



The Potential for Upscaling Kelp (*Saccharina latissima*) Cultivation in Salmon-Driven Integrated Multi-Trophic Aquaculture (IMTA)

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Integrated multi-trophic aquaculture (IMTA) has the potential of reducing open-cage fish farming impacts on the environment while also introducing new value chains. The aim of this study was to investigate the growth and composition of the kelp *Saccharina latissima* in salmon-driven IMTA, and to assess the spatial extent of the influence of salmon derived nitrogen in order to evaluate the upscaling potential for IMTA. *S. latissima* was cultivated 100, 200, and 1,000 m east and 1,000 m west of a 5,000 tons salmon farm in Western Norway from February to September 2013. The proportion of salmon derived nitrogen available for the kelp showed a clear decline with distance from the farm. Accordingly, the kelp cultivated near the salmon cages grew faster during the spring season, and growth rate decreased with increasing distance from the farm. A spatially explicit numerical model system (SINMOD), including compartments for dissolved nutrients and kelp growth, was tuned to the field data and used to investigate the potential for upscaling IMTA production. The model was used to introduce a new metric—the *impacted area IA*—for the areal effects of IMTA in terms of the increase in production by IMTA. The model showed that a 25 hectare kelp farm in the vicinity of the studied salmon farm could take up 1.6 of the 13.5 tons of dissolved inorganic nitrogen released during kelp cultivation, amounting to almost 12% of the ammonia released during the cultivation period from February to June. The 25 hectare kelp farm would have a production yield of 1,125 tons fresh weight (FW), being 60% more than that of a non-IMTA kelp farm, while a 20% increase of kelp FW could be obtained over a 110 hectare area in salmon-driven IMTA. To achieve an even mass balance, an area of approximately 220 ha⁻¹ would be needed to cultivate enough kelp to fix an equivalent of the nitrogen released by the fish.

Keywords: nitrogen, bioremediation, stable isotope, numerical modeling, Norway

INTRODUCTION

Norway is the world's largest producer of Atlantic salmon (*Salmo salar*) with a current production of 1.3 million tons per year (FAO, 2017). Atlantic salmon farming traditionally takes place in open sea cages and involves the use of nutrient rich formulated diets. One of the challenges of this open cage production are the waste discharges from fish excretion and feed loss. Even with a low feed conversion ratio of around 1.1 and relatively low losses of feed (3%), there is a substantial release of metabolic waste products such as feces and excreted inorganic nutrients in salmon aquaculture (Wang et al., 2012, 2013). It is estimated that about 60% of feed nitrogen and 70% of feed phosphorus is released into the environment as metabolic wastes. This means that the aquaculture industry fertilizes the Norwegian coast with 52,000 tons of nitrogen and 10,000 tons of phosphorus annually (Wang et al., 2012). Accumulation of effluents in the marine environment may lead to stimulation of phytoplankton growth and eutrophication of pelagic ecosystems, but this is highly dependent on the assimilative capacity of the area and on local hydrodynamics. The area of influence of effluents in surface water ecosystems is dependent on many factors including the biomass produced at the farm, farming practices, and especially environmental conditions such as local topography, water exchange rates, and water depth (Nordvarg and Håkanson, 2002). A full-scale fertilization experiment has suggested that the production capacity of Norwegian coastal waters is quite high (Olsen et al., 2014), and with the current releases of dissolved nutrients from Norwegian aquaculture, these effluents are unlikely to cause unacceptable degradation of surface water quality on a larger scale (Wang et al., 2012; Taranger et al., 2014). With the aim of increasing food production for a continually growing population, there is projected a five-fold increase in Norwegian salmon production by 2050 (Olafsen et al., 2012). By continuing to produce salmon in open sea cages, this increase in production will also lead to a substantial increase in the amount of dissolved nutrients released. In order to ensure a sustainable increase in production, it is important to determine how the potential influence of dissolved inorganic nutrients released by fish farming can influence pelagic ecosystems, and to find ways to mitigate possible negative impacts.

One of the possible methods to alleviate the effect of increased nutrient loading on the environment is by co-cultivating salmon with species of lower trophic levels. Around two thirds of the nitrogen waste from salmon farming is inorganic ammonia (Wang et al., 2013), which is taken up by primary producers such as phytoplankton and macroalgae. Macroalgae cultivated in the vicinity of salmon farms may utilize the dissolved inorganic nutrients released from salmon farms in open-water integrated multi-trophic aquaculture (IMTA) systems (Buschmann et al., 1996; Chopin et al., 2001; Wang et al., 2012, 2014; Handå et al., 2013). Co-producing salmon and kelp might therefore reduce the ecological influence of dissolved nitrogen released into the environment and at the same time produce valuable kelp biomass. Previous studies have shown that the use of kelp in IMTA with salmon has limited bioremediation potential as the kelp can only take up a part of the dissolved nutrients, but the

potential for increased biomass production is substantial (Broch et al., 2013).

In this paper, we assess the potential of salmon driven kelp cultivation in terms of increased production and nutrient extraction efficiency. As a basis for the assessment, kelp growth, and elemental composition along a gradient from a salmon cage farm off the coast of Sogn and Fjordane, Norway, was studied over one kelp growth season from deployment in February to September. A numerical model was used to assess the spatial extent of the influence of salmon derived nitrogen to evaluate the potential for upscaling IMTA.

MATERIALS AND METHODS

Experimental Design

The cultivation experiment took place at the Marine Harvest salmon farm at Flåtegrunnen (N 61°34.586, E 4°48.942), in the county of Sogn and Fjordane, Norway from February to September 2013 (Figures 1A,B). The farm consisted of eight circular cages of 150 m circumference, of which seven were used for salmon production. The eighth cage was used as one of four experimental stations situated 100 m (100 E; empty fish cage), 200 m (200 E), 1,000 m East (1,000 E), and 1,000 m West (1,000 W) of the farm (away from cage with fish) (Figure 1B). Based on previous literature, 1,000 m was expected to be outside of the impacted area and considered being a reference stations reflecting the natural situation (Merceron et al., 2002; Sanderson et al., 2008; Pitta et al., 2009). Bottom depths at the stations varied between 10 and 40 m, and the water column in the upper 10 m was always mixed due to tidal action.

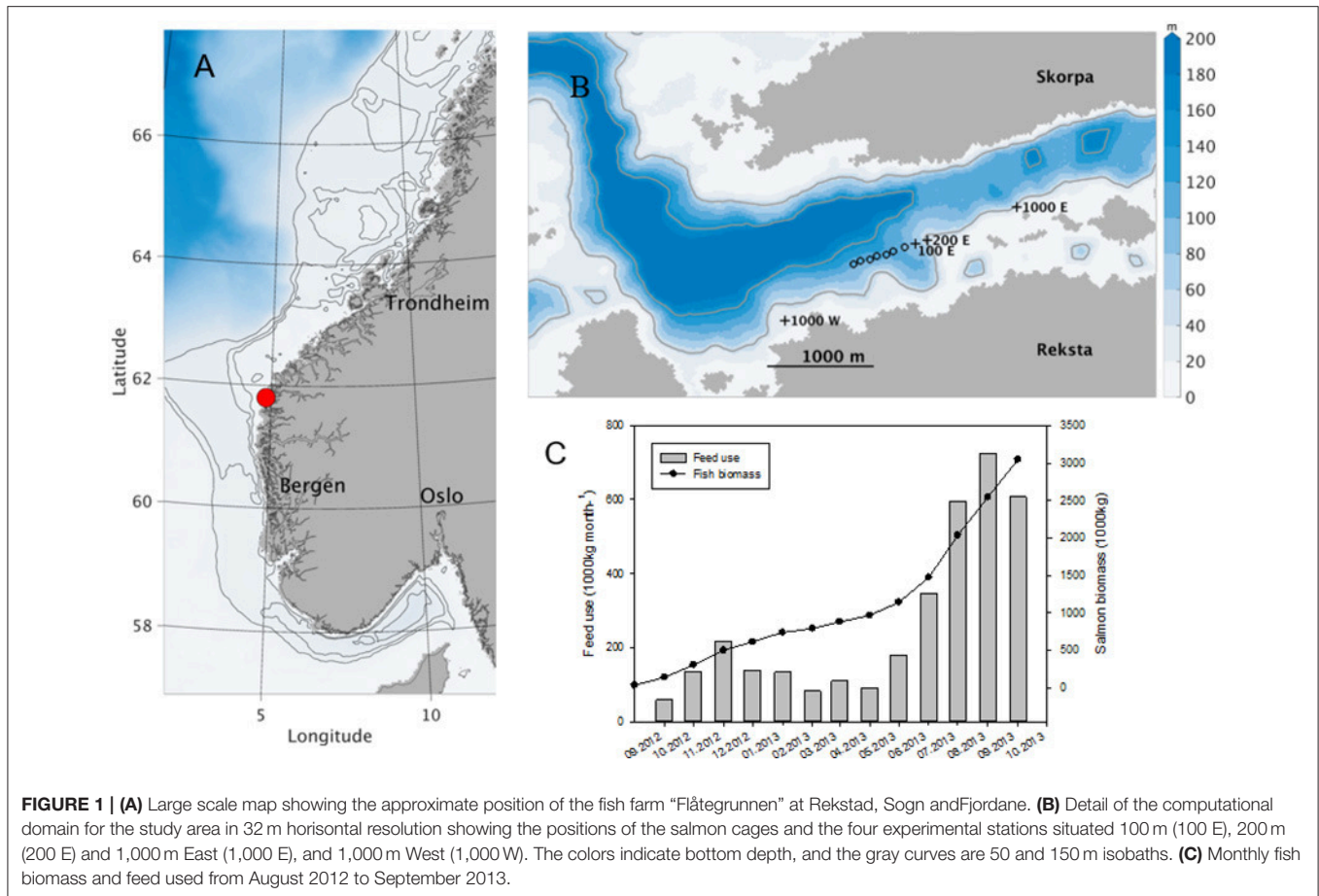
The westernmost station served as a reference station in this study. Horizontal carrier lines were aligned in the north-south direction and perpendicular to the alignment of the fish farm (East-West) at each station (Figure 1B).

Seedling Production and Deployment

Non-sporogenous individuals of *S. latissima* were collected close to the field site in September 2012. Meristematic tissue, stipes, and haptera were cut off and the remaining thalli were transferred to experimental tanks for sorus induction. When the sori were mature, the spores were released and seeded on 1.4 mm string, and to maintain genetic variability, spores from 10 different individual sporophytes were seeded. Seedlings were cultivated in flow-through incubators according to methods in Forbord et al. (2012) and Forbord et al. (2018). After 11 weeks in the hatchery the seeded string was twined onto 14 mm diameter ropes, and 20 ropes of 7 m length were deployed vertically with three meters distance from each horizontal carrier line at each station. The sporophyte length at deployment was on average 1 cm. Sampling of sporophytes throughout the season was predetermined for water depths of 2, 5, and 7 m below the surface based on a previous study showing best growth of kelp at 5 m depth in Mid-Norway (Handå et al., 2013).

Sampling

Two cruises were done to run water samples campaigns for the analysis of dissolved inorganic nitrogen (DIN: NO_2^- , NO_3^- ,



NH_4^+), one in April when feed use and fish biomass were low and one in September when feed use and fish biomass were high. Water samples were taken in triplicates from 5 m depth with a Niskin water collector. CTD-profiling (SAIV 200) showed that the water body was mixed down up to ~10–40 m (Jansen et al., 2018).

During each sampling month the length and width of seaweed of 15 randomly selected individual sporophytes from each of 6 ropes at each of three 1 m depth horizons at 2, 5, and 7 meters at each station was measured at each sampling (15 sporophytes \times 6 ropes \times 3 depths = 270 sporophytes per station, 4 stations, 4 samplings). Due to fouling related losses of sporophytes during summer we were only able to measure a minimum of 5 sporophytes per depth horizon and rope in few cases. The 1 m depth horizons 2, 5 and 7 meters relates to 1.5–2.5 m, 4.5–5.5 m, and 6.5–7.5 m below the surface, respectively. These horizons were marked on the ropes before deployment.

For chemical analysis five whole sporophytes were collected from 5 m depth at each of the 6 ropes per station and sampling date. The samples were stored at -80°C until freeze-drying. After freeze drying, the whole 5 sporophytes were milled into a powder to achieve a representative sample for the whole sporophyte and remove individual variability. Subsamples of this powder were used for further analysis of carbon and nitrogen concentrations as well as stable isotope signatures of carbon and nitrogen.

Salmon feed, seston and kelp tissue were sampled for stable isotope analysis when fish biomass was low in April and high in September to determine the contribution of salmon derived and naturally occurring nitrogen during the period of kelp cultivation.

Seston was sampled in triplicate at each of the four stations using a Niskin water collector and filtered on a pre-combusted Whatman GF/F filter. Salmon feed was sampled from the respective feed batch in use in April and September. Samples were frozen with dry ice, freeze dried, and stored at -80°C under nitrogen atmosphere until analysis. Stable isotope data was used to calculate one-isotope, two-source linear mixing models (reviewed by Boecklen et al., 2011).

Chemical Analysis

The concentration of total ammonia nitrogen (TAN) and nitrate in the water samples were determined in parallel with a Fluorescence Detector (DFL-10) (Kérouel and Aminot, 1997) and a flow analyser with O.I. Analytical cartridge Part A002603, respectively, according to Jansen et al. (2018). Samples of kelp, salmon feed, and filters with seston were analyzed at the Colorado Plateau Stable Isotope Laboratory at the Northern Arizona University (USA) for content of carbon and nitrogen and stable isotope ratios of $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$. The samples were weighed and packed in tin capsules before analyzing with a Carlo Erba NC2100

elemental analyser coupled with a Thermo-Electron Delta V Advantage isotope ratio mass spectrometer (IRMS), configured through a CONFLO III. Isotope ratios were calculated as follows:

$$\delta X \text{ sample}(\text{‰}) = [(R_{\text{sample}}/R_{\text{standard}}) - 1] \times 1000, \quad (1)$$

where $X = {}^{13}\text{C}$ or ${}^{15}\text{N}$ and R is ${}^{13}\text{C}/{}^{12}\text{C}$ ratio for carbon and ${}^{15}\text{N}/{}^{14}\text{N}$ for nitrogen. IAEA standards were used for carbon isotope nitrogen calibration.

Statistical Analyses

One-way ANOVA followed by a Holm-Sidak *post-hoc* tests were run on growth, % carbon, % nitrogen and C/N data using SigmaPlot 12.0 with a significance level of 0.05. To assess the influence of the fish farm on kelp growth rates we calculated the specific growth rate (μ). As maximum size of the kelp occurred at different sampling dates, we calculated the growth rate from deployment to maximum size. The influence of salmon farming on the relative contribution of inorganic nitrogen in kelp stemming from salmon farming or from natural sources was assessed by ANOVA followed by Holm-Sidak *post-hoc* test using the arcsin transformed proportion of salmon derived nitrogen calculated in isotope mixing models as the dependent variable. To assess the influence of distance from the farm on specific kelp growth rate and the proportion of natural and salmon derived nitrogen we fitted a 2-parameter hyperbolic decay model to the data, using salmon derived nitrogen from the isotope mixing models as the dependent variable and distance from the farm as the independent variable.

Stable Isotope Mixing Models

The relative contribution of nitrogen from natural sources and from salmon in the kelp was calculated using $\delta^{15}\text{N}$ signatures of seston and salmon feed as sources and of kelp from the four stations as mixture following Phillips and Gregg (2001). The stable isotope signature of nitrogen excreted by consumers is typically depleted compared to the ingested food sources and 1‰ was subtracted from the salmon feed $\delta^{15}\text{N}$ signatures (Steele and Daniel, 1978; Minagawa and Wada, 1984).

$$f_A = (\delta^{15}\text{N}_M - \delta^{15}\text{N}_B) / (\delta^{15}\text{N}_A - \delta^{15}\text{N}_B) \quad (2)$$

$$f_B = 1 - f_A \quad (3)$$

where $\delta^{15}\text{N}_M$, $\delta^{15}\text{N}_A$, and $\delta^{15}\text{N}_B$ represent the mean nitrogen signatures of the mixture M and sources A and B , respectively, and f_A and f_B are the proportions of A and B in M . Errors are expressed as variance. Due to heavy degradation of kelp in September and virtually no kelp growth throughout the summer, we omitted September from the analysis.

Mathematical Modeling of *S. latissima* Growth and Production Upscaling

The nested, coupled 3D hydrodynamic-ecological model system SINMOD (Wassmann et al., 2006; Broch et al., 2013) was used to simulate the general environmental conditions around Flåtegrunnen from February to June 2013. The model includes

variables for concentrations of phytoplankton (mmol N m^{-3}) and nutrients (NO_3^- , NH_4^+ -N) and is coupled with a growth model for *S. latissima*. The system can thus simulate dispersal and dilution as well as uptake of nutrients by phytoplankton and kelp. A model setup for the region around the farm at Flåtegrunnen in 32 m horizontal resolution was established (Figure 1). The nesting technique, forcing data and coupling of the models are detailed in Broch et al. (2013, 2017). Release rates of TAN (no distinction between NH_4^+ and NH_3 was made) from the fish farm were calculated from reported figures on feed usage (Figure 1C), using a conversion factor of $0.025 \text{ kg TAN-N released (kg dry fish feed used)}^{-1}$ (adapted from Wang et al., 2012; see also Broch et al., 2013). The *S. latissima* growth model (Broch and Slagstad, 2012; Broch et al., 2013) was updated and tuned to differentiate between uptake and use of NO_3^- and NH_4^+ (see Appendix).

Four simulations were run using the kelp-ecosystem model. The first two (one with and one without a fish farm effluent) were run in order to determine the spatial extent of the IMTA-effect. The last two (one with and the other without a fish farm effluent) were run in order to assess the increased growth and uptake potential of a large scale (25 ha) kelp installation west of the fish farm, with the same kelp densities as in the main simulations in Broch et al. (2013). The simulations were run from February to June as the fouling of the sporophytes until mid-June was negligible (Førde et al., 2016). After this point, the sporophytes were too heavily fouled and are therefore omitted from this study. For comparison with the simulation results, the frond area of the kelp was calculated as follows (Broch et al., 2013):

$$A = 0.75 LW \quad (4)$$

where A is the frond area, and L and W are the length and width of the kelp frond, respectively.

Quantification of the Spatiotemporal Influence of IMTA

A new metric, IA_P (Integrated Area, unit: area), was used to quantify the spatial extent of the influence of IMTA at a location. This term is applicable to any open IMTA aquaculture system. IA_P is the total area of the region where cultivation in IMTA leads to at least a production increase of $P\%$ in biomass or another variable of interest (e.g., accumulated carbon or nitrogen) compared with monoculture not associated to a salmon farm. The IA_P changes with time, location and cultivation variables. This metric is reminiscent of the area-effect curves presented e.g., in Zeldis et al. (2011), but relates to improved effect due to IMTA rather than effects on basic ecosystem variables. For example, an $IA_{50} = 10 \text{ ha}$ for dry weight in April for *S. latissima* cultures near a salmon farm, means that an area of 10 ha will experience a potential increase in dry weight of at least 50% in April relative to cultivation at the same location, but without the fish farm effluent. Based on the simulation results (see previous subsection), the IA_P was calculated as a function of P and time for the kelp-salmon IMTA system studied here.

RESULTS

Environmental Variables

During the experimental period from February to September 2013, the net increase in total salmon biomass at the farm was 2,250 tons, reaching a maximum value of 3,000 tons in September. The monthly feed use increased steadily over the summer and reached a peak value of 700 tons in August (Figure 1C). The surface water temperature ranged from 4°C in April to 15°C in September.

The NH_4^+ concentrations at 5 m depth ranged from 0.3 to 0.5 mmol m^{-3} in April and 1.5–3.4 mmol m^{-3} in September (Figure 2). While there were no differences in NH_4^+ concentrations between the stations in April, the concentrations found in September were generally higher, and significantly higher at the 100 E station (3.4 mmol m^{-3}) compared to the other stations (<2.0 mmol m^{-3}).

The average simulated pattern of dispersal and dilution (including uptake by phytoplankton) of released NH_4^+ from the fish farm was similar in April, June, and September, but with different average concentrations (Figure 2). A clearly

defined ellipsoidal plume of nutrients around the fish farm is apparent, indicating transport of NH_4^+ in both easterly and westerly directions. The average daily ranges of simulated NH_4^+ concentrations at the four stations were underestimated in September compared with the measured concentrations (Figure 2). The measured values were within the average simulated range at all stations in April and at the 100 E and 200 E stations in September.

Growth

Kelp growth did not differ between the three depths (2, 5, and 7 m). *S. latissima* sporophytes grew significantly faster ($P < 0.05$) at the 100 E station than at the other stations from February to June, and the sporophytes reached approximately twice the length and width compared to other stations at all depths (Figure 3), corresponding to a three times higher frond area. While the sporophyte length at the 100 E station decreased due to heavy fouling during the summer at all depths, kelp from the 200 E station kept growing, and grew significantly ($P < 0.05$) faster than kelp from both the 1,000 m stations, and in August reached

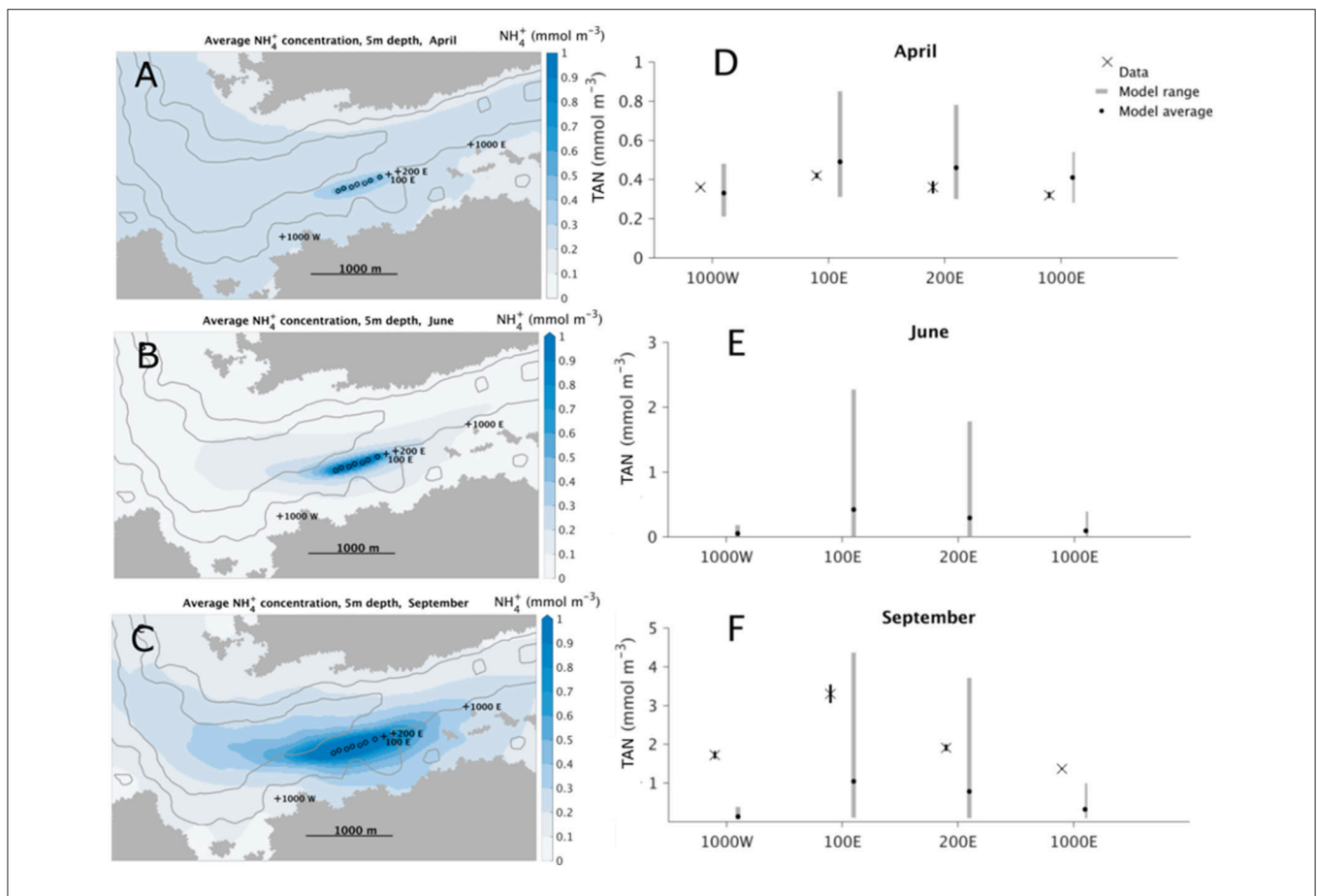


FIGURE 2 | Measure and modeled concentrations of TAN (mmol m^{-3}) in seawater at different stations in April, June and September 2013 (mean \pm se, $n = 3$). (A–C) Simulated spatiotemporal extent of the effluent and background NH_4^+ around the fish farm at Flåtegrunnen. (D–F) Simulated and measured NH_4^+ -concentrations at all the stations at 5 m depth for April (D) and September (F), and simulated concentrations in June (E). Black xs: measured concentrations; black dots: monthly mean simulated concentrations. Dark gray bars: range of simulated mean daily minimum to mean daily maximum concentrations with fish farm effluents. Light gray bars: range of simulated mean daily minimum to mean daily maximum concentrations without fish farm effluents.

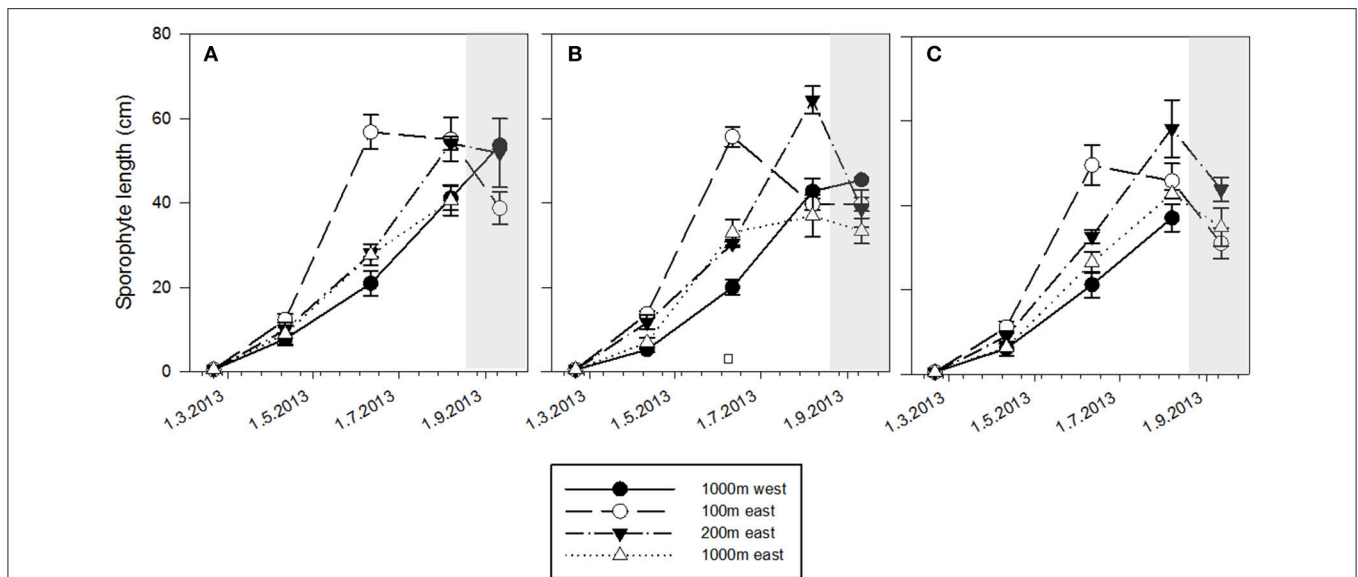


FIGURE 3 | Seasonal development of sporophyte length at 2 m (A), 5 m (B), and 7 m (C) depths at the four experimental stations (mean \pm standard error, $n = 6$). Gray shaded area indicates period of high bio-fouling.

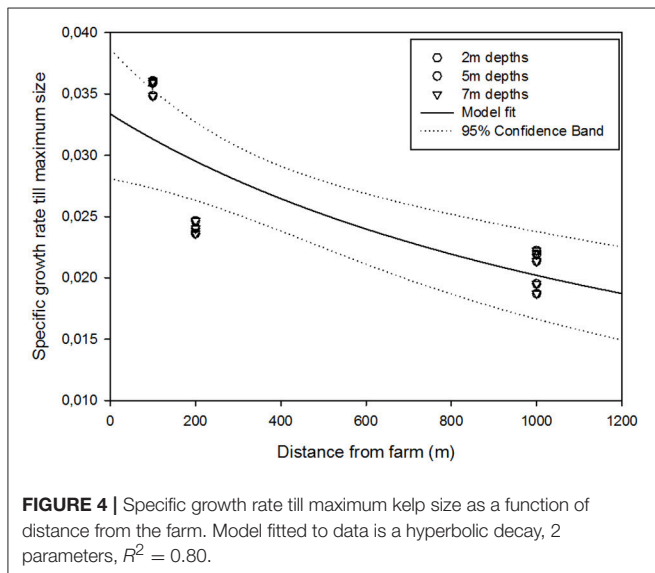


FIGURE 4 | Specific growth rate till maximum kelp size as a function of distance from the farm. Model fitted to data is a hyperbolic decay, 2 parameters, $R^2 = 0.80$.

comparable sporophyte lengths as the 100 E kelpid in June. The algae grown at the 1,000 m stations never reached comparable sizes. The reduction in plant size from August onwards was caused by heavy fouling by bryozoans, leading to almost complete losses of fronds by September. There was a clear gradient of decreasing growth rates with increasing distance from the farm (Figure 4), suggesting the main influence of the farm to be in the first 200 m from the farm.

Simulated frond areas at the four stations showed less spatial variation than the measured values (Equation 4), especially in April (Figure 5). When the simulated frond sizes were presented as ranges over an interval of time (± 1 week around the sampling

date), the model revealed much the same spatiotemporal growth pattern as the field estimates, especially in June (Figure 5). The simulated frond areas gradually decreased from the 100 E to the 200 E and 1,000 E stations, while the measurements of frond areas at 200 E and 1,000 E were approximately the same.

The IA_{30} -region for dry weight (DW) of individuals in June had a shape reminiscent of the contours of NH_4^+ -concentrations (Figure 2 and Figure 6). The IA_{30} -region is the region in which the simulated kelp dry weight (DW) yield increased by at least 30% from the scenario without fish farm effluents included to the one with effluents included. The region stretched further in a westerly than an easterly direction from the fish cages, covering an area of ~ 50 ha, including the fish farm.

The plots of the areas of the IA_P regions for different times and different values of P (the percentage increase) illustrate the increasing importance of the contribution of NH_4^+ from the fish farm as the natural background NO_3^- was being depleted (Figure 7). In April the area of the region of 20% or higher increase in DW by IMTA (the IA_{20} -region) was almost negligible, while in June this area covered 60–110 ha.

The total simulated biomass production of the hypothetical 25 ha IMTA kelp farm from February to June was about 1,125 tons FW, with a production ha^{-1} ranging from about 30 tons FW ha^{-1} furthest away to 130 tons FW ha^{-1} closest by the fish farm (Table 1). There was a simulated net uptake of 1.6 tons DIN-N, including TAN-N from the farm and TAN-N and NO_3^- from the ambient waters. By comparison, the effluent from the fish farm was ~ 13.5 tons TAN-N from February to June, so that the kelp culture extracted about 12% of the N-effluent during the cultivation period. The 25 ha IMTA farm had a 50% higher average nitrogen uptake, and a 60% higher DW yield compared to an identical farm without fertilization from a fish farm, while the FW yield was 100% higher in the IMTA than the non-IMTA case

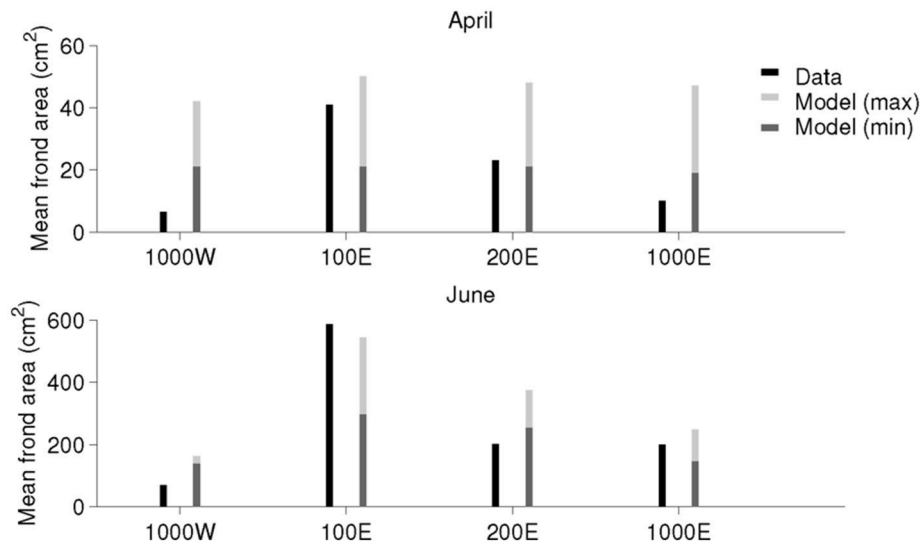


FIGURE 5 | Comparison between simulated and measured frond area values at the four stations (ordered from West to East) on 11th of April (**Top**) and 11th of June (**Bottom**). The black stems to the left represent the values estimated from measured frond length and width. The dark gray stems to the right represent the spatial average (3 model grid cells, representing the spatial extent of a kelp long line) of simulated frond areas at 5 m depth 7 days before the sampling dates (4 April and 4 June). The light gray stems represent the same spatial averages 7 days after the sampling dates (18 April and 18 June).

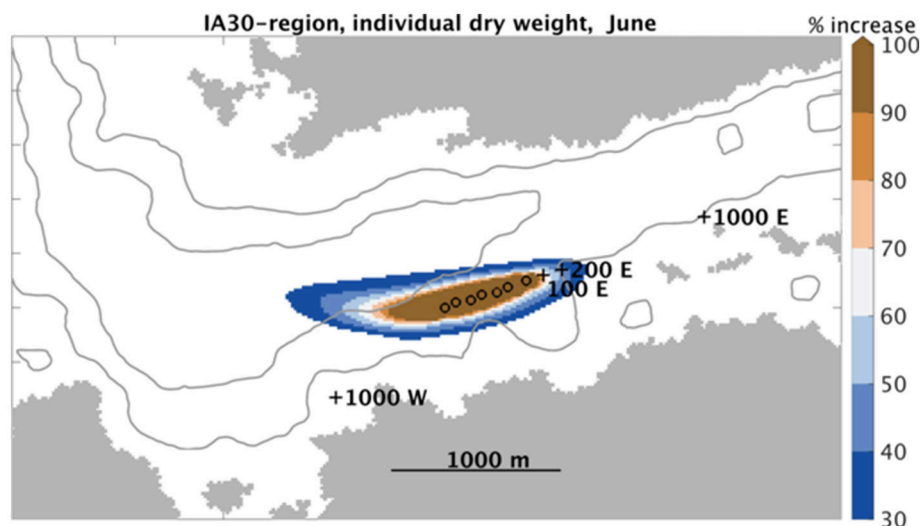


FIGURE 6 | IA_{30} for dry weight. Spatial extent of the region with a 30% (or greater) increase in individual sporophyte dry weight yields from a simulation scenario without to one with salmon farm effluents. Each model grid cell represents an area of $32\text{ m} \times 32\text{ m}$. $IA_{30} = 27\text{ ha}$ in this case.

(**Table 1**). We also simulated the area necessary to take up 13.5 tons of nitrogen, which equals the estimated N release from the farm during the same period. To achieve an even mass balance, an area of $\sim 220\text{ ha}^{-1}$ would be needed to cultivate enough kelp to fix an equivalent of the nitrogen released by the fish.

Isotope Mixing Models

The relative contribution of nitrogen released through salmon farming and natural inorganic N sources differed significantly between the stations (ANOVA, $p < 0.05$), with the highest

contribution ($\sim 40\%$ of salmon derived N) at the nearest station to the farm (100 m) being different from the two stations farthest from the farm (1,000 m, $\sim 10\%$), with the 200 m station falling in between (29%), not being different from either the 100 m or the 1,000 m stations (**Figure 8A**). More important than differences between individual stations is the general pattern that the relative contribution of salmon derived nitrogen utilized by the kelp decreased with increasing distance from the salmon cages (**Figure 8B**). Seston was only sampled in April and September, allowing calculating one-isotope, two-source linear

TABLE 1 | *Saccharina latissima* fresh weight (FW) and dry weight (DW) yields and nitrogen (N) and carbon (C)-uptake with and without IMTA.

Scenario	FW	DW	N	C
IMTA	45.5 (28.7–136.6)	5.4 (4.1–11.6)	0.06 (0.05–0.12)	1.7 (1.4–2.8)
Regular	22.6 (20.5–25.3)	3.4 (3.1–3.7)	0.04 (0.04–0.05)	1.2 (1.1–1.3)

Key figures were obtained from simulation of a 25 ha farm in the IA₃₀-zone assuming cultivation from mid-February to mid-June. Unit: t ha⁻¹. All figures are averages for the entire farm and have been rounded to the nearest 0.1 t, except values for nitrogen, rounded to the nearest 0.01 t. The figures in parentheses represent the range within the farm.

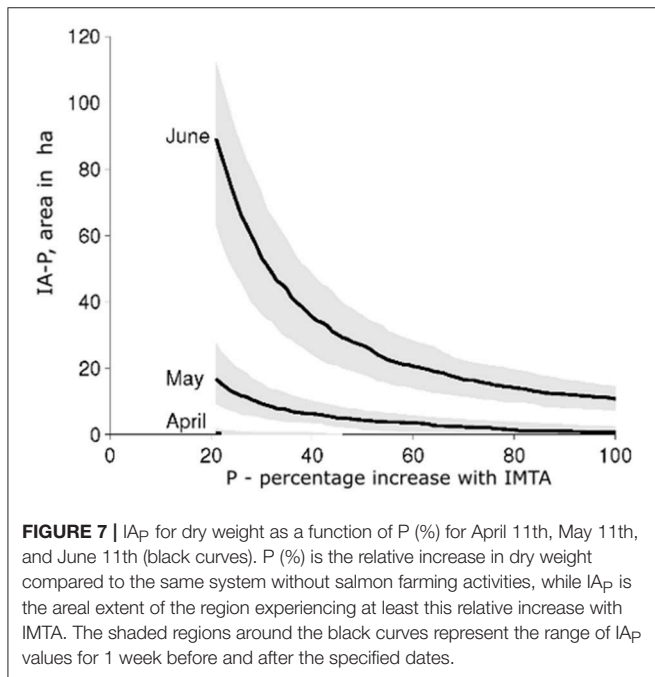


FIGURE 7 | IA_P for dry weight as a function of P (%) for April 11th, May 11th, and June 11th (black curves). P (%) is the relative increase in dry weight compared to the same system without salmon farming activities, while IA_P is the areal extent of the region experiencing at least this relative increase with IMTA. The shaded regions around the black curves represent the range of IA_P values for 1 week before and after the specified dates.

mixing models for these 2 months only. However, the kelp sporophytes were heavily fouled and degraded in September and no considerable growth took place at least from August to September. A meaningful use of stable isotope data as an indicator for the origin of nitrogen in kelp is based on incorporation of nitrogen, i.e., positive growth rates.

Composition

The fertilizing effect of the farm generally kept the elemental composition of the kelp more stable. The carbon concentration of the kelp tissue varied between 23 and 28% of the dry weight between the stations and within season (Figure 9), with significant differences between stations ($p < 0.05$). Within the period of positive growth, the 100 E station exhibited the lowest carbon content. The nitrogen contents in the kelp varied from 1 to 2% of the dry weight, reflecting spatiotemporal variability in nitrogen availability (Figure 9). All stations displayed a decrease in nitrogen contents until June, with a less pronounced decrease at the 100 E station ($p < 0.05$), indicating a less severe nitrogen limitation and the subsidizing role of the fish farm in a period with little or no nitrogen available from natural sources. The

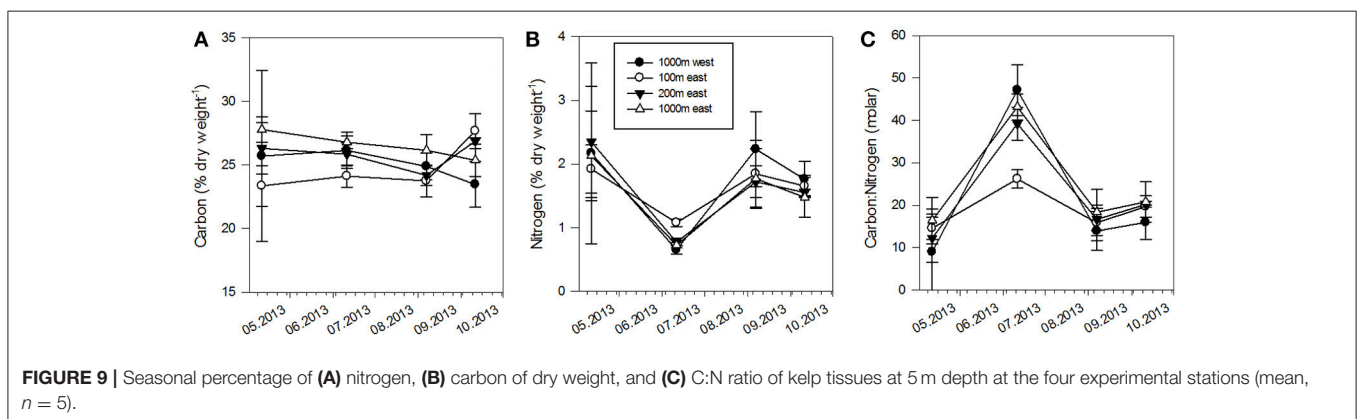
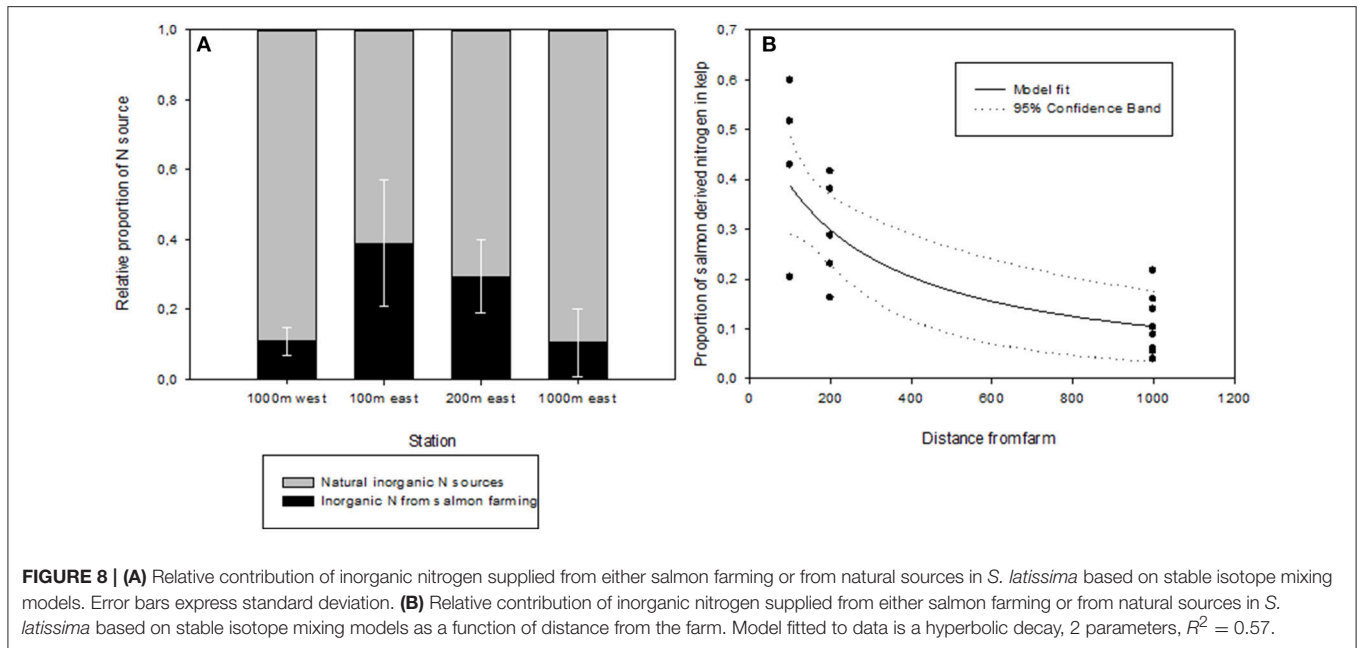
lower C/N ratios at the farm (Figure 9) were a result of lower carbon contents as well as higher nitrogen contents. The peak C/N ratios in June, showing the highest values at the 1,000 W station and lowest near the farm, again reflected the low nitrogen content at that time and the potential of nitrogen fertilization of the fish farm.

DISCUSSION

The present study examined the feasibility of IMTA combining *S. latissima* and *S. salar* in an open net cage system in western Norway. *S. latissima* sporelings were deployed at several distances away from a salmon farm and algal growth and composition were monitored over one cultivation season from February until September. Additionally, ammonia concentrations were measured and the contribution of salmon derived nitrogen to the total nitrogen pool of the plants was estimated. Finally, the potential increase of kelp production related to the nitrogen input from a fish farm was modeled to quantify the potential for upscaling of kelp (*S. latissima*) cultivation in salmon-driven IMTA.

The Norwegian coastal ecosystem is characterized by strong seasonal changes in light availability, temperature and nutrient availability. Typically a phytoplankton spring bloom, usually occurring in late March, early April in this part of Norway (Ibrahim et al., 2014), depletes dissolved inorganic nutrients. Such decreases in nitrogen availability can be traced in both micro- and macroalgae by a relative increase in cellular carbon contents and a decrease in nitrogen contents, and consequently increases in C:N ratios in kelp biomass (Gevaert et al., 2001). In the present study, a similar trend with reduced N-contents in kelp tissues farmed 1,000 m away from the salmon farm compared to that farmed closer to the salmon farm, illustrated the nitrogen subsidiary role of fish farming. The nitrogen contents of the sporophytes analyzed in April did not reflect the low DIN concentrations in the surrounding waters. This can be explained by surplus storage of nitrogen in the kelp tissue before the spring bloom when DIN concentrations are still high (Gevaert et al., 2001). As DIN concentrations change quickly in a bloom setting and the timing of the onset of the bloom may vary up to around 20 days from 1 year to another (Vikebø et al., 2012), ambient DIN concentrations were most likely higher just prior to our first sampling on the 8th of April. The nitrogen content of the kelp varied least at the 100 E station despite high growth rates, again suggesting a more constant nitrogen supply than at the other stations. Similar results have been reported by Sanderson et al. (2008), Handã et al. (2013), and Freitas et al. (2016).

Kelp size, elemental, and biochemical composition changes during the growth season (Lüning, 1990). Sharma et al. (2018) reported that sugars, which could serve to produce biofuels peaked earlier in the season, while a later harvest contained the higher proportions of amino acids, minerals, and phenolic compound. Schiener et al. (2015) also reported seasonal variability, where the highest yields of laminarin and mannitol coincided with the lowest yields in ash, protein, moisture and polyphenols. Clearly, the harvest strategy must be adjusted to



the product the kelp shall be converted to (human consumption, biochemicals, biomass). However, there always is a trade-off between little biomass and early harvest. This trade-off might be tipped toward an early harvest if the price of the product counterbalances the potential biomass loss due to early harvest. Potential higher value products might be young kelp sporophytes for human consumption, which tends to have a more tender and favorable texture matching consumer preference. In our study, the nitrogen concentrations showed less variation with season, documenting the fertilizing effect, meaning IMTA could provide more stable and reliable and ultimately predictable product quality, which would add an extra benefit to seaweed culture next to solely increased productivity.

Stable nitrogen isotopes have proved to be useful indicators for the detection of farm derived nitrogen in macroalgae, as they are incorporated into plant tissue (Carballeira et al., 2013). Costanzo et al. (2004), for example, showed a clear decrease of the influence of a shrimp farm in North East Australia with distance from

the farm using stable nitrogen isotopes of mangrove leaves. The gradient of the $\delta^{15}\text{N}$ signature of mangroves along the creek transporting the effluent of the shrimp farm clearly showed not only a dilution effect, but also that nitrogen originating from the shrimp farming operation was taken up along the creek, and hence were incorporated in plant tissue. This suggests that a precise positioning of macroalgae downstream of the nitrogen source in open-water IMTA systems can enable recycling of a part of the dissolved nutrients released from fish farms into the surrounding environment.

In the present study, salmon derived nitrogen was traceable in the kelp tissue by means of stable nitrogen isotopes, and the relative contribution decreased with increasing distance from the fish farm early in the season. Stable isotope analysis samples were taken only in April and September but not during the summer months. The kelp sporophytes were heavily fouled by September, and virtually no growth took place for at least 4 weeks prior the last sampling. Using mixing models to reveal the

relative contribution of nitrogen derived from either natural or farming sources to the nitrogen assimilated by kelp in September therefore provided little information. Non-growing organisms are unlikely to isotopically reflect their surroundings, as no or very little new tissue is generated, hence no new materials are incorporated. To further complicate the issue, isotopic ratios change drastically in starving organisms (e.g., Oelbermann and Scheu, 2002; Haubert et al., 2005; Aberle and Malzahn, 2007), and may therefore not reflect the nutrient sources anymore, hence violating one of the basic assumptions behind the use of stable isotopes as tracers for nutrient sources (Hobson et al., 1993).

The decreasing trend in salmon derived nitrogen in the sporophytes raises the question to which extent an aquaculture operation fertilizes the system. Surprisingly, we found a small, but detectable contribution of salmon derived nitrogen 1,000 m away from the farm. The literature suggests rather a more limited spatial influence. Marinho et al. (2015) showed increased production and N uptake in kelp (*S. latissima*) production 100–500 m away from a in a rainbow trout/blue mussel IMTA system compared to the reference site, which was 2,000 m away from the fish cages in a Danish fjord. However, they note that the fish did not release enough nitrogen to increase N concentrations near the fish cage above background levels. Sanderson et al. (2008) studied ammonium concentrations around salmon cages in north-west Scotland. The authors concluded that generally the N fertilization impact of a fish farm did not exceed 200–300 m and used station a 400 m away from the nutrient source as reference stations. Pitta et al. (2009) reported similar spatial trends in the Aegean Sea, where they concluded that nutrient effects were not visible anymore 500 m away from fish cages. Troell et al. (1997) found higher growth rates of a red algae cultured 10 m away from salmon cages in Chile, than algae grown 150 m and 1,000 m away from the fish farm. As growth and composition of kelp in that study did not differ between 150 and 1,000 m the authors concluded that in their study the spatial impact of the fish farm was small. Even smaller influenced areas were reported by Merceron et al. (2002) who noticed a spatial influence < 50 m of a trout cage in Cherbourg, France. However, Sanderson (2006) found indications for fish farm derived nitrogen 1,000 m away from a fish farm in wild kelp. Similar findings were reported by Abreu et al. (2009), who also found a fertilizing effect of a salmon farm in southern Chile. The authors report that *Gracilaria chilensis* utilized nitrogen derived from salmon farming at least 800 m away from the farm. Nordvang and Johansson (2002) even showed increased system productivity related to fish farms of up to 8 km², however, the latter study mainly dealt with semi-closed bays with limited water exchange. Clearly, the extent of the influence of a fish farm depends on the amount of nutrients released and the hydrographic conditions the fish farm is placed at, and the term “influenced” is not well-defined and can span from “traces of fertilizing effects can be found,” to “substantial fertilizing effect useful for IMTA.”

The measured field data was used to update the kelp model to quantify the potential for upscaling of kelp (*S. latissima*) cultivation in salmon-driven IMTA. The simulation results showed a development in *S. latissima* frond size similar to that

found by field measurements, though with a less pronounced difference between the stations (Figure 4). The simulations further indicated a greater potential for increased biomass in IMTA in the western than the eastern direction from the fish farm. The introduction of the IA-metric (Figure 6) is useful in this context because it focuses on the influenced area, independent of any specific current direction. The IA-metric integrates all factors like background nutrient availability, light intensity, temperature, water current speed, and the feeding intensity of the fed species (in this case Atlantic salmon) over time and is one measure of the potential efficiency of an IMTA system.

An efficiency measure in IMTA related to this is the increase in production by implementing IMTA compared to a monoculture system. The total fresh biomass production for a 25 ha virtual kelp farm in the IA₃₀ zone was 100% higher in the IMTA case than in monoculture without nutrients from salmon farming. By comparison, Sanderson et al. (2012) reported a 26% increase in *L. saccharina* productivity when grown in IMTA. However, the latter study reported a production capacity of about 220 t ha⁻¹ y⁻¹, which exceeds an average production of 45 t ha⁻¹ y⁻¹ modeled in our study. Troell et al. (1997) calculated similar high potential yields in an salmon/kelp-IMTA with a production potential of about 250 t wet weight ha⁻¹ y⁻¹. Clearly the increase in production due to IMTA and the production capacity depends on the trophic state of the system, where a lower natural nutrient background leads to higher increased in productivity.

From a bioremediation and mass balance perspective, the simulations indicated that a 25 ha kelp farm could take up almost 12% of the ammonia released during the cultivation period from February to June. This equals 0.47% taken up per ha, which corresponds well with the 0.41% per ha obtained in a previous study (Broch et al., 2013). Other model exercises (Hadley et al., 2016) revealed higher N extraction capacities of kelp IMTA systems, where the authors report an uptake potential of 30–80%, albeit without any specific mention of the area covered by the seaweed cultures. However, in the current study the yield would be larger if there would be a better temporal overlap in feed use in aquaculture (being high in the second half of a year) and the main kelp growth period. In general, the correct timing of seeding and harvest is of utmost importance for a successful extraction of fish farm released nitrogen, as stressed in several studies (e.g., Broch et al., 2013; Freitas et al., 2016; Buck et al., 2018).

CONCLUSIONS

The results suggest that direct uptake of nutrients in close vicinity to salmon farms can enhance growth and biomass production of *S. latissima* in Norwegian coastal waters. Macroalgae cultivated in IMTA contained more nitrogen suggesting higher protein content. Over the main growth season for macroalgae (February–June), a 25 ha *S. latissima* farm positioned close to a salmon farm could take up 12% of the 13.5 tons of inorganic nitrogen released from the salmon. The farm would have a potential average yield of 45.5 tons of *S. latissima* per hectare compared

to a kelp farm monoculture of 22.6 tons. Co-cultivating salmon and kelp in IMTA therefore has potential to produce more biomass, however, has also limited bioremediation effects when it comes to the uptake of released dissolved inorganic nutrients from salmon farms. This is especially an effect of the temporal mismatches between the release by fish and uptake rates by kelp.

AUTHOR CONTRIBUTIONS

JF, SF, AH, HJ, MB, HF, and AF were involved in planning, execution of the experiment, and writing of the manuscript. OB ran the modeling exercises and contributed to writing. JS and YO were involved in planning, and writing. AM was analyzing data and writing the manuscript.

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SUPPLEMENTARY MATERIAL

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Conflict of Interest Statement: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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