

# Differing criteria for setting crop group MRLs leads to non- harmonized MRLs - Current challenges and opportunities

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# What's the problem?

## **Crop Grouping**

is a well accepted and cost effective approach that facilitates the establishment of pesticide tolerances for both major and minor crops.



**Non-  
Harmonized  
MRLs**

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# Step 1 – Domestic Registration



## Fungi-Blaster®

*For effective treatment of Shot Hole disease in fruit trees*

	Cherry	6 field trials	Apricot; apricot, Japanese; capulin; cherry, black; cherry, Nanking; cherry, sweet; cherry, tart; Jujube, Chinese; nectarine; peach; plum; plum, American; plum, beach; plum, Canada; plum, cherry; plum, Chickasaw; plum, Damson; plum, Japanese; plum, Klamath; plum, prune; plumcot; sloe; cultivars, varieties, and/or hybrids of these
	Peach	9 field trials	
	Plum	6 field trials	

# Step 1 – Domestic Registration



Fungi-Res  
Cherry  
US  
3x100g/ha 0 PHI

Total number of data (n)	6
Percentage of censored data	0%
Number of non-censored data	6
Lowest residue	1.200
Highest residue	2.900
Median residue	2.550
Mean	2.283
Standard deviation (SD)	0.624
Correction factor for censoring (CF)	1.000

Proposed MRL estimate

- Highest residue	2.900
- Mean + 4 SD	4.780
- CF x 3 Mean	6.850
Unrounded MRL	<u>6.850</u>

Rounded MRL

7

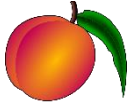
High uncertainty of MRL estimate.  
[Small dataset]

Residues (mg/kg)	n
1.2	1
1.9	1
2.5	1
2.6	2
2.9	1

**US  
MRL**

Cherry	7
Peach	
Plum	
<b>Stone Fruit</b>	

# Step 1 – Domestic Registration



Fungi-Res  
Peach  
US  
3x100g/ha 0 PHI

Total number of data (n)	9
Percentage of censored data	0%
Number of non-censored data	9
Lowest residue	0.500
Highest residue	2.700
Median residue	1.100
Mean	1.556
Standard deviation (SD)	0.872
Correction factor for censoring (CF)	1.000

Proposed MRL estimate

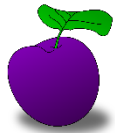
- Highest residue	2.700
- Mean + 4 SD	5.043
- CF x 3 Mean	4.667
Unrounded MRL	<u>5.043</u>
Rounded MRL	<u>5</u>

Residues (mg/kg)	n
0.5	1
0.8	1
0.9	2
1.1	1
2.2	1
2.4	1
2.5	1
2.7	1

**US  
MRL**

Cherry	7
Peach	5
Plum	
<b>Stone Fruit</b>	

# Step 1 – Domestic Registration



Fungi-Res  
Peach  
US  
3x100g/ha 0 PHI

Total number of data (n)	6
Percentage of censored data	0%
Number of non-censored data	6
Lowest residue	0.800
Highest residue	2.200
Median residue	1.300
Mean	1.333
Standard deviation (SD)	0.516
Correction factor for censoring (CF)	1.000

Proposed MRL estimate

- Highest residue	2.200
- Mean + 4 SD	3.399
- CF x 3 Mean	4.000
Unrounded MRL	<u>4.000</u>

Rounded MRL 4

High uncertainty of MRL estimate.  
[Small dataset]

Residues (mg/kg)	n
0.8	1
0.9	1
1.1	1
1.5	2
2.2	1

**US  
MRL**

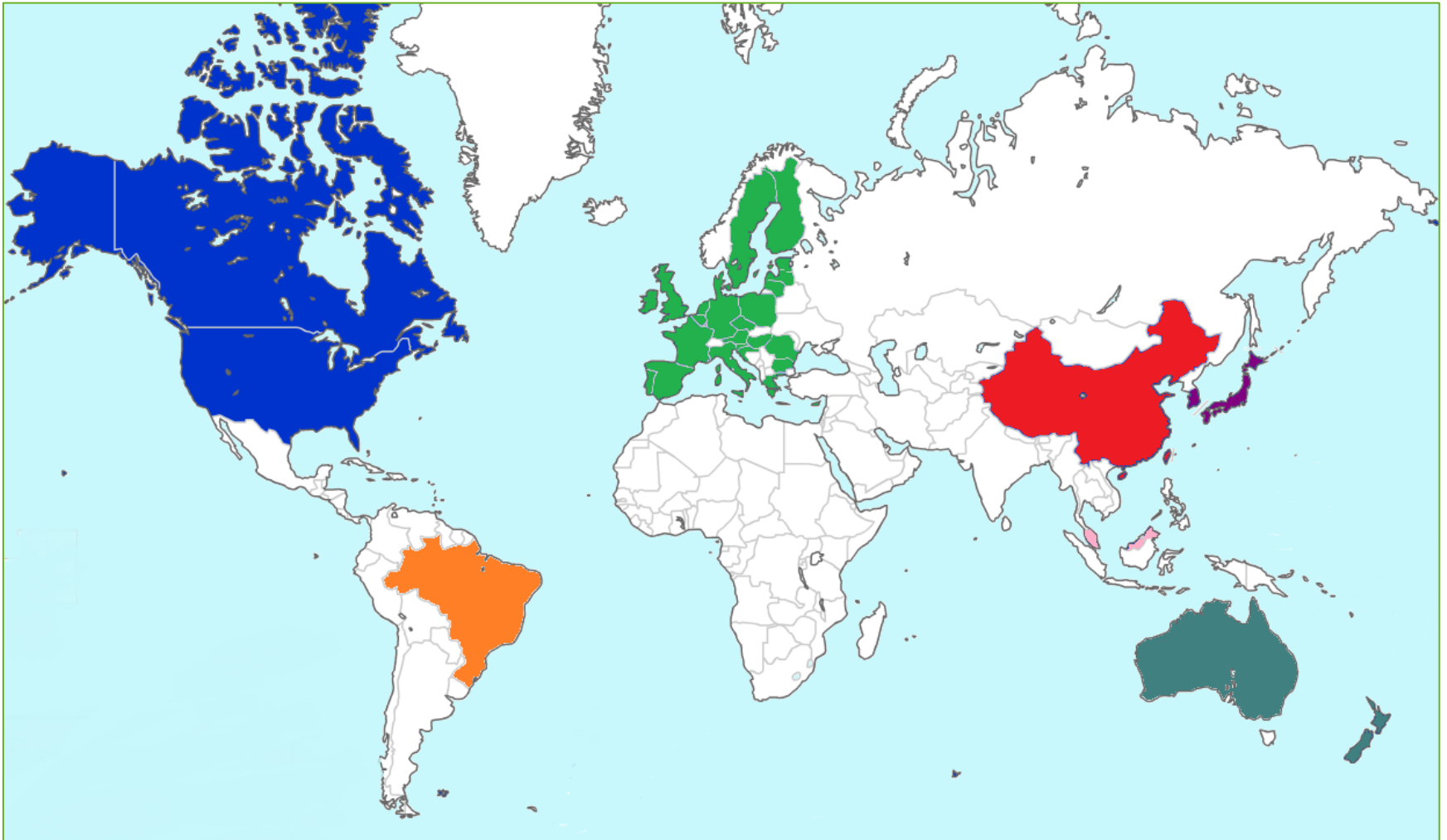
Cherry	7
Peach	5
Plum	4
<b>Stone Fruit</b>	<b>7</b>

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## Step 2 – Import Tolerances

### Crop Grouping Systems around the World



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# Crop Grouping – EU and US

broccoli or  
cauliflower

cabbage

mustard  
greens

broccoli and  
cauliflower

head cabbage

kale



## Brassica Vegetables





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# Crop Grouping – EU and US

tomato    pepper

tomato    pepper  
cucumber  
and melon    sweet corn



## Fruiting Vegetables



# EU: Guidelines on comparability, extrapolation, group tolerances and data requirements for setting MRLs

			nut”with the exception of coconuts (4 trials)	→	“Closed nuts”
(iii) Pome fruit	Apples Pears		Apple or pears (with a minimum of 4 annual trials)	→	Whole group
(iv) Stone fruit	Apricots Peaches (including nectarines and similar hybrids)		Peaches or apricots <b>(with a minimum of 4 trials on apricot)</b>	→	Nectarines, apricots, peaches
	Cherries		Sweet cherries	↔	Sour cherries
	Plums				
(v) Berries and small fruit					
(a) Table and wine grapes	Table grapes Wine grapes		Table grapes	↔	Wine grapes




**Major crops  
8 trials (each)**

# Challenge: different trial requirements in each country

*...could lead to denial of MRLs*

Fungi-Res	
Fungi-Res	
Fungi-Res Peach US 3x100g/ha 0 PHI	
Total number of data (n)	6
Percentage of censored data	0%
Number of non-censored data	6
Lowest residue	0.800
Highest residue	2.200
Median residue	1.300
Mean	1.333
Standard deviation (SD)	0.516
Correction factor for censoring (CF)	1.000
<u>Proposed MRL estimate</u>	
- Highest residue	2.200
- Mean + 4 SD	3.399
- CF x 3 Mean	4.000
Unrounded MRL	4.000
Rounded MRL	4
High uncertainty of MRL estimate. [Small dataset]	

	EU MRL
Cherry	7
Peach	5
Plum	4
<b>Stone Fruit</b>	

	Cherry	2+6 field trials
	Peach	9 field trials
	Plum	2+6 field trials




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# Challenge: different crop group extrapolations in different countries

## Fungi-Res MRLs

*...leads to different MRLs*

		<b>US MRL</b>	<b>EU MRL</b>
	Cherry	7	7
	Peach	7	5
	Plum	7	4

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# Step 3 – Codex MRLs

**Group 003 Stone fruits**

<u>Code No.</u>	<u>Commodity</u>
FS 0012	<b>Stone fruits</b> <i>Prunus</i> spp.
FS 0013	<b>Cherries</b> <i>Prunus cerasus</i> L.; <i>P. avium</i> L.
FS 0014	<b>Plums (including Prunes)</b> <i>Prunus domestica</i> L.; other <i>Prunus</i> spp and ssp.
FS 0240	<b>Apricot</b> <i>Prunus armeniaca</i> L.; syn: <i>Armeniaca vulgaris</i> Lamarck
FS 0241	<b>Bullace</b> <i>Prunus insititia</i> L.; syn: <i>Prunus domestica</i> L., ssp. <i>insititia</i> (L.) Schneider
FS 0242	<b>Cherry plum</b> <i>Prunus cerasifera</i> Ehrhart, syn: <i>P. divaricata</i> Ledebouer <i>P. salicina</i> Lindl., var. Burbank
FS -	<b>Chickasaw plum</b> , see Plum, Chickasaw
FS 0243	<b>Cherry, Sour</b> <i>Prunus cerasus</i> L.
FS 0244	<b>Cherry, Sweet</b> <i>Prunus avium</i> L.
FS -	<b>Damsons (Damson plums)</b> , see Plum, Damson
FS -	<b>Greengages (Greengage plums)</b> , see Plum, Greengage
FS 0250	<b>Japanese apricot</b> <i>Prunus mume</i> Siebold & Zucc.
FS 0245	<b>Nectarine</b> <i>Prunus persica</i> (L.) Batch, var. <i>nectarina</i>
FS -	<b>Mirabelle</b> , see Plum, Mirabelle
FS 0246	<b>Morello</b> <i>Prunus cerasus</i> L., var. <i>austera</i> L.



## Fungi-Res Residues

Residues (mg/kg)	n
1.2	1
1.9	1
2.5	1
2.6	2
2.9	1



Residues (mg/kg)	n
0.5	1
0.8	1
0.9	2
1.1	1
2.2	1
2.4	1
2.5	1
2.7	1



Residues (mg/kg)	n
0.8	1
0.9	1
1.1	1
1.5	2
2.2	1

# JMPR often uses statistical tests in their review process

	A	B	C	D	E	F	G	H
1	<b><u>Kruskal-Wallis Test</u></b>							
2								
3								
4								
5	Cherry	Peach	Plum	Group 4	Group 5	Group 6	Group 7	Group 8
6	1.2	1.1	0.8					
7	2.5	2.7	1.5					
8	2.6	2.5	1.5					
9	1.9	2.4	2.2					
10	2.9	0.9	0.9					
11	2.6	0.5	1.1					
12		2.2						
13		0.8						
14		0.9						
15								
16								
17								

N Groups	Chi-Sq. Value	DF	p-value
3	5.47	2	0.065

Group	N	Median	Mean of Ranks
Cherry	6	2.550	15.92
Peach	9	1.100	9.56
Plum	6	1.300	8.25

Statistical test indicates that there is no evidence that the residues are from different distributions

**Can combine datasets for MRL calculation**

# Challenge: same MRL calculator – different result



Fungi-Res  
Stonefruit  
Region / Country  
3 x 100g/ha 3D PHI

Total number of data (n)	21
Percentage of censored data	0%
Number of non-censored data	21
Lowest residue	0.500
Highest residue	2.900
Median residue	1.500
Mean	1.700
Standard deviation (SD)	0.787
Correction factor for censoring (CF)	1.000

Proposed MRL estimate




- Highest residue	2.900
- Mean + 4 SD	4.850
- CF x 3 Mean	5.100
Unrounded MRL	<u>5.100</u>
Rounded MRL	<u>6</u>

Residues (mg/kg)	n
0.5	1
0.8	2
0.9	3
1.1	2
1.2	1
1.5	2
1.9	1
2.2	2
2.4	1
2.5	2
2.6	2
2.7	1
2.9	1

# Challenge: different ways of combining residue data from representative crops

## Fungi-Res MRLs

*...leads to different MRLs*

		US MRL	EU MRL	Codex CXL
	Cherry	7	7	6
	Peach	7	5	6
	Plum	7	4	6



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# Opportunities

*Projects and efforts that will  
make a difference  
...to reduce the difference*



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# The International Crop Grouping Consulting Committee



The ICGCC has been working to harmonize crop groups internationally for more than 10 years

Chaired by US and the Netherlands  
Includes more than 200 crop experts and 35 countries

*Argentina, Australia, Bangladesh, Belgium, Brazil, Burkina Faso, Canada, Chile, China, Columbia, France, Germany, Guatemala, Honduras, Hungary, India, Israel, Japan, Kenya, Lebanon, Mali, Mexico, Morocco, New Zealand, Nicaragua, Nigeria, S. Korea, Senegal, South Africa, St. Kitts, Taiwan, Thailand, Trinidad, United Kingdom, United States*



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*Develop harmonized crop groups that are considered by EPA and CCPR for revision of the Codex Committee on Pesticide Residues (CCPR) Classification of Food and Animal Feeds.*



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*“Principles and Guidance for the Selection of Representative Commodities for the Extrapolation of MRLs”*



# EPA 5X Rule - Current approach or...

Fungi-Res

Fungi-Res

Fungi-Res  
Peach  
US  
3x100g/ha 0 PHI

Total number of data (n)	6
Percentage of censored data	0%
Number of non-censored data	6
Lowest residue	0.800
Highest residue	2.200
Median residue	1.300
Mean	1.333
Standard deviation (SD)	0.516
Correction factor for censoring (CF)	1.000
<u>Proposed MRL estimate</u>	
- Highest residue	2.200
- Mean + 4 SD	3.399
- CF x 3 Mean	4.000
Unrounded MRL	4.000
Rounded MRL	4

High uncertainty of MRL estimate.  
[Small dataset]

## MRL

Cherry	7
Peach	5
Plum	4
<b>Stone Fruit</b>	<b>7</b>



EPA uses the “5X Rule” to determine if representative crops can support a crop group.

The MRL for each rep crop is calculated separately and the highest MRL is used for the whole group.

# ... The statistical approach?

	A	B	C	D	E	F	G	H
1	<b><u>Kruskal-Wallis Test</u></b>							
2								
3								
4								
5	<b>Cherry</b>	<b>Peach</b>	<b>Plum</b>	<b>Group 4</b>	<b>Group 5</b>	<b>Group 6</b>	<b>Group 7</b>	<b>Group 8</b>
6	1.2	1.1	0.8					
7	2.5	2.7	1.5					
8	2.6	2.5	1.5					
9	1.9	2.4	2.2					
10	2.9	0.9	0.9					
11	2.6	0.5	1.1					
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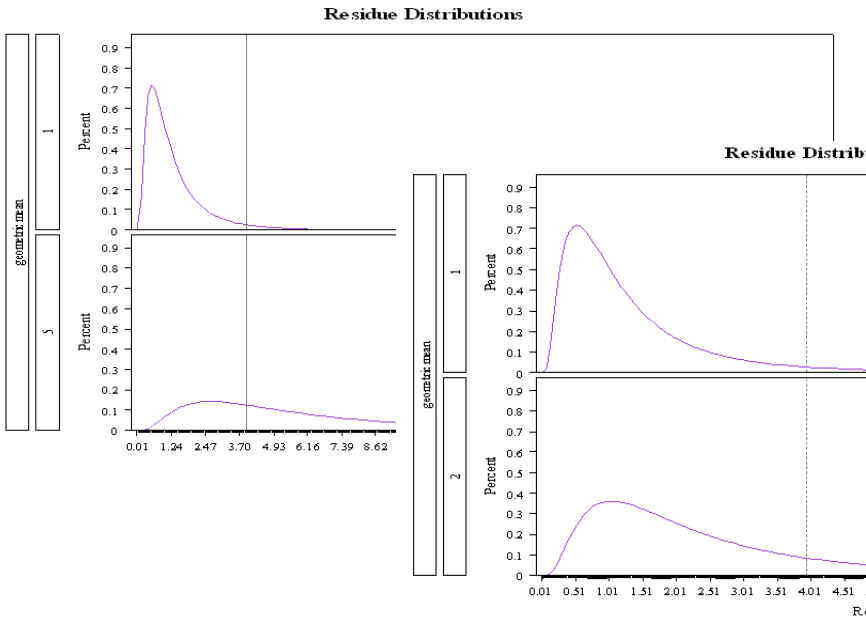
**Can combine datasets for MRL calculation**

<b>MRL</b>	
Cherry	
Peach	
Plum	
<b>Stone Fruit</b>	<b>6</b>

# Possible harmonization of MRL calculation procedures

The EPA is currently testing the performance of the **Kruskal-Wallis** test against the current **5X Rule** to see which approach is better at distinguishing residue data sets that are statistically similar / different.

**From a statistical viewpoint  
Initial results look very promising in favor of the K-W test**



Designed Max Ratio Factor (R)	Number of Groups (rep crops)	N Field Trials per group	Power to detect differences between groups when R=4		
			Kruskal Wallis	Max 5X	Med 5X
	2	5	0.63	0.38	0.36
		7	0.80	0.38	0.33
		10	0.93	0.36	0.31

Designed Max Ratio Factor (R)	Number of Groups (rep crops)	N Field Trials per group	Power to detect differences between groups when R=5		
			Kruskal Wallis	Max 5X	Med 5X
5	2	5	0.75	0.49	0.50
		7	0.89	0.50	0.50
		10	0.98	0.49	0.50
	3	5	0.66	0.60	0.56
		7	0.86	0.61	0.59
		10	0.97	0.60	0.58
	4	5	0.65	0.69	0.66
		7	0.85	0.69	0.66
		10	0.97	0.68	0.63
	5	5	0.66	0.78	0.73
		7	0.87	0.75	0.70
		10	0.97	0.74	0.69

0.49	0.43
0.49	0.42
0.46	0.38
0.60	0.54
0.58	0.50
0.56	0.44
0.69	0.60
0.65	0.55
0.64	0.50

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# What's the answer?



International  
harmonized crop groups  
Guidance on  
representative crops

Harmonized methods for  
calculating crop group  
MRLs

Crunchi-Wallis H test  
(degrees of freedom)

$$= \left[ \frac{12}{n(n+1)} \sum_{j=1}^c \frac{T_j^2}{n_j} \right] - 3(n-1)$$

n = sum of sample sizes in all  
Number of samples  
sum of ranks in the

Harmonized MRLs allow greater commodity trade among countries and open more markets for growers





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