

Working Group Information and Documentation

Chair: Christina Ligda

	<i>Tuesday, 2 May 2017</i>
	Arrival
	<i>Wednesday, 3 May 2017</i>
	Arrival
10:00 - 10:30	Welcome - Opening Objectives of the meeting - Report on the WG activities (C. Ligda)
10:30-11:15	Main recommendations from the Ad hoc action "Socio-economic and environmental parameters and their applicability into a tool to evaluate risks and trends "(E. Sturaro) Discussion
11:15 – 11:30	<i>Coffee break</i>
11:30 -12:30	ERFP ad hoc action improving communication tools : Report from meeting (14 -15 March, Ljubljana) (G. Osojnic) Discussion
12:30 – 13:15	Transboundary breeds in EFABIS: - What kind of information exists on in situ / ex situ conservation (Z. Ducheve) Discussion : Using the information (and risk status) to promote genetic diversity management between the countries
13:15-13:30	Election of Chair (<i>ERFP Secretariat</i>)
13:30 – 15:00	<i>Lunch</i>
15:00 – 15:45	Mountain breeds in Europe, and how EFABIS data can be used to assess their common characteristics, evolution etc. What are the gaps in existing data, how to analyze (E. Charvolin) Discussion
15:45-16:15	Update on DAD-IS situation (R. Baumung)
16:15– 16:30	<i>Coffee break (if needed)</i>
16:30 -17:15	DAD –IS Developments and European node / Reporting from WG members
17:15 – 17:45	CGRFA - Honeybee and pollinator fields (R. Baumung)
17:45 – 18:15	Workplan for 2017 -2018 (<i>to be approved in Tallinn</i>) (Chair)
19:30	<i>Dinner</i>

Thursday, 4 May 2017 – Joint with WG ex situ / TF in situ	
8:15-8:30	Welcome and Opening (<i>Host / S. Stojanovic</i>)
8:30-9:30	Report of the <i>Ad hoc</i> action: Development of the ERFP communication tools (<i>G. Osojnkik</i>)
9:30-10:00	Update on EUGENA Webportal development (<i>Z. Ducheve</i>)
10:00-10:30	Data to be collected for SDG indicator 2.5.1 on cryoconservation (<i>R. Baumung</i>)
10:30-10:45	Coffee break
10:45-11:15	Results of the IMAGE questionnaire/survey on genetic collections in Europe (<i>D. Duclos</i>)
11:15-12:30	IMAGE next steps – Access to gene bank data (<i>S.J. Hiemstra</i>)
Afternoon	Departure