16.00-21.00	Registration
	Opening Remarks
18.30-19.30	Opening Lecture: Fungus-host interactions during Candida
	colonisation and infection
	Alistair Brown, United Kingdom
19.30	Welcome Reception

SCIENTIFIC PROGRAM • TUESDAY, 19 SEPTEMBER

19.30	welcome Reception
SCIENTII	FIC PROGRAM • WEDNESDAY, 20 SEPTEMBER
Systems & Syn 08.30	thetic Biology Collateral sensitivity of antibiotic resistant microbes
00.50	Csaba Pál, Hungary
08.55	Oral presentation selected from abstracts
09.15	The role of xenogeneic silencers in (pro-) phage-host interaction Julia Frunzke, Germany
Towards Biotec	chnology – Metabolic Modeling & Genome/Enzyme Engineering
09.40	Lessons from (meta)genomes - Novel biocatalysts and compounds Karl-Erich Jaeger, Germany
10.05	Oral presentation selected from abstracts
10.25-11.00	Coffee Break
Towards Biotec	chnology – Metabolic Modeling & Genome/Enzyme Engineering
11.00	Microbial solvent formation revisited by comparative genome
11.00	analysis
	Anja Poehlein, Germany
11.25 11.45	Oral presentation selected from abstracts Oral presentation selected from abstracts
12.05	Novel halohydrin dehalogenases by database mining from
12.00	identification to protein engineering Anett Schallmey, Germany
12.30	Oral presentation selected from abstracts
12.50-13.30	Lunch Break
13.30-15.00	Poster Session I
Omics at its be	st – Global Gene Regulation & Proteome Dynamics
15.00	Regulatory RNAs in <i>Helicobacter pylori</i> and <i>Campylobacter jejuni</i> Cynthia Sharma, Germany
15.25	Oral presentation selected from abstracts
15.45	Regulatory network controlling sporulation and stress response
	in the enteropathogen Clostridium difficile
1010	Isabelle Martin-Verstraete, France
16.10 16.30	Oral presentation selected from abstracts
16.50-17:20	Oral presentation selected from abstracts Coffee Break
10.00 11.20	

	analysis
	Anja Poehlein, Germany
11.25	Oral presentation selected from abstracts
11.45	Oral presentation selected from abstracts
12.05	Novel halohydrin dehalogenases by database mining from
	identification to protein engineering
	Anett Schallmey, Germany
12.30	Oral presentation selected from abstracts
12.50-13.30	
13.30-15.00	Poster Session I
Omice at its be	st - Global Gene Regulation & Proteome Dynamics
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15.25	Oral presentation selected from abstracts
15.45	Regulatory network controlling sporulation and stress respo
20110	in the enteropathogen <i>Clostridium difficile</i>
	Isabelle Martin-Verstraete, France
16.10	Oral presentation selected from abstracts
16.30	Oral presentation selected from abstracts
16.50-17:20	Coffee Break
	st - Global Gene Regulation & Proteome Dynamics (continue
17.20	How to make a fruiting body: comparative genomics and
	transcriptomics in filamentous fungi
47.45	Minou Nowrusian, Germany
17.45	Oral presentation selected from abstracts

SCIENTIFIC PROGRAM • WEDNESDAY, 20 SEPTEMBER

18.05	Bacteriophage diversity in human bacterial diseases
	Martha Clokie, United Kingdom
18.30	Oral presentation selected from abstracts
18:50	End of 1st day

SCIENTIFIC PROGRAM • THURSDAY, 21 SEPTEMBER

MetaOmics &	Microbial Ecosystems	
08.30	The soil resistome: Impact of human activities on the diversity	
	and abundance of antibiotic resistance genes in the soil	
	Pascal Simonet, France	
08.55	Oral presentation selected from abstracts	
09.15	Mining and expressing biosynthetic gene clusters	
	from soil metagenomes	
	Mark Liles, United States	
09.40	Oral presentation selected from abstracts	
10.00	Activity-based discovery of novel enzymes from marine	
	environments	
	Peter Golyshin, United Kingdom	
10.25-11.00	Coffee Break	

Infectiongenomics & Host Microbe Interplay

inicononigenon	incodorigenomics & nost wicrobe interplay		
11.00	Pathogen heterogeneity in host tissue - impact on disease		
	progression and treatment		
	Dirk Bumann, Switzerland		
11.25	Oral presentation selected from abstracts		
11.45	Tba		
	Sebastian Suerbaum, Germany		
12.10	Oral presentation selected from abstracts		
12.30	Genomic approaches to invasive fungal infections		
	Oliver Kurzai, Germany		
12.55-13.30	Lunch Break		
13.30-15.00	Poster Session II		
Infectiongenon	Infectiongenomics & Host Microbe Interplay (continued)		

13.00	Overcoming nost (innate) infinitine barriers.	
	Pathogenic microbes use related survival strategies	
	Peter Zipfel, Germany	
15.25	Oral presentation selected from abstracts	
15.45	An OMICs view of physiology and pathophysiology of	
	Staphylococcus aureus	
	Uwe Völker, Germany	

Microbial Evolution, Population Genomics & Epidemiology

16.10	Bacterial within-host evolution and implications for
	transmission analysis
	Xavier Didelot, United Kindgom
16.35	Oral presentation selected from abstracts
16.55-17.25	Coffee Break

SCIENTIFIC PROGRAM • THURSDAY, 21 SEPTEMBER

Microbial Evolution, Population Genomics & Epidemiology (continue	
17.25	Bacterial pathogens from population biology to genomic epidemiology
	Sylvain Brisse, France
17.50	Oral presentation selected from abstracts
18.10	Oral presentation selected from abstracts
18.30	Oral presentation selected from abstracts
18.50	End of 2 nd day
19.30	Conference Dinner

SCIENTIFIC PROGRAM • FRIDAY, 22 SEPTEMBER

MetaOmics &	Microbial Ecosystems
08.30	Metagenomic analysis for crop production
	Trevor Charles, Canada
08.55	Oral presentation selected from abstracts
09.15	Sub polar sediment comparative metagenomics and applied
	ecosystem functions
	Sara Sjöling, Sweden
09.40	Oral presentation selected from abstracts
10.00	Driving forces of microbial community structure,
	diversity and function in the marine realm
	Bernd Wemheuer, Australia
10 25-11 00	Coffee Break

Omics at its be	st - Global Gene Regulation & Proteome Dynamics
11.00	Interactions between phage and bacteria studied on
	pan-genomic scales
	Rotem Sorek, Israel
11.25	Oral presentation selected from abstracts
11.45	Lessons learned from deep proteomics and
	proteogenomics of model bacteria
	Boris Macek, Germany
12.10	Oral presentation selected from abstracts
12.30	Lunch Break

Microbiome Host Interaction	
13.05	Oral presentation selected from abstracts
13.30	Post-genome analyses of a plant pathogenic bacterium
	Cornelius Schmidtke, Germany
13.55	Oral presentation selected from abstracts
14.15	Microbiome analysis of the human gut and the ocean
	Peer Bork, Germany
14.40	Closing remarks
14.50	End of conference