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The hepatitis C continuum of care among HIV infected individuals in EuroSIDA

<u>Sarah Amele</u>, Lars Peters, Jens D. Lundgren, Jürgen K. Rockstroh, Maryana Sluzhynska, Alexei Yakovlev, Alexandra Scherrer, Pere Domingo, Jan Gerstoft, Jean-Paul Viard, Robert Zangerle, Robert Flisiak, Sanjay Baghani, Matti Ristola, Clifford Leen, Elzbieta Jablonowska, Gilles Wandeler, Hans-Jürgen Stellbrink, Karolin Falconer, Antonella d'Arminio Monforte, Andrzej Horban and Amanda Mocroft

on behalf of the EuroSIDA study group

Presenter Disclosure Information

Sarah Amele

disclosed no conflict of interest.

Background

- Globally 2.3 million HIV/HCV co-infected, majority are injecting drug users (IDU)¹
- WHO goal of eliminating viral hepatitis as a public health threat by 2030¹ - HCV continuum of care (CoC) is an essential framework to monitor and evaluate progress in achieving these targets
- Also useful to identify leaks/breaks in the continuum that need to be addressed to ensure individuals transition through all stages and achieve sustained virologic response (SVR)
- More work is required to develop a standardised continuum for HCV infected people living with HIV (PLWH) to allow cross country or population comparisons

¹World Health Organization. Global hepatitis report, 2017. 2017. 62 p.

Aims

- To develop and evaluate a HCV continuum of care in HIV coinfected individuals across Europe at 1/1/2015
- Look at regional differences in the continuum
- Examine the proportion of individuals genotyped and with a fibrosis marker
- Describe factors associated with being HCV-RNA tested once already anti-HCV positive

EuroSIDA study

- Large prospective observational cohort study with over 22,000 HIV-positive individuals
- Inclusion criteria
 - HIV positive
 - Under follow-up at 1/1/2015
 - Anti-HCV positive before 1/1/2015
 - >16 years of age

Regions

• South: Greece, Israel, Italy, Portugal, Spain, Argentina

 Central West: Austria, Belgium, France, Germany, Luxembourg, Switzerland

 North: Denmark, Finland, Iceland, Ireland, Netherlands, Norway, Sweden, United Kingdom

Central East: Bosnia-Herzegovina,
 Croatia, Czech Republic, Hungary,
 Poland, Romania, Serbia, Slovakia,
 Slovenia

East: Belarus, Estonia, Georgia,
 Latvia, Lithuania, Russia, Ukraine

Methods - Definitions

Stage	Definition				
1: anti-HCV +ve	Anti-HCV positive, HCV-RNA positive, HCV genotyped or received HCV treatment before 1/1/2015				
2: Ever HCV-RNA tested	Ever HCV-RNA tested, HCV genotyped or received HCV treatment before 1/1/2015				
3: Currently HCV-RNA +ve	Most recent HCV-RNA test before 1/1/2015 was positive, HCV genotyped but not treated before 1/1/2015, started treatment for the first time after 1/1/2015 or first HCVRNA test result after 1/1/2015 is positive and never treated.				
4: Ever HCV-RNA +ve	Ever had a positive HCV-RNA test, received HCV treatment or HCV genotyped before 1/1/2015				
5: Ever received treatment	Started HCV treatment on or before 1/1/2015				
6: Treatment completed	Completed HCV treatment on or before 1/1/2015				
7: FU HCV-RNA available	HCV-RNA test after completing treatment (HCV-RNA test data included for duration of FU to allow for assessment of SVR)				
8: SVR	HCV-RNA negative test at least 12 or 24 weeks post treatment (for IFN-free and IFN-based therapy, respectively)				

Methods - Statistics

- Tested for regional differences within each stage of continuum
- Identify predictors of being HCV-RNA tested:

Anti-HCV+ve



HCV-RNA tested?

Logistic regression

Model adjusted for:

age, sex, ethnicity, region in Europe, CD4 count, HIV-RNA, previous use of cART, mode of HIV transmission, mode of HCV transmission, stage of liver fibrosis, hepatitis B co-infected and prior AIDS diagnosis

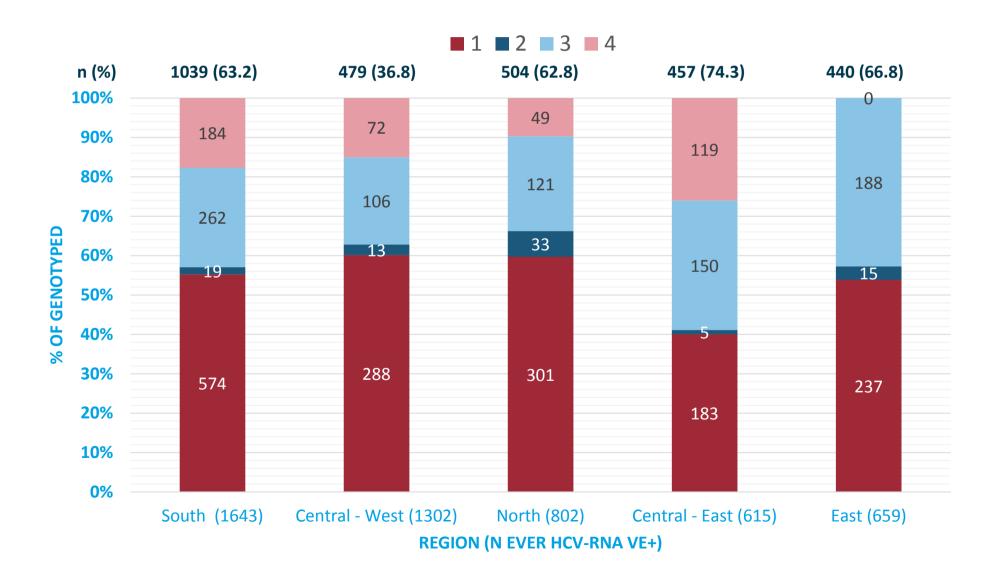
Characteristics at 1/1/2015

Region n (%)		Overall 6985 (100.0)	South 1910 (27.3)	Central - West 1614 (23.1)	North 966 (13.8)	Central - East 925 (13.2)	East 1570 (22.5)	
Varia	bles	%						
Gender	Male	71.6	72.5	75.1	78.4	72.0	62.7	
Ethnicity	White	88.3	94.3	68.2	82.3	98.6	99.4	
Fibrosis	<f3< th=""><th>74.7</th><th>73.9</th><th>80.9</th><th>69.5</th><th>69.4</th><th>75.7</th></f3<>	74.7	73.9	80.9	69.5	69.4	75.7	
	≥F3*	12.9	15.4	11.6	12.8	9.8	13.1	
HIV risk group	MSM	21.0	16.5	32.2	42.0	21.2	2.0	
	IDU	54.2	60.1	40.0	37.9	59.1	68.9	
cART	Yes	88.8	95.3	80.4	95.9	95.0	81.7	
		Median						
Age		47	50	51	51	41	37	
CD4 count (cells/mm³)		278	297	332	234	244	267	

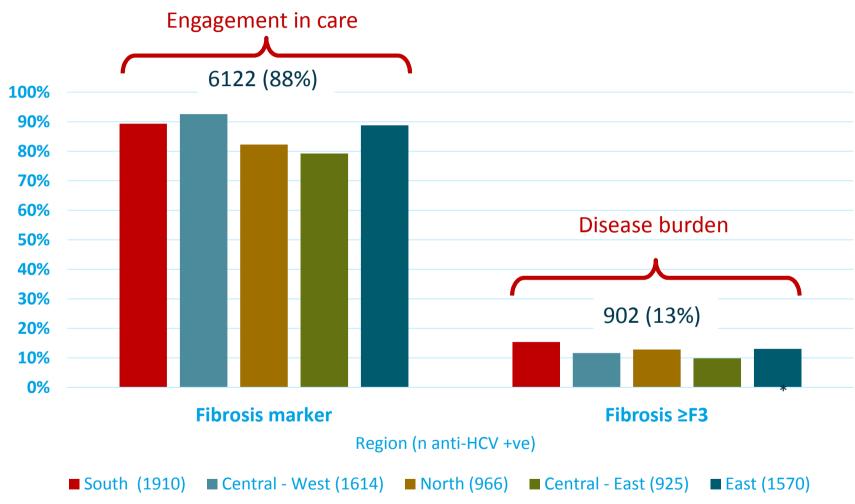
^{*}Either a biopsy (≥METAVIR stage F3), APRI (score >1.5), hyaluronic acid (>160ng/mL) or FibroScan (>9.5kPa) test during follow-up

Evidence of difference between regions for all variables (p<0.001)

HCV genotype



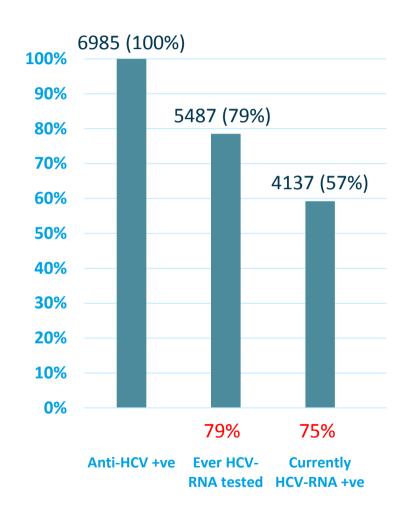
Fibrosis



^{*}Either a biopsy (≥METAVIR stage F3), APRI (score >1.5), hyaluronic acid (>160ng/mL) or FibroScan (>9.5kPa) test during follow-up

Evidence of difference between regions for all variables (p<0.001)

Overall CoC at 1/1/2015



RNA +ve treatment completed available

Treatment

FU HCV-RNA

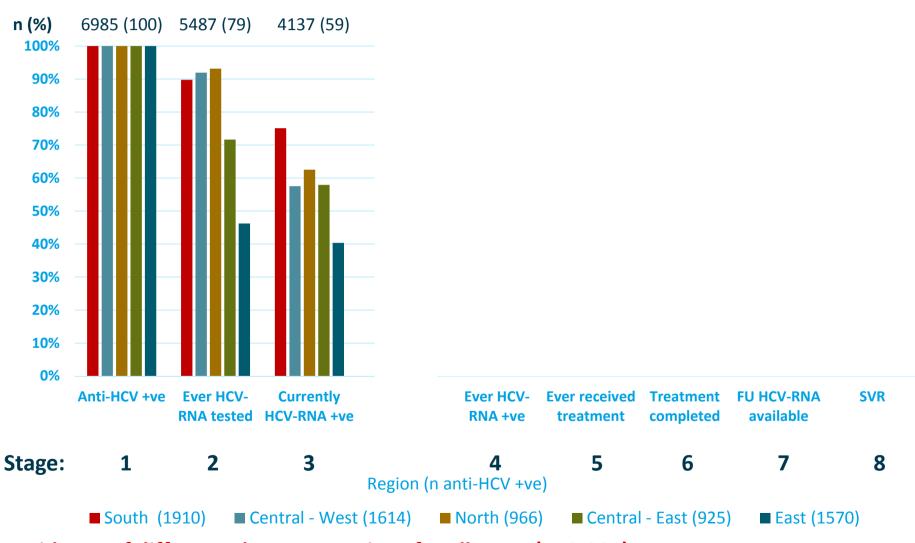
SVR

Ever received

Stage: 1 2 3 4 5 6 7 8

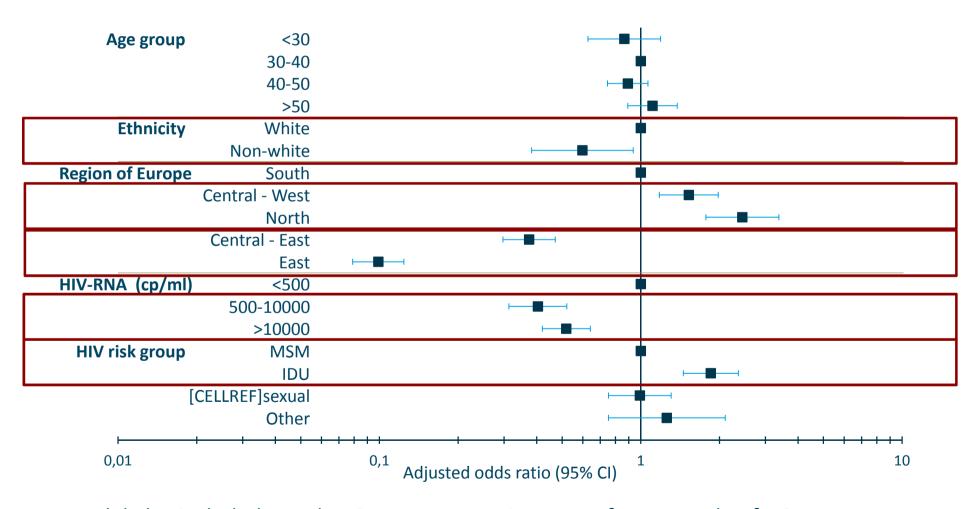
Ever HCV-

CoC by region at 1/1/2015



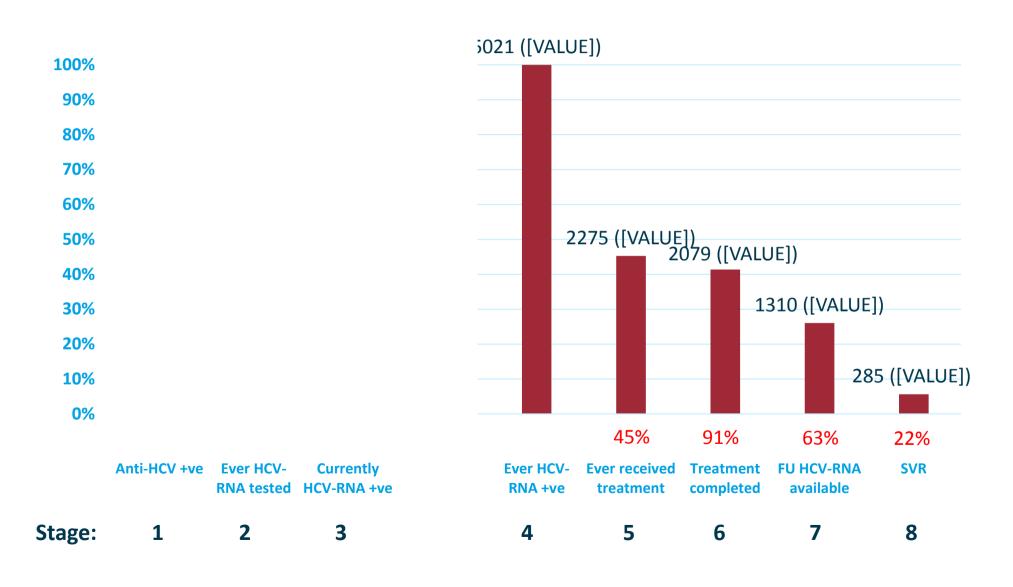
Evidence of difference between regions for all stage (p<0.001)

Odds of being HCV-RNA tested

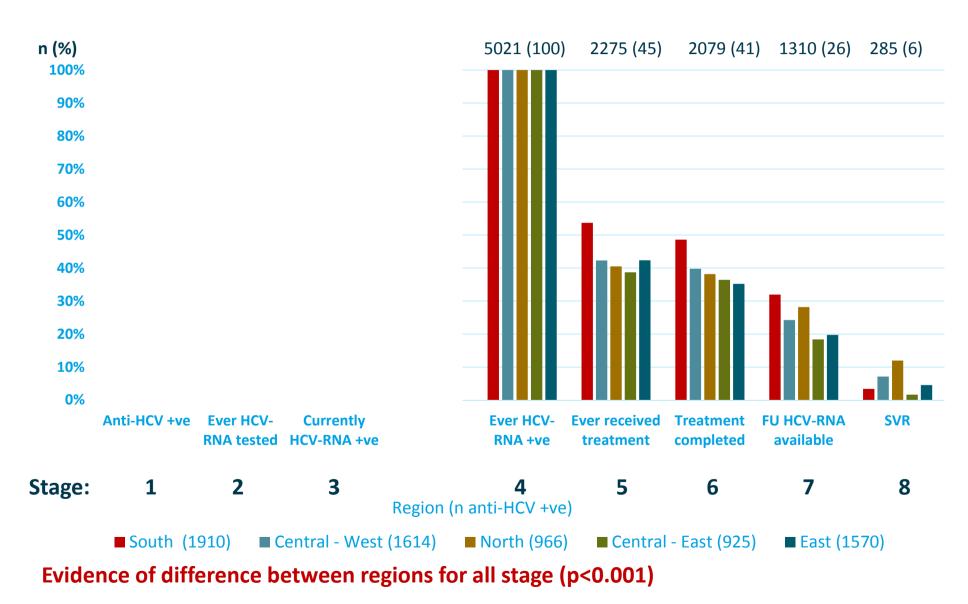


Model also included: gender, CD4 count, previous use of cART, mode of HCV transmission, stage of liver fibrosis, hepatitis B co-infection and prior AIDS diagnosis

Overall CoC at 1/1/2015



CoC by region at 1/1/2015

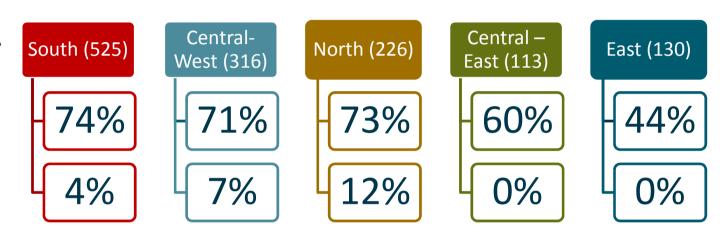


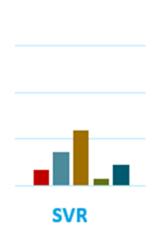
Low SVR?

Region (n FU HCV-RNA available):

Genotype 1 or 4
Genotyped

IFN-free regimen
FU HCV-RNA available





- High proportion of individuals with genotype 1 or 4 that are hard to treat with IFN based regimens.
- Very low number of individuals received
 IFN-free treatments

Limitations

- Cohort individuals not necessarily representative of all HCV infected individuals (vulnerable groups, incarcerated population etc. not included in cohort)
- Not everyone has a HCV-RNA measurement at 12/24 weeks after completing treatment
- CoC was examined at a fixed point in time and therefore may be different if repeated now
- Differences in treatment guidelines, access to care and patient management approaches within countries and regions
- Did not look at the whole continuum, undiagnosed population not estimated

Summary

- Significant proportion of individuals lost at all stages of our proposed HCV continuum, with large variations between regions of Europe
- Region, ethnicity, and HIV viral load found to affect the odds of being HCV-RNA tested after anti-HCV +ve test
- Large gap between testing and starting treatment likely due to multifactorial reasons, including access to treatment, lack of DAAs and patient characteristics

Conclusions

- The proportion of individual achieving SVR was low, in part due to our definitions but improvements in HCV treatment cannot be realised unless barriers to care are addressed
- Our proposed definitions would allow repeated analyses over time to monitor changes in the break points
- Further work on defining the HCV CoC is urgently needed

The EuroSIDA Study Group

The multi-centre study group, EuroSIDA (national coordinators in parenthesis).

South: Greece: (P Gargalianos), G Xylomenos, K Armenis, Athens General Hospital "G Gennimatas"; H Sambatakou, Ippokration General Hospital, Athens. Israel: (I Yust), D Turner, M Burke, Ichilov Hospital, Tel Aviv; E Shahar, G Hassoun, Rambam Medical Center, Haifa; H Elinav, M Haouzi, Hadassah University Hospital, Jerusalem; D Elbirt, ZM Sthoeger, AIDS Center (Neve Or), Jerusalem. Italy: (A D'Arminio Monforte), Istituto Di Clinica Malattie Infettive e Tropicale, Milan; R Esposito, I Mazeu, C Mussini, Università Modena, Modena; F Mazzotta, A Gabbuti, Ospedale S Maria Annunziata, Firenze; V Vullo, M Lichtner, University di Roma la Sapienza, Rome; M Zaccarelli, A Antinori, R Acinapura, M Plazzi, Istituto Nazionale Malattie Infettive Lazzaro Spallanzani, Rome; A Lazzarin, A Castagna, N Gianotti, Ospedale San Raffaele, Milan; M Galli, A Ridolfo, Osp. L. Sacco, Milan. Portugal: (L Caldeira), Hospital Santa Maria, Lisbon; K Mansinho, Hospital de Egas Moniz, Lisbon; F Maltez, Hospital Curry Cabral, Lisbon. Spain: (JM Gatell), JM Miró, Hospital Clinic Universitari de Barcelona, Barcelona; S Moreno, J. M. Rodriguez, Hospital Ramon y Cajal, Madrid; B Clotet, A Jou, R Paredes, C Tural, J Puig, I Bravo, Hospital Germans Trias i Pujol, Badalona; P Domingo, M Gutierrez, G Mateo, MA Sambeat, Hospital Sant Pau, Barcelona; JM Laporte, Hospital Universitario de Alava, Vitoria-Gasteiz. Argentina: (M Losso), M Kundro, Hospital JM Ramos Mejia, Buenos Aires.

Central West: Austria: (B Schmied), Pulmologisches Zentrum der Stadt Wien, Vienna; R Zangerle, Medical University Innsbruck, Innsbruck. Belgium: (N Clumeck), S De Wit, M Delforge, Saint-Pierre Hospital, Brussels; E Florence, Institute of Tropical Medicine, Antwerp; L Vandekerckhove, University Ziekenhuis Gent, Gent. France: (J-P Viard), Hôtel-Dieu, Paris; P-M Girard, Hospital Saint-Antoine, Paris; C Pradier, E Fontas, Hôpital de l'Archet, Nice; C Duvivier, Hôpital Necker-Enfants Malades, Paris. Germany: (J Rockstroh), Universitäts Klinik Bonn; G Behrens, Medizinische Hochschule Hannover; O Degen, University Medical Center Hamburg-Eppendorf, Infectious Diseases Unit, Hamburg; HJ Stellbrink, IPM Study Center, Hamburg; C Stefan, JW Goethe University Hospital, Frankfurt; J Bogner, Medizinische Poliklinik, Munich; G. Fätkenheuer, Universität Köln, Cologne. Luxembourg: (T Staub), R Hemmer, Centre Hospitalier, Luxembourg. Switzerland: (A Scherrer), R Weber, University Hospital Zurich; M Cavassini, University Hospital Lausanne; A Calmy, University Hospital Geneva; H Furrer, University Hospital Bern; M Battegay, University Hospital Basel; P Schmid, Cantonal Hospital St. Gallen.

North: Denmark: G Kronborg, T Benfield, Hvidovre Hospital, Copenhagen; J Gerstoft, T Katzenstein, Rigshospitalet, Copenhagen; NF Møller, C Pedersen, Odense University Hospital, Odense; L Ostergaard, Skejby Hospital, Aarhus, L Wiese, Roskilde Hospital, Roskilde; L N Nielsen, Hillerod Hospital, Hillerod. Finland: (M Ristola), I Aho, Helsinki University Central Hospital, Helsinki. Iceland: (M Gottfredsson), Landspitali University Hospital, Reykjavik. Ireland: (F Mulcahy), St. James's Hospital, Dublin. Netherlands: (P Reiss), Academisch Medisch Centrum bij de Universiteit van Amsterdam, Amsterdam. Norway: (DH Reikvam), A Maeland, J Bruun, Ullevål Hospital, Oslo. Sweden: (K Falconer), A Thalme, A Sonnerborg, Karolinska University Hospital, Stockholm; A Blaxhult, Venhälsan-Sodersjukhuset, Stockholm; L Flamholc, Malmö University Hospital, Malmö. United Kingdom: (B Gazzard), St. Stephen's Clinic, Chelsea and Westminster Hospital, London; AM Johnson, E Simons, S Edwards, Mortimer Market Centre, London; A Phillips, MA Johnson, A Mocroft, Royal Free and University College Medical School, London (Royal Free Campus); C Orkin, Royal London Hospital, London; J Weber, G Scullard, Imperial College School of Medicine at St. Mary's, London; A Clarke, Royal Sussex County Hospital, Brighton; C Leen, Western General Hospital, Edinburgh.

Central East: Bosnia-Herzegovina: (V Hadziosmanovic), Klinicki Centar Univerziteta Sarajevo, Sarajevo. Croatia: (J Begovac), University Hospital of Infectious Diseases, Zagreb. Czech Republic: (L Machala), D Jilich, Faculty Hospital Bulovka, Prague; D Sedlacek, Charles University Hospital, Plzen. Hungary: (J Szlávik), Szent Lásló Hospital, Budapest. Poland: (B Knysz), J Gasiorowski, M Inglot, Medical University, Wroclaw; A Horban, E Bakowska, Centrum Diagnostyki i Terapii AIDS, Warsaw; R Flisiak, A Grzeszczuk, Medical University, Bialystok; M Parczewski, K Maciejewska, B Aksak-Was, Medical University, Szczecin; M Beniowski, E Mularska, Osrodek Diagnostyki i Terapii AIDS, Chorzow; T Smiatacz, M Gensing, Medical University, Gdansk; E Jablonowska, E Malolepsza, K Wojcik, Wojewodzki Szpital Specjalistyczny, Lodz; I Mozer-Lisewska, Poznan University of Medical Sciences, Poznan. Romania: (R Radoi), C Oprea, Spitalul Clinic de Boli Infectioase si Tropicale: Dr. Victor Babes, Bucuresti. Serbia: (D Jevtovic), The Institute for Infectious and Tropical Diseases, Belgrade. Slovenia: (J Tomazic), University Clinical Centre Ljubljana, Ljubljana.

East: Belarus: (I Karpov), A Vassilenko, Belarus State Medical University, Minsk, VM Mitsura, Gomel State Medical University, Gomel; D Paduto, Regional AIDS Centre, Svetlogorsk. Estonia: (K Zilmer), West-Tallinn Central Hospital, Tallinn; Jelena Smidt, Nakkusosakond Siseklinik, Kohtla-Järve. Georgia: (N Chkhartishvili) Infectious Diseases, AIDS & Clinical Immunology Research Center, Tbilisi Latvia: (B Rozentale), Infectology Centre of Latvia, Riga. Lithuania: (V Uzdaviniene) Vilnius University Hospital Santaros Klinikos, Vilnius; R Matulionyte, Centro poliklinika, Vilnius, Vilnius University Hospital Santaros Klinikos, Vilnius. Russia: (A Panteleev), O Panteleev, St Petersburg AIDS Centre, St Peterburg; A Yakovlev, Medical Academy Botkin Hospital, St Petersburg; T Trofimora, Novgorod Centre for AIDS, Novgorod, I Khromova, Centre for HIV/AIDS & and Infectious Diseases, Kaliningrad; E Kuzovatova, Nizhny Novgorod Scientific and Research Institute of Epidemiology and Microbiology named after Academician I.N. Blokhina, Nizhny Novogrod; E Borodulina, E Vdoushkina, Samara State Medical University, Samara. Ukraine: A Kuznetsova, Kharkov State Medical University, Kharkov; G Kyselyova, Crimean Republican AIDS centre, Simferopol; M Sluzhynska, Lviv Regional HIV/AIDS Prevention and Control CTR, Lviv.

EuroSIDA Steering Committee: J Gatell, B Gazzard, A Horban, I Karpov, M Losso, A d'Arminio Monforte, C Pedersen, M Ristola, A Phillips, P Reiss, J Lundgren, J Rockstroh, A Scherrer, Aho I, Rasmussen LD, Svedheim V, Wandaler G, Pradier C, Chkartishvili N, Matulionyte R, Oprea C, Kowalska J, Begovac J, Miro J, Guaraldi G. Chair: J Rockstroh. Study Co-leads: A Mocroft, O Kirk. EuroSIDA staff: Coordinating Centre Staff: O Kirk, L Peters, AH Fischer, A Bojesen, D Raben, D Kristensen, K Laut, JF Larsen, D Podlekareva. Statistical Staff: A Mocroft, A Phillips, A Cozzi-Lepri, L Shepherd, S Amele, A Roen, A Pelchen-Matthews.

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