





Biomolecular Characterization of *E. granulosus*Larval Form in Sardinian Patient Affected by Cystic Echinococcosis: Project Results

Santucciu C, Mastrandrea S, Peruzzu A, Profili S, Porcu A, Fancellu A, Carta A, Bagella G, Fiamma M, Bonelli P, Piseddu T, Masu G, Masala G

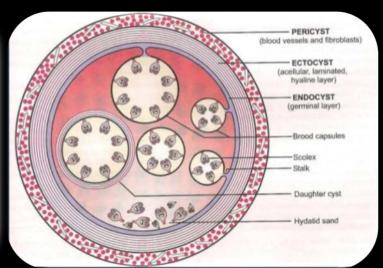
OIE and NRL for Echinococcosis, IZS della Sardegna, Italy

presenting author: Dr. Cinzia Santucciu email: cinzia.santucciu@izs-sardegna.it



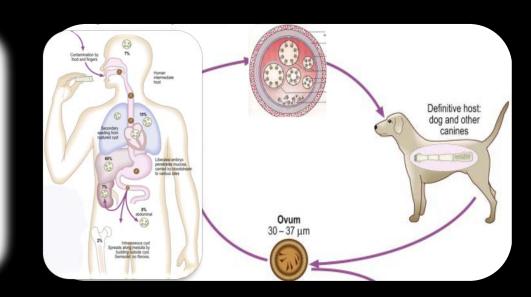


Cystic Echinococcosis (CE) is a neglected zoonotic disease caused by the larval stage of the tapeworm *Echinococcus* granulosus sensu lato (s.l.).





The parasite life cycle typically includes intermediate hosts, usually ungulates, and definitive hosts, domestic and wild canids, harboring the adult worm.



Aim of the project

- Improve the knowledge of CE
- Increase the data flow (e.g. ERCE)
- •Standardize the diagnostic and therapeutic paths and the follow up of CE patients on the base of OMS Guide Lines

34 patients with symptoms referable to CE from Sassari and Nuoro Hospital Wards were investigated 15 females 19 males **Imagine techniques** 12 negative to CE cysts 23 CE cysts positive 21 liver /2 lungs Serum analysis: ELISA Echinococcus IgG (DRG) **Echinococcus WB IgG (LDBIO)** 14 positive sera 20 negative sera

CE Pos cases were included in the European Register of Cystic Echinococcosis (ERCE)

7 patients were undergone to surgery hydatids excised: HU1-HU7



WHO classification

cysts stadium:

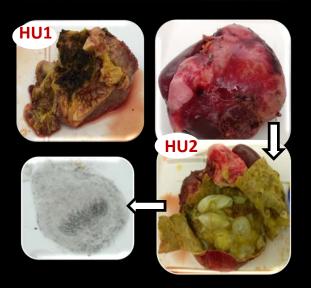
active - CE2

cysts stadium:

transitional CE3b

cysts stadium:

inactive - CE5

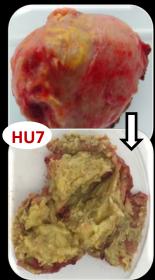












..... 7 hydatids surgically excised the external and internal structures were inspected by naked eye and a microscope

cysts	protoscolices
HU1	non-viable
HU2	non-viable
HU3	non-viable
HU4	non-viable
HU5	no
HU6	no
HU7	no



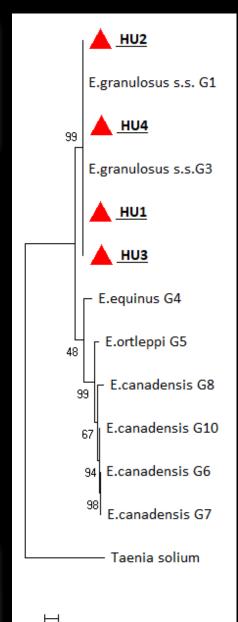


....biomelocular characterization: genotyping of larval form

PCR *E.g.s.s.* [primers: **Calreticulin** (**Cal**) gene of 1001 bp (F5':CAATTTACGGTAAAGCAT-3' R5':CCTCATCTCCACTCTCT-3')].

cysts	PCR E.g.s.s.	genotype
HU1	<i>E. granulosus</i> ss	G3
HU2	<i>E. granulosus</i> ss	G1
HU3	E. granulosus ss	G3
HU4	E. granulosus ss	G1
HU5	neg	/
HU6	neg	/
HU7	neg	/

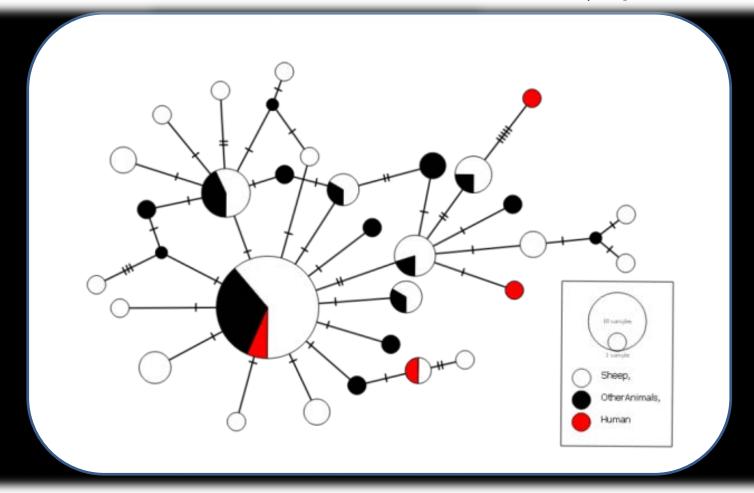
Maximum Likelihood Phylogenetic tree build on a dataset comprising 4 human isolates (HU1, HU2, HU3, HU4) of *E. granulosus* from this study and other reference sequences of *E. granulosus* s.l. retrieved from GenBank.



0.20

....biomelocular characterization: sequencing analysis of larval form

•PCR for sequencing: **Cytochrome Oxidase I (COX1)** (F5'-TTTTTTGGCCATCCTGAGGTTTAT-3' R5'-TAACGACATAACATAATGAAAATG-3') for].



Haplotype networks calculated on 83 DNA sequences of *E. granulosus* isolated in Sardinia from different animal species.

Thanks for your attention!!!