Australia New Zealand Food Standards Code – Amendment No. 94 – 2007

Food Standards Australia New Zealand Act 1991

Preamble

The variations set forth in the Schedule below are variations to Standards in the *Australia New Zealand Food Standards Code* published by the National Health and Medical Research Council in the *Commonwealth of Australia Gazette*, No. P 27, on 27 August 1987, which have been varied from time to time.

These variations are published pursuant to section 23A of the *Food Standards Australia New Zealand Act 1991*.

Citation

These variations may be collectively known as the *Australia New Zealand Food Standards Code* – Amendment No. 94 – 2007.

Commencement

These variations commence on 11 October 2007.

Note: These variations were published in the Commonwealth of Australia *Food Standards Gazette* No. FSC 36 on 11 October 2007.

SCHEDULE

- [1] **Standard 1.3.3** is varied by –
- [1.1] *omitting from clause 1 the definition of* EC [number], *substituting* –

EC number (Enzyme Commission number) means the number which the Enzyme Commission uses to classify the principal enzyme activity.

[1.2] omitting from the Table to clause 14, cupric citrate on a bentonite base, substituting –

Cupric citrate	Removal of sulphide compounds	GMP
	from wine	

[1.3] *omitting the* Table to clause 15, *substituting* –

Table to clause 15

Enzyme	Source
Lipase, triacylglycerol EC 3.1.1.3	Bovine stomach; salivary glands or forestomach of calf, kid or lamb; porcine or bovine pancreas
Pepsin EC 3.4.23.1	Bovine or porcine stomach
Phospholipase A ₂ EC 3.1.1.4	Porcine pancreas

Thrombin	Bovine or porcine blood
EC 3.4.21.5	
Trypsin	Porcine or bovine pancreas
EC 3.4.21.4	

[1.4] omitting the Table to clause 16, substituting –

Table to clause 16

Enzyme	Source
α–Amylase EC 3.2.1.1	Malted cereals
β-Amylase EC 3.2.1.2	Sweet potato (<i>Ipomoea batatas</i>) Malted cereals
Actinidin EC 3.4.22.14	Kiwifruit (Actinidia deliciosa)
Bromelain EC 3.4.22.4	Pineapple stem (Ananas comosus)
Ficin EC 3.4.22.3	Ficus spp.
Papain EC 3.4.22.2	Carica papaya

[1.5] omitting the Table to clause 17, substituting –

Table to clause 17

Enzyme	Source
α–Acetolactate decarboxylase	Bacillus amyloliquefaciens
EC 4.1.1.5	Bacillus subtilis
	Bacillus subtilis, containing the gene for α–Acetolactate
	decarboxylase isolated from Bacillus brevis
Aminopeptidase	Aspergillus oryzae
EC 3.4.11.1	Lactocococcus lactis
α–Amylase	Aspergillus niger
EC 3.2.1.1	Aspergillus oryzae
	Bacillus amyloliquefaciens
	Bacillus licheniformis
	Bacillus licheniformis, containing the gene for α -Amylase
	isolated from Geobacillus stearothermophilus
	Bacillus subtilis
	Bacillus subtilis, containing the gene for α -Amylase isolated
	from Geobacillus stearothermophilus
	Geobacillus stearothermophilus
β-Amylase	Bacillus amyloliquefaciens
EC 3.2.1.2	Bacillus subtilis
α–Arabinofuranosidase	Aspergillus niger
EC 3.2.1.55	
Carboxyl proteinase	Aspergillus melleus
EC 3.4.23.6	Aspergillus niger
	Aspergillus oryzae
	Rhizomucor miehei
Carboxylesterase	Rhizomucor miehei
EC 3.1.1.1	
Catalase	Aspergillus niger
EC 1.11.1.6	Microccocus luteus

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Invertase	Saccharomyces cerevisiae
EC 3.2.1.26	buccharoniyees cerevisiae
Lipase, monoacylglycerol EC 3.1.1.23	Penicillium camembertii
Lipase, triacylglycerol EC 3.1.1.3	Aspergillus oryzae Aspergillus oryzae, containing the gene for Lipase, triacylglycerol isolated from Fusarium oxysporum Aspergillus oryzae, containing the gene for Lipase, triacylglycerol isolated from Humicola lanuginosa Aspergillus oryzae, containing the gene for Lipase, triacylglycerol isolated from Rhizomucor miehei Candida rugosa Hansenula polymorpha, containing the gene for Lipase, triacylglycerol isolated from Fusarium heterosporum Mucor javanicus Penicillium roquefortii Rhizopus arrhizus Rhizomucor miehei Rhizophus niveus Rhizophus oryzae
Lysophospholipase EC 3.1.1.5	Aspergillus niger
Maltogenic α-amylase EC 3.2.1.133	Bacillus subtilis containing the gene for maltogenic α-amylase isolated from Geobacillus stearothermophilus
Metalloproteinase	Aspergillus oryzae Bacillus amyloliquefaciens Bacillus coagulans Bacillus subtilis
Mucorpepsin EC 3.4.23.23	Aspergillus oryzae Aspergillus oryzae, containing the gene for Aspartic proteinase isolated from Rhizomucor meihei Rhizomucor meihei Cryphonectria parasitica
Pectin lyase EC 4.2.2.10	Aspergillus niger
Pectinesterase EC 3.1.1.11	Aspergillus niger Aspergillus oryzae, containing the gene for pectinesterase isolated from Aspergillus aculeatus
Phospholipase A ₁ EC 3.1.1.32	Aspergillus oryzae, containing the gene for phospholipase A_1 isolated from Fusarium venenatum
Phospholipase A ₂ EC 3.1.1.4	Streptomyces violaceoruber
3-Phytase EC 3.1.3.8	Aspergillus niger
4-Phytase EC 3.1.3.26	Aspergillus oryzae, containing the gene for 4-phytase isolated from Peniophora lycii
Polygalacturonase or Pectinase multicomponent enzyme EC 3.2.1.15	Aspergillus niger Aspergillus oryzae Trichoderma reesei
Pullulanase EC 3.2.1.41	Bacillus acidopullulyticus Bacillus amyloliquefaciens Bacillus licheniformis Bacillus subtilis Klebsiella pneumoniae
Serine proteinase EC 3.4.21.14	Aspergillus oryzae Bacillus amyloliquifaciens Bacillus halodurans Bacillus licheniformis Bacillus subtilis

Transglucosidase EC 2.4.1.24	Aspergillus niger
Transglutaminase EC 2.3.2.13	Streptomyces mobaraensis
Urease EC 3.5.1.5	Lactobacillus fermentum
Xylose isomerase EC 5.3.1.5	Actinoplanes missouriensis Bacillus coagulans Microbacterium arborescens Streptomyces olivaceus Streptomyces olivochromogenes Streptomyces murinus Streptomyces rubiginosus

[1.6] *omitting the* Editorial note, *immediately following the* Table to clause 17, *substituting* –

Editorial note:

Bacillus amyloliquefaciens is a separate species from Bacillus subtilis.

Aspergillus niger group covers strains known under the names Aspergillus aculeatus, A. awamori, A. ficuum, A. foetidus, A. japonicus, A. phoenicis, A. saitor and A. usamii.

Trichoderma reesei also known as Trichoderma longibrachiatum.

Kluyveromyces marxianus – former names Saccharomyces fragilis and Kluyveromyces fragilis.

Kluyveromyces lactis – former name Saccharomyces lactis.

Rhizomucor miehei – former name Mucor miehei.

Microccocus luteus – former name Micrococcus lysodeikticus.

Paenibacillus macerans – former name Bacillus macerans.

Talaromyces emersonii – former name Penicillium emersonii.

Klebsiella pneumoniae – former name *Klebsiella aerogenes*.

Streptomyces mobaraensis – former name *Streptoverticillium mobaraensis*.

Humicola lanuginosa also known as Thermomyces lanuginosus.

Mucor javanicus also known as *Mucor circinelloides* f. *circinelloides*.

Penicillium roquefortii also known as Penicillium roqueforti.

Hansenula polymorpha also known as Pichia angusta.

Geobacillus stearothermophilus – former name Bacillus stearothermophilus.

4-Phytase also known as 6-phytase.

[2] *Standard 1.4.2* is varied by –

[2.1] *omitting from* Schedule 1 *all entries for the following chemical* –

Coumaphos

[2.2] inserting in Schedule 1 –

AZIMSULFURON	
AZIMSULFURON	
EDIBLE OFFAL (MAMMALIAN)	*0.02
EGGS	*0.02
MEAT (MAMMALIAN)	*0.02
MILKS	*0.02
POULTRY, EDIBLE OFFAL OF	*0.02

POULTRY MEAT RICE	*0.02 *0.02
PROHEXADIONE-CAL	CIUM
SUM OF THE FREE AND CONJUGA	ATED FORMS OF
PROHEXADIONE EXPRESSED AS I	PROHEXADIONE
APPLE	*0.02
EDIBLE OFFAL (MAMMALIAN)	*0.05
MEAT (MAMMALIAN)	*0.05
MILKS	*0.01

[2.3] omitting from Schedule 1 the chemical residue definition for the chemical appearing in Column 1 of the Table to this sub-item, substituting the chemical residue definition appearing in Column 2-

COLUMN 1	COLUMN 2
THIABENDAZOLE	COMMODITIES OF PLANT ORIGIN:
	THIABENDAZOLE
	COMMODITIES OF ANIMAL ORIGIN: SUM OF
	THIABENDAZOLE AND 5-
	HYDROXYTHIABENDAZOLE, EXPRESSED AS
	THIABENDAZOLE

[2.4] omitting from Schedule 1 the foods and associated MRLs for each of the following chemicals –

INDOXACARB	
INDOXACARB	
WINE GRAPES	1
PACLOBUTRAZOL	
PACLOBUTRAZOL	
ASSORTED TROPICAL AND SUB-	*0.01
TROPICAL FRUITS – INEDIBLE	
PEEL [EXCEPT AVOCADO]	
PROCYMIDONE	
PROCYMIDONE	
BEANS [EXCEPT GREEN BEANS]	T10
TETRACHLORVINPHOS	
TETRACHLORVINPHOS	
LEAFY VEGETABLES	2

[2.5] inserting in alphabetical order in Schedule 1, the foods and associated MRLs for each of the following chemicals –

AZOXYSTROBIN	
AZOXYSTROBIN	
BROCCOLI	T0.5
BRUSSELS SPROUTS	T0.5
CAULIFLOWER	T0.5
EGGS	*0.01
POULTRY, EDIBLE OFFAL OF	*0.01
POULTRY MEAT	*0.01

	BIFENTHRIN BIFENTHRIN	
TARO		T*0.05

CHLOROTHALONIL	
COMMODITIES OF PLANT ORIGIN:	
CHLOROTHALONIL	
COMMODITIES OF ANIMAL ORIGIN: SUN	1 OF
CHLOROTHALONIL AND 4-HYDROXY-2,	
TRICHLOROISOPHTHALONITRILE METAB	,
EXPRESSED AS CHLOROTHALONIL	
ASPARAGUS	T*0.1
ASPARAGUS	1 0.1
CYPERMETHRIN	
CYPERMETHRIN, SUM OF ISOMERS	
LEEK	T0.5
SHALLOT	T0.5
SPRING ONION	T0.5
DIFENOCONAZOLE	
DIFENOCONAZOLE	
BEETROOT	T0.2
ETHEPHON	
ETHEPHON	
MANGO	T10
OLIVES	T5
WHEAT	T1
ETOXAZOLE	
ETOXAZOLE	
GRAPES	T0.5
PEAR	T0.2
STONE FRUITS	T0.5
INDOXACARB	
INDOXACARB	
DRIED GRAPES	2
GRAPES	0.5
LEAFY VEGETABLES [EXCEPT	5
LETTUCE, HEAD]	
LINSEED	T0.5
SAFFLOWER SEED	T0.5
STRAWBERRY	T1
MCPA	
MCPA	
RHUBARB	*0.02

MERCHANICA	
METHOMYL	
SUM OF METHOMYL AND METH	
HYDROXYTHIOACETIMIDATE ('MET	
OXIME'), EXPRESSED AS METHO	MYL
SEE ALSO THIODICARB	
TARO	T1
PACLOBUTRAZOL	
PACLOBUTRAZOL	
ASSORTED TROPICAL AND SUB-	*0.01
TROPICAL FRUITS – INEDIBLE	
PEEL [EXCEPT AVOCADO AND	
MANGO]	
MANGO	T1
PROCYMIDONE	
PROCYMIDONE	
BROAD BEAN (DRY)	T10
BROAD BEAN (GREEN PODS	T10
AND IMMATURE SEEDS)	
COMMON BEAN (DRY)	T10
COMMON BEAN (PODS AND/OR	T3
IMMATURE SEEDS)	13
IIVIIVII (TOKE SEEDS)	
PROPICONAZOLE	
PROPICONAZOLE	
ASPARAGUS	T*0.1
	1 0.1
PYMETROZINE	
PYMETROZINE	
PODDED PEA (YOUNG PODS)	0.3
(SNOW AND SUGAR SNAP)	0.5
(SNOW THE SOUTH SIVIL)	
TEBUCONAZOLE	
TEBUCONAZOLE	
ASPARAGUS	T*0.02
710171101000	1 0.02
THIABENDAZOLE	
COMMODITIES OF PLANT ORIGI	'M'
THIABENDAZOLE	11.
	SUM OF
COMMODITIES OF ANIMAL ORIGIN: SUM OF	
THIABENDAZOLE AND 5-	
HYDROXYTHIABENDAZOLE, EXPRES	SED AS
THIABENDAZOLE SWEET DOTATO	T0.05
SWEET POTATO	T0.05

 $[1.6] \quad \textit{omitting from Schedule 1, under the entries for the following chemicals, the maximum residue limit for the food, substituting -$

AZOXYSTROBIN	
AZOXYSTROBIN	
PEANUT	0.05
PEANUT OIL, CRUDE	0.1
ETHEPHON	
ETHEPHON	
BARLEY	1

GLUFOSINATE AND GLUFOSINATE-	
AMMONIUM	
SUM OF GLUFOSINATE-AMMONIUM, N-ACETYL	
GLUFOSINATE AND 3-[HYDROXY(METHYL)-	
PHOSPHINOYL] PROPIONIC ACID, EXPRESSED	
AS GLUFOSINATE (FREE ACID)	
COTTON SEED 3	

GLYPHOSATE	
SUM OF GLYPHOSATE AND	
AMINOMETHYLPHOSPHONIC ACID (AM	PA)
METABOLITE, EXPRESSED AS GLYPHOS	ATE
COTTON SEED	15
IMIDACLOPRID	
SUM OF IMIDACLOPRID AND METABOLITES	
CONTAINING THE 6-	
CHLOROPYRIDINYLMETHYLENE MOIE	TY,
EXPRESSED AS IMIDACLOPRID	
BANANA	0.5
CITRUS FRUITS	2
QUINOXYFEN	
QUINOXYFEN	
DRIED GRAPES	2
GRAPES	0.6

THIAMETHOXAM	
COMMODITIES OF PLANT ORIGIN:	
THIAMETHOXAM	
COMMODITIES OF ANIMAL ORIGIN: SUN	A OF
THIAMETHOXAM AND N-(2-CHLORO-THIAZOL-	
5-YLMETHYL)-N'-METHYL-N'-NITRO-	
GUANIDINE, EXPRESSED AS THIAMETHO	OXAM
SUNFLOWER SEED	*0.02
TRIFLOXYSULFURON SODIUM	
TRIFLOXYSULFURON	
SUGAR CANE	*0.01
UNICONAZOLE-P	
SUM OF UNICONAZOLE-P AND ITS Z-ISC	MER
EXPRESSED AS UNICONAZOLE-P	
AVOCADO	0.5

[3] Standard 2.2.1 is varied by omitting from clause 1 the definition of meat pie, substituting –

meat pie means a pie containing no less than 250 g/kg of meat flesh.

[4] Standard 4.5.1 is varied by omitting from the Table to clause 4, cupric citrate on a bentonite base, substituting –

Cupric citrate	
Cupite citiate	