

Summary & PowerPoint

Addressing Emerging Concerns Related to Safety of Wheat-Based Products

Summary to follow.

Learning Objectives

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Presentation Time

Monday, February 25, 2019 2:10 pm - 2:45 pm

Session Breakout 3





Indicators and Pathogens in Wheat Grain

Prevalence of indicators and pathogenic microorganisms in wheat grains in different geographic regions

	Geographic Origin ¹								
Microorganisms	Australia	Australia	Australia	Great Britain	United States	Hungary	Algeria		
Escherichia coli	NR	NR	2 (50)	NR	NR	NR	NR		
Coliforms	ND	93 (58)	NR	NR	NR	NR	NR		
Salmonella spp.	NR	0.5 (412)	2 (50)	NR	NR	NR	NR		
Bacillus cereus	NR	81.0 (58)	4 (50)	NR	NR	NR	NR		
Reference	Eyles et al (1989)	Berghofer et al (2003)	Eglezos (2010)	Seiler (1986)	Manthey et al (2004)	Peles et al (2012)	Riba et al (2008)		

¹ Values are percentages of positive samples. Sample sizes are provided in parentheses. NR = not reported; ND = not detected.

Microbial load of wheat milled products:							
Wheat milled	Average Microbial Counts (log CFU/g) ×						
products	<u>APC y</u>	Coliform	Generic E. coli	Eb ^y	Yeasts	Molds	
Incoming wheat	5.5 ± 0.2 a	1.0 ± 0.9 b	< 1.0 ± 0.0 ^z	3.2 ± 1.2 a	2.7 ± 0.9 a	2.9 ± 0.6 a	
Dry clean wheat	5.3 ± 0.3 a	1.8 ± 1.7 ab	< 1.0 ± 0.0 ^z	3.9 ± 1.1 a	2.5 ± 1.1 a	2.8 ± 0.6 a	
Tempered wheat	5.4 ± 0.4 a	1.3 ± 1.2 b	< 1.0 ± 0.0 ^z	3.7 ± 0.8 a	3.3 ± 1.2 a	2.6 ± 0.4 a	
1 st Break	4.9 ± 0.7 ab	1.9 ± 1.7 ab	< 1.0 ± 0.0 ^z	3.4 ± 0.7 a	1.8 ± 1.5 a	3.1 ± 0.5 a	
3 rd Break	4.5 ± 0.7 ab	2.3 ± 0.4 ab	< 1.0 ± 0.0 ^z	3.4 ± 1.1 a	2.1 ± 0.6 a	2.5 ± 0.3 a	
5 th Break	4.8 ± 0.8 ab	3.0 ± 0.1 ab	< 1.0 ± 0.0 ^z	3.6 ± 0.4 a	2.9 ± 1.4 a	2.9 ± 0.4 a	
1 st Middlings	3.5 ± 0.4 b	2.4 ± 0.3 ab	< 1.0 ± 0.0 ^z	2.7 ± 0.5 a	1.8 ± 0.6 a	2.5 ± 0.1 a	
3 rd Middlings	3.9 ± 0.3 ab	2.3 ± 0.2 ab	< 1.0 ± 0.0 ^z	2.3 ± 0.5 a	1.6 ± 1.4 a	2.6 ± 0.1 a	
5 th Middlings	4.9 ± 1.3 ab	2.8 ± 0.5 ab	< 1.0 ± 0.0 ^z	3.2 ± 0.2 a	2.2 ± 0.9 a	2.9 ± 0.9 a	
Straight-grade flour	3.9 ± 0.5 ab	2.1 ± 0.5 ab	< 1.0 ± 0.0 ^z	2.2 ± 0.7 a	1.6 ± 0.5 a	3.0 ± 0.6 a	
Bran	5.3 ± 0.2 a	3.2 ± 0.5 ab	< 1.0 ± 0.0 ^z	3.4 ± 1.0 a	2.9 ± 1.1 a	2.6 ± 0.6 a	
Shorts	5.1 ± 0.4 ab	3.0 ± 0.5 ab	< 1.0 ± 0.0 ^z	3.6 ± 0.8 a	2.7 ± 0.6 a	2.8 ± 0.4 a	
Germ	5.1 ± 0.5 ab	3.7 ± 1.0 a	$< 1.0 \pm 0.0^{2}$	4.0 ± 1.1 a	2.6 ± 1.0 a	3.0 ± 1.1 a	

Indicators and Pathogens in Wheat Flour

Prevalence of indicators and pathogenic microorganisms in wheat flour in different									
geographic regions									
	Geographic Origin ¹								
Microorganisms	Turkey	Australia	Australia	United States	United States	Pakistan			
Escherichia coli	50.7 (142)	1.4 (71)	0.7 (300)	12.8 (3350)	0.7 (2921)	NR			
Coliforms ²	NR	1.0 - 1000 (71)	NR	1.2 (1477)	1.6 (3,688)	3.0 - 4.0 (150)			
Salmonella spp.	NR	NR	ND (150)	1.3 (3040)	0.14 (4358)	NR			
Bacillus cereus	4.2 (142)	93 (71)	ND (350)	NR	NR	NR			
C. perfringens	9.9 (142)	NR	NR	NR	NR	NR			
Reference	Aydin et al (2009)	Berghofer et al (2003)	Eglezos (2010)	Richter et al (1993)	Sperber (2007)	Batool et al (2012)			

¹ Values are percentages of positive samples. Sample sizes are provided in parentheses.

² Expressed as most probable number per gram (MPN/g). NR = not reported

Microbial Load on Equipment Surface

Microbial load associated with selected equipment used to process the wheat and handle the resulting flour.

	Average Microbial Counts (log CFU/10 cm ²) ^x							
Sample Location	APC ^y	Coliform	Generic <i>E.</i> coli	Eb ^y	Yeasts	Molds		
Cleaning and tempering equipmen								
Water addition system (Turbolizer)	7.0 ± 1.6	4.3 ± 1.2	$< 1.0 \pm 0.0$ ^z	4.8 ± 0.7	3.9 ± 1.8	3.3 ± 0.5		
Milling and sifting equipment								
Inside walls of grinding machines	4.3 ± 0.7	1.7 ± 0.5	$< 1.0 \pm 0.0$ ^z	2.1 ± 0.4	1.4 ± 1.1	3.3 ± 0.7		
Walls of sifter boxes	3.3 ± 1.0	0.7 ± 0.9	$< 1.0 \pm 0.0$ ^z	1.5 ± 1.4	0.8 ± 1.0	2.2 ± 0.8		
Flour handling and storage equipment								
Twin screw conveyors	3.7 ± 0.1	1.5 ± 0.5	$< 1.0 \pm 0.0$ ^z	2.1 ± 0.3	2.1 ± 0.2	2.9 ± 0.1		
Product scales	5.3 ± 1.0	1.6 ± 0.6	$< 1.0 \pm 0.0$ ^z	1.4 ± 1.2	1.4 ± 1.2	4.7 ± 1.1		
× Values represents average ± standard deviation. ^y APC, aer [∞] bic plate count; Eb, <i>Enterobacteriaceae.</i>								

 z Values are below the limit of detection of 1.0 log CFU/10 cm².

