

# Welcome!

## **Swiss Galaxy Workshop** (workflow management / reproducible research)

### **3<sup>rd</sup> Oct. 2012, Bern**

Status of the project

Developments

Interface to other systems

Extensions

Best practice

***<http://wiki.g2.bx.psu.edu/Events/Switzerland2012>***

# Thank You!

all the speakers

Dave Clements and the support  
from the Galaxy Project

Peter Kunszt and the support  
from SystemsX.ch



# aim of today's workshop

promoting Galaxy in Switzerland (and Europe)

establishing a 'central Europe' Galaxy Community

evaluating Galaxy / comparison with other systems



a special Welcome to all Non-Galaxiens

# who is attending

(based on the 38 original registrations)

<b>Galaxy Users</b>	<b>17</b>
<b>Galaxy Admins</b>	<b>16</b>
<b>Tool Developers</b>	<b>12</b>
<b>Non-Galaxiens</b>	<b>4</b>
<b>no answer</b>	<b>11</b>

# what is Galaxy used for

(based on the 38 original registrations)

<b>NGS analysis</b>	<b>22</b>
<b>Microarray analysis</b>	<b>3</b>
<b>High Throughput Screening</b>	<b>6</b>
<b>Proteomics</b>	<b>3</b>
<b>Data management</b>	<b>12</b>
<b>Set-theory / interval operations</b>	<b>7</b>
<b>Text/File manipulation</b>	<b>15</b>
<b>other</b>	<b>0</b>
<b>no answer</b>	<b>11</b>

# morning schedule

9:50 - 10:30

Dannon Baker

10:30 - 11:00

BREAK

11:00 - 11:15

Mark Robinson

11:15 - 11:30

Tim Roloff

11:30 - 11:45

*cancelled*

11:45 - 12:00

Manuel Kohler

12:15 - 13:30

LUNCH

# afternoon schedule

<b>13:30 - 13:45</b>	<b>Nikolai Vazov</b>
<b>13:45 - 14:15</b>	<b>Jelle Scholtalbers</b>
<b>14:15 - 14:30</b>	<b>Mario Emmenlauer</b>
<b>14:30 - 15:00</b>	<b>BREAK</b>
<b>15:00 - 15:15</b>	<b>Grainne Kerr</b>
<b>15:15 - 15:30</b>	<b>Björn Grüning</b>
<b>15:30 - 15:45</b>	<b>Sveinung Gundersen</b>
<b>15:45 - 16:10</b>	<b>conclusion/panel discussion</b>

