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CONFERENCE PROGRAMME

Tuesday, 6 March 2012

08:30 - 08:50	Bus pickup from downtown stops and train station
08:30 - 09:15	Registration <i>ATC Main Entrance</i>
09:15 - 09:30	Opening Remarks (Seán O'Donoghue) <i>Klaus Tschira Auditorium</i>
09:30 - 10:30	Keynote: Integrating Visualization and Analysis (Georges Grinstein) <i>Klaus Tschira Auditorium</i>
10:30 - 11:00	Coffee Break <i>ATC Foyer</i>

Visualizing Single Genome Data

Session Chair: James Procter
Klaus Tschira Auditorium

11:00 - 11:30	Assembly for NGS (Cydney Nielsen)
11:30 - 12:00	Browsers for medical genomics (Michael Brudno)
12:00 - 12:30	Visualizing functional genomics (Jim Robinson)
12:30 - 12:37	BioVis Honourable mention (Hendrik Strobel)
12:37 - 13:00	Fast-Forward for Poster Session 'A'
13:00 - 15:00	Lunch Break + Poster Session 'A' <i>ATC Foyer</i>

Visualizing Transcript Data

Session Chair: John Quackenbush
Klaus Tschira Auditorium

15:00 - 15:30	2D & 3D Structures of RNA (Yann Ponty)
15:30 - 16:00	Visualizing Transcriptional Regulation & Abundance (Kay Nieselt)
16:00 - 16:30	Alternative Splicing (Jernej Ule)
16:30 - 17:00	Coffee Break <i>ATC Foyer</i>

VIZBI Art & Biology Evening

Organisers: Bang Wong & Rikke Schmidt Kjrgaard

17:00 - 18:00	Outreach Keynote: Communicating Biology Visually (David Goodsell) <i>Klaus Tschira Auditorium</i>
18:00 – 19:00	Pre-dinner drinks (<i>Sponsored by our Silver Partners Autodesk</i>) <i>ATC Foyer</i>
19:00 – 22:00	Dinner <i>EMBL Canteen</i>
21:00	1st bus departs for downtown
22:00	Last bus departs for downtown

Wednesday, 7 March 2012

09:00 - 09:10 Bus pickup from downtown stops and train station

09:45 - 10:00 Announcements (Organisers)
Klaus Tschira Auditorium

Visualizing Proteins & Complexes

Session Chair: Gaël McGill

Klaus Tschira Auditorium

10:00 - 10:30 Protein Folds & Folding (Valerie Daggett)
10:30 - 11:00 Visual Exploration of Protein Dynamics (Ivet Bahar)
11:00 - 11:30 Mesoscale Visualization (Graham Johnson)
11:30 - 12:00 Coffee Break
12:00 - 12:07 BioVis contest - Biologists' favourite (Ryo Sakai)
12:07 - 12:30 Fast-forward for Poster Session 'B'

12:30 - 14:30 Lunch Break + Poster Session 'B'
ATC Foyer

Visualizing Cellular Systems

Session Chair: Nils Gehlenborg

Klaus Tschira Auditorium

14:30 - 15:00 Protein Networks (Lars Juhl Jensen)
15:00 - 15:30 Biological Pathways (Alex Pico)
15:30 - 16:00 Cellular Image Data (Jason Swedlow)
16:00 - 16:30 Coffee Break
16:30 - 17:30 Keynote: Human Computer Interfaces (Jessie Kennedy)
17:30 - 18:00 Fast-forward for Poster Session 'C'

18:00 - 20:00 Dinner + Poster Session 'C'
ATC Foyer

20:00 Bus to all downtown stops

20:30 Get together at Restaurant *Kulturbrauerei*
Leyergasse 6, 69117 Heidelberg
Bus Stop Neckarmünzplatz

Thursday, 8 March 2012

- 08:30 - 08:40 Bus pickup from downtown stops and train station
09:15 - 09:30 Announcements (Organisers)
Klaus Tschira Auditorium
09:30 - 10:30 Keynote: Visual Design Principles (Katy Börner)
Klaus Tschira Auditorium
10:30 - 11:00 Coffee Break
ATC Foyer

Visualizing Organism Data

Session Chair: Richard Baldock
Klaus Tschira Auditorium

- 11:00 - 11:30 Developmental Anatomy (Fons Verbeek)
11:30 - 12:00 Neural Anatomy (Gene Myers)
12:00 - 12:30 Physiology & Function (Marco Viceconti)
12:30 - 12:37 BioVis contest - experts favorite (Günter Jäger)
12:37 - 13:00 Fast-forward for Poster Session 'D'

13:00 - 15:00 Lunch Break + Poster Session 'D'
ATC Foyer

Alignments & Phylogenetics

Session Chair: Chris Creevey
Klaus Tschira Auditorium

- 15:00 - 15:30 Multiple Sequence Alignments (Teresa Attwood)
15:30 - 16:00 Metagenomics & Phylogeography (Daniel Huson)
16:00 - 16:30 Population Structure & Dynamics (Philippe Lemey)
16:30 - 17:00 Closing Remarks (Organizing Committee & Session Chairs)

17:00 - 17:30 Coffee Break
ATC Foyer

17:30 Buses depart for downtown, train station & airport
EMBL Bus Stop

FAST-FORWARD AND POSTER PRESENTATIONS

A01

Structural Kinome Inhibitor Library

Chandan Sona and V Badireenath Konkimalla

A02

Expression in strains forming disulfide bonds of phospholipases-D of Brown Spider.

Matheus Regis Belisário, Adriano Marcelo Morgon, Luiza Helena Gremski, Waldemiro Gremski, Andréia Senff Ribeiro, Olga Meiri Chaim, Silvio Sanches Veiga

A03

Quasar - peptide measurement efficiency prediction

Markus-Hermann Koch; Julian Uszkoreit, Stefan Lorocho, Katja Kuhlmann, Martin Eisenacher, Hanna Diehl, Raphael Voges, Marco Oldiges, Helmut E. Meyer, Christian Stephan

A04

Visualization of cells using cryo electron tomography

Roman Koning

A05

Visual Analysis of Uncertainties within Biochemical Reaction Networks

Corinna Vehlow, Jan Hasenauer, Andrei Kramer, Julian Heinrich, Nicole Radde, Frank Allgöwer, Daniel Weiskopf

A06

ZMap: Efficient and Informative Presentation of High Volume Genomic Data

Malcolm Hinsley, Gemma Barton, Ed Griffiths

A07

A spatial and temporal modeling study of the tumor and its microenvironment

Naamah Bloch, David Harel

A08

Interactive Visual Comparison of Multiple Phylogenies

Sebastian Bremm, Martin Heß, Kay Hamacher, Tatiana von Landesberger

A09

Blixem; a viewer for multiple gapped, pairwise alignments on a genomic sequence.

Jeremy Henty, Gemma Barson

A10

KoriBlast: interactively explore NGS datasets from a global perspective.

Alain Meil, Patrick Durand

A11

Visualization of Arabidopsis Interactome

Julia Dmitrieva, Laurence Lins

A12

Visualizing multi-dimensional genomics data using interactive heatmaps in Gitools

Michael P Schroeder, Christan Pérez-Llamas, Abel Gonzalez-Perez & Nuria Lopez-Bigas

A13

Fungal Genome Collection

C Anderson, SD Harris, EN moriyama

A14

Identification and visualization of transient binding pockets in proteins

Daria B. Kokh, Stefan Henrich, Paul Czodrowski, Friedrich Rippmann, Rebecca C. Wade

A15

Vizualization for Logic Modeling tools

M.P. van Iersel, E.J. Gonçalves, C. Chaouiya, J. Saez-Rodriguez

A16

The Open Microscopy Environment: Open Image Informatics for the Biological Sciences

Josh Moore, Jason R. Swedlow, and the OME Consortium

A17

Visual Analysis of RNAseq Data to Detect Overlapping Genes in Bacterial Genomes

S. Simon, D. Oelke, R. Landstorfer, K. Neuhaus, D. A. Keim

A18

Disturbed networks in stage III and IV melanomas

Sarah-Jane Schramm, Anna E Campaign, Simone S Li, David C Y Fung, Vivek Jayaswal, Chi Nam Ignatius Pang, Richard A Scolyer, Yee Hwa Yang, Marc R Wilkins and Graham J Mann

A19

Re-designing SRS 3D and Reflect

Christian Stolte, Seán O'Donoghue

B01

Towards an almost complete, highly reliable protein-protein interaction network of yeast

Kubra Karagoz, Kazım Yalçın Arga

B02

Genome sequencing and variant analysis of a Pakistani individual

M. Kamran Azim, Yong Zhang, M. Iqbal Choudhary, Zhixiang Yan, Chuanchun Yang, Asifullah Khan, Huma Asif, Yasmeen Rashid

B03

Meander: Using Hilbert Curves to visually explore structural variations

Georgios A. Pavlopoulos, Alejandro Sifrim, Ryo Sakai, Jan Aerts

B04

HyperBalls on GPU: Dynamic Visualization of Molecular Structures

M. Chavent, A. Tek, Z. Lu, M. PiuZZi, B. Laurent, M. Baaden

B05

Proposed Web Interface to the Next Generation of SWISS-MODEL

Marco Biasini, Konstantin Arnold, Stefan Bienert, Andrew Waterhouse, Torsten Schwede

B06

Automated 3D layout for SBML network diagrams

Mark M. Shovman, James Bown

B07

Biological Network Visualisation

Kai Xu

B08

PDBe widgets - compact and reusable visualizations of biomacromolecular data

S. P. Gore, G. van Ginkel, S. Velankar, G. J. Kleywegt

B09

Transcriptome Viewer

Johannes Wollnik

B10

On Visualizing Fragmented Peptide Mass Spectra in Proteomics using PEAKPLOT

Christian Panse, Jonas Grossmann, Bertran Gerrits, Ralph Schlapbach

B11

Effectiveness analysis of different binarization algorithms in histology

Vladyslav Luzin, Vladimir Ovcharenko, Anton Yeryomin

B12

FvNano : An Interactive Molecular Dynamics platform

Marc PiuZZi, Alex Tek, Matthieu Chavent, Nicolas Férey, Matthieu Dréher, Ahmed Turki, Sébastien Limet, Sophie Robert, Bruno Raffin, Marc Baaden

B13

Visualisation of Cross-link/MS data

Colin Combe; Salman Tahir; Lutz Fischer; Zhuo Angel Chen; Juan Zou; Adam Belsom; Lauri Peil; Jimi-Carlo Bukowski-Wills; Juri Rappsilber

B14

Tracing cooperativity: 3D-electron microscopy of Limulus polyphemus hemocyanin

Christoph Kuehne, Frank Depoix, Jürgen Markl

B15

MedSavant: accelerating the discovery of disease-causing genetic variants

Marc Fiume, Andrew Brook, Eric Smith, and Michael Brudno
Marc Fiume, Andrew Brook,
Eric Smith, and Michael Brudno

B16

GRAPE RNAseq Analysis Pipeline Environment and Browser

D. González, M. Röder and R. Guigó

B17

Visualization of freshwater ostracod anatomy

Renate Matzke-Karasz, Radka Symonova, Libor Morkovsky

B18

NetTrapper: An interactive tool to view and explore network features and models

Evan Molinelli, Martin Miller, Arman Aksoy

C01

The Use of Bioinformatics Tools in the Analysis of Toxins from the Brown-Spider Venom

Adriano M. Morgon, Matheus R. Belisário, Gabriel O. Meissner, Larissa Vuitika, Dilza Trevisan-Silva, Andrea S. Ribeiro, Luiza H. Gremski, Olga M. Chaim, Waldemiro Gremski, Silvio S. Veiga

C02

3D-EM of Molluscan Rhogocytes

Maria Kokkinopoulou and Jürgen Markl

C03

Omix - a highly customizable visualization tool for multi-omics data from life sciences

Peter Droste, Wolfgang Wiechert, Katharina Nöh

C04

Visualization and quality assessment of de novo genome assemblies

Oksana Riba-Grognuz, Laurent Keller, Laurent Falquet, Ioannis Xenarios and Yannick Wurm

C05

The PROFILE Commander Software - From Measurement to Presentation

Michael Kohl, Maïke Ahrens, Hagen Meckel, Julian Uszkoreit, Michael Turewicz, Martin Eisenacher, Dominik A. Megger, Barbara Sitek, Helmut E. Meyer, Christian Stephan

C06

Visualizing Molecular Dynamics using VMD and Maya

Samuel Hertig & Viola Vogel

C07

Visual Analysis for 3D Electron Microscopy Data

Ingvar Lagerstedt, Ardan Patwardhan, Gerard Kleywegt

C08

Quips – Bringing biomacromolecular structures to a non-specialist audience

M. F. Symmons, S. P. Gore, M. J. Conway, G. van Ginkel, S. Velankar, G. J. Kleywegt

C09

VIPER: Visual Interactive Pedigree ExploreR

Trevor Paterson, Martin Graham, Jessie Kennedy, Andy Law

C10

Interactive Visual Analysis of Biological Data

Cagatay Turkay, Julius Parulek, Helwig Hauser

C11

FISH Oracle: a web server for visualization of DNA copy number data in a genomic context

Malte Mader, Ronald Simon, Sascha Steinbiss, Stefan Kurtz

C12

Reveal - Visual eQTL Analytics

Günter Jäger, Florian Battke, Corinna Vehlow, Julian Heinrich, Kay Nieselt

C13

Clustering biomolecular structures by residue contact similarity

João PGLM Rodrigues, Mikaël Trellet, Christophe Schmitz, Adrien SJ Melquiond, Alexandre MJJ Bonvin

C14

Intramuscular architecture of the human back muscles

Heiko Stark, Rosemarie Fröber, Nadja Schilling

C15

UnityMol: Visualization with a game engine

Zhian Lv, Alex Tek, Matthieu Chavent, Marc Baaden

C16

Automatic CNV Scoring Tool for unpaired sample analysis with Affymetrix SNP-Chips

Sven Haag, Daniel Nowak, W.-K. Hofmann, Markus Gumbel

C17

The Vesicle Builder - A Plugin for the CELLmicrocosmos MembraneEditor

Björn Sommer, Yan Zhou

C18

Visualization and Analysis of Very Large Network Graphs from High Dimensional Data

Thanos Theo and Tom Freeman

D01

TBDB: Publishing TB Systems Biology Data in an Interactive Genomic Context

Peter Sisk, Christian Stolte, Thomas Abeel, Matthew Peterson, Jeremy Zucker, Antonio Gomes, Anna Lyubetskaya, Tige Rustad, Kyle Minch, Irina Glotova, David Sherman, James Galagan

D02

Visual Analysis of the Digital Embryo

Jens Fangerau, Burkhard Höckendorf, Jochen Wittbrodt, Bernhard X. Kausler, Fred A. Hamprecht, Heike Leitte

D03

PhenoTimer: connecting time-resolved phenotypes

Maria Secrier, Reinhard Schneider

D04

Using dynamic simulations to better visualize signaling networks

Megan Riel-Mehan, Kevan Shokat

D05

Effect of carbon nano-particles on the flowering

Khushbu Kumari, Kishore CS Panigrahi

D06

A framework for pan-genome analysis and visualization

Leonid Zaslavsky, Vyacheslav Chetvernin, Boris Fedorov, Boris Kiryutin, Alexandre Souvorov, Igor Tolstoy, Tatiana Tatusova

D07

Nucleolar self-assembly as modeled by Monte Carlo simulations of free diffusion

Michaela Blazikova, Ivan Novotny, Petr Herman, Jan Malinsky

D08

Realtime genome browsing in Chipster

Petri Klemelä, Alekski Kallio, Taavi Hupponen, Massimiliano Gentile, Rimvydas Naktinis, Eija Korpelainen

D09

PresentaBALL - A Web Technology based Presentation Extension of BALL

Sabine C. Mueller, Stefan Nickels, D. Stoeckel, A.K. Dehof, H.P. Lenhof, Andreas Hildebrandt

D10

Pipit: Visualising Genomic Structural Variation

Ryo Sakai, Matthieu Moisse, Joke Reumers, Alejandro Sifrim, Raf Winand, Georgios A. Pavlopoulos, Andrew Vande Moere, Jan Aerts

D11

3D Warping and Image Registration in the EMAGE Database

S. Venkataraman, Bill Hill, Zsolt Husz, Chris J. Armit, Dominic Rannie, Richard A. Baldock

D12

SBGN-ED - working with the Systems Biology Graphical Notation

Tobias Czauderna and Falk Schreiber

D13

Visualizing time-dependent gene expression pattern in an animal model of drug addiction

Oliver Staehlin, Stephanie Perreau-Lenz, Wolfgang Sommer

D14

Leeds Virtual Microscope & Paramorama Visualization System

Roy Ruddle, Darren Treanor, A Johannes Pretorius

D15

PhysiIllustration

Ivan Viola

D16

Stochastic Single-Cell Simulation: The External Signal

Martin Falk, Thomas Ertl

D17

StratomeX - Integrative Visualization of Tumor Subtypes in Cancer Genomics Data Sets

Alexander Lex, Marc Streit, Hans-Joerg Schulz, Christian Partl, Dieter Schmalstieg, Peter J. Park, Nils Gehlenborg

D18

uPy: a ubiquitous Python API with biological modeling applications enables ePMV & autoFill

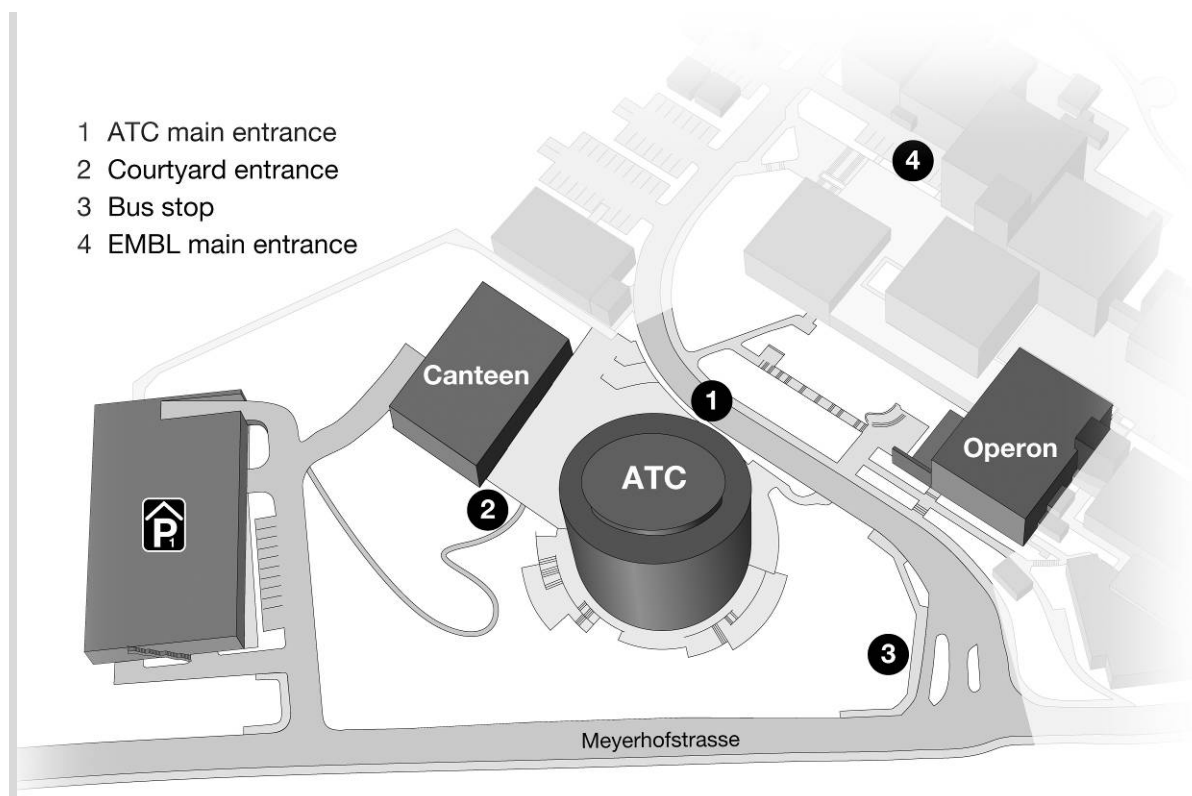
Ludovic Autin, Graham Johnson, Johan Hake, Arthur Olson, Michel Sanner

Y02

Nuclear Pore Transport

Martin Falk

MAP OF EMBL CAMPUS



The conference will take place at the **EMBL Advanced Training Centre (ATC)**.

While on campus, please wear your conference badge at all time for security purposes.

HEALTH & SAFETY NOTES



Do not smoke in any EMBL building.



Eating and drinking is prohibited in the Auditorium and all laboratories.



Do not enter any restricted areas or the laboratories unless instructed to do so.



If first aid is required ...

- A first aid room (room 10- 205) is located next to the registration desk in the ATC.
- Dial the **Emergency number 222** from any internal phone. Where is the accident? What happened? How many injured?
- Please report all accidents to the conference officer at the registration desk.



In case of fire...

- Press the nearest fire alarm button or the **Emergency number 222** from any internal phone.
- A loud fire alarm will go off if an evacuation is required. On hearing the alarm leave the building immediately following the escape route and fire wardens. Do not use the lifts.

CONFERENCE BUS SCHEDULE

EMBO Conference Series: Visualizing Biological Data (VIZBI 2012)

Date	Time	from	to
06 March 2012			
	08:15	Main Station	EMBL
	08:15	Neckarmünzplatz	EMBL
	08:20	Peterskirche	EMBL
	08:25	Crowne Plaza/ back entrance	EMBL
EMBL - Downtown			
	21:00	EMBL	downtown, all stops
	22:00	EMBL	downtown, all stops
07 March 2012			
Downtown - EMBL			
	09:00	Main Station	EMBL
	09:00	Neckarmünzplatz	EMBL
	09:05	Peterskirche	EMBL
	09:10	Crowne Plaza/ back entrance	EMBL
EMBL - Downtown			
	20:00	EMBL	downtown, all stops
08 March 2012			
Downtown - EMBL			
	08:30	Main Station	EMBL
	08:30	Neckarmünzplatz	EMBL
	08:35	Peterskirche	EMBL
	08:40	Crowne Plaza/ back entrance	EMBL
EMBL - Downtown			
	17:30	EMBL	downtown, all stops
	17:30	EMBL	Frankfurt International Airport
Main Station	The main train station is west of the town centre. The EMBL shuttle bus will stop in front of McDonald's to the left of the main entrance.		
Neckarmünzplatz	This is the main tourist bus stop on north-eastern edge of the town centre. The EMBL shuttle bus will stop in the drop-off/boarding area.		
Peterskirche	This is a public bus stop on the south-eastern edge of the town centre. The bus stop is in the Friedrich-Ebert-Anlage just above Peterskirche (St. Peter's church). The EMBL shuttle bus will stop slightly before the regular bus stop.		
Crowne Plaza	The hotel is on south-western corner of the town centre (Kurfürstenanlage/ Adenauerplatz). The EMBL shuttle bus will stop at the back of the hotel next to the Lufthansa Airport Shuttle stop.		

Please make sure that you are at the bus stop 5 minutes before the departure time.

The buses are with the Hoffmann Reisen company



PUBLIC BUS

A one way bus fare costs 2.20 Euro to be paid in cash to the bus driver.

Bus 19 / SF

Departs from the EMBL bus stop just outside the Advanced Training Centre to Heidelberg "*Bismarckplatz*".

Bus 19/SF (SpezialFahrt)			
Morning		Afternoon	
Departure Bismarckplatz*	Arrival EMBL	Departure EMBL	Arrival Bismarckplatz*
		14:11	14:30
06:33	06:49	16:11	16:30
07:33	07:49	17:11	17:30
08:33	08:49	18:11	18:30
09:33	09:49	19:11	19:30
<p>*Bismarckplatz - is the main bus and tramway stop in Heidelberg city centre.</p> <p>Please note: this bus does not run at the weekend or public holidays.</p>			

Bus 39 (schedule available on the info board)

Departs from the Max-Planck-Institut für Kernphysik "*Kernphysikalisches Institut*" or "*Bierhelderhof*" to Heidelberg "*Bismarckplatz*".

Route One to bus stop: The quickest way to get to the bus stop at the Max-Planck-Institut is by walking uphill from the Advanced Training Centre, past the EMBL main entrance, until you come to an iron gate. Go through the gate and simply follow the road until you have crossed the Max-Planck-Institute campus. After passing through another gate you are on the main road where the bus passes. The bus stop is across the road (slightly left). The walk from the ATC to the bus stop takes approximately 12 minutes.

Please note: You need a Staff Access Card to go through the gates between EMBL and MPI – on a week-day, an EMBL or MPI employee will let you through sooner or later, but at the weekend or a public holiday you might not be able to get through. In this case it is advisable to take Route Two to "*Bierhelderhof*" bus stop.

Route Two to bus stop: To reach the "*Bierhelderhof*" bus stop, depart EMBL via the Meyerhofstrasse, turn right onto Bierhelderhofweg, continue past the dairy farm to the top of the hill and take the next right just after a private property. The bus stop can be found at the beginning of this road. The walk from the ATC to the bus stop takes approximately 20 minutes.

TAXI

You can also call a local taxi on: **Taxi Centrale** +49 (0) 6221/302030

CAR

If you are travelling by car we recommend parking in the car park P1, at the beginning of Meyerhofstrasse. The car park is free of charge.

GPS Address: Meyerhofstrasse 1, 69117 Heidelberg

TRAIN

For train travel times check the German Rail website. www.deutschebahn.de
(Timetable to Frankfurt International available on the info board)

COACH TO FRANKFURT INTERNATIONAL AIRPORT

Thursday, 8 March 17:30

EMBL Bus Stop – Frankfurt International Airport

A coach ticket costs 25 Euro and is available at the registration desk from the second day of the conference.

AIRPORT TRANSPORT

Private Taxi Airport Transfer

TLS Transfer & Limousine-Service

Tel: +49 (0)6221-770077

Fax: +49 (0)6221-770070

<https://www.tls-heidelberg.de/en/>

York Airport Service

Tel: +49 (0) 6226 990135

E-mail: york-airport-service@t-online.de

PMJ Fahrservice

<http://www.pmj-fahrservice.de>

info@pmj-fahrservice.de

Tel.: 06223 / 7399020, Mobile: 0163 / 7232348

Lufthansa Bus Service (schedule available on the info board)

Tel: +49 (0) 6152-97690-99 for reservations

(Monday - Friday 7.00 – 18.00, Saturdays 8.00 – 13.00)

Email: info@frankfurt-airport-shuttles.de

<http://www.transcontinental-group.com/en/frankfurt-airport-shuttles>

The bus runs once an hour, seven days a week between Terminal 1, Arrival Area, Hall B, Exit B4-5, Frankfurt International Airport and downtown Heidelberg Crowne Plaza Hotel, Kurfürstenanlage 1–3, Heidelberg (city centre).

You do not need to be a Lufthansa passenger in order to use this service.

The shuttle has 8 seats and therefore we recommend early booking in order to reserve a seat. The fares are:

	Lufthansa Passengers	Other
One Way Ticket	20 EUR	22 EUR
Return Ticket	38 EUR	40 EUR

FAST-FORWARD AND POSTER SESSION INFORMATION

In order to guarantee the smooth running of **your Fast-Forward Session**, please make sure you sit in the numbered seat reserved for you in the front rows.

Your Fast-Forward Session seat number and your poster board number correspond to the number in your ID letter-number combination, e.g.

A 01

The poster boards are located in the Foyer in the Advanced Training Centre. The sizes of our poster boards are 1.05 m wide x 2.30 m high. Pins for fixing the posters are available on the board. If you need additional pins, please ask at the registration desk.

CATERING

All meals and coffee breaks are included in the registration fee. Our catering staff will prepare a wide variety of vegetarian meals, meat and fish dishes, soups, pasta, fresh fruit and vegetables, as well as a variety of desserts.

PLEASE WEAR YOUR BADGE AT ALL TIMES WHEN SERVING YOURSELF.

NO FOOD OR DRINKS ARE ALLOWED IN THE AUDITORIUM.

NO FOOD OR DRINKS ARE ALLOWED ON THE RAMPS OF THE HELIX.

Coffee breaks will take place in the Foyer.

Lunch will be served in the Foyer.

Dinner will be served in the EMBL Canteen on Tuesday 6 March and in the Foyer on Wednesday 7 March.

QUESTIONNAIRES & BADGES

Please help us with the organisation of future conferences by giving us your **feedback** on this meeting. After the meeting we will be sending the questionnaire link per mail to all participants.

Please help us **to be green** by returning the reusable **plastic holder of your badge** at the end of the conference at the registration desk.

PHOTOGRAPHY

During the conference an EMBL Photographer will be taking photographs. If you would not like to appear in these, please inform the photographer or a member of the Course and Conference Office.

EMBL MERCHANDISE

EMBL Merchandise is available during weekdays. If you are interested in purchasing an EMBL souvenir (products presented in the glass display in the registration area), please ask at the registration desk for more information.

USEFUL TELEPHONE NUMBERS

Internal Numbers:

EMBL's Advanced Training Centre registration desk	8752
Course & Conference Office	8797/ 8908
Switchboard/Security	8100
Ambulance	112

CONFERENCE TELEPHONE FAX & E-MAIL

The conference **telephone number** where you can be reached or where messages can be left for you is: +49 6221 387 8752.

The conference **fax number** is: +49 6221 387 8158. Faxes or messages can be picked up at the registration desk.

WiFi is available everywhere on the premises (no password required, just log on as ATC Guest). It is possible to access your emails via the terminals in the E-Lounge on Helix A.

CONTACT: CONFERENCE OFFICER

If you have any further questions please contact the conference officer at the registration desk. There will be a member of the conference office at the registration desk throughout the entire conference.



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USEFUL GERMAN EXPRESSIONS

Hello	Hallo
Goodbye	Auf wiedersehen
Good morning	Guten Morgen
Good afternoon	Guten Tag
Good evening	Guten Abend
Good night	Gute Nacht
I'm sorry	Tut mir leid
Excuse me...	Entschuldigen Sie
How are you?	Wie gehts?
I'm fine thanks. And you?	Mir geht es gut , danke, und dir/Ihnen?
What is your name	Wie heisst du? Wie heissen Sie?
My name is	Ich heisse
Do you speak English	Sprechen sie Englisch?
I don't understand	Ich verstehe nicht
Please speak more slowly	Können sie bitte langsamer sprechen
Thank you	Danke schön
Where is the toilet?	Wo ist die Toilette?
Please call me a taxi	Bitte rufen sie mir ein Taxi
How do I get to....?	Wie komme ich zum/zur.....?
A beer/two beers please	Ein Bier/zwei Bier bitte?
A glass of red/white wine please	Ein Glas Rot?Weisswein bitte
The menu, please	Die Speisekarte, bitte
Is there a local speciality?	Gibt es eine Spezialität aus dieser Gegend?
I'm Vegetarian	Ich bin Vegetarier
It was delicious	Es war hervorragend
The bill, please	Die Rechnung, bitte
I have a headache	Ich habe Kopfschmerzen
I have a sore throat	Ich habe Halsschmerzen
My stomach hurts	Ich habe Magenschmerzen
I'm allergic to	Ich bin allergisch gegen
I need a doctor who speaks English	Gibt es einen Arzt, der Englisch spricht?

Enjoy your stay in Heidelberg!

