life-long gluten-free diet is the only safe and efficient treatment available. A diet is called gluten free only if the gluten content is less than 20 ppm.

The pooled global prevalence of serology confirmed CD is 1.4%, while biopsy confirmed prevalence is only 0.7%. Children are more affected (0.9%) than the adults (0.5%) (Singh *et al.*, 2018). In India, the serology confirmed CD toxicity has been reported more in northern and north-western parts as compared to southern parts in India. This may be because of higher consumption of wheat in northern parts of India. For example, the mean daily intake of wheat is 455g/day in North India and only

25g/day in Southern parts of India (Ramakrishna *et al.*, 2016). CD toxicity is mainly caused by the presence of glutamine and proline rich specific epitopes in gluten protein. *In vitro* and *in vivo* studies in rats and humans have shown that a 33-mer peptide from gliadin (LQLQPFQPOLPYQPOLPYQPOLPYQPOLPYQPQPF) is not digestible by gastric, pancreatic, and intestinal brush-border membrane endoproteases (Shan *et al.*, 2002). This and similar peptides have been identified as the main stimulators of the inflammatory response. Development of wheat varieties with low or very less immunogenicity (low amount of CD toxic epitopes) can lead to reduced toxicity load in human beings.

Wheat germplasm has wide variation and there are chances of screening out genotypes with less immunogenicity (Spaenij-Dekking *et al.*, 2005). The identified genotypes can be used in breeding in reducing CD toxic epitopes in high yielding varieties. Many biotechnological and genetic engineering techniques like genome editing by CRISPER/Cas9, downregulation of gliadin genes by RNAi have also been used to remove the toxic sequences from the wheat (Malamgoda *et al.*, 2017). But the main issue with the removal of such sequences is to maintain the baking quality of the modified wheat genotypes. Recently, Pilolli *et al.*, (2019) presented a comprehensive approach for characterization of durum wheat genotypes to identify low gluten, lower toxicity along with conserved rheological properties and yield. Lately, enzyme therapy techniques have also been used to inactivate the antigenic protein just before food preparation (Scherf *et al.*, 2018). The use of biodiversity, biotechnology and breeding may

Table 1 Wheat varieties released in different years used for the study.

Before 1960		1961-80		1981-2000		After 2000	
Variety	Year of Release	Variety	Year of Release	Variety	Year of Release	Variety	Year of Release
A -206	1954	DURGAPURA-65	1974	AJANTA	1983	COW(W)I	2006
BAXI-288-18	1952	HD-1941	1972	CPAN-1676	1984	DBW-17	2007
C-281	1955	HD-1949	1973	DL-153-2	1985	GW-322	2002
C-591	1934	HD-2177	1980	GW-40	1981	HD-2888	2006
GULAB		HD-2189	1980	HD-2329	1985	HI-1531	2006
JOB-666		HI-385	1976	HI-977	1988	HPW-251	2008
KENPHAD-25	1951	HP-1102	1980	HS-240	1989	K0307	2007
KHARCHIA L		HYB-65	1976	K-8020	1987	K-9644	2000
NI-345	1955	K-68	1974	KRL-1-4	1990	MP-4010	2003
NP-12V	1910	KALYANSONA	1969	LOK-1	1982	RAJ-4037	2004
NP-4	1905	KHARCHIA-65	1970	NARMADA112	1982	VL-804	2002
NP-52	1920s	LERMA ROJO	1969	NIAW-34	1997	KRL-210	2012
NP-721	1949-50	NI-5439	1975	PBW-343	1996	DBW-39	2010
NP-761	1949-50	NI-917	1973	PBW-65	1987	DPW-621-50	2011
NP-771		PV-18	1969	RAJ-3765	1996	MACS-6222	2010
NP-809	1954	SONALIKA	1969	UP-2121	1986	VL-892	2008
NP-830	1958-62	SONORA-64	1967	VL-616	1986	DBW-88	2014
RAJ-4125		UP-262	1978	WH-291	1985	WH-1080	2011
RIDLEY	1954	WH-147	1978	WH-542	1992	HD-2967	2011
WH-331		WL-711	1977	HUW-206	1985	HS-490	2009

help to develop grains that have a low or zero content of immunotoxic sequences, but with reasonable baking quality. ICAR-IIWBR, Karnal has initiated a study to identify wheat varieties with low CD toxicity which can be used in breeding program.

For this study, 80 wheat varieties released in India during different periods were used (Suppl Table 1). These were grown at ICAR-IIWBR farm during the crop season 2017-18. Two genotypes each of rice and maize were purchased from the local market and used as negative controls. The flours were prepared using Cyclotec mill (FOSS) with 0.5mm screen. The alcohol soluble prolamins were extracted in 60% ethanol and quantified by Bradford Method. The dilutions of the antigen, primary antibody (Pierce Gliadin peptide Antibody, 4F3, 1:4000) and secondary antibody (Pierce Goat Anti-Mouse IgG, peroxidase conjugated, 1:10000) were standardized using the standard checker board method. Equal amount of extracted protein of each sample was subjected to

the standardized Indirect ELISA protocol as used by Gregorini *et al.* (2009). The ELISA results were adjusted with the total prolamins content of each sample and the relative reactivity against the antibody was calculated. ANOVA and Tukey's comparison test were performed using XLSTAT. The ELISA results were further confirmed by western blotting using same antibody.

Monoclonal antibodies have been used in many studies to show the presence of CD toxic sequences in protein fractions of cereals like wheat, barley and oats (Spaenij-Dekking *et al.*, 2005; Gregorini *et al.*, 2009; Ribeiro *et al.*, 2016). In this study monoclonal antibody against the -gliadin peptides p58-73 (KLQPFPPQPELPYPQPQ) containing a core region reported for CD toxicity was used for the indirect ELISA for screening the wheat genotypes for the presence of these toxic sequences. The presence of CD toxic epitopes was also confirmed by western blotting.

Around 3-fold variability (2.1-6.0) was observed for this specific (Glia - 2) CD toxic epitope in wheat varieties across

the periods (Table 2). The results showed that most of the varieties developed before 1960 had significantly low antibody reactivity (3.25). However, after 1960 varietal differences were observed but overall no significant change observed in antibody reactivity in the wheat varieties released in last 60 years in India. Some wheat varieties with low reactivity have been identified across the periods (JOB-666, BAXI-288-18, NI-345, NP 830, C 591, Khenpad 25, HD 2329, HI 977, HS 240, K 9644). But further confirmation is required in the second year and also with antibodies against different epitopes. Some other studies have also reported that the ancient wheat

Table 2 Relative antibody reactivity against CD toxic epitopes in wheat varieties released during different periods.

Period	Before 1960	1961-1980	1981-2000	After 2000
Range	2.1 - 4.9	3.1- 5.2	2.8 - 6.0	2.5- 5.8
Mean*	3.25b	4.10a	3.98ab	4.03a

*Mean of 20 wheat varieties. Means with the same letter in the same row are not significantly different (P<0.05)

varieties have less number of CD toxic epitopes and these might be better to be used in the diets of people suffering from celiac disease (van den Broeck *et al.*, 2010). Contrasting studies suggesting that the modern wheat varieties instead of ancient varieties have fewer CD toxic epitopes (Colombo and Gregorini, 2012; Prandi *et al.*, 2017) have also been reported. While Malalgoda *et al.* (2018) reported that there is no association between the cultivar release year and the amount of immunogenic epitopes and -gliadin. Ribeiro *et al.* (2016) have predicted that breeding has not contributed to the prevalence of CD immune stimulatory epitopes. Mohan Kumar *et al.* (2017) reported that the tetraploid wheat varieties are less immune reactive as compared to the hexaploids. Shewry (2018) reviewed the studies on CD and found that wide range of variation exists in reactions among genotypes of all species. However, it was pointed that these studies used only one to four antibodies which target only a small proportion of the total number of celiac epitopes in gluten proteins.

The presence of CD epitopes was also confirmed by the western blotting. Representative samples with low and high relative reactivity based on ELISA result were selected. Equal quantity of these samples were used for SDS-PAGE and further detected with the 4F3 antibody.

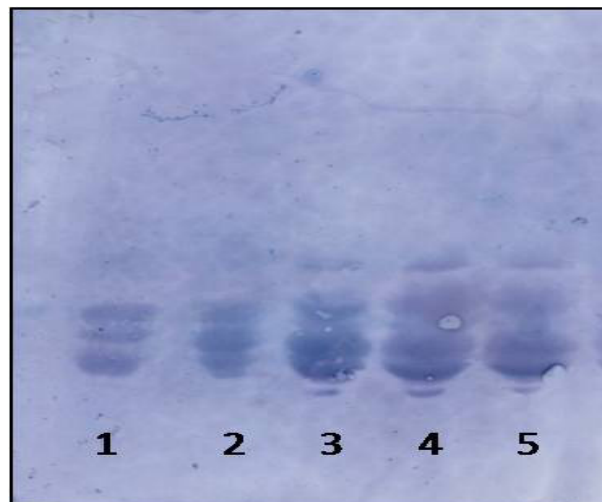


Fig 1. Immunoblotting of gliadin extracts with 4F3 monoclonal antibody. 1 and 2 – low antibody reactivity; 3, 4 and 5 – high antibody reactivity.

The intensity of bands clearly showed the relative reactivity of the wheat genotypes (Fig 1). Other studies have also reported such confirmation of ELISA results by western blotting (Gregorini *et al.*, 2009; Spaenij-Dekking *et al.*, 2005).

The conflicting results on the immunogenicity of ancient and modern wheat varieties highlights the need to screen the large wheat germplasm available including different wheat species, wild relatives and synthetics. The identified genotypes with low CD toxicity can be used in breeding programmes for reducing CD toxic epitopes in high yielding varieties. This study is a very small initiative where only one monoclonal antibody against few epitopes has been used. More monoclonal antibodies can be used in future studies to target more CD toxic epitopes. The most important challenge while developing a new genotype with low CD toxicity either by breeding or genetic engineering will be to keep the baking and processing quality of the wheat intact. This will bring less CD toxic wheat products in food chain with consequences of reduced celiac disease.

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