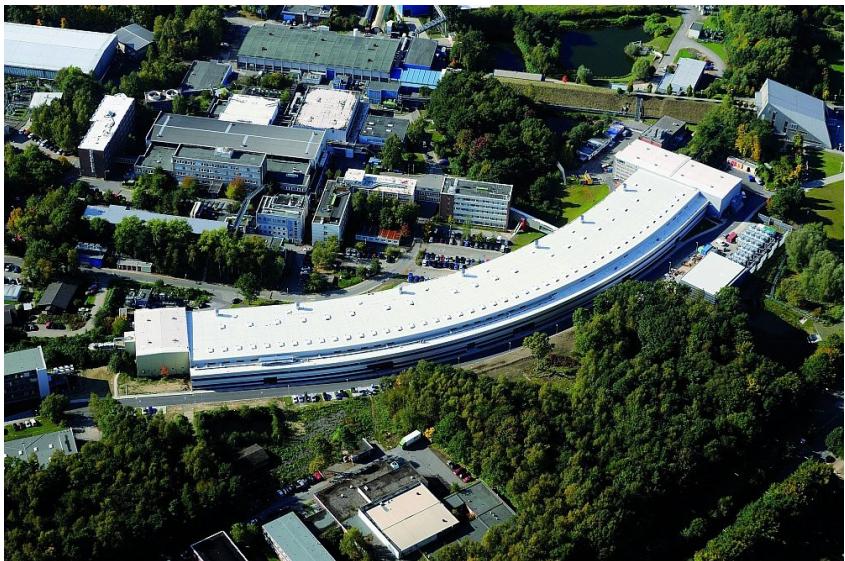


IT Infrastructure at DESY

The IT infrastructure for the PETRA III synchrotron

Eugen Wintersberger
IT infrastructure at DESY
BNL, 26.05.2016

The PETRA III synchrotron at DESY



Diverse Environment

- **Various kinds of research**
- **Different techniques**
- **Multiple kinds of analysis**

Beamline status: user operation

1. High resolution dynamics beamline
2. Hard x-ray diffraction beamline for time-resolved studies
3. Micro- and nanofocus x-ray scattering
4. Variable polarization XUV
5. HZG/DESY: Imaging
6. Hard x-ray micro/nano probe
7. HZG: High energy materials science
8. High resolution diffraction
9. Resonant scattering and diffraction
10. Coherence applications
11. MPI/HZIR/DESY: Protein crystallography and Bio-Imaging
12. EMBL: BioSAXS
13. EMBL: Macromolecular crystallography I
14. EMBL: Macromolecular crystallography II

Deutsches Elektronen-Synchrotron

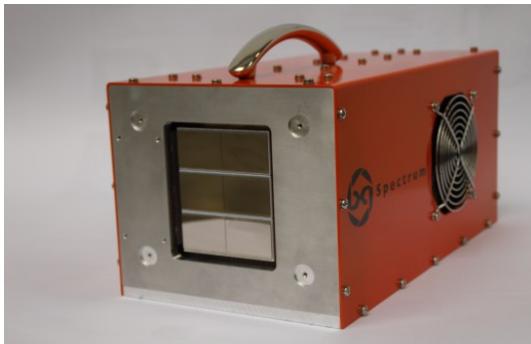
Helmholtz-Zentrum Geesthacht Centre for Materials and Coastal Research
European Molecular Biology Laboratory



Detectors used at PETRA III beamlines



Dectris Pilatus and Eiger systems



Lambda systems
• 750k – 2.7GB/s
• 2M – 3x2.7GB/s



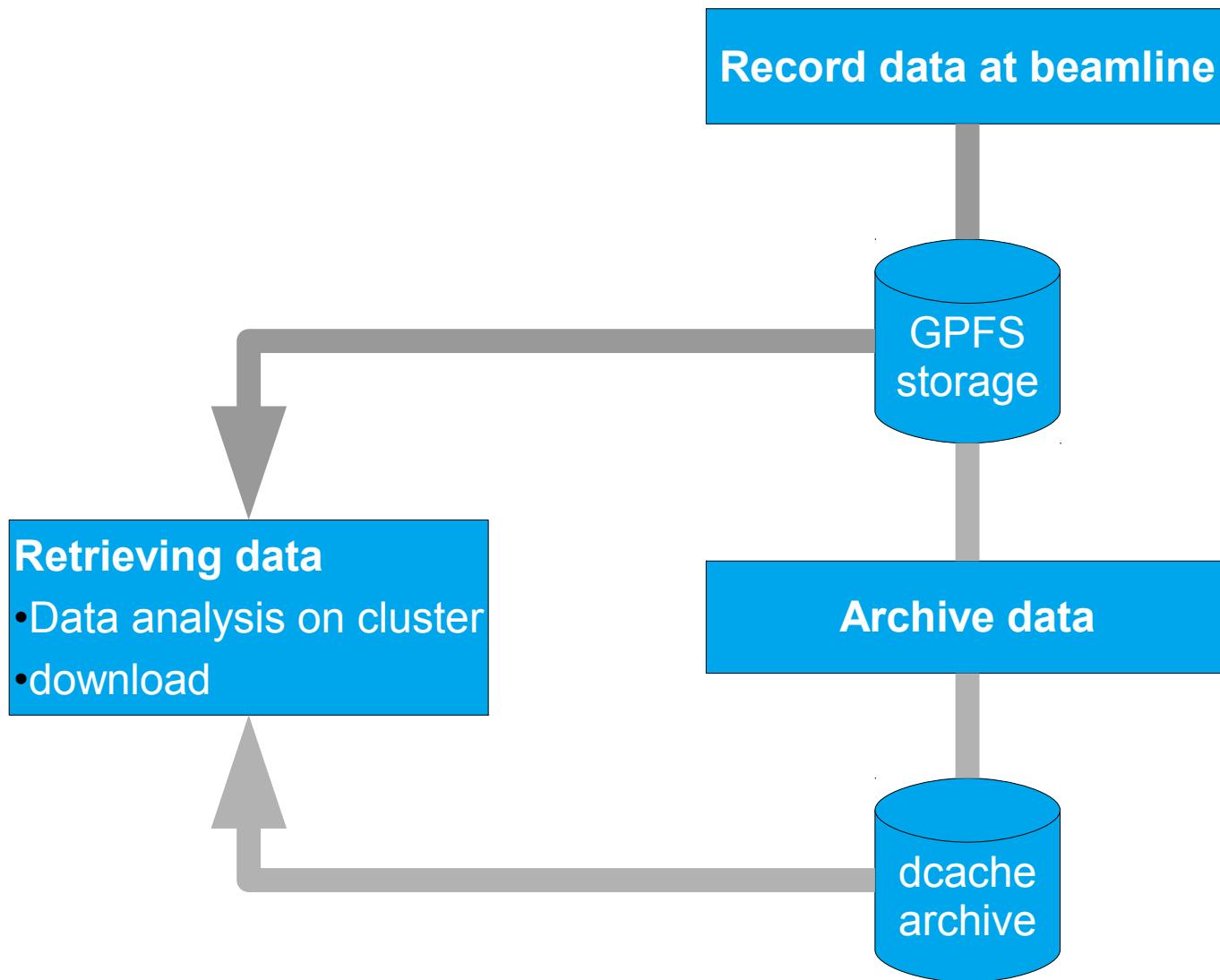
PCO Edge (~1GB/s) and Perkin Elmer (240MB/s) systems

Network Computing center – synchrotron facility

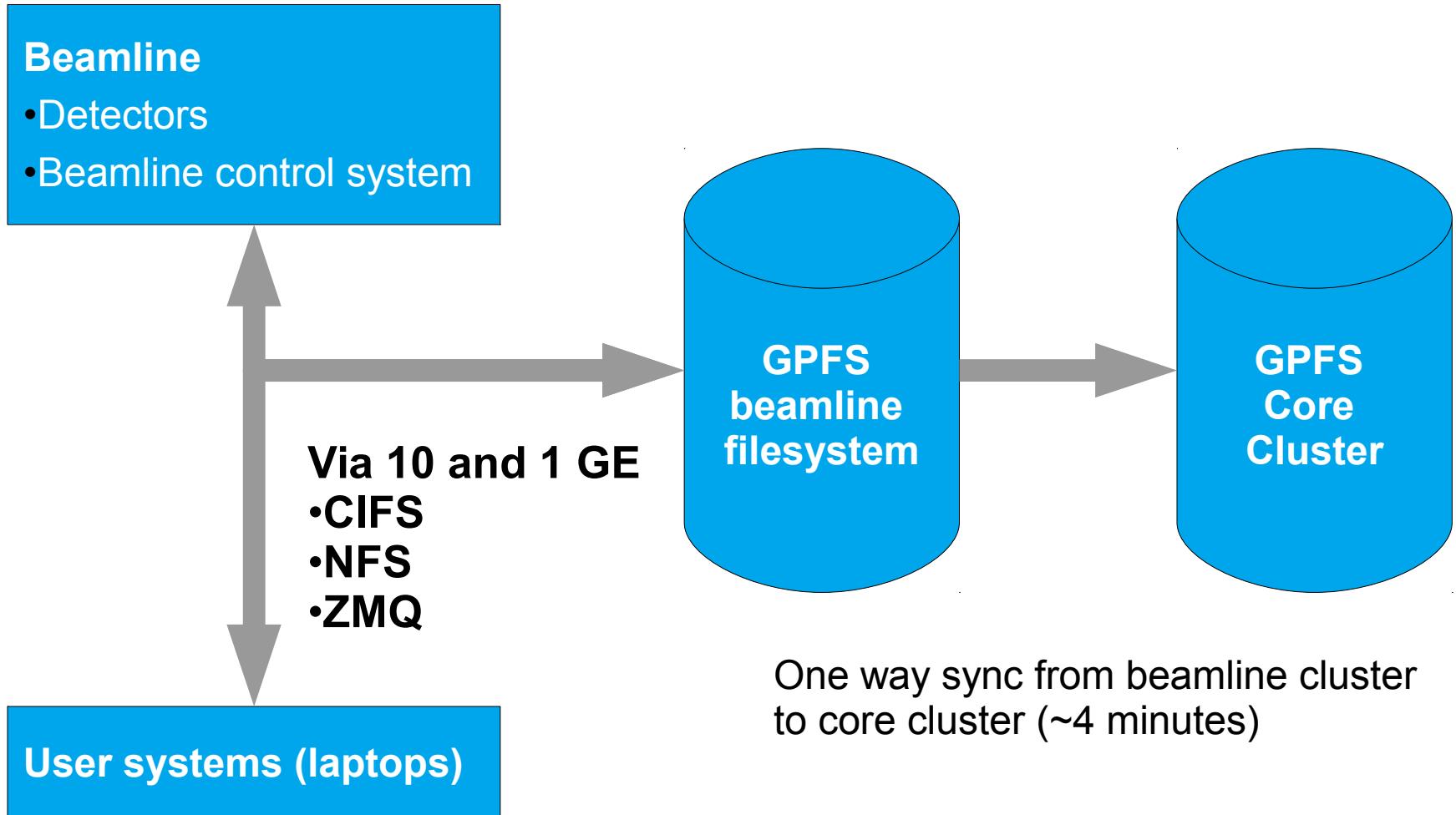


- 3 Arista switches with 2x40GE (fibre) each in the hall
- Total bandwidth 240 GE
- Can double bandwidth easily

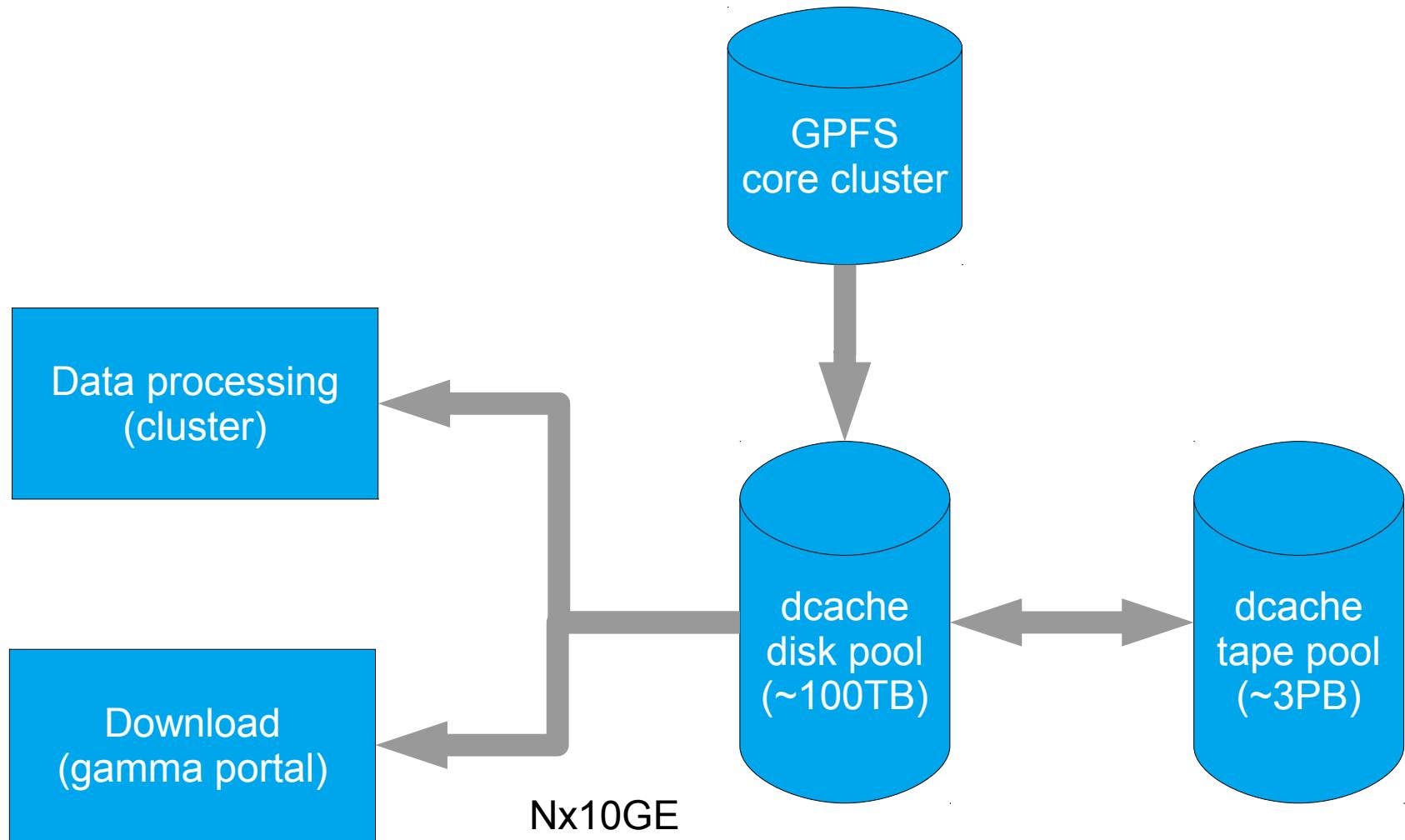
Data lifecycle



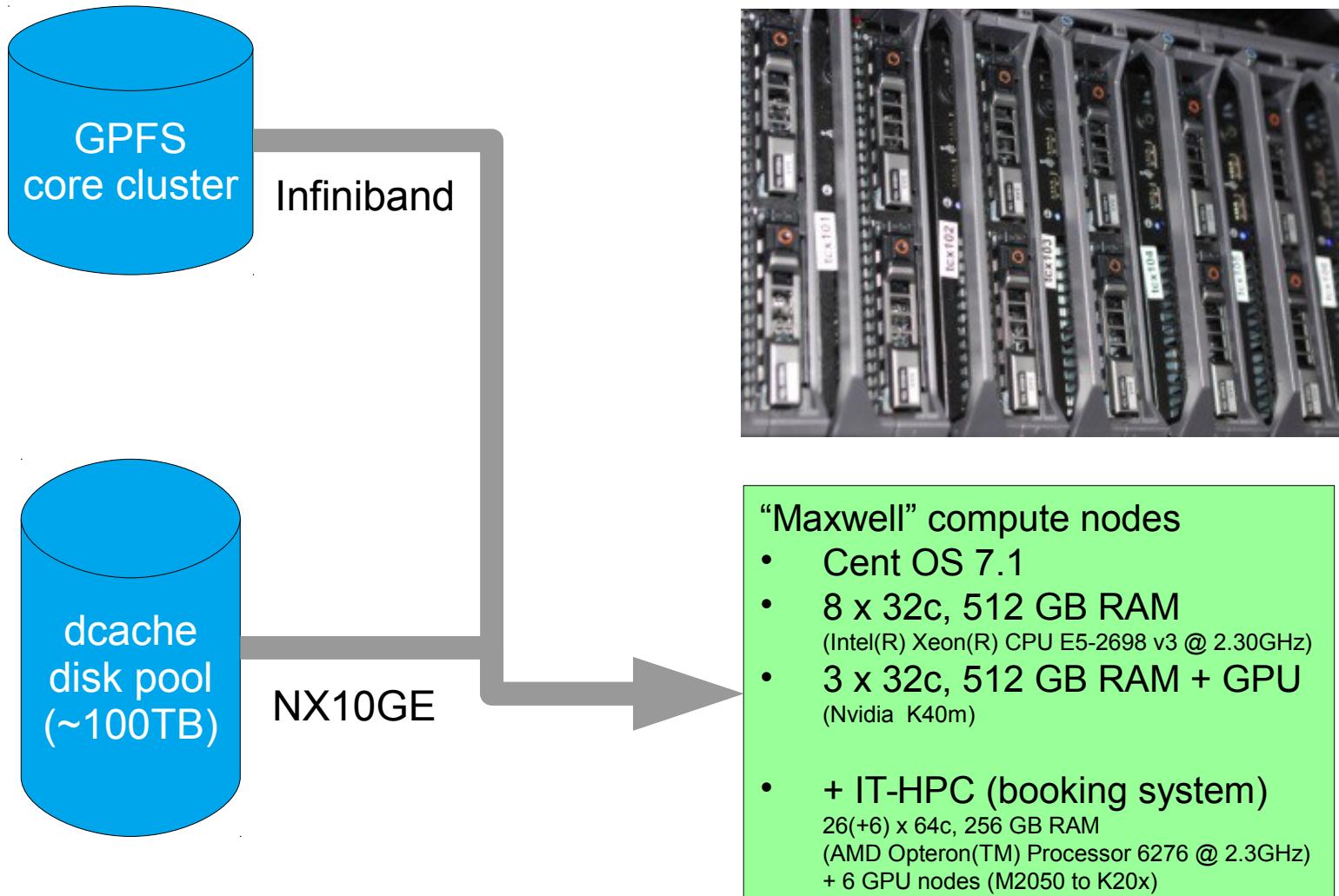
Beamline storage



Long time archive (dcache)



Supercomputing and storage



Data transfer to user facilities: Gamma-Portal

The screenshot displays two Mozilla Firefox browser windows side-by-side.

Left Window (Details): This window shows the "Browse archive / Details" interface. It includes a sidebar with links to Home, Browse archive, -Details (which is highlighted in orange), Staging status, and Migration status. The main area shows a table of archived datasets:

Archived Directory	Format	Size in Mb	Archived Date
experiment3	tar	0.01	06-FEB-2013
experiment3/kupfer	tar	0.01	06-FEB-2013
experiment3/protein1	tar	0.01	06-FEB-2013

Buttons for Stage and Download are at the bottom, along with a note: "* Stage means to copy multiple datasets to DESY disk space".

Right Window (edit_door_users): This window shows the "edit_door_users" interface. It has sections for Home, Browse archive, -Users/ACLs (highlighted in orange), Staging status, and Migration status. The main area is titled "Browse Archive / Users" and shows a table of users:

Door Account	Lastname	Firstrname	User Role	Access Permission	Create Date	Change Date
rothkrc	Rothkirc	Andre	applicant	download/stage	27-FEB-2013 15:14	27-FEB-2013 15:14
wroblewski	Wroblewski	Thomas	leader	download/stage	27-FEB-2013 15:14	27-FEB-2013 15:14

Buttons for Save, Delete, and Cancel are present. Below the table are sections for "ADD DOOR USER TO EXPERIMENT FOR DATA ACCESS" (with a "Door account" dropdown and "Add User" button) and "ACL MANAGEMENT" (with "Update Acls" and "Show Acls" buttons). A status bar at the bottom indicates "ROTHKIRC ADMIN en".

- Different views
- Locate data by keyword search in metadata
- Download & stage
- Project leader & applicant can manage ACLs