

## Microbial control of live/dead zooplankton ratio in Sevastopol Bay

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### Abstract

To explain higher fraction of live zooplankton in heavily polluted and eutrophic Sevastopol Bay comparing with cleaner adjacent waters, a hypothesis has been proposed and tested experimentally that more intensive bacteria-driven decomposition of dead organisms in the bay reduced their pool and, as a result, increased the live-to-dead zooplankton ratio. In the experiment, a heat-killed batch culture of the copepod *Calanipeda aquaedulcis* was used as a substrate for decomposition by natural microbial communities from the waters of different pollution status. Bacterioplankton abundance and *in situ* decomposition rate of copepod carcasses were shown to be about 3-fold higher in the bay ( $1.3 \times 10^6$  cells ml<sup>-1</sup> and 0.13 day<sup>-1</sup>, respectively) while an approximation of zooplankton non-predatory mortality rates gave equal values for both the sites (about 0.046 day<sup>-1</sup>). These findings call for revising the ways of interpreting the results of zooplankton viability assays in their relation to water pollution status.

**Key words:** zooplankton, viability, mortality, copepod, carcasses, decomposition, microbial hotspots, Sevastopol bay.

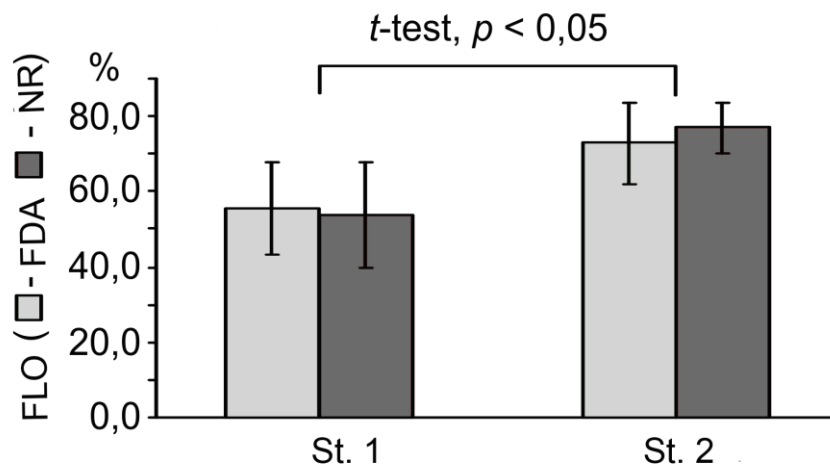
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### Introduction

Zooplankton have been widely shown to suffer from non-predatory mortality owing to starvation, diseases, injuries, parasites, harmful algal blooms, environmental stresses (Carpenter et al., 1974; Murtaugh, 1981; Byron et al., 1984; Burns, 1985; Ianora et al., 1987; Kimmerer, McKinnon, 1990; Hall et al., 1995; Delgado, Alcaraz, 1999; Gomez-Gutierrez, 2003; Tang et al., 2006; Dubovskaya, 2009; Elliott, Tang, 2011). However, a linkage between the mortality (and its derivative – live/dead organisms ratio) and pollution status of aquatic environments is still poorly understood.

Recent studies of live and dead planktonic copepods in Sevastopol bay and adjacent coastal waters (Litvinyuk et al., 2011) have revealed a surprising pattern: average annual fraction of live organisms (FLO) in highly polluted and eutrophic waters of the bay was significantly higher than one in the cleaner offshore area (Fig. 1). This result contradicted the conclusions made earlier by other authors (e. g. Pavlova et al., 2001; Pavlova, Melnikova, 2005) who explained higher necrozooplankton abundances and low FLO by adverse environmental conditions. In this study, bacterial degradation of copepod carcasses was considered a key factor controlling the ‘dead pool’ and, hence, the value of FLO in the studied area. Higher microbial activity in the polluted bay was hypothesized to accelerate copepod decomposition and, as a result, shift the ratio of live-to-dead organisms to higher values. The hypothesis was tested experimentally and a simple

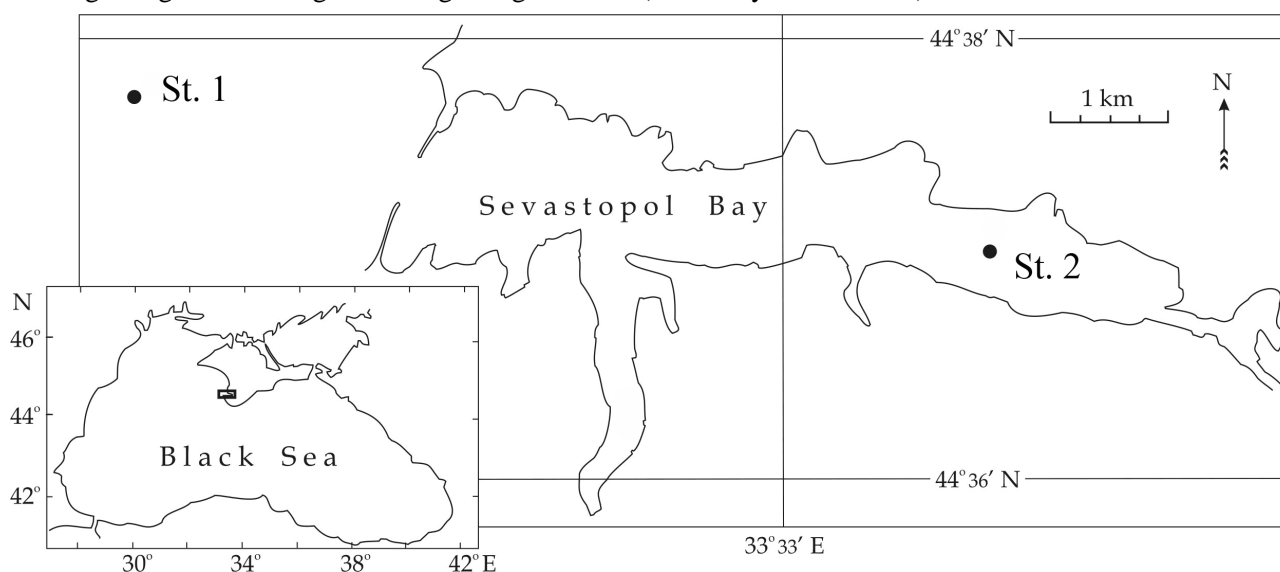
model was used to understand how the pool of dead organisms forms in the water column and how the balance between their inflow (via non-consumptive mortality) and outflow (via decomposition) is maintained in time.



**Figure 1.** Fluorescein diacetate- (FDA) and neutral red (NR) -based estimates of the average annual FLO in the open coastal waters (St. 1 in this study) and the polluted bay (St. 2 in this study) in 2010 – 2011. Calculated from the data presented in Litvinyuk et al. (2011).

## Material and Methods

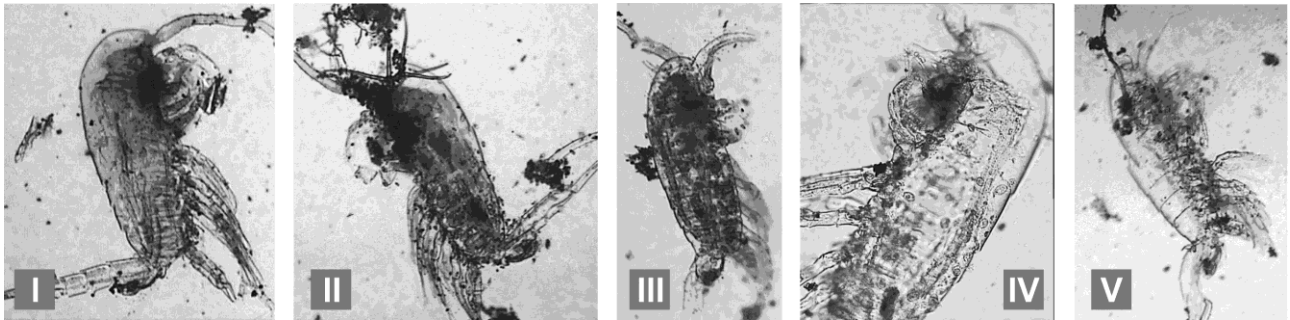
To estimate microbial abundances and conduct decomposition experiments *in situ*, seawater samples were collected at two stations in Sevastopol Bay and adjacent, open waters (Fig.2). The water area was characterized by pronounced gradients of water salinity, contamination and eutrophic state from the mouth of the Black River in the apex of the bay (close to St. 2) to the bay mouth and the open sea outside the bay (St. 1). The concentration of biogenic elements in the river outflow was 2 to 10 times higher than in the bay (Ovsyanyi et al., 2007), resulting in formation of stable concentration gradients of nitrite-N (from  $0.9 \mu\text{g L}^{-1}$  at St. 1 to  $2.3 \mu\text{g L}^{-1}$  at St. 2), nitrate-N ( $3.2$  and  $13.3 \mu\text{g L}^{-1}$ , respectively), ammonium-N ( $17.4$ – $23.1 \mu\text{g L}^{-1}$ ), phosphate-P ( $7.9$ – $8.8 \mu\text{g L}^{-1}$ ), and silica ( $57$ – $81 \mu\text{g L}^{-1}$ ) (V. Gubanov, unpublished data). The average E-TRIX eutrophication indices (Vollenweider, 1998) were 4.7 (St. 1) and 5.5 (St. 2) (Gubanov et al., 2015). The degree of chronic contamination by petroleum products and heavy metals increased from the open sea outside the bay to the bay center. Thus, the content of oil hydrocarbons in the bottom sediments of St. 1 was  $\sim 30 \text{ mg } 100 \text{ g}^{-1}$ , increasing to  $180 \text{ mg } 100 \text{ g}^{-1}$  at St. 2 (Osadchaya et al., 2004).



**Figure 2.** Sampling sites in Sevastopol Bay and adjacent coastal waters.

The rates of degradation and decomposition of dead ‘model’ organisms (the copepod *Calanipeda aquaedulcis* Kritchagin, 1873) by natural microbial communities from marine waters of different pollution status (St. 1 vs St. 2) were measured experimentally and compared. A batch culture of the copepods fed microalgal mixture (Bacillariophyceae: *Phaeodactylum tricornutum*; Chlorophyceae: *Chlorella vulgaris*, *Dunaliella salina*; Dinophyceae: *Prorocentrum cordata*, *Prorocentrum micans*; Prymnesiophyceae: *Isochrysis galbana*). The copepods were killed by heat (85 °C for 10 min.), trapped on 100-µm nylon mesh, rinsed and resuspended in native seawater freshly collected at the two stations (June 2012) and prefiltered through 100-µm nylon mesh to remove mesozooplankton. The final concentration of the dead organisms in the experimental flasks was about 300 ind. L<sup>-1</sup>. The flasks were prepared in triplicates for each of the locations, submersed to the sea (down to about 2-m depth) and incubated under *in situ* temperature (22 °C) and light conditions for 4 days that was an expected time period for decomposing the bulk of the carcasses (Tang et al., 2006). Then, decomposition of the copepods was examined by light microscopy.

130 carcasses of nauplii, copepodites and adults were analyzed visually and photographed in every replicate for identifying visual signs of disintegration and decomposition of the carcasses and classifying the extent of their degradation to the five categories: I – no visual signs of any body injuries or internal decomposition, II – partial postmortal decomposition of the body with light ‘spots’ and cavities, III – distinct visual signs of internal decomposition of up to 50% of the body volume, IV – body injuries and more than 50% internal decomposition, cephalothorax deformation and degradation, V – fragmented and/or empty carcasses with no tissue inside (Fig. 3). The carcasses at the stage V were defined as completely decomposed.



**Figure 3.** Stages of decomposition of copepod carcasses. Explanations are in the text.

To compare average annual bacterioplankton abundances at St. 1 and St. 2, seawater samples were collected in the surface layer bimonthly over 2010-2011. These data and the data on zooplankton FLO in Sevastopol Bay published earlier (Litvinyuk et al., 2011) were both obtained simultaneously at the same stations (as a part of Sevastopol Bay Time Series), thus allowing their inter-comparison.

Bacterial abundances in the samples and in the experimental flasks were estimated by flow cytometry. Counts were performed using a Beckman Coulter flow cytometer (Cytomics FC 500) equipped with an air-cooled blue laser (15 mW, 488 nm) and the standard filter setup. Aliquots of 1ml water samples previously fixed with formaldehyde (2% final conc.) were stained with SYBR Green I (Molecular Probes Inc.), following the procedures described by (Marie et al., 1997; Gasol, Giorgio, 2000). SYBR-Green I fluorescence at the green channel FL1 (525 nm) was considered proportional to intracellular nucleic acid content and interpreted as a measure of bacterial cell-specific metabolic activity according to (Servais et al., 2003). Consequently, the integral metabolic activity of the community was estimated as  $FL1 \times N$  (in relative units), where  $N$  is the total bacterial abundance in the sample.

To analyze the field and experimental results, a simplified model (ignoring sedimentation) has been used, describing dynamics of live and dead organisms and the balance between their pools:

$$dN_L/dt = (\mu - m) \times N_L \quad (1),$$

$$dN_D/dt = m \times N_L - d \times N_D \quad (2),$$

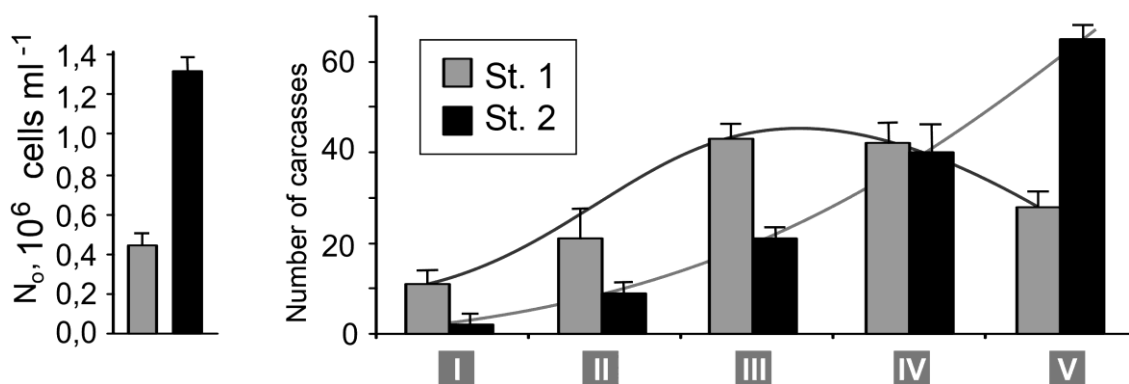
$$FLO = (N_L/(N_L+N_D)) \quad (3),$$

where  $N_L$  and  $N_D$  are abundances of live and dead organisms, respectively;  $\mu$  is specific growth rate;  $m$  is non-consumptive mortality rate;  $d$  is decomposition rate of zooplankton. Under steady-state conditions ( $\mu = m$ , constant  $N_L$ ) in the model population, FLO depends on  $N_D$  and, hence, on the ratio between the non-consumptive mortality rate ( $m$ ) of live zooplankton (inflow to the pool of dead organisms) and the decomposition rate ( $d$ ) of copepod carcasses (outflow from the pool). In the decomposition experiment, the equation 2 was used to calculate  $d$  at  $N_L = 0$ ,  $\mu = 0$ ,  $m = 0$ , and  $FLO = 0$ .

Statistical analyses were performed using software STATISTICA ver. 10 (StatSoft, Inc., USA).

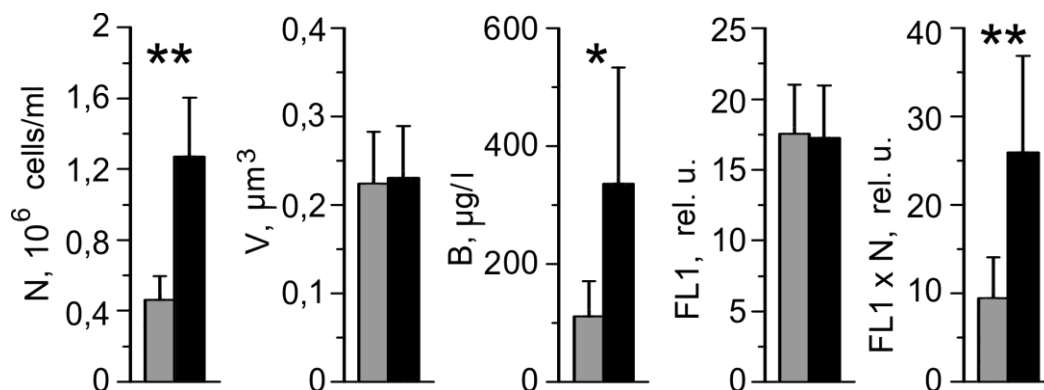
## Results and Discussion

By fourth day of the exposition, about 20% (St. 1) and 50% (St. 2) of the copepods were decomposed completely, reaching the stage V (Fig. 4). Frequency distributions of the copepod decomposition stages differed significantly in the locations, indicating faster process in the bay. The rates of decomposition ( $d$ ) approximated from these data amounted  $0.05 \text{ day}^{-1}$  in the bay (St. 1) and  $0.13 \text{ day}^{-1}$  outside it (St. 2), thus supporting the hypothesis of microbial control of live/dead zooplankton ratio in Sevastopol bay.



**Figure 4.** Initial bacterial abundances in the experiment ( $N_0$ , left plot) and frequency distribution of the copepod decomposition stages on the fourth day of exposition at St. 1 and 2 (right plot). Means and standard deviations are presented.

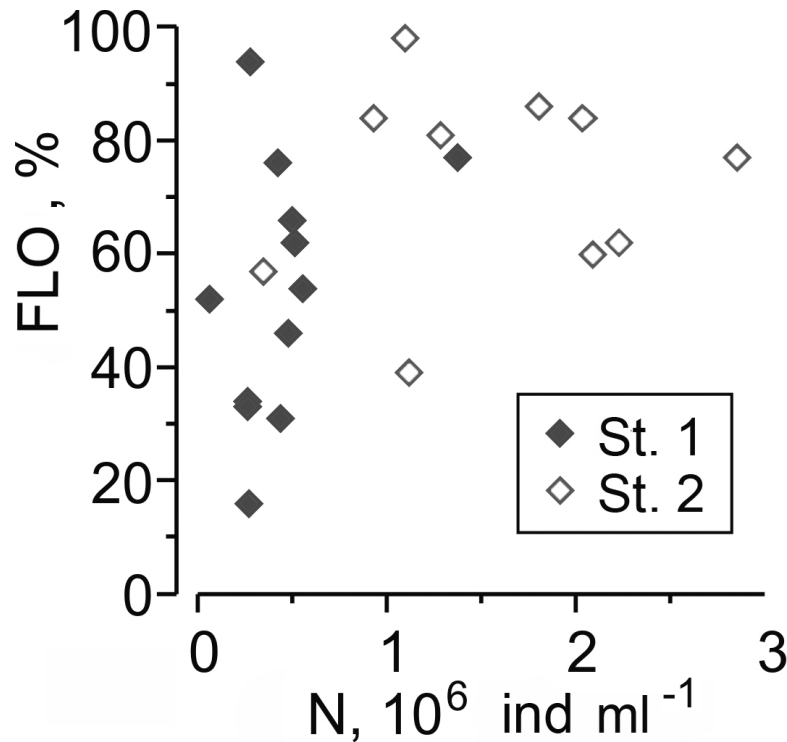
Disintegration and decomposition of zooplankton carcasses was earlier shown to be driven by external, not internal, bacteria (Harding, 1973; Lee and Fisher, 1992; Reinfelder et al., 1993), hence, indicating a direct link between FLO and microbial activity in the water column. The initial bacterial abundances in the experimental flasks differed significantly,  $0.4 \times 10^6 \text{ cells ml}^{-1}$  at St. 1 versus  $1.3 \times 10^6 \text{ cells ml}^{-1}$  at St. 2 (Fig. 4). Approximately the same difference between the locations maintained over the year – average annual abundance of bacterioplankton proved to be about 2.5-fold higher in Sevastopol Bay as against the adjacent waters (Fig. 5).



**Figure 5.** Bacterioplankton average annual (2010 - 2011) abundance ( $N$ ), cell volume ( $V$ ), biomass ( $B$ ), intracellular nucleic acids ( $FL1$ ) and integral metabolic activity ( $FL1 \times N$ ) ( $\pm 95\%$  CI) at St. 1 (grey) and St. 2 (black). Significant differences are marked (\*  $p < 0.05$ , \*\*  $p < 0.01$ ).

According to the seasonal microbiological data obtained in 2010-2011, no significant difference between the stations has been revealed in terms of bacterial cell volumes and intracellular nucleic acid contents (FL1 signal). However, the integral metabolic activity measured as  $FL1 \times N$  was 2.5- to 3-fold higher in the bay community (Fig. 5).

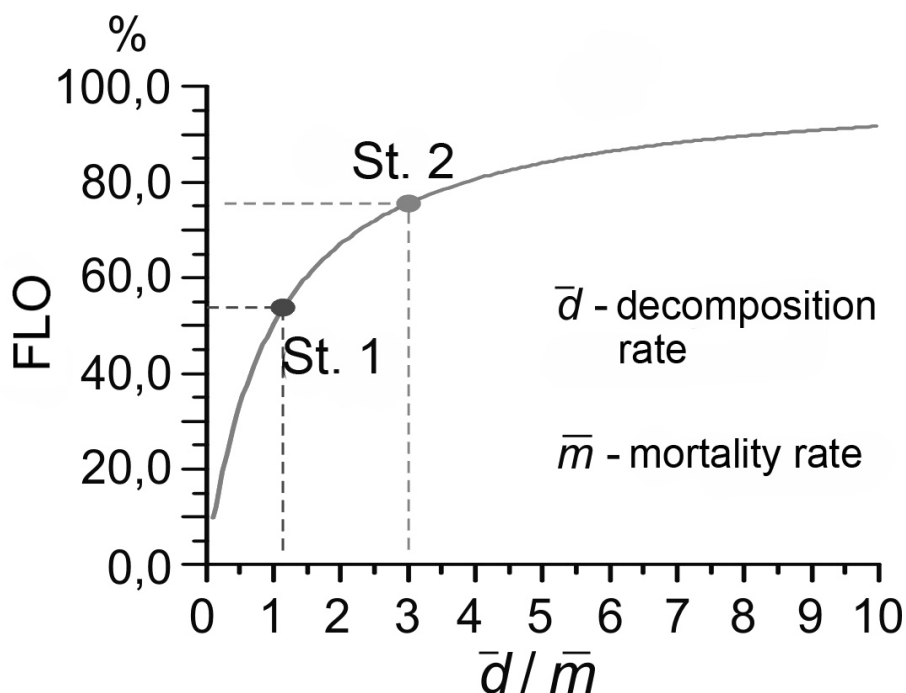
Thus, both the experimental results and microbiological data have provided an evidence of a linkage between FLO and bacterioplankton. There was an increase in FLO at higher bacterial abundances and high FLO variability at fewer bacteria (Fig. 6). The latter seemed to be a result of contribution of other important factors to the FLO control (like sedimentation or elimination by detritivores).



**Figure 6.** Fraction of live organisms (FLO) in zooplankton versus bacterioplankton abundance (N). Data on FLO (2010-2011) are from Litvinyuk et al. (2011).

Given constant  $N_L$  and equality of  $\mu$  and  $m$  (losses are compensated for by reproduction) under stationary conditions, the value of FLO depends only on the  $d/m$  ratio irrespective of the initial  $N_L$ ,  $m$  and  $d$  (Fig. 7). The dependence has allowed a calculation of the  $d/m$  ratios from the field data on the FLO annual averages (54% and 77%, Litvinyuk et al., 2011), which amounted about 1.2 and 3.0 at St. 1 and 2, respectively (Fig. 6). Thus, according to the model, the rates of copepod mortality and decomposition were roughly equal in the open waters while in the bay, decomposition occurred three times faster. Combining these calculations with the experimental results gives nearly equivalent rates of zooplankton non-consumptive mortality at the St. 1 and 2 ( $0.045$  and  $0.047 \text{ day}^{-1}$ , respectively) which are within the range ( $0.01$  to  $0.1 \text{ day}^{-1}$ ) commonly reported for marine zooplankton (see review in Tang et al., 2014).

In summary, it can be concluded from the present findings that higher FLO values observed in more polluted and eutrophic marine coastal waters were not necessarily a result of higher zooplankton mortality but rather a consequence of higher abundance of heterotrophic bacteria and more intensive bacteria-driven decomposition of dead organisms in the detrital food web. Unexpectedly, the rates of non-consumptive mortality of copepods in polluted Sevastopol Bay were approximated to not differ significantly from those in the less polluted adjacent waters. Thus, results of zooplankton viability assays should be interpreted carefully in their relation to water pollution status. Higher pollution status of an area does not necessarily imply higher zooplankton mortality and/or lower FLO values.



**Figure 7.** Fraction of live organisms (FLO) as a function of the decomposition-to-mortality ratio ( $\bar{d}/\bar{m}$ ) in the model under steady-state conditions (mortality and specific growth rates are balanced,  $\mu = m$ ) and projections of natural zooplankton communities (St. 1 and 2) onto the model curve.

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