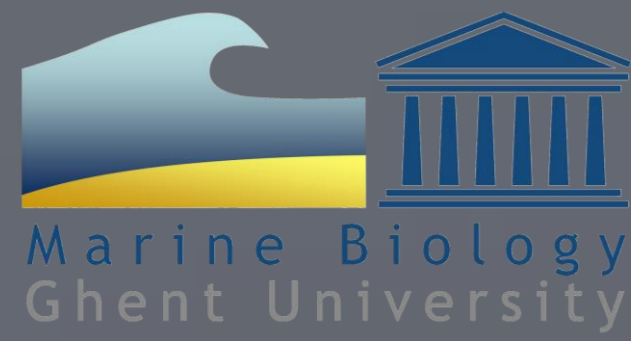


SUBSAMPLING ACCURACY IN BEAM TRAWL CATCHES



Samuel Van de Walle^{1,2}, Jochen Depestele^{1,2}, Hans Polet¹, Kris Van Craeynest¹ & Magda Vincx²

¹Institute for Agricultural and Fisheries Research, Animal Sciences – Fisheries ILVO, Biological Environmental Research, Ankerstraat 1, 8400 Oostende, Belgium

²Ghent University, Biology Research Group, Marine Biology Section Krijgslaan 281 (Campus Sterre – S8), B-9000 Gent, Belgium



Figure 1: Catch Sorting in Progress

Introduction

In commercial beam trawling data on fish catches is routinely available, while composition of invertebrate discards is ignored. However, from an ecosystem perspective such data is equally important yet is highly labor intensive to acquire. Subsampling the discards, and accepting the error this inherently imposes, is the only option. Here we investigate these error rates, which will allow for more accurate data or estimates in discard composition.

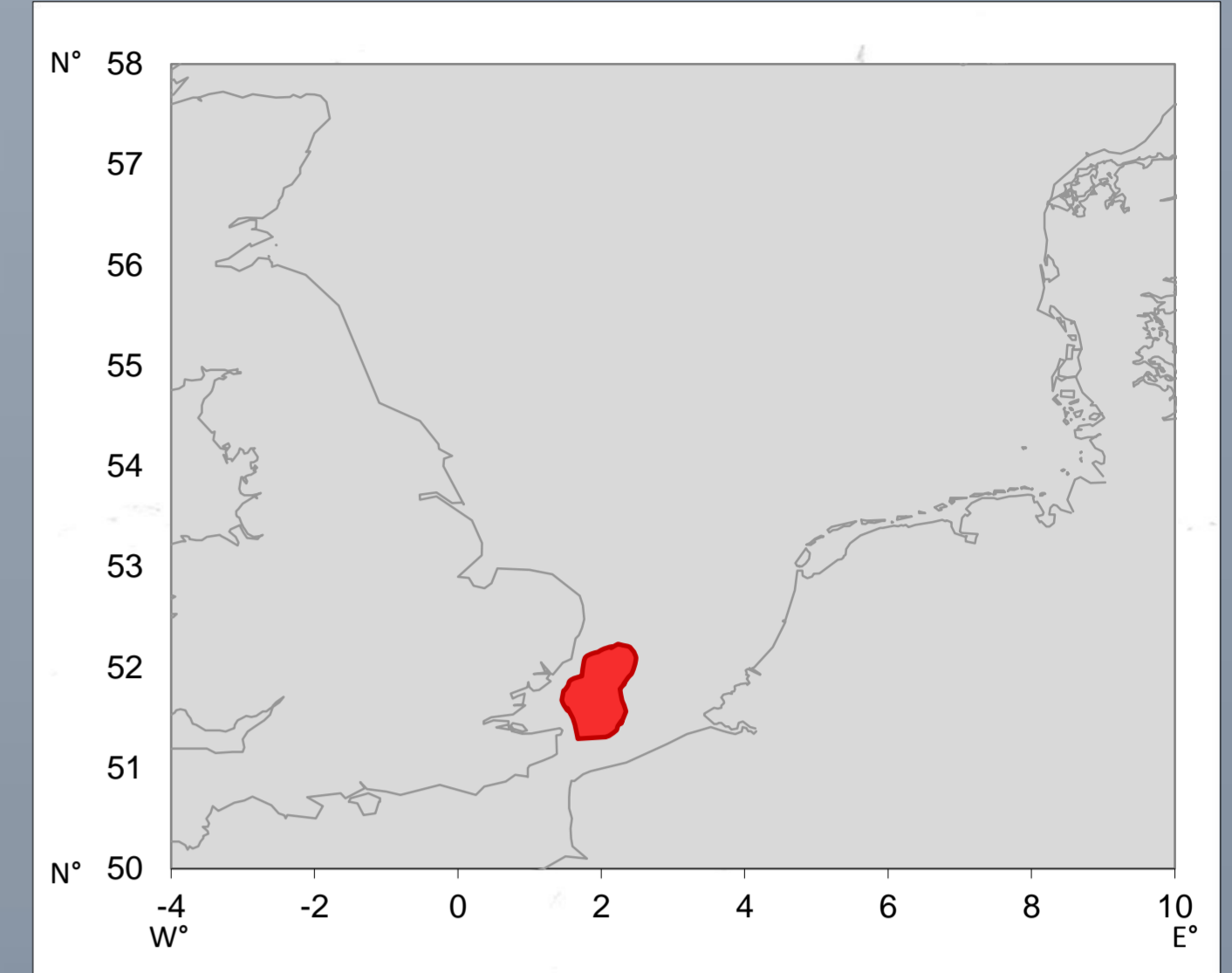


Figure 2: Trawling Location

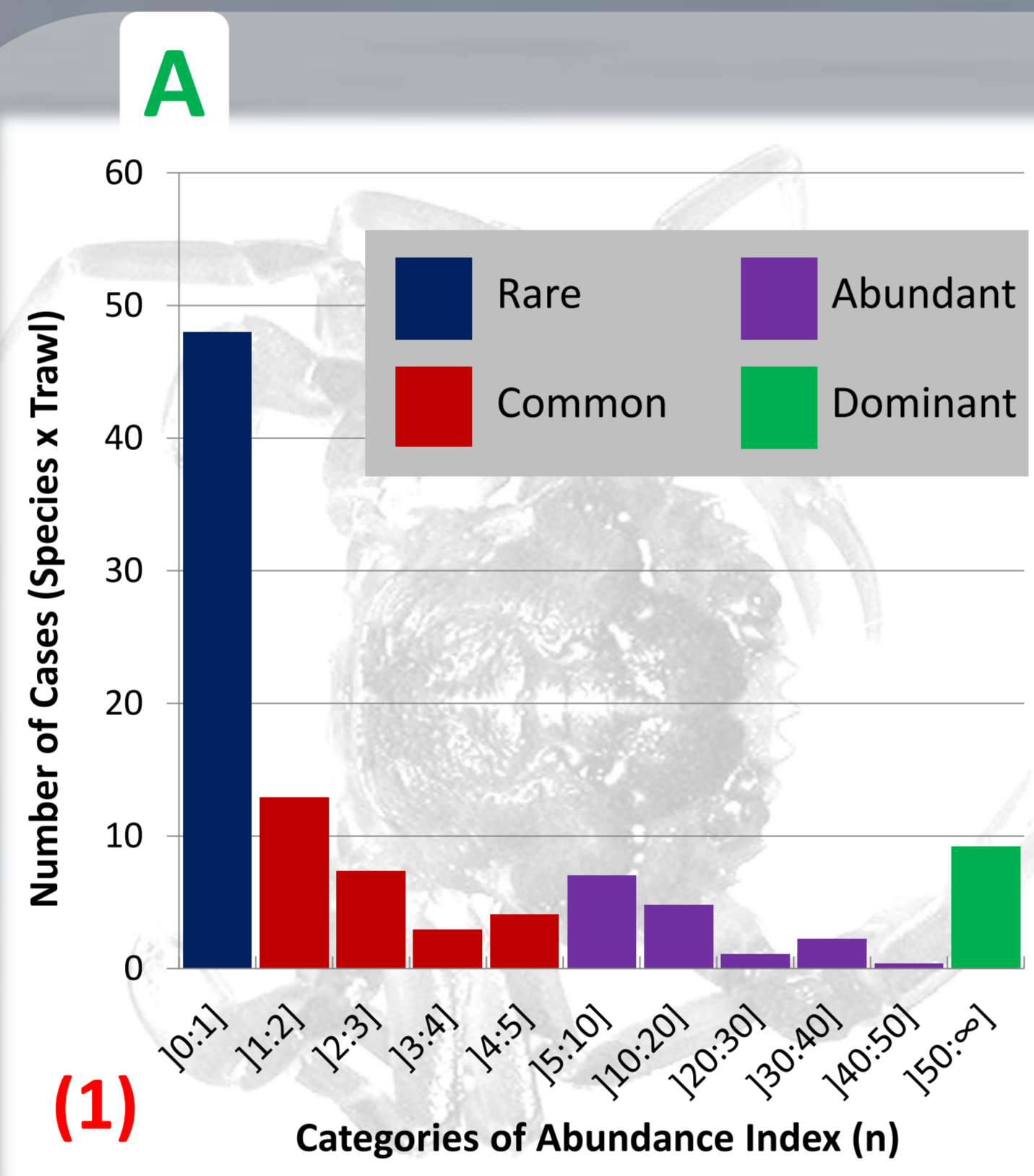
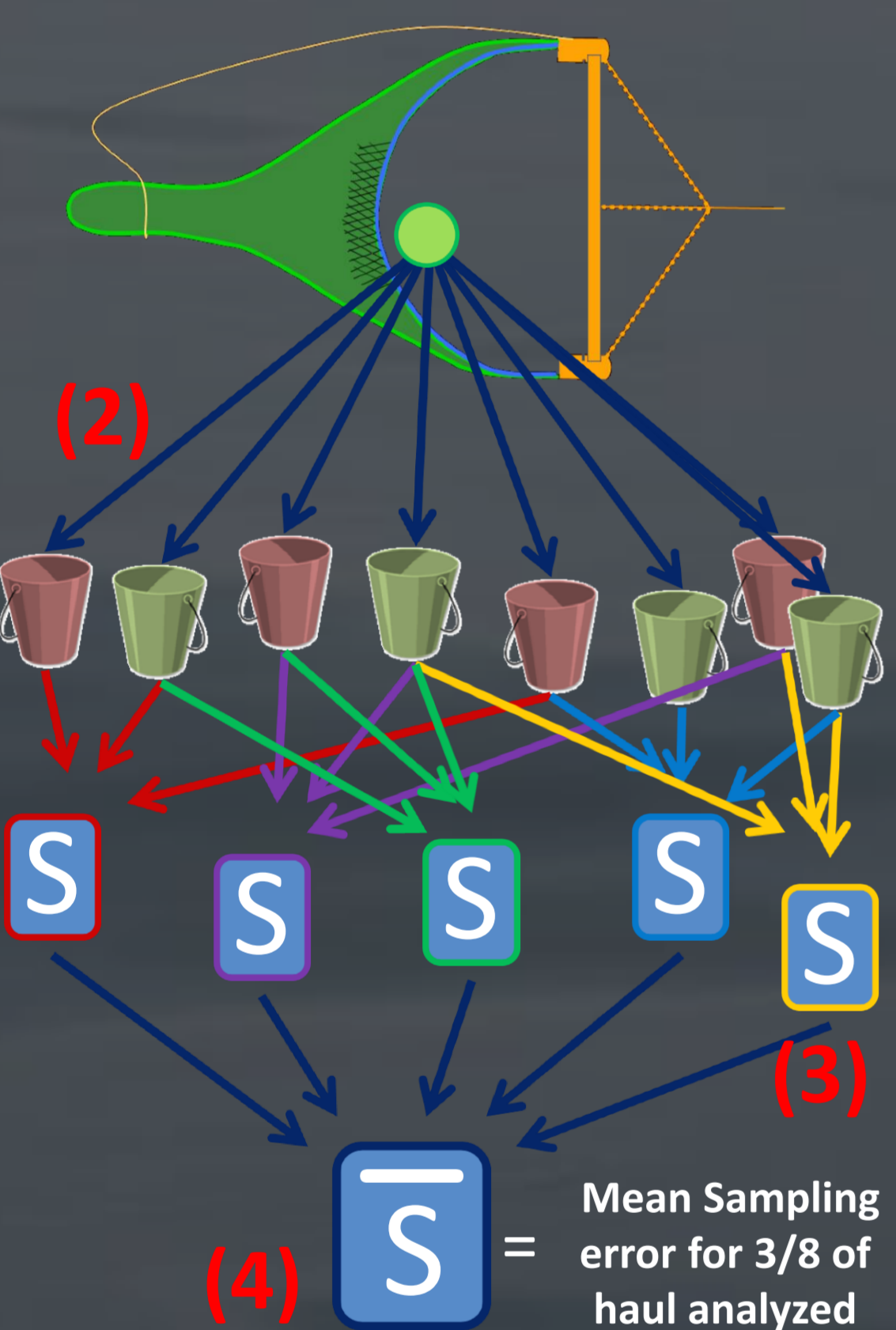
Materials & Methods

Twelve trawls were performed, in February 2009 (F1-F3), February 2010 (F4&F5), April (A1) and November 2011 (N1-N6) at the location in Fig. 2. Every species x trawl combination was assigned to an abundance category (1) based on an abundance index (n). Hauls were subdivided into 10L buckets (2) and individuals were identified. Different numbers of buckets (from 1 to all) from a trawl were used to simulate different subsample sizes. For a large number of random recombination's of buckets, a sampling error (S) was calculated (3) and from all these a mean error and a confidence interval was derived (4). A similar approach was then used to calculate the mean number of species in a subsample of a certain size.

$$n = 10 * (T/W)$$

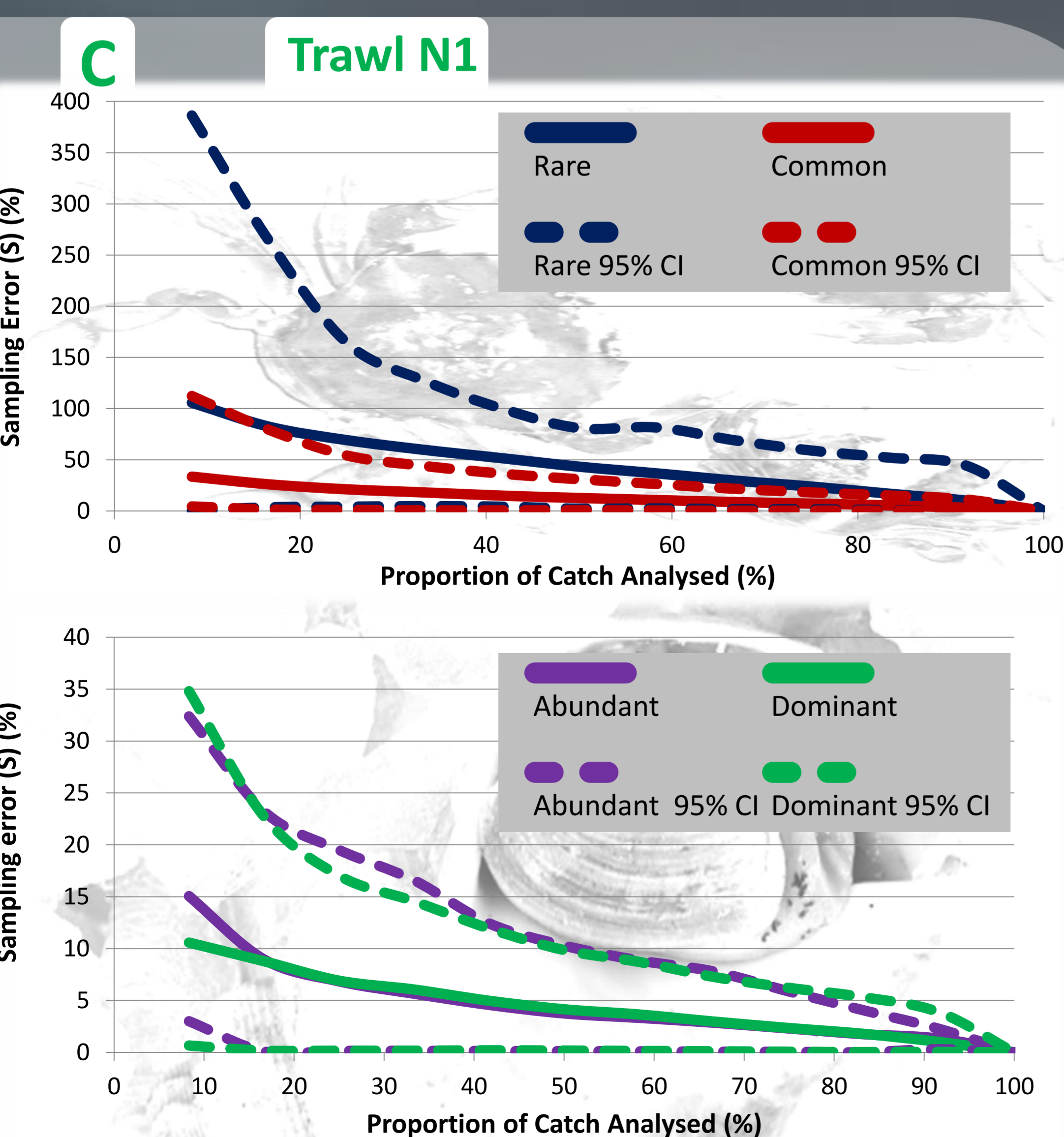
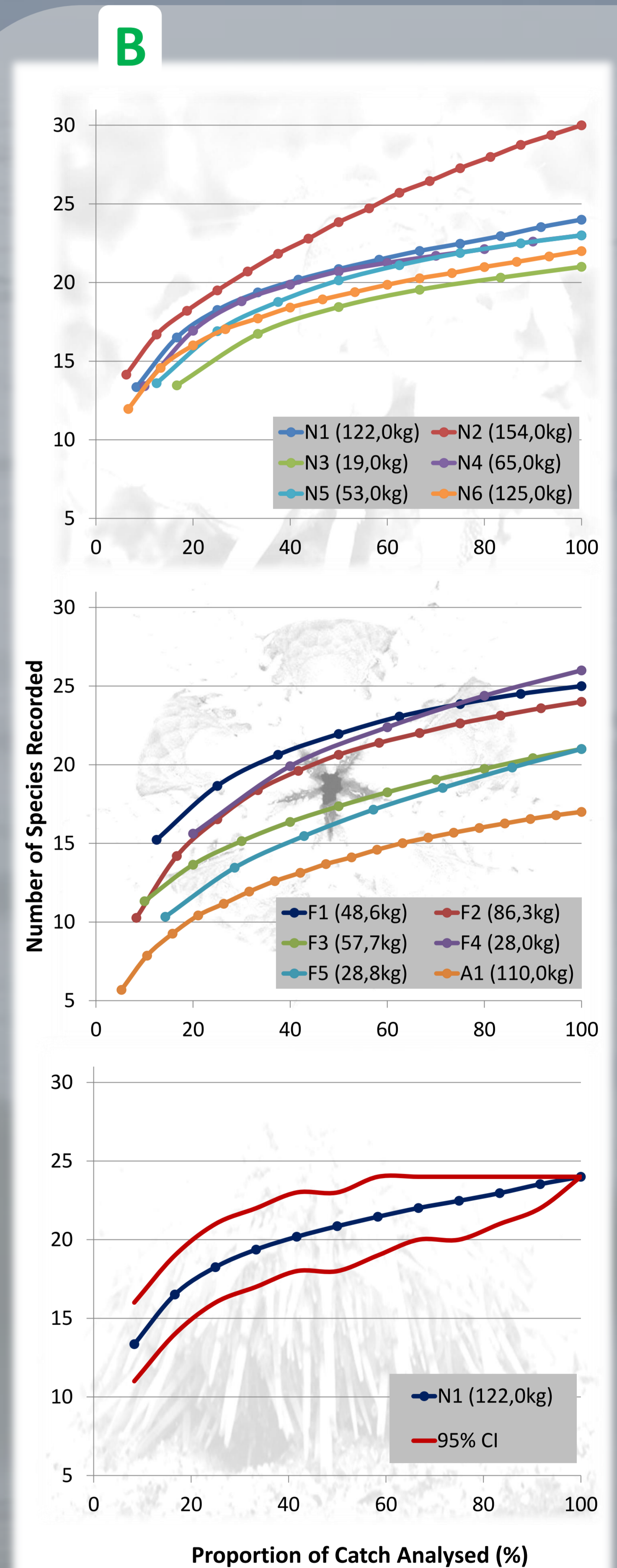
$$S = \left| \frac{a/p - T}{T} \right|$$

p = proportion of catch analyzed
a = Species abundance in subsample
T = Species abundance in haul
W = Total weight of haul



Species	Individuals	%
<i>Asterias rubens</i>	24068	47,19
<i>Echinidea</i> sp.	21324	41,81
<i>Buccinum undatum</i>	901	1,77
<i>Pagurus bernhardus</i>	675	1,32
<i>Liocarcinus depurator</i>	636	1,25
<i>Liocarcinus holsatus</i>	608	1,19
<i>Aphrodita aculeata</i>	416	0,82
<i>Palaemoninae</i> sp.	318	0,62
<i>Necora puber</i>	247	0,48
<i>Ophiura ophiura</i>	236	0,46
...
Totaal	51001	

Table 1: Top Abundant Species



Discussion

(A) Species Abundances

All hauls were highly dominated by starfish and sea urchins. A majority of species was found to be rare.

(B) Species Richness

An average of 23 species per trawl was found. Despite large differences in catch weight there was comparatively little variation in number of species. Most trawls follow a similarly shaped curve regardless of their weight or number of species. For N1 the confidence interval showed a constant range over all subsample sizes.

(C) Sampling Error

Results for N1 are given as example, other trawls revealed similar patterns. Average error decreased with increasing subsample size. Rare species showed the highest error rates while common species had intermediate error rates. For abundant & dominant species errors were up to 10 times lower and surprisingly similar to each other. This indicates that after a certain abundance threshold is reached (around 5 ind./10kg), error rate is unlikely to drop much further, even at highest abundances.

Future Prospects

Currently most results are represent all trawls individually. In the next step, we aim to produce overall results for all trawls combined. Interpolation and/or a general linear mixed model approach should allow for this, while also accounting for the complex interdependence of all data points.

Corresponding author:

+32(0)472 31 94 88 samuel.vandewalle@outlook.com