Future challenges in cephalopod research*

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37 Abstract

Cephalopods (Mollusca: Cephalopoda) play an important role as keystone invertebrates 38 in various marine ecosystems as well as being a valuable fisheries resource. At the 39 World Malacological Congress, held 21-28 July 2013 in Ponta Delgada, Azores, 40 Portugal, a number of cephalopod experts convened to honour the contribution of the 41 late Malcolm R. Clarke, FRS (1930-2013) to cephalopod research. Endorsed by the 42 Cephalopod International Advisory Council (CIAC), the meeting discussed some of the 43 major challenges that cephalopod research will face in the future. These challenges were 44 identified as follows: 1. To find new ways to ascertain the trophic role and food web 45 links of cephalopods using hard tissues, stable isotopes and novel concepts in theoretical 46 ecology; 2. To explore new approaches to the study of cephalopod morphology; 3. To 47 further develop cephalopod aquaculture research; 4. To find new ways to ascertain 48 cephalopod adaptation and response to environmental change; 5. To strengthen 49 cephalopod genetics research; and 6. To develop new approaches for cephalopod 50 fisheries and conservation. The present contribution presents brief reviews on these 51 topics, followed by a discussion of the general challenges that cephalopod research is 52 bound to face in the near future. By contributing to initiatives both within CIAC and 53 54 independent of CIAC, the principle aim of the article is to stimulate future cephalopod research. 55

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58 Introduction

There is a long tradition of researchers striving to predict what might happen in the 59 future. Until recently, most research related to biodiversity and conservation has been 60 performed in an unsystematic manner (Sutherland & Woodroof, 2009, Sutherland et al., 61 2013). However, an assessment of future research opportunities, knowledge gaps and 62 new areas of science coonstitutes the first step in identifying and communicating 63 hypotheses and insights for the future (Rands et al., 2010, Sutherland et al., 2010). 64 Prognoses for marine ecosystems are becoming increasingly important, because of the 65 66 threats that have been emerging in recent decades and which require urgent scientific attention. These threats include global climate change, ocean warming, sea level rise, 67 biodiversity loss, overfishing, ocean acidification and expanding hypoxia (Pauly, 1998, 68 Pauly et al., 1998, Pauly et al., 2003, Orr et al., 2005, Rockstrom et al., 2009, Turner et 69 70 al., 2009). Addressing these challenges asks for greater synergy between research, management and policy, and it will be important to inform researchers and funding 71 72 agencies as to where their efforts might best be focused.

Cephalopods (Mollusca: Cephalopoda) are widely recognized as playing a 73 74 pivotal role in many marine ecosystems, both as predators and prey (Clarke, 1996, 75 Piatkowski et al., 2001, Boyle & Rodhouse, 2005). Furthermore, cephalopod fisheries have been increasing steadily in recent decades and it is likely that more species will be 76 commercially exploited in the future (FAO, 2005). As marine biologists whose research 77 is focused on cephalopods, our aims include a better understanding of cephalopod 78 biology and ecology and the role of these organisms in marine ecosystems, identifying 79 patterns and mechanisms, quantifying changes at different scales, recognizing problems 80 and testing potential solutions (e.g. related to conservation, fisheries management and 81 aquaculture). Except for nautiluses, cephalopods have a short life span, rapid growth, 82 83 and semelparous maturation patterns (Boyle & Rodhouse, 2005). These life cycle traits may have positive or negative effects on cephalopod species in relation to 84 85 environmental change, as cephalopods can be both sensitive (in terms of rapid response) and resilient (in terms of recovery) to phenomena such as overfishing or climate 86 variability and change (Pecl & Jackson, 2008, Rosa & Seibel, 2008, André et al., 2010, 87 Pierce et al., 2010, Hoving et al., 2013, Rodhouse, 2013). However, exactly how these, 88 and other, phenomena affect cephalopods is not yet fully understood. Therefore, a broad 89 discussion of these issues can be valuable in providing guidance for future directions of 90 91 cephalopod research.

The 2013 World Malacological Congress, held 21-28 July in Ponta Delgada, 92 93 Azores, Portugal, brought together a number of cephalopod experts to participate in a symposium focusing on the role of cephalopods in the world's oceans. The symposium 94 was held in honour of the late Malcolm R. Clarke, FRS (1930-2013), and was endorsed 95 by the Cephalopod International Advisory Council (CIAC). Malcolm Clarke himself 96 97 had initiated the CIAC in 1981, and the council was officially founded two years later as an international forum to encourage research on cephalopods, promote international 98 collaboration in cephalopod science, and to provide an official body to answer the 99 100 increasing number of questions about cephalopods, particularly those related to cephalopod fisheries (Hochberg & Hatfield, 2002). In the present article, cephalopod 101 102 experts, including present and former members of CIAC, working in specific fields and 103 at different organisational scales, ranging from a species perspective to the ecosystem 104 level, discuss some of the challenges that cephalopod research will face in the future. The individual sections provide brief reviews of topics in cephalopod research that 105 106 deserve further attention.

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108 1. New ways to ascertain the trophic role and food web links of cephalopods

Studying cephalopods in the world's oceans using top predators as biological samplers:
where are we heading? (José C. Xavier)

Knowledge on cephalopods, particularly those from oceanic waters that are not 111 commercially caught, largely originates from analyses of stomach contents collected 112 from their natural predators, such as toothed whales, seals, seabirds, sharks and teleost 113 fish (Clarke, 1996). This is, because current methods for direct sampling, especially of 114 oceanic squid, are still inefficient (Clarke, 1977, Xavier et al., 2007, Hoving et al., in 115 press). Therefore, an essential tool in the study of cephalopod remains found in predator 116 117 stomachs is the identification and measurement of their chitinized upper and lower beaks (Clarke, 1986, Cherel et al., 2004, Xavier & Cherel, 2009, Xavier et al., 2011), 118 119 and, to a lesser extent, the morphological and molecular analysis of soft tissues in case 120 these should still be available (Pierce & Boyle, 1991, Barrett et al., 2007, Karnovsky et al., 2012). 121

However, the analysis of hard tissues can be biased. For instance, a recent study showed that the ratio of upper to lower beaks in diet samples from top predators varied significantly during one year as well as between years. This bias was larger in some cephalopod species than in others, resulting in the underestimation of the relative importance of some species in data derived from this approach (Xavier et al., 2011).
This can result in an under- or over-estimation of relative cephalopod abundance and
suggests that it is essential to count both (i.e. lower and upper) beaks in stomach content
analyses. Furthermore, in instances where there is a consistent bias (>30%), all beaks
should be identified, and the higher quantity of beak type should be considered to
reconstruct the cephalopod component of the diet by mass (Santos et al., 2001, Xavier et al., 2011).

In samples collected from predators that tend to retain material, it is of 133 134 importance to separate old and fresh material during the initial sorting process in order to obtain a qualitative assessment of the degree of erosion of the material as well 135 136 (Piatkowski & Pütz, 1994, Cherel et al., 2000, Xavier et al., 2005). These components 137 can then be analysed separately, as required, and the results compared. In general, more 138 effort should be put into describing upper beak morphology to aid identification (Clarke, 1962, Imber, 1978, Pérez-Gándaras, 1983, Wolff, 1984, Kubodera & 139 140 Furuhashi, 1987, Lu & Ickeringill, 2002, Xavier & Cherel, 2009), to measuring upper beaks in diets, and to developing regressions or allometric equations for estimating 141 142 cephalopod mass based on both lower and upper beak measurements. Indeed, for numerous species no allometric equations are yet available, which is why scientists have 143 to rely on equations from closely related species. In addition, various allometric 144 equations were produced based on a limited number and size range of cephalopod 145 specimens. Therefore, more material must be collected, particularly from cephalopod 146 natural predators or by research as well as commercial fishing vessels. 147

Malcolm Clarke emphasized the importance of additional ship time devoted to 148 cephalopod research, as well as the need for the development of better capture methods 149 (Xavier et al., 2007, Hoving et al., in press). Many cephalopods are fast-swimming 150 organisms and therefore only the small or less-mobile specimens are usually captured 151 (Clarke, 1977). This dilemma still holds true, despite a long history of sampling. In 152 153 order to maximize the success rate of capturing bigger specimens, larger nets and 154 modified net gear (e.g. underwater lights) have been developed to attract cephalopods into the nets (Clarke & Pascoe, 1997, Clarke & Pascoe, 1998, Clarke, 2006). However, 155 new techniques are required to enhance the catch ratio of poorly-known cephalopod 156 species in the world's oceans in order to complement the work already being carried out 157 on the feeding and foraging ecology of cephalopod predators. 158

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160 Stable isotopes, hard tissues and the trophic ecology of cephalopods (Yves Cherel) 161 Stable isotopes (δ^{13} C and δ^{15} N) have recently emerged as new efficient intrinsic 162 markers of the trophic ecology of cephalopods (Jackson et al., 2007), and pioneer 163 investigations (Takai et al., 2000, Cherel & Hobson, 2005) have lead to a steady 164 increase in the use of the method over the last ten years (Navarro et al., 2013). In this 165 section, attention will be paid to the most recent findings, methodological issues and 166 perspectives on the use of these tools on hard tissues of cephalopods.

In contrast to soft tissues (e.g. mantle), hard tissues (i.e. beaks, gladii, statoliths, 167 168 and eye lenses) are metabolically inactive structures that grow continuously by accretion of new molecules with no turnover after synthesis. Consequently, these 169 170 structures retain molecules laid down throughout the lives of cephalopods, and their δ^{13} C and δ^{15} N values thus integrate the feeding ecology of individuals over their 171 lifetime. Indeed, various parts of hard tissues have different isotopic signatures. For 172 example, δ^{13} C and δ^{15} N values of the tip of the wings and anterior tip of the gladius (i.e. 173 174 the most recently synthesized parts of lower beaks and gladii, respectively) integrate the 175 feeding ecology prior to capture (Cherel & Hobson, 2005, Hobson & Cherel, 2006, 176 Cherel et al., 2009a). Gladii have the advantage over beaks that their growth increments are larger, better defined and easier to sample along the longitudinal proostracum axis 177 178 (Cherel et al., 2009a). Furthermore, in the most recent part of the gladius, assuming 179 increments are daily, a day-by-day picture can also be established which can directly be related to body size (as gladii length is approximately the same as the dorsal mantle 180 181 length; Graham Pierce, unpublished data).

Stable isotopes from hard structures have two practical advantages and one 182 methodological disadvantage. Firstly, measuring the isotopic signature of serially 183 184 sampled beaks and gladii presents the unique opportunity to reconstruct the foraging history of individuals. For example, $\delta^{15}N$ profiles of beaks from *Architeuthis dux* 185 suggest an ontogenetic shift early in life (Guerra et al., 2010), and sequential isotopic 186 values along gladii of *Dosidicus gigas* highlight contrasted individual foraging 187 strategies (Ruiz-Cooley et al., 2010, Lorrain et al., 2011). In the same way, the only 188 189 published investigation on concentric eye lens layers reveals variations in δ^{13} C and δ^{15} N 190 values at fine temporal scales, indicating substantial variability in squid feeding patterns (Hunsicker et al., 2010a). Secondly, the combination of the stable isotope techniques 191 192 with the use of predators as biological samplers, and cephalopod identification using

external features of accumulated beaks in predators' stomachs (Clarke, 1986, Xavier &
Cherel, 2009) allows information to be gathered on poorly known species. This method
has already revealed new trophic relationships and migration patterns together with the
trophic structure of deep-sea cephalopod assemblages (Cherel & Hobson, 2005, Cherel
et al., 2009b).

However, a main problem with using δ^{13} C and δ^{15} N values of hard structures is 198 199 that biological interpretation is confused by differences in biochemical composition between hard and soft tissues. Beaks and gladii contain not only protein but also chitin 200 201 (Hunt & Nixon, 1981, Rubin et al., 2010), a modified polysaccharide that contains impoverished ¹⁵N nitrogen (Schimmelmann, 2011). The presence of chitin explains why 202 hard tissues have consistently much lower δ^{15} N values than soft tissues (Cherel et al., 203 204 2009a). Moreover, the ratio of chitin to protein varies within beaks, with the undarkened, darkening and darkened parts of beaks containing decreasing amounts of 205 206 chitin (Rubin et al., 2010). Chitin content is thus likely to be different between individual beaks (e.g. small, undarkened versus large, darkened beaks), and the gladius 207 is richer in chitin than darkened beaks (Hunt & Nixon, 1981). This particular issue is 208 209 analogous to that arising from the different fractionation apparent in lipids compared to other components of soft tissues. Three different approaches enable the 'chitin effect' to 210 211 be dealt with, namely the use of isotopic correction factors between hard and soft tissues (Hobson & Cherel, 2006, Cherel et al., 2009a), the removal of chitin and measuring 212 213 stable isotopes on amino acids. Determining the stable isotope ratios of chemically extracted proteins from hard tissues has not yet been performed, but a more promising 214 way is to measure δ^{15} N values of amino acids resulting from protein hydrolysis. 215 Selecting appropriate source and trophic amino acids (e.g. phenylalanine and glutamic 216 acid, respectively) allows quantification of both δ^{15} N baseline levels and the trophic 217 position of consumers relative to the baseline [i.e. the δ^{15} N signature of source amino 218 219 acids (e.g. phenylalanine) does not increase along the food chain, while that of trophic amino acids (e.g. glutamic acid) does - hence the $\delta^{15}N$ difference between trophic and 220 source amino acids is a direct estimation of the trophic position of an organism]. This 221 222 approach was recently used on cephalopod hard tissues, including cuttlefish cuttlebone (Ohkouchi et al., 2013) and squid gladii (Ruiz-Cooley et al., 2013) and has the potential 223 224 to depict previously unknown trophic relationships, habitat use and migration patterns 225 of cephalopods in marine ecosystems.

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227 Population dynamics of cephalopods under a trophic relations (as well as age and
228 growth) context: possible future research (Marek R. Lipinski)

- 229 The present-day population dynamics of cephalopods are still largely at the descriptive,
- natural-history stage. The best summaries of current knowledge were given by Boyle
 and Boletzky (1996), Boyle and Rodhouse (2005) and Rodhouse et al. (in press).

The known data on cephalopod population dynamics that has been widely and 232 comprehensively quantified may serve a practical purpose in the management of 233 234 fishable stocks of selected species of octopods, cuttlefish and squid (Rodhouse et al., in press). This part, however, largely ignores the general theoretical framework of 235 population dynamics in ecology (described by Turchin (2003)), based on predator-prey 236 interactions. However, there are efforts aimed at incorporating predator-prev 237 238 relationships in sustainable resource management (Bowman et al., 2000, Overholtz et al., 2000, Overholtz et al., 2008, Tyrrell et al., 2008, Tyrrell et al., 2011). Hence, 239 240 predation should be considered in setting Maximum Sustainable Yield (MSY) for a fishery, including those fisheries exploiting squid (such as *Doryteuthis pealeii* and *Illex* 241 242 *illecebrosus*). When this is done, MSY is usually considerably smaller. Model inputs are 243 usually based on stomach contents analysis, and actual consumption is calculated 244 (subject to some assumptions). The underlying reasoning is that a lack of this resource in the diet of predators will be to the detriment of these predators. 245

However, this may not correspond to reality. This approach usually assumes 246 either a specialist character of these predators, or at best, a hyperbolic response of a 247 generalist predator, according to a model where dN/dt = rN(1-N/k)-gN/d+N [with N: 248 population density; r: per capita rate of population change; k: carrying capacity 249 250 (logistic); d: half-saturation constant (hyperbolic); g: total killing rate by generalist predators; h: half saturation constant (sigmoid)] (Turchin, 2003). However, cephalopods 251 are opportunistic generalists both as a predators and in turn are preyed upon by 252 253 generalist predators themselves (preved upon by fishes, birds, mammals and 254 cephalopods), giving a sigmoid response to predation according to the model dN/dt = $rN(1-N/k)-gN^2/h^2+N^2$. These two scenarios are illustrated in Figure 1, where solid lines 255 represent per capita growth rates of the prey population in the absence of predators, 256 whilst dashed curves represent per capita death rate of prey as a result of predation. 257 258 Numbers correspond to the specific cases (out of many possible). In the hyperbolic scenario (Fig. 1A), case 1 refers to the total extinction of prev as a result of predation. 259

260 Case 2 is the situation where only very high density of prey secures the end equilibrium 261 (and hence the survival of prey). Case 3 ends with survival of the prey population regardless what happens, therefore predators have a minimum impact upon their prey. 262 263 In the sigmoid scenario (Fig. 1B), case 1 refers to an equilibrium where prey densities 264 are low (survival is probable); case 2 represents three equilibria and therefore the final 265 result depends upon the initial conditions, but all of them are likely to be stable (survival of prey is probable in most situations); and in case 3 equilibrium is reached at 266 high prey densities, therefore survival is even better than in case 1. 267

Given the above, future work should apply theoretical ecology models to real cephalopod populations, and only then should feed into well-intentioned resource management. This is not happening as of yet, simply because it is a complicated task. Cephalopods will require the development of a 'multi-opportunistic links model' compatible with other findings of theoretical ecology. This model may be useful for fisheries management only if the required parameters can be obtained or assessed in practice.

A second case study considered here is on cephalopod age, ageing, longevity 275 276 and growth from a population dynamics perspective. This field also has a background in 277 theoretical ecology (Turchin, 2003), with its emphasis on changing ages, different 278 average longevity (and ultimately, growth parameters) with change between subsequent generations, and on the influence these changes may have on oscillating numbers of 279 280 individuals in a population. Here, in contrast to the field of predator-prey relationships, theoretical ecology feeds into practical applications (Quinn & Deriso, 1999), although 281 282 the focus is somewhat different. Cephalopods, however, have not yet been the subjects of thorough studies in this discipline. This is, because it is felt that some fundamental 283 284 problems in understanding population structuring have not yet been resolved. There is a 285 relative abundance of age data, but a paucity of studies using these data to model 286 population structure based either on generations (for theoretical purposes) or to 287 construct suitable keys (e.g. age-length) for stock assessment and management analyses.

Therefore, there is a need for new research and more data. Firstly, there is a requirement for physiological studies on the interpretation of age marks (mostly biomineralization studies) to construct true instead of biased validation procedures. Secondly, no one has so far adequately addressed Daniel Pauly's paradoxon regarding the metabolic limitation of squid growth (Pauly, 1998): according to him, large squid cannot grow quickly due to their energetic requirements, which goes against the age 294 readings of squid statoliths (supported by aquarium observations), which in turn support 295 the inference that large squid do grow quickly. However, a good start to reconciling 296 these contradictory data was made by O'Dor and Hoar (2000). Thirdly, studies of 297 cephalopod growth are required, that will combine a theoretical ecology approach 298 (suffering at the moment from an assumption of non-overlapping generation cycles) (Turchin, 2003), a wealth of matrix models (Quinn & Deriso, 1999), and a solid 299 physiological basis (which is lacking at the moment). It is to be hoped that the state of 300 the art, presently fragmented into these three areas (Arkhipkin & Roa-Ureta, 2005, 301 302 André et al., 2009, Keyl et al., 2011, Semmens et al., 2011, Zavala et al., 2012), will 303 improve in the future.

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305 2. New approaches to the study of cephalopod morphology (Elizabeth K. Shea, 306 Alexander Ziegler)

307 Comparative morphology is an essential, yet increasingly rare specialty in organismic 308 biology. The slow pace of work for detailed analysis and the current lack of a centralized repository for morphological data contribute to the widely-cited 'taxonomic 309 310 impediment' that contemporary biodiversity research is facing (Crisci, 2006, de 311 Carvalho et al., 2007). Due to the lack of open access to structural data, molecular 312 methods (e.g. barcoding) are rapidly supplanting morphology in systematics and taxonomy research, resulting in a one-sided discussion about invertebrate relationships 313 314 and evolution. Rejuvenating morphological research through the development of online repositories for morphological data will provide new avenues of inquiry that would 315 316 contribute substantively to systematic and taxonomy research. In addition, morphology would become a more accessible contributor to large interdisciplinary research 317 initiatives such as the Census of Marine Life (Decker & O'Dor, 2003) or major online 318 319 compilations of organismic data such as the Encyclopedia of Life (Wilson, 2003). In 320 this section, we highlight several new and developing digital techniques that have the 321 potential to expedite morphological work, and which could encourage a shift in focus 322 from data acquisition to data analysis, consequently resulting in a more rapid and regular research output in cephalopod morphological research. 323

324 Cephalopod beak identification is notoriously difficult, but an in-depth
325 understanding of beak morphology is critical to stomach content analyses, as well as a
326 for an understanding of predator-prey dynamics (Clarke, 1986, Xavier & Cherel, 2009).
327 Three-dimensional (3D) anaglyph images constitute an alternative to complex line

drawings or photographs (Xavier & Cherel 2009). Richard E. Young is in the process of
building a collection of such images, archiving them on the Tree of Life Web Project
website (http://www.tolweb.org/notes/?note_id=4541). The upper and lower beaks of
over 140 species from all major clades have so far been analyzed (Young, 2009). In
addition, new hybrid approaches such as rotational SEM could be used to image
miniscule morphological features such as statoliths, sucker dentition or cartilaginous
strips and tubercules at very high resolutions and in 3D (Cheung et al., 2013).

In addition, robotic microscopy systems developed for applications in pathology 335 336 permit rapid digitization of histological sections on a large scale and at high resolutions (Al-Janabi et al., 2012). Such systems could be used to digitize and catalogue 337 338 histological data on cephalopod neuroanatomy, such as, for example, the John Z. Young slide collection deposited at the National Museum of Natural History (Washington, DC, 339 340 USA). The resulting tomographic image stacks can be aligned using semi-automatic and automatic algorithms (Eliceiri et al., 2012), and can subsequently be made accessible as 341 342 full 3D datasets in online repositories. These image stacks would then become a baseline of information that permits directly connecting past research (Young, 1971) to 343 344 present compilations (Nixon & Young, 2003), as well as future studies.

345 Furthermore, non-invasive scanning techniques such as magnetic resonance imaging (MRI), computed tomography (CT), or micro-computed tomography (μ CT) 346 now allow analyzing whole specimens from the millimetre to the metre scale (Walter et 347 al., 2010). Following dataset acquisition, specialized (but often open source) software 348 can be used to virtually dissect the scanned specimen in real-time and in 3D (Ziegler & 349 350 Menze, 2013). While MRI is particularly suitable for soft tissue imaging (Ziegler et al., 2011a), the X-ray-based techniques CT and µCT constitute the methods of choice for 351 352 hard part imaging (Ziegler et al., 2010). However, specimen state (in vivo or ex vivo), 353 scanning medium (e.g. air, ethanol, formalin, water), scanning time (minutes to hours), dataset resolution (nm to μ m), as well as scanning cost per specimen (up to many 354 355 hundreds of US-\$) may vary considerably and primarily depend on the system used.

³⁵⁶ Due to the dominance of soft tissues, cephalopods constitute suitable candidates ³⁵⁷ for MRI scanning (Ziegler et al., 2011a). For example, 3D MRI datasets can be used to ³⁵⁸ visualize internal organs in their natural context (Fig. 2A, B). In contrast, mineralized ³⁵⁹ tissues present in cephalopods (e.g. eye lenses, beaks, statoliths, shells) can be rapidly ³⁶⁰ visualized using CT or μ CT (Fig. 2C-E). However, whole specimen staining using ³⁶¹ electron-dense elements such as iodine or tungsten (Metscher, 2009, Kerbl et al., 2013) also allows analyzing soft tissues of smaller cephalopod specimens using µCT (Fig. 2 F,
G).

Current online projects such as The Digital Fish Library (Berquist et al., 2012) 364 or The Digital Morphology website (http://digimorph.org/) provide a good starting point 365 366 for the web-based hosting of morphological data and constitute potential infrastructures for future efforts in cephalopod research. In addition, dissemination of complex 367 biological structures is still carried out primarily in the form of 2D publications (Ziegler 368 et al., 2011b), although interactive 3D models based, for example, on the ubiquitous 369 370 portable document format (PDF) have been integrated into electronic publications already for several years (Ruthensteiner & Heß, 2008, Kumar et al., 2010). Nonetheless, 371 372 continued development of such approaches is required in order to adapt them, for example, to mobile devices. Furthermore, 3D printing is poised to become an important 373 374 tool in the communication of complex biological structures, whether in research or in teaching (Kelley et al., 2007, Ziegler & Menze, 2013). 375

376 In general, digital morphological techniques permit shifting the workload from data acquisition to data analysis, which will open new avenues of research both across 377 378 and within cephalopod species. Previously collected, well-identified, and data-rich museum specimens could form the backbone of a large-scale, non-invasive scanning 379 380 program (Ziegler, 2012). Apart from developing a collection of 3D datasets that can be accessed in the form of a digital museum collection, the novel, high-throughput 381 scanning techniques described above provide new opportunities for a variety of 382 cephalopod specimens. For example, scanning of bulk-collected, commercially-trawled 383 384 cephalopods could be employed to answer long-standing questions of character variation within species (Vecchione et al., 2005). Reared cephalopods such as Sepia 385 officinalis could be used for in vivo experiments, where images taken before and after a 386 387 stimulus would be required. Furthermore, specimens too valuable for dissection (e.g. 388 holotypes) can now be scanned with virtually no impact on the specimen and be made fully accessible online in 3D. 389

Cephalopods constitute a small-enough class of molluscs that an effort to
digitally scan one representative from each genus or species would constitute a realistic
goal, and one that should be pursued in parallel to molecular barcoding (Strugnell &
Lindgren, 2007). A concise, user-friendly, widely-disseminated, morphological
infrastructure that parallels ongoing efforts to barcode all cephalopod species would

render cephalopods not just a group with multiple model organisms, but also a modelclade for systematic and taxonomy research.

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398 **3.** Challenges in cephalopod culture (Roger Villanueva, Erica A. G. Vidal)

399 Experimental approaches have been an important tool for understanding fundamental principles of cephalopod life cycles, physiology and behaviour, thus providing the basis 400 for pilot commercial culture of some species. A recent publication summarizes modern 401 culture techniques used for the most common cephalopod species (Iglesias et al., 2014). 402 403 Another publication focuses on four species which are highlighted as cephalopod culture models for which there are comprehensive data available, primarly because they 404 405 are frequently used by researchers around the world, namely Sepia officinalis, 406 Sepioteuthis lessoniana, Octopus maya and O. vulgaris (Vidal et al., 2014). These four 407 species show versatile characteristics for culture, such as fast growth and high food conversion rates. In addition, these species mate and spawn in captivity, laying eggs 408 409 that, with the exception of *O. vulgaris*, produce large hatchlings. These biological features make them suitable candidates as experimental laboratory animals with a 410 411 potential for aquaculture. However, nearly all zootechnical aspects related to the culture of these species still require improvement and need to be adapted for closely related 412 species from different geographic regions. 413

At present, most of our knowledge on cephalopod culture techniques relies on 414 415 shallow water species. This is due to the relatively easy access to this group of cephalopods, most of them with commercial interest, and to the ease of reproducing the 416 417 characteristics of coastal waters in the laboratory. In contrast, techniques for the maintenance of oceanic or deep sea cephalopods remain virtually unexplored. In 418 particular, little experimental work has been directed towards deep sea octopods (Wood 419 et al., 1998, Hunt, 1999), oceanic squids (O'Dor et al., 1977, Bower & Sakurai, 1996, 420 Hunt, 1999, Bush, 2012, Hoving & Robison, 2012, Villanueva et al., 2012), or polar 421 422 species (Daly & Peck, 2000). However, as research efforts in the open ocean, the deep 423 sea, and polar regions are bound to increase around the world in the near future, methods for the study of captured cephalopods from these regions will be needed to 424 obtain new information on their life cycles and ecology. 425

High-priority research targets in cephalopod culture are the development of
sustainable artificial foods and the control of reproduction (Villanueva et al., 2014).
Littoral cephalopods are carnivorous and require food rich in protein to maintain their

vigorous metabolism, as well as high quality lipids rich in essential fatty acids, 429 430 phospholipids and cholesterol to sustain their fast growth. Recent efforts to obtain artificial foods have shown promising results (Rosas et al., 2008, Rosas et al., 2013, 431 432 Martínez et al., 2014). However, a major challenge will be to develop a sustainable 433 artificial diet independent from fisheries products, completely formulated from plant 434 sources, and in addition supporting good survival and growth, as is now a reality for some marine carnivorous fishes (Watson et al., 2013). The study of feeding dynamics of 435 delicate planktonic paralarvae of cephalopods should also become a priority, because it 436 437 would enable the commercial culture of octopod species such as O. vulgaris, which produce small eggs (Iglesias et al., 2007, Villanueva & Norman, 2008). For example, an 438 439 adapted, enriched Artemia protocol would be desirable to feed planktonic octopods or squids - recent work is currently shedding light on this aspect (Guinot et al., 2013). 440

441 A further area of development required to facilitate cephalopod culture is the control of reproduction and an understanding of the effects of maternal condition on egg 442 443 quality and offspring competence. Currently, egg masses are collected from the field, 444 are obtained by spontaneous spawning in aquaria, or stem from *in vitro* fertilization. As 445 cephalopods are semelparous and often have a natural spawning period restricted to a 446 few months in the year, researchers currently need to adapt their laboratory studies, 447 timing and experimental protocols to the natural sexual maturation period of the target species. The development of methods to accelerate or retard sexual maturation and 448 449 spawning in aquaria will open new experimental possibilities and will be particularly useful to the planning and development of commercial culture. The influence of light 450 451 intensity and photoperiod on sexual maturation has been studied in a few cases (Richard, 1971, Zúñiga et al., 1995) and, if extended, could open new opportunities for 452 453 the control of reproduction. Furthermore, in cephalopod culture, the development of 454 ethical guidelines that aim to reduce pain, suffering and stress are strongly encouraged and should be based on the 3Rs principle, i.e. replacement, refinement and reduction 455 456 (Mather & Anderson, 2007, Moltschaniwskyj et al., 2007, Andrews et al., 2013, Fiorito 457 et al., 2014).

Finally, genetic intervention has already been applied to other metazoans in culture in order to enhance production of cultured animals and to tackle challenges in culture (Hulata, 2001). Such an approach can be expected to have the potential for taking cephalopod culture to the next level. Important new research topics in this respect would be genomic sequencing or studies looking for genes that code for particular traits or that govern protein expression. For example, it would be interesting to identify thegenes responsible for desirable broodstock features, control of sexual maturation,

465 growth, immunology, and pathology.

466

467 4. New ways to research cephalopod adaptations and responses to environmental468 change

469 Cephalopods and climate change (Paul G. K. Rodhouse)

The effects of global climate change will include warming of the atmosphere and the 470 471 oceans, intensification of ocean currents, more frequent and intense extreme weather events, retreat of sea ice in the polar regions, reduction in the depth of the oxygen 472 473 minimum layer and reduced seawater pH (Raven et al., 2005). These physical changes 474 will drive changes in marine ecosystems, which are predicted to reduce biodiversity, 475 although they will not necessarily reduce overall primary and secondary production. However, these effects will not be uniform. Currently, warming of the atmosphere is 476 477 most intense in Alaska, Siberia, and the Antarctic Peninsula. In addition, warming of the ocean surface and upper layers in the vicinity of the Antarctic Peninsula has been 478 479 reported by Meredith and King (2005).

480 Because cephalopods are poikilotherms, they could be expected to physiologically respond to ocean warming. Warming will increase growth rate (subject 481 to food availability and sufficient water oxygen), shorten life span, and increase 482 turnover, which in turn might drive changes in life history parameters (Pecl & Jackson, 483 484 2008). This will only happen if the species do not shift their distribution in response to warming in order to remain within their present thermal environment. However, there is 485 evidence that some species expand their distribution when facing a warmer environment 486 (Zeidberg & Robison, 2007, Golikov et al., 2013). 487

488 Furthermore, many cephalopods, especially the oegopsid squids, produce planktonic paralarvae, which, by definition, are transported by ocean currents and have 489 been shown in some species to be dependent on mesoscale structuring in the ocean to 490 491 complete their planktonic phase (Bakun & Csirke, 1998, Dawe et al., 2000). Such species are likely to be affected by changes in oceanic circulation, the effects of which 492 may be positive or negative. For example, small changes in large-scale circulation are 493 unlikely to affect Antarctic squid, but changes in mesoscale oceanography may have a 494 495 significant impact (Rodhouse, 2013).

Extreme local events such as storms or basin-scale events such as the El Niño 496 497 Southern Oscillation or North Atlantic Oscillation, which are predicted to be intensified 498 by global climate change, will influence changes in populations (Hoving et al., 2013). 499 Basin-scale events are known to drive variability in the recruitment and abundance of 500 species, including Illex argentinus (Waluda et al., 1999), I. illecebrosus (Dawe et al., 501 2000), and Dosidicus gigas (Waluda et al., 2006). Intensification of such events might be deleterious and/or advantageous to these species, but there are currently no models 502 503 which can predict likely outcomes.

In the polar regions, changes in sea ice may cause changes in the distribution of some species, but there are no species known to be dependent on sea ice as, for instance, is the Antarctic krill *Euphausia superba* (Murphy et al., 2007, Constable et al., in press, Xavier & Peck, in press). In these high latitudes, changes in ocean ecology driven by retreating sea ice may have a greater effect on cephalopod populations than the direct effect of ice retreat.

510 At least two cephalopod species, D. gigas and Vampyroteuthis infernalis, are 511 associated with the oxygen minimum layer, where they descend to during daylight 512 (Robison et al., 2003, Rosa & Seibel, 2008, Hoving & Robison, 2012). These two 513 species are physiologically adapted to survive the low oxygen tension of the oxygen minimum layer, and probably enjoy the selective advantage of avoiding active water-514 breathing predators in this zone. Depending on how widespread this habit is among 515 pelagic cephalopods, changes in the oxygen minimum layer associated with global 516 climate change will have effects on other species (Bograd et al., 2008, Stramma et al., 517 2008, Keeling et al., 2010, Gilly et al., 2013). 518

519 Furthermore, all cephalopods possess calcarious statoliths, while some possess 520 larger mineralized structures such as an external shell (e.g. nautiluses) or an internal 521 shell (e.g. cuttlefish). Although there is some evidence that cuttlefish are pre-adapted to 522 ocean acidification (Gutowska et al., 2008), there is still a need for more data on the 523 effects of reduced ocean pH on cephalopods.

Cephalopods evolved from an ancestral mollusc in the Cambrian. They have
survived major extinction events at the end of the Palaeozic and at the end of the
Mesozoic, and have thrived in spite of competition from fishes (Packard, 1972,
Rodhouse, 2013). Although some cephalopod groups such as ammonites and belemnites
became extinct in geological time, the coleoids have survived and radiated. Their life
history traits have adapted them for ecological opportunism and provide them with the

potential to quickly evolve in response to new selection pressures (Murphy et al., 1994,
Murphy & Rodhouse, 1999, Hoving et al., 2013). There is therefore reason to believe
that these characteristics will enable cephalopods to evolve under global climate change,
enabling them to avoid becoming extinct, and ultimately giving rise to new forms
adapted to a new 'greenhouse world'.

535

536 Physiological adaptations of cephalopods to environmental change (Rui Rosa)

Coastal marine ecosystems are warming at a higher rate than most other 537 538 ecosystems (MacKenzie & Schiedek, 2007). Because many coastal organisms already live close to their thermal tolerance limits (Helmuth et al., 2006), ocean warming is 539 540 expected to negatively impact their performance and survival. Cephalopods are some of the most adaptable marine organisms, capable of adjusting their biology (and life 541 542 cycles) according to the prevailing environmental conditions (Boyle & Rodhouse, 2005, Hoving et al., 2013). Yet, although their short life spans and great life history plasticity 543 544 allow them to respond rapidly to new climate regimes, ocean warming may cause serious biological impairments to the more vulnerable early ontogenetic stages, namely 545 546 shorter embryonic periods and an increased likelihood of premature hatching (Rosa et 547 al., 2012b, Rosa et al., 2014). Future changes in ocean chemistry are also expected to pose particular problems. Cephalopods possess statoliths that may be reduced and 548 abnormally shaped (with increased porosity) under hypercapnia (Kaplan et al., 2013). It 549 550 is also noteworthy, that along with the rise of pCO_2 in the embryo (combined with a drop in pH and pO_2), the current record of oxygen tension below critical pO_2 values 551 552 reveals that the harsh (i.e. hypoxic and hypercapnic) conditions inside cephalopod egg capsules are expected to be magnified in the future (Rosa et al., 2013a). Such 553 554 environmental conditions may promote untimely hatching and smaller post-hatching 555 body sizes (Table 1), thus challenging survival and fitness.

In the last few decades, marine hypoxia has become a major ecological concern 556 557 (Diaz & Rosenberg, 2008). Surprisingly, some squids that were thought to be driven 558 from hypoxic areas due to anatomical and physiological constraints (e.g. Dosidicus gigas) instead seem to benefit from expanding hypoxia (Rosa et al., 2013b). 559 Nonetheless, the synergistic impact of these climate-related factors (i.e. hypoxia, global 560 warming, and ocean acidification) is expected to compress the habitable night-time 561 depth range of these vertically migrating squid species due to unfavorable high 562 563 temperature and decreasing pH at the ocean surface (Rosa & Seibel, 2008).

At macroecological scales, a species distribution model (SDM) linked to the 564 565 field of conservation physiology may help to explore future changes in the global patterns of cephalopod diversity. However, the reliability of SDM-based predictions 566 567 needs to be improved, because models often lack a physiological underpinning and rely 568 on assumptions that may be unrealistic under global climate change. For instance, 569 additional information on the limits of thermal tolerance [e.g. maximum critical temperature (CTMax), lethal temperature at which 50% of the sample population dies 570 (LT₅₀)] will improve our ability to predict the effects of climate change on the present 571 572 distribution patterns of cephalopods (Rosa et al., 2008a, Rosa et al., 2008b, Rosa et al., 573 2012a).

574

575 **5.** Future cephalopod research in genetics (Jan M. Strugnell)

576 The volume of research that contains cephalopod genetic sequences has increased markedly over the last 20 years, in particular thanks to the decreasing costs of molecular 577 578 sequencing. Prices are now sufficiently low for sequencing to become an attractive research tool for scientists representing a range of disciplines, including fisheries 579 580 science, systematics, or neuroscience and developmental biology. The next exciting 581 wave of genetic research on cephalopods is approaching as the first cephalopod genomes are being sequenced. Genome sequencing of at least ten cephalopod species is 582 583 currently underway, representing a broad range of taxonomic groups, including Octopus vulgaris, O. bimaculoides, Hapalochlaena maculosa, Sepia officinalis, Doryteuthis 584 pealeii, Euprymna scolopes, Idiosepius paradoxus, I. notoides, Architeuthis dux, and 585 Nautilus pompilius (Albertin et al., 2012). Obtaining high-quality whole genome 586 sequences of cephalopods will stimulate new inquiries by providing a wide range of 587 research opportunities in which a reference genome is required, as well as in the 588 589 interpretation of the genomes themselves.

590 However, the sequencing of cephalopod genomes is not without its challenges, 591 and early work has shown cephalopod genomes to be large and to contain many 592 repeated regions, making sequence assembly difficult (Albertin et al., 2012). In 593 addition, at least one whole genome duplication event has been suggested to have occurred during the evolution of the Cephalopoda (Hallinan & Lindberg, 2011), which 594 may further complicate assembly. Nonetheless, important lessons in sequencing whole 595 molluscan genomes have been learned through sequencing of the few whole molluscan 596 597 genomes that exist to date (i.e. Lottia, Aplysia, and Biomphalaria). But, although bestpractice methods of sequencing and assembly are being implemented (Albertin et al.,2012), the task will not be trivial.

In addition, annotation of cephalopod genomes will likely prove to be a 600 601 significant challenge as well. Part of the annotation process for a novel genome 602 typically involves de novo gene prediction, a task that is known to be difficult and error-603 prone (Albertin et al., 2012, Yandell & Ence, 2012). Large taxonomic distances exist 604 between cephalopods and taxa with well-annotated animal genomes, which will increase the difficulties of annotation. Therefore, the sequencing of corresponding 605 606 transcriptome data will be essential to supplement any de novo predictions, because it definitively identifies regions of the genome that are transcribed, and thus can help to 607 608 identify boundaries between genes through differences in transcript abundance.

609 Despite these initial difficulties, the sequencing of the first cephalopod genome 610 holds great promise for improving our understanding of the evolution and function of this fascinating group of marine organisms. Completely sequenced genomes will 611 612 provide researchers with the ability to thoroughly study the function of different genes and also to investigate evolutionary relationships, not only within cephalopods, but also 613 614 more broadly within molluscs and lophotrochozoans. In addition, whole genomic data 615 of cephalopods will open up fields of research that have to date largely been unavailable 616 or subject to only a handful of studies. Such research areas include epigenetic modification, RNA editing and microRNAs (Albertin et al., 2012). 617

The development of a cephalopod model organism (possibly Idiosepius due to 618 its small size) will allow focused studies of the development of the cephalopod body 619 620 plan. This will facilitate investigation and understanding of many morphological features characteristic of cephalopods that are commonly suggested to be 'vertebrate-621 622 like', such as complex eyes, well-developed brains and highly differentiated vascular 623 and neuroendocrine systems. As such, research of the evolution and development of 624 these features, facilitated by whole-genome data, may not only provide further insight 625 into cephalopod evolution, but also into the evolution of man (depending on whether the 626 similarity of vertebrate structures is superficial or based on genuine homology).

627

628 6. Challenges in cephalopod fisheries and conservation

629 The future trends in cephalopod fisheries (Graham J. Pierce)

630 Historically, cephalopod fisheries have been less important in the northeast Atlantic

631 compared to much of the rest of the world (Caddy & Rodhouse, 1998, Hunsicker et al.,

632 2010b), despite a strong tradition of cephalopod consumption in southern Europe. 633 However, a combination of declines in other fishery resources has led to an increase in 634 directed cephalopod fishing as well as increased attention from fishers, national 635 governments, and fisheries organizations such as the International Council for the 636 Exploitation of the Sea. In Europe, therefore, the short-term trend is likely to be an 637 increased effort in cephalopod fishing, extending exploitation to currently under- or unexploited species, coupled with novel implementation of formal stock assessment and 638 regulated fishing policies. However, it is fairly unlikely that existing stocks can absorb a 639 640 substantial increase in fishing pressure (Royer et al., 2002) and past experience shows 641 that the unpredictable nature of cephalopod abundance tends to discourage commercial 642 fishery interests (Young et al., 2006).

643 These remarks can be generalised to world cephalopod fisheries in the sense that 644 landings have been increasing (at least until around 2005), new species have assumed high importance (notably Dosidicus gigas in the eastern Pacific) and evidence is already 645 646 being seen of overexploitation is some areas (Pierce and Portela, 2014). A key issue will be understanding the rise (and fall) of important cephalopod fisheries, especially those 647 648 of ommastrephid squids such as *Todarodes pacificus*, *Illex argentinus* and *D*. gigas. While we suspect that environmental sensitivity is one key to understanding 649 650 population trajectories, effects of overexploitation may at least partially explain some of the spectacular crashes like that of the T. sagittatus fishery off Norway in the mid-651 1980s. As suggested above, global climate change may have a range of impacts on 652 cephalopod populations and may result in a shift in the relative importance of fisheries 653 654 and environment in controlling population dynamics.

Cephalopod culture, especially for Octopus spp. (Iglesias et al., 2014), may help 655 656 to fill the growing demand for cephalopods in Europe and its export markets. Relevant 657 recent developments in cephalopod culture include in vitro fertilization (Villanueva et al., 2011). Nonetheless, artisanal fisheries will remain important, and are increasingly in 658 659 need of assessment and management that is appropriate to the small scale of the 660 fisheries and the particular biological features of the resource species. However, perhaps 661 the biggest question mark concerns whether exploitation of deep sea cephalopod resources is capable of expansion. Malcolm Clarke, among other cephalopod scientists, 662 suggested that there are vast resources of oceanic squids in the world. His assessment 663 was based on the estimated amount of food needed to sustain the world's sperm whale 664 665 population (Clarke, 1996, Santos et al., 2001). This potential resource presents an

enticing opportunity for fisheries, but others have cast doubt on the large abundance of
such species. In addition, a practical challenge relates to palatability, although fishery
companies are currently developing processing methods for ammonium-rich squid
tissues to permit their marketing as food products.

670 Fisheries management and governance in Europe is currently undergoing a 671 revolution, with the implementation of an integrated ecosystem assessment and management approach as part of the reform of the Common Fisheries Policy, while at 672 the same time looking ahead to a future integrated marine management, in which 673 674 fisheries are simply one of many relevant sectors. The move towards an ecosystem approach to fisheries is of course not unique to Europe. However, the steep increase in 675 676 data requirements (compared to single species assessments) presents a real obstacle, 677 especially in a period of economic recession; thus, alternative approaches based on 678 indicators and expert judgement are also likely to be needed. In this context, the Marine Strategy Framework Directive (MSFD) of the European Union is relevant, as it focuses 679 680 on the development of indicators of ocean health. At least in the United Kingdom, there are plans to develop cephalopod indicators for the MSFD. As a final note, cephalopod 681 682 waste from fishery processing, and cephalopod species of lesser interest for human 683 consumption, may be increasingly used in animal feedstuffs, fertilizers (Fetter et al., 684 2013), or other industrial products such as pharmaceuticals.

685

686 *Cephalopod conservation (A. Louise Allcock)*

Assessing the conservation status of a wide range of cephalopod taxa reveals just how 687 little is known about many species. Studies carried out for the International Union for 688 the Conservation of Nature (IUCN) Red List, focusing on different higher cephalopod 689 690 taxa (e.g. Sepiida, Oegopsida, Cirrata), have found that between about 50 and 75% of species in these higher taxa are 'Data Deficient' (Kemp et al., 2012). Many species are 691 692 known from just a few specimens, so that little is known about their biology and 693 ecology. In some cases, we can conclude that species meet the IUCN category of 'Least 694 Concern' simply because their very wide geographic distribution and high fecundity with planktonic dispersal means that they are unlikely to be impacted across their entire 695 distribution range, despite the possible existence of local threats, so the lack of data is 696 actually under-reported. 697

In particular, data are lacking for cirrate octopods. These cephalopods arepotentially long-lived, are slow to reach maturity and have low fecundity (Collins &

700 Villanueva, 2006). Opisthoteuthis, the most shallow cirrate genus, is characterized by a 701 close association to the benthos, and is therefore the genus most affected by commercial 702 deep sea trawling. O. chathamensis was considered 'Nationally Critical' on the New 703 Zealand Red List (Freeman et al. 2010) and Collins and Villanueva (2006) suggested 704 that populations of other species may already have declined as a result of deep sea 705 trawling. However, a lack of specific population data and information on fisheries 706 impacts will likely prevent many potentially vulnerable species being listed in a category other than 'Data Deficient'. Therefore, one of the future challenges for 707 708 cephalopod biologists is to improve the quality and consistency of population estimates 709 for all cephalopod species, particularly those subjected to direct or indirect 710 anthropogenic impacts, including fishing.

711 Taxonomic issues may also prevent the actual vulnerability of a species from 712 being reflected in its conservation assessment. Recent dramatic declines in the size of the Sepia apama population in the upper Spencer Gulf (South Australia) have been well 713 714 documented (Hall, 2008, Hall, 2010), but attempts to have this population listed as 'Critically Endangered' under Australia's Environment Protection and Biodiversity 715 716 Conservation Act 1999 failed (Anonymous, 2011), apparently because the population had not been formally described as a distinct species, despite little evidence of it inter-717 breeding with other populations (Anonymous, 2011). However, a temporary localised 718 ban on fishing was enacted in 2013. S. apama was assessed as 'Near Threatened' on the 719 720 IUCN Red List (Barratt & Allcock, 2012), but this assessment considered the whole range of the species, as is normal practice. The IUCN assessment notes that "If the 721 722 population in the upper Spencer Gulf is shown to be a separate species then the Spencer Gulf species would be assessed as Endangered.". 723

724 Conservation efforts for Nautilus are similarly hindered. The slow growth and 725 low fecundity of nautiluses (Dunstan et al., 2011) make them vulnerable to fishing pressure and several overfished populations have crashed (Dunstan et al., 2010). The 726 very wide distribution range reported for N. pompilius suggests that threats are likely to 727 728 be local, until one considers recent genetic data. For example, molecular phylogenetic work (Bonacum et al., 2011, Sinclair et al., 2011, Williams et al., 2012) indicates that N. 729 pompilius comprises several distinct phylogenetic species. This suggests that the impact 730 of fisheries is far more likely to lead to species extinctions than previously thought. 731 However, descriptions of individual species within the *N. pompilius* species complex 732 733 and accurate information on the range of these species are required if conservation

- ristings are to reflect the perceived vulnerability to anthropogenic impacts. Therefore,
- ensuring that all cephalopod species are accurately described, and that species
- complexes and cryptic species are distinguished, constitutes an essential future
- range for cephalopod conservation.
- 738

739 **Discussion**

Cephalopods will continue to attract scientific interest, particularly in the fields of 740 physiology, genetics, ecology and fisheries. Furthermore, the traditional scientific 741 742 disciplines of taxonomy and morphology are currently being rejuvenated by the application of new technologies. Studies on cephalopods will continue to range from the 743 744 organismic level (e.g. physiology, behaviour), to the species level (e.g. taxonomy, 745 systematics, population dynamics, distribution, abundance), and finally to the ecosystem 746 level (e.g. fisheries, biodiversity, conservation). In addition, new cephalopod research is 747 emerging on issues such as global climate change and ocean acidification or habitat and 748 food-web modelling.

Cephalopods constitute an important trophic link between the lower levels of 749 750 food webs and top predators (Young et al., 2013). About 800 species of extant 751 cephalopods have been described, but we only have sufficient data to understand the life 752 history (e.g. distribution, habitat, feeding ecology, reproductive biology) for approximately 60 species (Jereb & Roper, 2005, Jereb & Roper, 2010, Jereb et al., 753 754 2014). Therefore, taxonomists and geneticists must increasingly work together to ensure that specimen data uploaded to databases are based on correctly identified specimens. 755 756 The combination of molecular genetics, DNA barcoding, and digital morphological techniques offers new ways to resolve numerous outstanding issues in cephalopod 757 758 taxonomy and evolution. In this context, an increase in molecular work is of particular 759 importance, because the lack of transcriptomic and genomic information, for example, 760 has limited advances in neurobiology research, where cephalopods act as model 761 organisms (Zhang et al., 2012).

Cephalopods have several interesting traits, which make them suitable model organisms for broad evolutionary research. For instance, they have one of the largest size range of any metazoan class and could therefore become model species for studying metazoan growth and metabolism. Furthermore, cephalopods show a remarkable diversity of life history traits and a better understanding of evolutionary relationships among cephalopods would help to determine the plasticity of these traits or could reveal simple switches between individual strategies. In addition, because of
the presence of mineralized structures or the planktonic early life stages, most
cephalopod species may be highly sensitive to global climate change and/or ocean
acidification, because of the presence of mineralized structures or the planktonic early
life stages. Hence, cephalopods should be increasingly used as model organisms to
predict the effects of global warming on ocean life (Hanlon et al., 1989, Rodhouse,
2013).

In addition, a quantitative PCR approach should finally allow reliable 775 776 identification of cephalopod species as prey. Also, because top predators are still a major source of information on cephalopods, novel techniques in trophic research such 777 778 as the analysis of stable isotopes, DNA, or fatty acids as well as 3D imaging will 779 complement the data obtained by conventional means (Jarman et al., 2004, Barrett et al., 780 2007, Karnovsky et al., 2012). These latter techniques have suffered due to a decline in taxonomists actually able to perform this type of work (Pearson et al., 2011). 781 782 Furthermore, the use of ecological tracers, especially fine-scale analyses of tracer molecules within informative structures such as statoliths, beaks or shells will offer new 783 784 insights into stock structuring and individual life history (Cherel & Hobson, 2005, 785 Cherel et al., 2009a, Ramos & Gonzalez-Solis, 2012). In addition, recent improvements 786 in specimen tagging now allow studying movements of cuttlefish and squid (Gilly et al., 2006, Semmens et al., 2007, Wearmouth et al., 2013). If tag weight could be further 787 reduced and some attachment issues resolved, this approach might be extendible to 788 smaller cephalopod species or earlier developmental stages. 789

790 Future research should certainly also focus on the ecology of cephalopod species, particularly for those species with immediate commercial fishery interest. As a 791 792 result of the increasing international capacity to explore deeper environments, deep sea 793 cephalopods will attract the attention of fisheries and research. For example, the 794 increasing amount of deep sea imagery calls for creative solutions to compiling and 795 using such data. Advanced and more complete morphological data will improve our 796 ability to identify specimens based on photographic records alone. For the well-known 797 commercial cephalopod species, long-term monitoring and the establishment of marine protected areas will be the primary focus of discussion in cephalopod conservation. 798 Furthermore, the usage of the continental shelf slope, deep sea, and oceanic areas by 799 800 numerous pelagic predators and cephalopods is a further area that will receive attention 801 from conservationists (Harris et al., 2007, Game et al., 2009, Tancell et al., 2012). In

order to catch fast-swimming cephalopods, efforts should be channeled into the use of
more efficient nets that allow catching sub-adult or adult stages of the larger species.
Incorporating such research foci into major multidisciplinary projects could become
essential for success in obtaining funding.

806 At present, the effects of global climate change, linked with acidification, 807 warming, and expanding hypoxia, perhaps represent the biggest threat to certain species of cephalopods, but also constitute a challenge to researchers, policymakers, and society 808 at large. From a scientific point of view, one of the greatest challenges in this discipline 809 810 will be to discriminate between the effects of global climate change and fisheries on cephalopod populations. In this regard, experimentation has always been an important 811 812 approach to resolving open questions in cephalopod research. From a technological 813 perspective, cephalopod culture should be further developed to meet challenges such as 814 the development of sustainable artificial foods or the control of reproduction and genetic manipulation. In addition, the successful maintenance of deep sea and oceanic 815 816 cephalopods in captivity would be a major step forward to understanding their life 817 cycles and would contribute to assessing the potential impact of fisheries targeted at 818 other species in their habitats. Such research efforts would also constitute an important 819 contribution to cephalopod conservation efforts (Hoving et al., in press).

820 Finally, collaboration, in particular between scientific disciplines, is essential for 821 tackling some of the big scientific challenges the world is currently facing. Early career scientists, such as the CIAC Young Researchers group, should make ample use of 822 novel, digital approaches to networking, communication, and collaboration. Social 823 824 media, along with digital repositories as well as new data and research sharing protocols, will continue to facilitate international and interdisciplinary research on 825 826 cephalopods and related scientific areas. Furthermore, education and outreach initiatives 827 are bound to follow suit, resulting in the increased dissemination of cephalopod science 828 to a wider audience.

829

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- 1342
- 1343
- 1344 Figures

1345Figure 1. Different per capita growth rates of prey, according to the presence or absence

of predators. (A) Hyperbolic response. (B) Sigmoid response. See the main text forexplanation. Modified from Turchin (2003), with copyright permission from Princeton

- 1348 University Press.
- 1349

1350 Figure 2. Analysis of cephalopod specimens using non-invasive imaging techniques.

- 1351 The two- and three-dimensional visualizations shown here are based on a MRI dataset
- 1352 of a whole museum wet specimen of *Bathypolypus arcticus* (A, B), a µCT dataset of the
- 1353 dry shell of *Spirula spirula* (C-E), and a µCT dataset of a tungsten-stained wet specimen
- 1354 of *Idiosepius pygmaeus* (F, G) (dataset courtesy of Brian D. Metscher).

1355

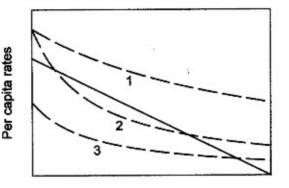
1356 Tables

1357 Table 1. Responses of different cephalopod life stages to ocean acidification.

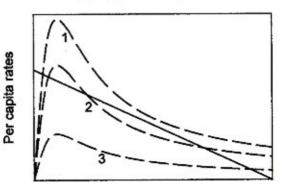
Species	Life stage	Ecological	Effect	Reference
		parameters		
C	E ush mare		T 1 4	(Daga at al
Sepia	Embryo	<i>p</i> CO ₂ 400–1650	Lower survival rate;	(Rosa et al.,
officinalis		ppmv; pH 8.0-7.5;	metabolic	2013a)
		18 and 22°C	depression;	
			premature hatching;	
			lower hypoxic	
			threshold	
Sepia	Juvenile	<i>p</i> CO ₂ 705–6068	4% daily increase in	(Gutowska et
officinalis		ppmv; pH 8.0-7.1;	body mass; mass of	al., 2008,
		16-17°C	calcified cuttlebone	Gutowska et
			increased 5- to 7-	al., 2010)
			fold	
Loligo	Embryo	<i>p</i> CO ₂ 424–1680	Lower survival rate;	(Rosa et al.,
vulgaris		ppmv; pH 8.0-7.5;	shorter mantle	2014)
		13, 15, 17, and 19°C	length; premature	
			hatching; greater	
			incidence of	
			abnormalities;	
			metabolic	
			depression; lower	
			thermal limit;	
			increased heat shock	
			response	

Doryteuthis	Paralarva	<i>p</i> CO ₂ 390–2200	Increased time of	(Kaplan et
pealeii		ppmv; pH 8.0-7.3;	hatching; shorter	al., 2013)
		20°C	mantle length;	
			statoliths with	
			reduced surface area;	
			abnormally shaped	
			statoliths with	
			increased porosity	
			and altered crystal	
			structure	
Dosidicus	Juvenile	рСО2 380-1000	Metabolic	(Rosa &
gigas		ppmv; pH 8.0-7.5;	depression; reduced	Seibel, 2008)
		10°C	aerobic scope;	
			reduced level of	
			activity	

(a) Hyperbolic response



(b) Sigmoid response



Prey density

