

InfoVeg - Bug #3609

subsampling typo? 11-C-309

11/07/2008 04:13 PM - Michael Lee

Status:	Resolved	Start date:	11/07/2008
Priority:	Normal	Due date:	
Assignee:	Michael Lee	% Done:	0%
Category:	DataFix	Estimated time:	0.00 hour
Target version:	2009-June	Spent time:	0.00 hour
Bugzilla-Id:	3609		
Description			
011-0C-0309 is listed in a way in the file that I think the 20% and 10% subsample should be something else, but I'm not sure what exactly. Probably requires a look at the original datasheet.			
Related issues:			
Blocks InfoVeg - Bug #3611: Newell R module = supersample?			In Progress 11/07/2008

History

#1 - 03/20/2009 08:46 AM - Michael Lee

This is another Claire plot where the supersampling was split into an R module. Here, it is for all TREE species, presumably. Only trees are recorded in R.

Solution to this depends on the solution to 3611, which is the same weird issue.

#2 - 03/20/2009 08:51 AM - Michael Lee

I change my mind. There is a diagram on the tree sheet showing the location of the supersample. It is clearly 2 m along the far side of module 1:

[MOD2][MOD1][2m super sample]
(diagram not to scale)

Therefore, I think the best solution is to move the supersample entirely to the module #1 and have module 2 be treated as 100% sample, and 1 as 120% supersample for trees only.

#3 - 03/20/2009 09:11 AM - Michael Lee

This has been fixed, archive version 1.1.24

#4 - 03/27/2013 02:23 PM - Redmine Admin

Original Bugzilla ID was 3609