

1 **Evaluation of wetted surface area of commercial ships as biofouling habitat flux to the United States**

2
3 **Running title:** Flux of underwater ship surface habitats

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15 **ABSTRACT**

16 Commercial ships inadvertently transfer vast numbers of living organisms beyond their evolutionary
17 ranges, sometimes resulting in invasions of distant marine habitats. Biofouling on ship hulls translocate
18 organisms that cling to the undersides and interstices of ships that function as hard substrate habitat for
19 biota. Because biofouling accumulates over space and time continually, it poses risk to all ports visited.
20 To better understand the potential magnitude of the biofouling vector in the United States, we compiled
21 information on ship-specific dimensions as well as actual arrival histories of the fleets of ships calling at
22 U.S. ports (2011-2014) in an effort to calculate wetted surface area (WSA) flux to the U.S. The annual
23 mean flux of WSA from overseas bioregions to the U.S. is $333 \text{ km}^2 \cdot \text{y}^{-1}$. An additional $177 \text{ km}^2 \cdot \text{y}^{-1}$ of WSA
24 moves among the eight distinct biogeographic regions of the lower 48 United States. We confirm that
25 over 90% of all global marine bioregions (120 of 132 identified by IUCN) are visited by commercial ships
26 within five port calls of arriving to the U.S. Our analysis is the first ever to quantify the extent of WSA
27 flux among global marine bioregions and underscores the urgent need for management approaches and
28 technologies that will reduce associated invasion risks.

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30 **Key words:** aquatic nuisance species, biofouling, hull fouling, marine invasive species, non-indigenous
31 species, vector

INTRODUCTION

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Biofouling of commercial ship hulls has long been of concern to industry due to the adverse effects fouled hulls have on drag and fuel use (e.g., Callow & Callow 2002, Schultz et al. 2009, 2011). Further, the transfer of living organisms that cling to the hull or occupy niche spaces of ships outside the hydrodynamic slipstream and shear forces (e.g., sea chests, thruster tunnels, etc.) is widespread and recognized as a dominant vector for the transfer and introduction of marine species beyond their natural, evolutionary ranges (Ruiz et al. 2000, Carlton 2001, Fofonoff et al. 2003, Minchin 2006). Despite its operation over millennia, ship biofouling is a potent contemporary source of biological invasions throughout the world, with diverse management and policy approaches underway to minimize associated ecological and economic impacts (e.g., IMO 2011, Davidson et al. 2016).

Historically, wooden hulled vessels moved slowly and spent more time in port than modern-day ships (Carlton 1985, Hewitt et al. 2009). These ships were especially prone to biofouling and even structural damage. Damages from boring animals such as the molluscan shipworm (*Teredo navalis*) were dangerous and costly to the maritime industry due to the destruction of hulls and wooden pilings and support structures in harbors and marinas (Atwood & Johnson 1924, Carlton 1985). Biofouling and the widespread transport of marine biota associated with ship hulls remains active today and may actually be increasing in recent time for multiple reasons. First, some effective biocides have had unintended and adverse environmental consequences and are being discontinued, including the global ban on tributyl and organotin (Nehring 2001, Hewitt et al. 2009, IMO 2001). Second, the magnitude of shipping has been increasing over time, with vessels becoming more numerous and larger, increasing opportunities for biofouling colonization and transfers.

Ships' underwater surfaces, like other submerged artificial and natural hard substrates, serve as habitat for an enormous array of benthic marine species that settle, grow, and reproduce. Indeed, ships can be likened to mobile islands that shift habitat and associated species from place to place (Godwin

56 2003). The extent or proportion of a ship's hull that is fouled is related to a range of factors, including
57 voyage routes and biological source regions, vessel operational profiles, maintenance schedules, and the
58 amount of colonizing organisms that can remain attached to in-service ships. However, to date, few
59 papers describe the extent and diversity of biota on submerged surfaces of contemporary commercial
60 ships (Gollasch 2002, Davidson et al. 2009, Inglis et al. 2010, Sylvester et al. 2011), especially in the
61 context of global shipping. As a result, accurate predictions of species assemblages and their flux around
62 the world are lacking (Ashton et al. 2016).

63 Understanding the magnitude of biofouling on ships as a mechanism for species transfer is
64 important but complex because of the overlapping influence of factors related to biological colonization
65 of ships' surfaces (and subsequent introduction in ports of call) and shipping behavior (Inglis et al. 2010).
66 Furthermore, the effect of oceanic passage or shifts in parameters such as temperature and salinity
67 when transiting inter-ocean corridors, such as the Suez and Panama Canals, on hull biota is not yet fully
68 understood. Similarly, biological responses to changing environmental conditions associated with
69 emerging shipping routes such as Arctic passages (Miller and Ruiz 2014) are not yet known. Thus,
70 despite the significant investment of ship owners in maintaining clean hulls (\$5 billion per year and
71 growing) of tens of thousands of ships plying the world's oceans, there are still many fundamental gaps
72 in knowledge about biofouling communities associated with the wide diversity of vessel types and
73 routes involved in global trade.

74 The magnitude of biofouling species transfers is related to both biological content of vessels and
75 number of vessels arriving for a particular time period and geographic location. For the latter
76 component, wetted surface area (WSA) is a common standardized measure of the maximum submerged
77 surface area of a ship, and it represents the area of that ship that can potentially be colonized by marine
78 organisms (Davidson et al. 2016; Moser et al. 2016). Despite not being a direct biological measurement,
79 WSA provides a useful proxy for understanding the potential for ships to transfer biofouling organisms

80 and how this is partitioned by vessel type, source or recipient regions, and time. The extent of biofouling
81 that occupies WSA is highly variable and somewhat idiosyncratic, and is influenced by time since dry-
82 docking, operational history, maintenance and a range of other factors (Inglis et al. 2010). This study
83 estimated annual flux of total WSA to marine bioregions of the U.S. from both vessels arriving from
84 overseas (OS) bioregions as well as vessels engaged in coastwise (CW) traffic that transit U.S. bioregion
85 boundaries. In addition, we compared the relative contribution of different vessel types and different
86 geographic source regions to the total annual flux. Finally, we estimated the WSA for niche areas by
87 vessel type, for OS and CW vessel traffic. Our analysis illustrates clearly that OS WSA exposure to U.S.
88 bioregions is uneven, suggesting important geographic components to vessel biofouling invasion
89 opportunity.

90 **MATERIALS AND METHODS**

91 Using formalized naval architecture formulas, we calculated the WSA for a variety of prevalent
92 ship types and modeled the relationship of WSA to an independent parameter (Net Register Tonnage) to
93 determine WSA for individual ships. Combining data of ship arrivals (2011-2014) to the United States
94 with WSA values for multiple ship types, calculated using the relationship between WSA and NRT, we
95 were able to estimate the annual flux of WSA to ports in the U.S. Using a commercially available
96 database of ship identity and statistics for the world fleet, IHS Fairplay World Register of Ships
97 (<http://www.ihs.com/products/maritime-information/ships/world-register.aspx>), we collated
98 dimensional measurements for six ship types (Bulkers, Tankers, Passenger, Container, Roll-on Roll-Off
99 [RORO], and General Cargo). We then calculated WSA for several thousand individual ships of each
100 category using the naval architecture formula of Van Maanen & Van Oossanen (1988). An independent
101 dimension available for all ships, net register tonnage (NRT), was used to regress NRT to each vessel's
102 corresponding WSA, yielding regression models that predict WSA from NRT for each ship type; this step
103 was necessary, because the WSA of all ships arriving to the U.S. could not be calculated directly, as all

104 ships' dimensional measurements were not available. This model was then applied to the population of
105 ships arriving to U.S. ports according to geographic arrival region, last port region, and ship type in
106 2011-2014, as identified by the U.S. Department of Homeland Security's National Vessel Movement
107 Center (NVMC). We confined our regression models to vessels of 100 NRT or greater to reflect arrivals
108 by ships of this size and greater. The dataset of 373,833 arrivals included, in some cases, multiple
109 arrivals to U.S. ports by a single vessel.

110 To evaluate WSA flux among global bioregions, and specifically to understand flux to U.S.
111 regions, we estimated WSA of individual ships and paired these with actual voyage histories and arrivals.
112 Importantly, ships connect geographically and biologically separated segments of the globe (i.e., ships
113 move among marine bioregions that have evolved in various degrees of isolation) and move some
114 organisms well beyond their native ranges. For this reason, we chose to bin individual arrivals according
115 to the marine bioregion to which each ship arrived. We used the International Union for Conservation of
116 Nature (IUCN) marine biogeographic regions (Kelleher et al. 1995), which includes eight regions in the
117 contiguous U.S. (i.e., lower 48 states, see Figs 1 & 2). By definition, all OS arrivals to the U.S. cross
118 between separate marine bioregions, so all were considered in this analysis as potential vectors for
119 species transfers. In contrast, only CW traffic that moved between separate bioregions were considered
120 potential invasion vectors for the purpose of this analysis, although we recognize invasions can occur
121 within bioregions (especially as newly established invaders spread). For each of the 373,833 OS and CW
122 arrivals considered here, the last ports of call and arrival ports were assigned to IUCN marine
123 biogeographic regions. This approach enabled us to differentiate vessel transits that crossed *between*
124 bioregions from those that remained *within* a single bioregion, and in this manner, each transit's
125 potential for transferring invasive species among unique biogeographic regions containing distinct
126 assemblages of marine species was assessed.

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128 **WSA calculation and regressions**

129 WSA was calculated for a subset of ships in each ship-type category as follows (Van Maanen &
130 Van Oossanen 1988):

131
$$WSA = L(2T + B)C_M^{0.5} \left(0.4530 + 0.4425C_B - 0.2862C_M - 0.003467\frac{B}{T} + 0.3696C_{WP} \right) + 2.38\frac{A_{BT}}{C_B}$$

132 Where:

133 L = length overall

134 T = average molded draft

135 B = breadth

136 C_M = midship coefficient

137 C_B = blocking coefficient

138 C_{WP} = waterplane coefficient

139 A_{BT} = cross-sectional area of bulbous bow (calculated as a percentage of the immersed area of midship)

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141 The coefficients and bulb area percentages for different vessel types are published in Van Maanen &
142 Van Oossanen (1988). Although length should technically be the waterline length of the hull, such data
143 are not available for the commercial fleet, and so length overall was used. For each ship type, the WSA
144 per ship was regressed on an independent univariate measure of ship size (NRT), and the models with
145 the best fit were selected.

146

147 **WSA Flux**

148 The four years of ship arrivals data was used to estimate the annual flux of WSA to the U.S.,
149 expressed as $\text{km}^2 \cdot \text{y}^{-1} \pm 1 \text{ SD}$). Additionally, we evaluated the global source regions and associated
150 magnitudes of WSA flux to each U.S. biogeographic region using four-year totals. Our analyses focused
151 on commercial ocean-going ship types that are capable of making long haul passages across oceans or

152 CW transits (i.e., Bulker, Container, Passenger, RORO, Tanker, and General Cargo). Some ships, for which
153 information was not widely available (e.g., Military, Tug/Barge), were excluded from this analysis. This
154 approach provides the first ever estimate of WSA flux to the United States (or elsewhere) and across
155 distinct bioregions. Two-way ANOVAs were used to test for differences in WSA by ship type and
156 receiving coastal region for both OS and CW flux. WSA was log-transformed prior to analysis to conform
157 to ANOVA assumptions.

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159 ***Niche Area Flux***

160 Mean annual niche area flux was estimated for each ship type investigated, both for OS and CW
161 arrivals. Niche areas consist, broadly, of rudders, propellers, propeller shafts, thrusters, sea-chest grates,
162 bilge keels, and dock-block surfaces for each ship type. Niche area was calculated by multiplying mean
163 annual WSA flux by a ship-type-specific multiplier developed by Moser et al. (in review) in an analysis
164 that quantified niche area by ship type, based on the global commercial shipping fleet. Moser et al. (in
165 review) concluded that an area equal to 10.2% of the WSA of the global fleet could be attributed niche
166 areas. Of this percentage, about half or $\approx 5\%$ of total WSA is included in the WSA calculations (e.g., dry
167 dock strips) and $\approx 5\%$ is additional submerged surface area not included in WSA calculations. Applying
168 the niche area multipliers to ship type WSA gives a conservative estimate of niche area flux, since on
169 average, it discounts niche area by approximately 5%. Nevertheless, applying these across the fleet of
170 vessel types that call at U.S. ports highlights the relative extent of niche WSA, that is, the areas with the
171 highest probability of biofouling occurrence (i.e., hot spots for fouling, Coutts & Dodgshun 2007,
172 Davidson et al. 2016). Based on Moser et al. (in review), the following multipliers were used to estimate
173 the fraction of total WSA attributable to niche areas that are likely hotspots for biofouling organisms:
174 Bulker (7%), Container (9%), General Cargo (9%), Passenger (27%), RORO (15%), and Tanker (8%).

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199 **Overseas Flux.** In the United States, the Gulf of Mexico coast (CAR-I) received by far the greatest
200 annual OS WSA flux of any biogeographic region, 152 km², approximately 46% of total OS WSA, more
201 than double that of the next busiest bioregion (the Mid-Atlantic region, NA-ET3), and more than 27
202 times the WSA flux observed in the Northeastern states (NA-ET2; Fig. 1). The Gulf of Mexico, Mid-
203 Atlantic, and Southeastern U.S. (CAR-VII) receive a combined 246 km² (74% of total annual OS flux).
204 Southern California (NEP-VI) had the highest flux of OS WSA of the Pacific Coast bioregions, raising the
205 OS WSA flux of the top four bioregions to 292 km² per year (88% of total annual OS flux). The
206 contribution of each ship type to annual WSA arrival varied greatly among U.S. bioregions (Fig. 1).

207 **Coastwise Flux.** By comparison, the mean annual WSA flux across bioregions of the U.S. (i.e., CW
208 traffic) was 177 km², an area similar in size to Washington, D.C. (Fig. 2). CW flux is dominated by four
209 bioregions (CAR-VII, NEP-V, NA-ET3, CAR-I) whose combined WSA exposure is 139 km², 79% of the mean
210 annual CW WSA flux (Fig. 2). The Gulf of Mexico, Southeastern U.S., and Mid-Atlantic regions account for
211 58% of total CW WSA flux. In most cases, OS WSA flux exceeded CW flux in each bioregion, with the
212 exceptions of Central California (NEP-V) and the Southeastern U.S. (CAR-VII) which receives 330% and
213 119% more CW than OS WSA, respectively.

214 If compared at the coastal scale, the Gulf of Mexico coast (CAR-I) receives 46% of all OS WSA flux
215 but only 14% of CW WSA. The East and West coasts are more balanced in WSA flux: East coast receives
216 30% OS and 46% CW flux; West coast receives 24% OS and 30% of CW WSA flux (Figs 1 & 2).

217 **Flux by Ship type.** In most recipient bioregions there were marked shifts in the ship types that
218 dominate OS and CW traffic patterns (Figs 1 and 2). For example, in Southern California (NEP-VI) there
219 was a strong shift from 70% (OS) to 33% (CW) arrivals by Container ships accompanied by a parallel
220 increase in Tanker arrivals from 15% (OS) to 48% (CW). The inverse relationships exist for Containers and
221 Tankers arriving to Central California ports (NEP-V). When compared at the national scale, the
222 relationship of OS and CW fluxes also varied by ship type. WSA flux from Containers and Roll-on Roll-off

223 vessels were evenly balanced between OS and CW WSA flux, compared with all other ship types for
224 which OS flux far outstrips CW flux (Fig 3.).

225 There were 13,820 unique vessels engaged in trade with U.S. ports during the timeframe
226 studied (Table 2). Less than 2% of vessels were engaged in exclusive trade among U.S. ports, whereas
227 48% arrived exclusively from OS, and 50% operate in both OS and CW modes. A large majority of
228 Containers (79%) and ROROs (86%) operate both OS and CW, but the ratio of 'Both':'OS' for these ship
229 types was 4.1 and 8.0, respectively, while all other ship type had ratios of <2.0 (Table 2). Bulkers are only
230 ship type dominated by exclusive OS trade (67% of unique vessels).

231 Two-way ANOVAs indicated that there were highly significant differences in mean annual
232 overseas WSA among vessel types, but that these differences were not equal among the regions ($F_{34,139} =$
233 $73.03, p < 0.001$). A similar statistical pattern was confirmed for CW WSA flux, ($F_{35,143} = 71.55, p < 0.001$).
234 These results reinforce the fact that strong differences in WSA flux exist among U.S. bioregions and that
235 different assemblages of ship types serve them.

236 **Niche Area Flux.** Figure 3 also describes the estimated quantity of WSA corresponding to niche
237 areas, surfaces that are widely recognized as more prone to biofouling than submerged portions of the
238 hull (Coutts & Taylor 2004, Davidson et al. 2016). Overseas arrivals by Passenger vessels, although
239 representing just 13% of overall WSA flux, comprised 33% of the total niche area arriving to U.S. ports
240 from OS. By comparison, the OS flux of WSA by Containers and Tankers was more than two times
241 greater than that of Passenger vessels, but the flux of niche area attributable to these ship types was
242 roughly two-thirds that of Passenger vessels. This disparity was directly related to the prevalence of hull
243 thruster tunnels and other niche areas on Passenger vessels, estimated as 27% of total WSA (Moser et
244 al., in review). Based on differences in design, namely the extent of niche area, some ship types may be
245 more readily fouled and thus potent vectors of invasive biofouling species. However, owing to
246 differences in operations and trade patterns, ship types with similar OS WSA and niche area flux (e.g.,

247 Containers and Tankers) may pose very different threats to CW transport of ship biofouling species (Fig.
248 3).

249 **Geographic Source Richness and Global Connectedness.** As ships move from place to place,
250 their hulls are exposed to ever-increasing varieties of biota. Thus, knowing the voyage history of ships
251 provides insight into the diversity of bioregions visited prior to calling at a recipient port of interest. By
252 compiling the geographic history of arrivals to U.S. bioregions, we were able to compare the cumulative
253 source bioregion richness for each recipient region. Figure 4 plots total bioregion richness accumulated
254 over four yrs for each U.S. receiving bioregion by both one and five LPOCs. There is substantial variability
255 in richness among U.S. receiving bioregions with one LPOC (62.8 ± 6.38 bioregions). The Gulf Coast (CAR-
256 I) is directly connected to more than twice as many source bioregions as Puget Sound (NEP-III). There
257 was strong convergence among recipient regions when connecting to five LPOC (101.5 ± 2.25 per
258 receiving bioregion; mean and SE). The combined total LPOC-5 richness for the contiguous US was 120
259 bioregions and the world is comprised of 132 bioregions under the IUCN scheme. As a whole, the US is
260 sampling 91% of the world's coastal bioregions.

261 Mapping source bioregions accumulated by ships prior to arrival in U.S. bioregions across four
262 yrs enabled us to visualize how each region of the U.S. is connected to the rest of the globe. Strong
263 regional differences are especially apparent when evaluating the most recent last port (LPOC-1; Figure
264 S.1., Supporting Information).

265 When the arrival history of ships was traced back beyond the immediate last port of call (e.g.,
266 from one to five last ports) the domain of global source/donor regions expands for all U.S. recipient
267 bioregions (Fig. S.2, Supplemental Information). For example, connections to Southeast Asia, Australasia,
268 and the Indian Ocean accumulate for all 8 U.S. recipient bioregions, but the North Atlantic remains a
269 relatively minor source of WSA for West coast ports and bioregions. As voyage route history is more
270 fully accounted for, last ports of call accumulate, a much greater range of biotic source regions are

271 'sampled' by ships, and greater numbers of inter-ocean transits result in accumulated exposures with
272 potential physiologically stressful conditions (e.g., Panama Canal (warm water, low salinity exposure),
273 the Suez Canal (warm water, marine-hypersaline exposure).

274 Our investigation sought to characterize and quantify both the extent and geographic nature of
275 WSA flux across distinct marine biogeographic regions. Unlike ballast water (BW)—the most widely
276 recognized transport vector associated with commercial ships, which has a growing variety of
277 technologies and procedures by which invasion risk can be minimized (e.g., Minton et al. 2005, Miller et
278 al. 2011, Davidson & Simkanin 2012)—there are no adopted technologies or procedures that are both
279 highly effective and broadly used to prevent ship biofouling mediated invasions. In fact, explicit
280 management of ship biofouling to prevent marine invasions is only emerging at present despite the long
281 history of hull coating use (e.g., anti-fouling and foul-release) by shippers to promote vessel
282 performance and fuel efficiency (Dafforn et al. 2011, Davidson et al. 2016). Hull coatings represent the
283 focal point of the shipping industry's biofouling management approach, which has had inadvertent
284 benefits for invasion prevention, but modern ship biofouling remains a potent vector of marine invasive
285 species (e.g., Inglis et al. 2010, Ruiz et al. 2011) and biofouling management for explicit biosecurity
286 purposes is being promoted more broadly around the world (IMO 2011, Wells & Booth 2012, New
287 Zealand Government 2014).

288 A ship that arrives to a U.S. port, or any port of the world for that matter, may pose near-zero
289 risk of species introduction from ballast water (e.g., if no ballast water is discharged into a port, there is
290 no risk of associated BW introduction); however, as ships move from place to place, there is continual
291 opportunity for colonization of the submerged surfaces and a concomitant opportunity for biofouling
292 species to disembark to the surrounding environment. Biota may be dislodged through accidental
293 contact with a wharf or from the forces exerted on thruster tunnel inhabitants when thrusters are
294 activated. Alternatively, and more commonly, animals and seaweeds may reproduce at any port during

295 the 'opportunity window' that aligns with their reproductive life cycle. Moreover, stress-induced
296 spawning may be an important mechanism of release from ships when organisms are exposed to highly
297 variable and sometimes barely tolerable conditions that trigger spawning events in ports (Minchin &
298 Gollasch 2003). In the absence of perfect anti-fouling solutions, ships experience the continual
299 development and replacement (due to competition and death) of species assemblages that essentially
300 integrate across the places they visit. In this sense, ship biofouling is a concatenation vector (C. Hewitt
301 pers. comm.), perhaps more so than ballast water even though individual ballast tanks can contain
302 water from different ports. Conversely, ships may leave a trail of biota in their wake, reflective of their
303 particular voyage history, as has been postulated for certain voyage routes as explanations of both
304 marine and terrestrial species distributions (Darling et al. 2012, Gotzek et al. 2015).

305 It was clear from our study that the total OS WSA flux was nearly double that of CW WSA flux
306 (333 km² vs. 177 km²). Interestingly, this value is similar in magnitude to the estimated total WSA of the
307 entire global fleet of active commercial ships (\approx 120,000 commercial ships in the world fleet with
308 estimated 325 km² WSA, Moser et al. 2016). Nevertheless, some marine bioregions receive far more OS
309 flux than CW flux (e.g., Gulf of Mexico, Southern California, and the Mid-Atlantic) while others were
310 dominated by CW flux (e.g., the Central/Northern California coast). These patterns no doubt directly
311 reflect underlying commerce patterns to and from these regions, related to hub-and-spoke, point-to-
312 point, short-sea, and pendulum models of trade and transport geography (Rodrigue et al. 2013). Such
313 patterns imply that the conditions experienced by ship biofouling organisms will be quite different,
314 depending on voyage history characteristics like route, voyage length, potential exposure to divergent
315 environmental conditions, and overall differences in biota being mixed. Furthermore, overlaid on all of
316 these parameters is climate change, which will further complicate these relationships.

317 By definition, all ship arrivals to the U.S. from OS artificially connect unique marine
318 biogeographic regions and thus pose some risk of transferring invasive biofouling species (and

319 populations) to coastal habitats of the United States. Likewise, ship voyages emanating from U.S.
320 bioregions threaten to export North American biota to other parts of the world. However, it is critical to
321 recognize that CW ship transits can transfer species among distinct bioregions, either by mixing U.S.
322 natives across U.S. biogeographic boundaries or by secondary movement of an introduced invasive from
323 one bioregion to another. It is also important to note that the same individual ship can engage in an OS
324 arrival and subsequently conduct one or more inter-bioregion CW arrivals, as is the model voyage
325 pattern for certain Containers (e.g., a pendulum voyage route). In this case, OS biofouling from a trans-
326 or inter-oceanic source can be transferred to the original port of entry *and* adjacent bioregional ports,
327 while the OS and domestic bioregional biotas can coexist on the same vessel.

328 A recent study by Moser et al. (in review) based on >120,000 commercial vessels that were part
329 of the active global fleet (1999-2013) concluded that the equivalent of 10% of the total available WSA
330 was comprised of niche areas. These are locations where biota accumulate most often on ships'
331 submerged surfaces, typically areas outside the laminar hydrodynamic slipstream, and they are
332 locations that can be more difficult to maintain than hull surfaces (Coutts & Taylor 2004) and where
333 biofouling does not impose an immediate cost or performance penalty to ship operators (Davidson et al.
334 2016). Although some areas, such as dry dock strips and bilge keels are included in WSA calculations,
335 other niche areas (e.g., propellers, rudders, sea chests, thrusters and thruster tunnels, etc.) are in excess
336 of typical WSA estimates, providing an additional $\approx 5\%$ of area that can be colonized by biofouling.
337 Further, the extent and make-up of niche area is highly dependent on ship type, ranging from niche
338 areas that are 7% of total WSA for Bulklers to 27% for Passenger ships. Because these niche areas tend to
339 foul more readily, they are an elevated concern from an invasion risk and biosecurity perspective.
340 Although we did not formally parse the fleet of vessels visiting U.S. ports into its proportional hull and
341 niche area representation, applying niche area percentages of overall WSA for mean annual fluxes by
342 ship type provides a conservative estimate of biofouling hot spots associated with vessel flux in the U.S.

343 WSA acts as the potential surface area available for colonizers, but the actual or realized percent
344 cover of colonized space is likely highly variable and often linked to niche area hotspots for macro-
345 organisms (Coutts & Taylor 2004, Davidson et al. 2009). Although larger studies (n > 30 ships) of
346 biofouling on modern shipping have occurred in recent years (Inglis et al. 2010, Thomason 2010,
347 Sylvester et al 2011), the amount of ship sampling that has occurred is not commensurate with the scale
348 of the phenomenon and the range and complexity of factors that influence biofouling on ships. As such,
349 the degree to which WSA acts as a proxy measure for biofouling is not known. WSA is probably a more
350 accurate proxy measure for the extent of microbial organism transfers that are traversing the globe in
351 biofilms, however. Despite anti-fouling technology, biofilms develop quickly on ships' surfaces and
352 sterile surfaces are not possible, so biofilms develop to varying degrees across their surfaces.
353 Fundamental questions regarding microbial biogeography remain unanswered (Hughes Martiny et al.
354 2006), however, and an even greater dearth of micro-organism investigations from ships' submerged
355 surfaces (compared to macro-organism studies) hinders our understanding of biofilm composition,
356 diversity, extent, and transfer on ships (Hunsucker et al. 2014. Leary et al. 2014).

357 How biofouling organisms from coastal environments withstand prolonged exposure to pelagic
358 conditions during ocean crossings is not well documented. Likewise, the potential detrimental or lethal
359 effects of temperature and salinity on exposed biofouling organisms that pass through the Panama or
360 Suez Canals is not yet established. The expansion of these canals will amplify flux of ships (and WSA)
361 between oceanic bodies, but is also expected to strongly affect the strength of connections among ports
362 (Muirhead et al. 2015, Galil et al. 2015). Both conditions increase the opportunity for species transfer;
363 however, passage through physiologically stressful environments may actually serve to filter or kill some
364 fraction of the taxa on the undersides of vessels. If such a mechanism actually operates, then the
365 frequency with which any given vessel traverses the Panama or Suez Canals may affect its degree of
366 biofouling and by extension, its risk for moving invaders. Riverine and atypical salinity ports can exert

367 similar disturbances on ship biofouling, and analyses of port network structure and environmental
368 match among ports provide a useful basis for spatial comparisons of invasion risk (Kaluza et al. 2010,
369 Keller et al. 2011).

370 As ships begin opting for Arctic routes, such as the Northern Seaway and Northwest Passage as a
371 shortcut between oceans (Miller 2014), biofouling species will be increasingly exposed to cold water
372 without drastic changes in salinity. It is altogether possible that mortality and moribundity to biofouling
373 organisms will differ among inter-ocean corridors. Indeed, the spread of marine organisms between the
374 North Pacific and North Atlantic via the Arctic Ocean has already been documented (Reid et al. 2007). If
375 Arctic routes prove less destructive to ship biofouling than historically active inter-oceanic passages, the
376 prospect for accelerated exchange of species across oceans and invasions of the Arctic seems likely
377 (Miller and Ruiz 2014). Floerl (2014) has suggested that Arctic nations consider adopting a biosecurity
378 approach based on that developed by New Zealand to reduce ship-mediated marine invasions.

379 **WSA and BW Vector Contrast.** Both WSA and BW represent vectors that transfer entire species
380 assemblages (acting as ‘habitat vectors’), and therefore interesting to compare and contrast. When
381 mean annual OS WSA is plotted against mean annual discharge of OS ballast water to the U.S. (BW; NBIC
382 2016), there is a strong positive linear correlation between the two: mean annual BW Discharge ($10^6 \cdot \text{m}^3$)
383 $= 0.3668 \cdot \text{mean annual WSA (km}^2\text{)}$, $r^2 = 0.9182$). The characteristics of ship biofouling and ballast water
384 vectors offer a contrast in biotic uptake by an external and internal vector of ships; ship biofouling
385 accumulates throughout the inter-dry-docking period of each ship, which may result in a higher level of
386 bioregion sampling *per ship* via biofouling than ballast water (the sediment accumulating in ballast tanks
387 and microbial biofilms on ballast tank walls notwithstanding). These patterns underscore that invasion
388 opportunity related to commercial shipping is strikingly uneven across the coastal United States. Indeed,
389 the volume of unmanaged BW (either via open ocean BW exchange or onboard BW treatment)
390 discharged to the Gulf of Mexico is more than three-fold greater than unmanaged BW discharge to the

391 rest of the nation (NBIC Annual Report 2014). Federal regulations have required BW management from
392 ships discharging water from OS since 2004 (33 Code of Federal Regulations 151.2005), but uneven
393 compliance with such regulations have left large regions of the coastal U.S. under-protected from BW-
394 borne invaders.

395 **Management Implications.** The limitations to effective BW management in the United States
396 and globally (Miller et al. 2011) are minor compared to those associated with biofouling of commercial
397 ships. Although there are IMO Guidelines concerning best practices for ship biofouling and invasive
398 species (IMO 2011), and an emerging number of regional and state programs (McClay et al. 2015), U.S.
399 federal regulations are incomplete with respect to biofouling management for biosecurity purposes such
400 that their influence on the behavior of ship operators is unclear and their impact on invasion risk
401 reduction is likely minimal. Currently New Zealand is the only nation state to have proposed an explicit
402 policy for biofouling management and (soon-to-be) mandatory biofouling standards for invasion risk
403 reduction (New Zealand Government 2014). Other countries have also proposed IMO-style guidelines,
404 while some regions and states, particularly those with sensitive or highly-valued marine environments,
405 impose mandatory rules governing levels of biofouling on arriving vessels (McClay et al. 2015). In the
406 U.S., and most of the world, dry-docking frequency typically occurs on three to five-year intervals, as
407 suggested in the International Convention for Safety of Life at Sea and for ship classification purposes
408 (Takata et al. 2011). Dry-docking is aimed primarily at inspection, maintenance, and repair of a ship's
409 submerged surfaces, equipment, and processes for structural and insurance reasons, and is nearly
410 always accompanied by hull cleaning and re-application of coatings (typically anti-fouling or foul-release
411 coatings). While periodic cleaning will reduce the risk of invasions and often resets this vector's biota to
412 zero (until a biofilm is formed), efficacy of biofouling management strategies will depend on frequency
413 of dry-dock visits, appropriate matching of coatings to operational profiles, and the quality of cleaning

414 and coating application. For all coating types, but especially for foul-release coatings, the application
415 process is quite prescribed and must be precise to ensure proper efficacy over time.

416 In-water cleaning (IWC) of underwater hull surfaces by divers or robotic cleaning technologies
417 are management approaches designed to scour commercial ship hulls of their developing biological
418 communities between dry-dock visits. Oftentimes, reduced vessel performance triggers IWC, like a
419 downturn in propulsion efficiency and increased fuel costs. Such reactive cleaning can result in
420 depositing non-native species into recipient environments. More frequent pro-active IWC may impede
421 the development of extensive and biodiverse fouling communities, thereby reducing the risk of ship
422 biofouling introductions. However, until hurdles regarding the incidental release of both viable biota and
423 toxicants into local waters (McClay et al. 2016) are overcome, IWC may remain a response rather than a
424 preemption of biofouling problems. Both pose serious environmental concerns that mandate
425 improvements to the technology and practice before IWC can be truly effective for a broad range of
426 vessels.

427 Forward-looking management that reaches beyond issues of ship performance and addresses
428 marine bioinvasions seems prudent (Davidson et al. 2016), especially for a marine invasion vector as
429 enduring and globally potent as ship biofouling (Darwin 1854, Hewitt & Campbell 2010). Biofouling has
430 lagged behind BW management but is emerging as an international policy issue. Whether international
431 guidelines or a few nascent management and regulatory programs will generate sufficient awareness
432 and environmental concern to gain widespread attention from policy-makers in the U.S. and elsewhere
433 is unclear. Given the scale of unfettered WSA in motion across Earth's oceans, combined with rapidly
434 expanding and new shipping routes, the strength of port-to-port connections and the structure of
435 shipping networks is certain to change, bringing with it changes to invasion risk worldwide.

436

437 **ASSOCIATED CONTENT**

438 **Supporting Information.** Analysis and mapping of the biogeographic histories of ships that arrived to
439 U.S. bioregions (2011 – 2014), based on their immediate and last five ports of call. Comparison of WSA
440 and BW flux to recipient U.S. bioregions.

441

442

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648 **Table 1.** Relationships between net registered tonnage (NRT) and wetted surface area (WSA) for six
649 commercial ship types. Regressions are based on first order polynomial relationships with sample sizes
650 (N) of 1709 to 8005. Mean WSA \pm 1 SEM by ship types are included.

651
652 **Table 2.** Number of unique vessels calling at U.S. ports (13,820 during 2011-2014) by ship type,
653 according to the type of trade each is involved in (Coastwise [CW], Overseas [OS], and Both).

654

655 **Figure 1.**
656 Arrivals of ships' WSA from overseas sources. Average (\pm 1 SEM) annual WSA entering each U.S.
657 bioregion from overseas ports is shown for the nation (central panel; average 336.45 km² to the U.S. per
658 year). The mean annual (\pm 1 SEM) contribution of each of six ship types to each bioregion's total is
659 shown in the eight perimeter panels. B- Bulkers; C- Container ships; P- Passenger ships; R- Roll-on Roll-
660 off (Auto) Carriers; T- Tankers; G- General Cargo ship types.

661
662 **Figure 2.**
663 Arrivals of ships' WSA from coastwise voyages. Average annual WSA entering each U.S. bioregion from
664 bioregion crossing coastwise traffic is shown for the nation (central panel). The average coastwise flux
665 among U.S. bioregions was 178.42 km². The mean annual (\pm 1 SEM) contribution of each of six ship types
666 to each bioregion's WSA total is shown in the eight perimeter panels. Ship type codes are the same as
667 Fig. 1 y-axes differ from Fig. 1.

668
669 **Figure 3.**
670 Mean annual WSA and niche area flux to U.S. ports from overseas and domestic bioregions. Ship type
671 codes are the same as Fig. 1

672
673 **Figure 4.** Total number of source bioregions accumulated across 4 yrs by the collective ports in each U.S.
674 receiving bioregion from the immediate single last port of call (LOPC, grey) and 5 LOPC (gray + white).
675 The mean bioregion richness from 5 LPOC (\pm 1 SEM) = 101.5 \pm 2.25 source bioregions.