**caArray 2.x Status Meeting Minutes**

**Date:** January 19, 2011

**Attended By:**

|  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- |
| x | Juli Klemm | x | Rashmi Srinivasa |  | Vanessa Caldwell |
|  | George Komatsoulis |  | Daniel Kokotov | x | Jill Hadfield |
| x | Matt Tiller | x | Don Swan |  | Anand Basu |
| x | Quy Phung |  | Sriram Kalyanasundaram |  | Zhong Li |
| x | Xiaopeng Bian |  | Gavin Brennan |  | Zhou Ji |
| X | Maureen Colbert |  | Yeon Choi |  | Michael Honig |
|  | Jia Liu | x | Andrea Johnson |  | Jianguo “David” Lee |
| X | Maki Duncan |  | Cuong Nguyen |  | Jessica Cheng |
| x | John Scott | x | Jacob Mensah |  | Ken Smith |

The running action item list and past meeting minutes are available at:

<https://gforge.nci.nih.gov/plugins/scmsvn/viewcvs.php/trunk/docs/project_management/internal_meetings/?root=caarray2>

### **2.4.0.1 Status**

The problem importing copy number data is fixed but a new issue was filed with a larger Netherland data.

### Original issue found in 2.4.0.1: Array-1812: Netherlands Copy Number data failing to import (verified by QA)

New issues:

### Array-1792: Investigate Large Netherlands data (Copy Number (MAGE-TAB Data Matrix)) files fail on import

### Array-1811: Multiple associated array designs causes copy number data to be Imported (not parsed)

Array-1796: need decision either to include the fix in a new upgrader or provide the db script instead. We will wait till we receive a response on the forum, but we will likely only provide a db script to fix the problem for this particular institution.

Array-1791: issue with GenePattern. We cannot reproduce their problem.

Array-1722: Curation team will reload the data to a new experiment. This script will be very complex touching multiple tables.

### **2.5.0 Planning**

* Iteration 31: 2.5.0\_M4, 2.4.0.2\_M2

### **Systems**

Update

### **Documentation**

Update

### **Support / Knowledge Center**

**Update**

### **Other**

Feature request to validate so that you cannot have multiple hybes associated with a single cel file. Array-1813 was created to capture this request. We will need to discuss and consider the feature request.