MEMORANDUM


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Background

ICI has submitted residue data on numerous crops, including field and sweet corn, wheat, sorghum, soybeans, peanuts, sunflowers, broccoli, cabbage, tomatoes, hops, onions and lettuce. These crops and relevant processed commodities were analyzed for lambda-cyhalothrin, its epimer (an isomer pair that is always present at low concentrations), and three metabolites: PP890, 3-PBAcid and 3-PBAAlcohol. Our most recent memo is dated 11/19/92.

In the present submission, ICI has assembled all the collected residue data in one volume and calculated averages for each crop. Such information should be quite useful in determining anticipated residues. However, present policy is to first conduct a DRES analysis using tolerances. If the reference dose is exceeded, exposure is then calculated with anticipated residues, and the submitted information can be used at that time.

ICI has also submitted a proposal for conducting residue analyses based on patterns observed in previous analyses. The company notes the following:
2

1. Maximum metabolite residues (PP890, 3-PBAcid and PBAAlcohol) are always low and/or substantially less than the maximum regulated residues (lambda-cyhalothrin plus its epimer).

2. With one exception, metabolite residues are always below the limit of determination (<0.01 ppm) when the combined regulated residues are ≤0.04 ppm. In one field trial with wheat, residues of lambda-cyhalothrin and epimer were <0.01 but the level of PP890 was 0.02 ppm.

3. Average metabolite residues are always low (≤25%) relative to the combined residues of lambda-cyhalothrin and epimer.

The company therefore proposes the following program for future analyses:

1. All samples will be analyzed for lambda-cyhalothrin and epimer.

2. For commodities with maximum combined residues of parent and epimer ≥0.05, metabolite analyses will be conducted on samples from each trial having the three highest combined parent and epimer residues. [We assume that this means that from all the field trials on one RAC, metabolite analyses will be carried out on samples from three trials.]

3. For commodities with maximum combined parent and epimer residues ≤0.04 ppm, no metabolite analyses will be conducted.

For processed commodities, ICI notes that:

1. With the exceptions of grain dust [not really a processed commodity] and tomato pomace, all residues of parent, epimer and metabolites are low.

2. With the exception of crude corn oil, peanut soapstock and soybean soapstock, PP890 and combined 3-PBAcid and 3-PBAAlcohol residues are always ≤ to the lambda-cyhalothrin residues.

ICI therefore proposes that:

1. All samples would be analyzed for parent and epimer.

2. Metabolites would be analyzed in soapstock samples and in commodities with maximum combined parent and epimer levels of ≥0.05 ppm; otherwise no metabolite analyses
would be carried out.

**CBTS Comment**

CB I and TB I have agreed that, at present, only the parent and epimer need appear in tolerance expressions for lambda-cyhalothrin. This conclusion was based on probable toxicities of the metabolites as well as low concentrations. The residue data summary for the raw agricultural commodities does indicate that when concentrations of parent lambda-cyhalothrin are low, metabolite concentrations are lower. We therefore agree with ICI’s proposal to analyze for metabolites only when parent + epimer concentration exceeds 0.05 ppm.

On the other hand we recommend that metabolite analyses be continued for processed commodities, at least for human food items. Data from seven completed processing studies do indicate that even when parent is present at low levels, metabolites -- usually PP890 -- may be also present at low but quantifiable levels. For example, in tomato juice, parent + epimer concentration was 0.02 ppm, PP890 concentration was 0.02 ppm and PBAcid + PBAcohol concentration was 0.01 ppm. Should the tolerance expression for lambda-cyhalothrin be reassessed, such analytical data would be useful. Such reassessment might occur, for example, if data from new field trials show significant concentrations of metabolites and a general DRES analysis becomes necessary.

cc: SF, RF, Circu., Mike Flood, E. Haeberer, PP#7F3560, PP#2F4109, PP#2F4414.