NOTE

Cetacean strandings in Italy: an unusual mortality event along the Tyrrhenian Sea coast in 2013

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ABSTRACT: An unusual mortality event involving cetaceans, mainly striped dolphins *Stenella coeruleoalba* (Meyen, 1833), occurred along the Tyrrhenian Sea coast of Italy during the first 3 mo of 2013. Based on post-mortem analyses carried out according to body condition on 66 dolphins (54% of stranded animals), several hypotheses to explain the causes of this mortality event were proposed. Although no definitive conclusions can be drawn, dolphin morbillivirus was deemed the most likely cause, although other infectious agents (including *Photobacterium damselae damselae* and herpesvirus) or environmental factors may also have contributed to this recent mortality event.

KEY WORDS: Cetacean \cdot Unusual mortality event \cdot Dolphin \cdot Morbillivirus \cdot Marine mammal \cdot Stranding

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INTRODUCTION

Cetacean unusual mortality events (UMEs) refer to a marked rise in the number of strandings as compared with historical records for the same location and period (WGMMUME 2006). In the Mediterranean Sea, 3 different dolphin morbillivirus (DMV) dieoffs have occurred in the past 25 yr, each involving 100s—if not 1000s—of striped dolphins *Stenella coeruleoalba* (Meyen, 1833). The first was particularly dramatic and occurred between 1990 and 1992 (Domingo et al. 1992, Van Bressem et al. 2001) fol-

lowed by a second one, also involving pilot whales *Globicephala melas* (Traill, 1809), between 2006 and 2008 (Fernández et al. 2008, Raga et al. 2008, Keck et al. 2010). A recent epidemic struck the coast of Spain (Soto et al. 2011) in 2011, before moving eastward and reaching Italian waters (Soto et al. 2011, Mazzariol et al. 2012, Rubio-Guerri et al. 2013a,b).

More recently, an exceptional number of cetacean strandings, 10- to 12-fold the 25 yr average recorded for the same time period (Fig. 1), along the Tyrrhenian coast of Italy between January and March 2013 prompted diagnostic post-mortem investigation.

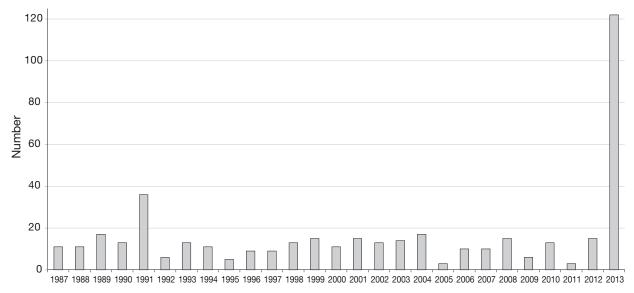


Fig. 1. Cetacean strandings recorded along the Tyrrhenian Sea coast of Italy (January to March quarter, 1987 to 2013)

METHODS AND RESULTS

A total of 122 stranded cetaceans were reported by the National Database on Cetacean Strandings (http://mammiferimarini.unipv.it/index_en.php), including 96 striped dolphins *Stenella coeruleoalba* (Meyen, 1833), 7 bottlenose dolphins *Tursiops truncatus* (Montagu, 1821), 1 fin whale *Balaenoptera physalus* (Linnaeus, 1758), 1 pilot whale *Globicephala melas* (Traill, 1809) and 3 Risso's dolphins *Grampus griseus*

(G. Cuvier, 1812). The species could not be determined in other 14 individuals (Fig. 2).

Age was estimated from total body length (TBL) in 80 striped dolphins (42 other specimens were directly disposed of by local health authorities due to poor preservation status), since direct techniques (e.g. counting growth layers in teeth and pectoral fin X-rays) were impractical because of the high number of stranded animals. Two standard approaches (Di Meglio et al. 1996, Marsili et al. 2004) for Mediterranean striped dolphins determined that 77 animals (96%) were younger than 20 yr of age, with a TBL shorter than 202 cm.

Post-mortem investigations of 66 specimens (54% of the stranded animals) were performed by various laboratories that used a standard sampling protocol according to the body conservation code (see Table 1) (Geraci & Loundsbury 2005). Tissues (brain, lung, heart, kidney, liver, lymph node, spleen and skin) were sampled, frozen at

-80°C for microbiological and biomolecular investigations to identify the main cetacean pathogens (DMV, *Brucella* spp., *Toxoplasma gondii*, *Photobacterium damselae* subsp. *damselae*, herpesvirus) and preserved in 10% neutral-buffered formalin for histopathological and immunohistochemical (IHC) analyses.

Diagnostic investigations were performed whenever possible, according to the preservation conditions of the stranded animals. Logistic difficulties

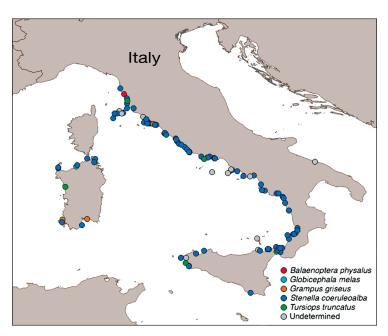


Fig. 2. Geographical distribution of recent cetacean strandings along the Italian coastline (1 January to 30 March 2013)

				Biomolecular investigations					Bacteriology		OC pollutants		
Condition		No. of animals	% of animals	DM No. tested	%	T. go No. tested	%	Herpe No. tested	svirus %	No. tested	%	No. tested	%
Code 1	Live animal	4	6.35										
Code 2	Fresh carcass	16	25.40	57	95	52	87	22	37	50	96	12	23
Code 3	Moderate decomposition	32	50.79										
Code 4	Advanced decomposition	8	12.70							Not		No	ot
Code 5	Mummified/ skeletal remains	3	4.76	Not feasible					feasible		feasible		

Table 1. Body condition score of 63 stranded cetaceans according to Geraci & Loundsbury (2005) (data not available for 3 animals) and diagnostic analysis. DMV: dolphin morbillivirus; *T. gondii: Toxoplasma gondii*; OC: organochlorine

sometimes precluded thorough examination due to the large geographic area involved in this UME and the unexpectedly high number of cetaceans found stranded. Most carcasses (63%) showed a moderateto-advanced degree of post-mortem autolysis, with empty stomachs and high loads of a range of parasites distributed throughout the body.

Biomolecular analyses for DMV (Raga et al. 2008, Di Guardo et al. 2010, Romano et al. 2011) and *Toxoplasma gondii* (Alba et al. 2013, Vitale et al. 2013) were performed in most animals triaged with codes from 1 to 4, while culture-based and molecular investigations for bacterial pathogens, including *Brucella* spp. (Alton et al. 1988, Carter 1992, OIE 2008) and *Photobacterium damselae* subsp. *damselae* (Holt et al. 1994, Osorio et al. 2000, Lozano-Leon et al. 2003, Labella et al. 2011, Woo & Bruno 2011), were attempted in animals triaged with codes from 1 to 3.

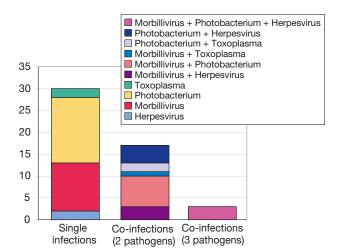


Fig. 3. Pathogens detected in necropsied cetaceans

Furthermore, single laboratories performed other related analyses, such as those against herpesvirus (VanDevanter et al. 1996, Rose 2005), which were mainly carried out on selected individuals with codes from 1 to 4 stranded along the coasts of Latium and Tuscany (22 animals, 37%) and environmental organochlorine (OC) pollutant determinations performed on a sample of 12 animals (23%) with codes from 1 to 3 stranded along the coasts of Tuscany and Sicily (Table 1).

DMV was detected by means of reverse transcription-polymerase chain reaction (RT-PCR) in 24 out of 57 animals tested (42%), mainly on samples from the brain or lungs. IHC analysis (Uchida et al. 1999) performed on animals that were RT-PCR-positive for DMV detected morbilliviral antigen in the brain of 1 dolphin. Other pathogens were frequently identified concurrently (Fig. 3).

Photobacterium damselae subsp. damselae was isolated from 31 out of 50 animals (62%), mainly from the brain, liver, lungs and lymph nodes. Toxoplasma gondii was detected by biomolecular analysis in 5 out of 52 animals (9.62%), mainly from the lungs and central nervous system (CNS), but without related pathological changes. Herpesvirus was detected in 12 out of 22 animals (54%), mainly from the CNS, lungs and spleen. No characteristic histopathological changes generally attributed to herpesvirus infection were found. No evidence of Brucella spp. was detected in any of the tissues.

Histopathology revealed chronic inflammation in several organs (CNS, lung, kidney, liver, spleen) of almost all the animals. Furthermore, follicular depletion and hyalinosis in lymphoid tissues suggested impairment of the immune system, and eosinophilic

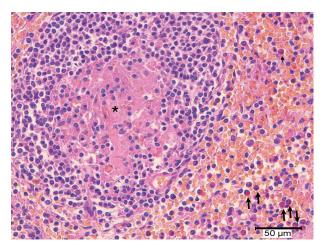


Fig. 4. Stenella coeruleoalba. A spleen probe from a dolphin morbillivirus-infected striped dolphin. Note the follicular hyalinosis (*) represented by abundant eosinophilic amorphous material filling the empty space due to severe lymphoid depletion. Furthermore, eosinophils can be noted scattered throughout the surrounding red pulp (arrows). Hematoxylin and eosin, scalebar = 50 μm

inflammatory infiltration was suggestive of a systemic reaction to parasites (Fig. 4). Finally, fibrinoid effusion and necrosis and/or hemolysis or hemorrhage in the spleen, lymph nodes, liver and kidneys were suggestive of acute septic shock and were often associated with the presence of pathogens in many animals.

Screening for OC pollutants, including hexachlorobenzene (HCB), dichlorodiphenyltrichloroethane (DDT) and polychlorobiphenyls (PCBs), was performed in blubber, muscle and liver of 12 animals according to Environmental Protection Agency (EPA) Method 8081/8082, with modifications (Marsili &

Focardi 1997). Total PCBs were quantified as the sum of individual congeners (IUPAC Nos. 95, 101, 99,151, 144, 135, 149, 118, 146, 153, 141, 138, 178, 187, 183, 128, 174, 177, 156, 171, 202, 172, 180, 199, 170, 196, 201, 195, 194, 206), while total DDTs were calculated as the sum of pp'DDT, op'DDT, pp'DDE, op'DDE, pp'DDD and op'DDD. Toxicological stress was evaluated using a theoretical model (Marsili et al. 2004) that estimates hazardous levels of OC pollutants at a canonical variable value > 0.47 (Table 2).

DISCUSSION

The microscopic findings, high parasite burdens and infections all point to immunocompromise of the cetaceans involved in this UME, which is particularly noteworthy because it involved young individuals belonging to a single species and occurred within a very short period of time and within a well-defined area. Furthermore, the immune system and its response may have been impaired by co-existing DMV infection and high tissue levels of OC pollutants in some of the stranded animals, making them more susceptible to secondary infections by other pathogens.

Morbillivirus is considered the most pathogenic viral agent for cetaceans. Endemic infections with milder lesions and lighter pathogen loads have been described in the western Mediterranean (Rubio-Guerri et al. 2013b); however, DMV displays a 'cyclic behavior', with seasonal outbreaks in this area. Indeed, the agent tends to 'reappear' as a population's anti-viral immune response decreases, leaving animals more susceptible to clinical disease.

Table 2. Levels of hexachlorobenzene (HCB), dichlorodiphenyltrichloroethane (DDT) and polychlorobiphenyls (PCBs) (ng g^{-1} dry wt) in the blubber of 12 cetaceans. Toxicological stress as evaluated according to the model proposed by Marsili et al. (2004). **Bold** entries denote animals with hazardous levels of organochlorines (canonical variable [CAN] value > 0.47). DMV: dolphin morbillivirus; CNS: central nervous system

ID	НСВ	DDTs	PCBs	CAN value	Pathological findings
22	301.4	63493.0	85625.0	0.17	Parasites
23	179.5	25512.4	59594.4	-0.24	Parasites
31	847.1	445582.0	471013.0	5.52	DMV (CNS, lungs); parasites
57	50.7	3162.8	10096.2	-0.81	Photobacterium damselae (multiple organs)
58	234.2	225558.6	215694.5	2.12	DMV (lungs)
59	417.1	133704.0	159855.7	1.19	DMV (heart); Photobacterium damselae (lungs)
60	1287.9	97955.9	92773.9	0.39	Parasites
62	232.2	233717.7	318458.6	3.14	DMV (CNS, heart, lungs)
63	478.2	163043.3	178431.9	1.49	Photobacterium damselae (multiple organs)
64	200.3	83689.0	110450.6	0.50	DMV (CNS)
65	349.7	96539.1	140581.9	0.84	Photobacterium damselae (multiple organs)
66	473.7	11718.1	35953.3	-0.53	Photobacterium damselae (multiple organs)

Despite the lack of a range of characteristic morphologic changes in the tissues from the DMVinfected animals, biomolecular investigations yielded clear evidence for DMV infections in 42% of those tested, supporting the hypothesis that DMV played an important role in this mortality outbreak. Given the geographical range of the strandings and the preservation status of the carcasses, the animals likely belonged to cetacean groups living in the southern Tyrrhenian Sea. The only DMV outbreak-the largest one to datereported for this area occurred in 1990 to 1992, whereas the 2 DMV outbreaks in 2006 to 2008 and 2010 to 2011 occurred in the western and northern Mediterranean basins (Di Guardo et al. 1995, Van Bressem et al. 2001, Di Guardo & Mazzariol 2013). Most of the animals in this UME were probably <20 yr of age, not previously exposed to DMV and, therefore, likely lacked adequate immunity; immunity to infection gradually tends to wane with the absence of recurrent infections in a population. Also, the level of anti-viral immunity in these animals or their relatives and conspecifics could have been impaired by other immunosuppressive agents, such as the high tissue levels of OC pollutants found in the stranded animals. Furthermore, because both DMV and OCs are known to impair host immune response, this could have led to secondary infections by other pathogens in the DMVinfected dolphins. Morbillivirus and herpesvirus infections have been reported to co-occur within the Mediterranean striped dolphin population, with co-infected animals frequently showing morbilliviral- but not herpesviral-related lesions (Bellière et al. 2010).

Usually considered an opportunistic pathogen, Photobacterium damselae subsp. damselae was identified in 62% of the stranded dolphins. Although some strains are known to be highly pathogenic for laboratory mammals under experimental conditions (Rivas et al. 2013), a clear association between this bacterium and mass mortality events or epidemic outbreaks in aquatic mammals has never been demonstrated. Little is known about the pathogenic significance of this microbial agent for free-ranging cetaceans. Equally critical is that few studies to date have investigated the entire viral genome of isolates recovered from morbillivirus-infected cetaceans. Future studies are therefore needed to characterize virus- and host-related factors in host-DMV interaction, as well as climate change-related factors contributing to the 'cyclic' occurrence of DMV epidemics in the Mediterranean.

No definitive conclusions on the causes of this cetacean UME can be drawn. Nonetheless, the effects possibly exerted by DMV infection on the immune system of the infected cetaceans may have contributed to this UME, in association with such other well-known immunosuppressive agents as OC contaminants. Other undetermined, unknown, or known environmental factors, such as reduced prey availability due to competition with human fishing activities, may have augmented the lethal effects of non-infectious or infectious agents such as *Photobacterium damselae damselae* or herpesvirus.

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