

Programm

Freitag, 20. Oktober

13.00 – 14.00	Get together & Lunch	
14.00 – 15.30	SESSION 1 / Begrüßung <u>G. Cario</u> , M. Stanulla, M. Schrappe <i>High interleukin 15 expression characterizes childhood acute lymphoblastic leukemia with involvement of the central nervous system</i>	Chair: W.-D. Ludwig [T09]
	<u>S. Breit</u> , M. Muckenthaler, A. Kulozik <i>Correlation of NOTCH1 mutational status and mRNA expression profiles in childhood precursor T-cell lymphoblastic leukemia</i>	[T11]
	<u>P. Rhein</u> , R. Kirschner-Schwabe, R. Spang, L. Karawajew, C. Hagemeier, W.-D. Ludwig, K. Seeger <i>Glucocorticoid therapy targets proliferation, differentiation and Bcl-2 dependent survival signaling in ALL blasts</i>	[T10]
	<u>J. Duyster</u> <i>Die genetische Grundlage für die Empfindlichkeit der Ph+ALL gegenüber dem Kinaseinhibitor Imatinibmesylat</i>	[T14]
15.30 – 16.00	Kaffeepause	
16.00 – 17.00	SESSION 2 <u>S. Bohlander</u> , A. Krause, P. Greif <i>The CALM/AF10 fusion: Identification of target genes and important protein interactors</i>	Chair: R. Kirschner-Schwabe [T15]
	<u>C. Buske</u> <i>Acute myeloid leukemia is propagated by a leukemic stem cell with lymphoid characteristics in a mouse model of CALM/AF10 positive leukemia</i>	[T17]
	<u>R. Marschalek</u> <i>Gene expression profiling of AF4 knock out cells</i>	[T13]
17.00 – 17.20	Präsentation AFFYMETRIX <u>M. Prüss</u> , E. Funk <i>Microarrays der nächsten Generation: GeneChip Exon und Tiling Arrays</i>	
17.20 – 18.00	Freie Diskussion / Ausblick NGFN-3	
20.00	Dinner im „Gourmet Sud“ (Selmaplatz 1-2 in 14163 Berlin-Dahlem)	

Samstag, 21. Oktober

09.30 – 11.00	SESSION 3	Chair: C. Hagemeier
	<u>F. Rücker, L.Bullinger</u> [T06] <i>Parallel analyses disclose novel genomic imbalances in acute myeloid leukaemia with complex karyotypes</i>	
	<u>A. Neubauer, S. Teichler, N. Hagner</u> [T18] <i>Genomic analysis of retinoic acid signalling in acute myeloid leukemia</i>	
	<u>C. Hömme, C. Müller-Tidow, H. Serve</u> [T07] <i>PML-RARalpha induced chromatin modifications regulate leukemogenic genes and functional networks</i>	
	<u>A. Peer Zada, G. Behre</u> [T16] NN	
11.00 – 11.15	Kaffeepause	
11.15 – 13.00	SESSION 4	Chair: R. Spang
	<u>M. Muckenthaler, A. Kulozik</u> [T11] <i>A sensitive array for microRNA expression profiling (miChip) based on locked nucleic acids (LNA)</i>	
	<u>C. Lottaz, R. Spang</u> [T12] <i>The ALC-WWW-Platform Diskussion der Meta-Analysen der Genexpressionsdaten</i>	
ab 13.00	Abschließendes gemeinsames Mittagessen	