



# Migration and habitat use of the landlocked riverine Atlantic salmon *Salmo salar* småblank

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**Abstract** “Småblank” is the only population of Atlantic salmon in Europe living its entire life in the river habitat without performing migrations to the sea or lakes. Home range size and habitat use were compared between an unregulated tributary and hydro-power regulated parts of the main river during 2014–2017. In total, 140 småblank were tagged with radio transmitters and tracked for 4–5 months. Tagged fish tended to stay within small areas: the average 50% home range was 1123 m<sup>2</sup> and the average 95% home range was 4416 m<sup>2</sup>. Mean distance between the furthermost positions for individual fish during tracking was 242 m, and the fish did not migrate between different stretches of the river. The small home range and non-migrating behaviour may explain why småblank are divided into several genetically distinct populations with limited gene flow. The results

showed that småblank were far more stationary than what is known for other landlocked salmon populations. Use of moderate or high water velocities (> 50 s<sup>-1</sup>) areas with coarse river substrates implies a vulnerability to anthropogenic effects such as water diversion, dams and weirs.

**Keywords** Home range · Movement · Namsblank · Landlocked · Relict salmon

## Introduction

Anadromy, with spawning and rearing in freshwater and feeding migrations to the sea, is the most common life history strategy for Atlantic salmon (*Salmo salar* Linnaeus, 1758). Non-anadromy occurs either as a reproductive strategy predominately in males that mature as parr in freshwater (Thomaz et al., 1997), or entire populations can be freshwater-resident, with both males and females remaining in freshwater throughout their life cycle (Klemetsen et al., 2003; Hutchings et al., 2019). Many of these freshwater-resident populations are landlocked—unable to perform feeding migrations between freshwater and the sea due to physical barriers such as waterfalls or dams—but some freshwater-resident populations have potential access to the sea (Behnke, 1972). Freshwater-resident populations of Atlantic salmon occur in both northeast North America and Europe (Kazakov,

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1992; Ozerov et al., 2010; Hutchings et al., 2019). Many landlocked populations have migrations between river and downstream lakes, and a large fraction of these populations have declined due to anthropogenic impacts, and some populations have become extinct (Ozerov et al., 2010; Hutchings et al., 2019).

The greatest threats to the persistence of landlocked salmon are found in Europe (Hutchings et al., 2019), including Norway, where the status of landlocked salmon is poor. Historically, there were at least four landlocked populations in Norway, but two were extirpated and a third came close to extirpation due to hydro-power development and acidification. The latter population (“bleke” from the Byglandsfjorden area, southern Norway) is now restored after large-scale stockings from a gene bank facility. The fourth population, called småblank (Berg, 1953), is regarded as having the least secure conservation status of all remaining landlocked populations in terms of extinction risk (Hutchings et al., 2019). Småblank is the only river-resident population in Europe, with no feeding migrations to lakes (Berg, 1984b). A few populations in North America have adapted a similar river-resident strategy (Behnke, 1972). Populations that do not utilize lakes are particularly vulnerable to negative impacts from anthropogenic changes in the river.

Habitat fragmentation and diminished population abundance can reduce genetic diversity and increase extinction risk. This is most likely the case for småblank, which has a lower genetic variation than anadromous Atlantic salmon, and is a unique endemic island population of Atlantic salmon (Bourret et al., 2013; Sandlund et al., 2014). It is in a precarious situation due to a variety of anthropogenic impacts on its restricted habitat area: reduced water discharge, hydro-power dams and weirs causing fragmentation and a change from lotic to lentic conditions, introduced species, and hybridization with anadromous Atlantic salmon after the construction of fishways (Sandlund et al., 2014; Karlsson et al., 2018). In addition, the regulated water flow of the main river contributes to sediment packing of the stony substratum and reduced shelter availability (Sandlund et al., 2014), which is important for growth and abundance of juvenile salmon (Finstad et al., 2007). Thus, maintaining population size, avoiding further habitat fragmentation and reducing negative impacts of river regulation are important for protecting this population.

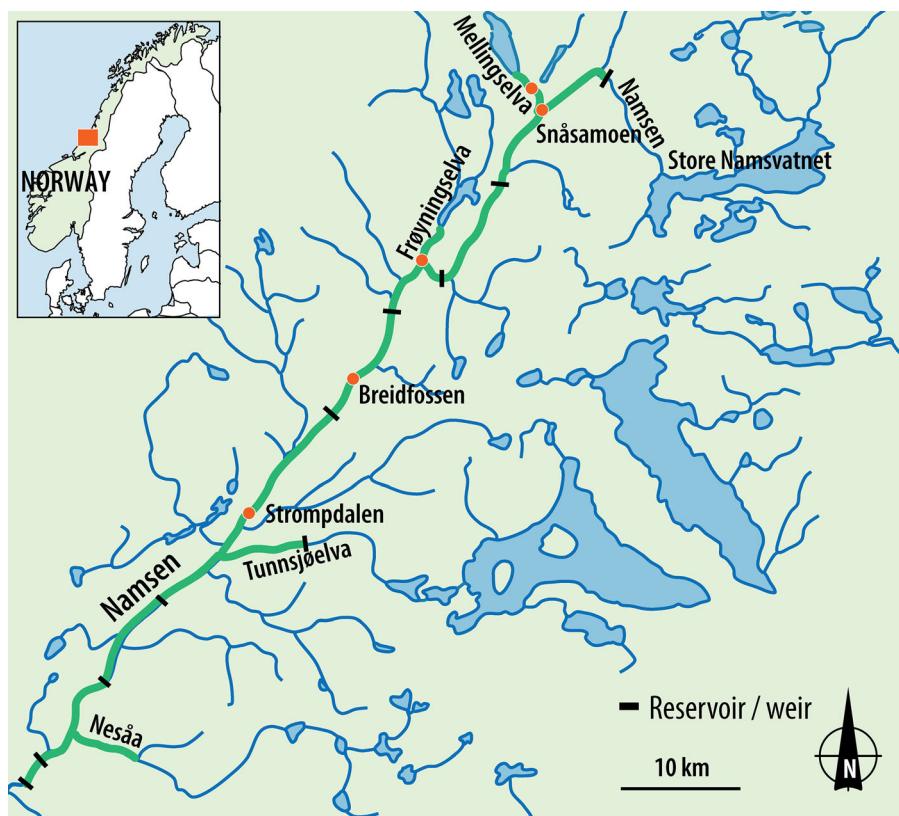
To improve the conservation status, it might also be necessary to restore suitable habitat in the degraded parts of the watershed.

The habitat use of stream-dwelling salmonids provides vital information on their life history. In contrast to landlocked salmon that migrate into lakes as part of their life cycle, little is known on the movement patterns and habitat use of individuals of river-resident Atlantic salmon populations. However, they differ genetically (Bourret et al., 2013) and in many aspects of their life-history (Sandlund et al., 2014) to anadromous Atlantic salmon, so it is also likely that they may differ in habitat use to anadromous Atlantic salmon. Identifying essential habitat is crucial when evaluating conservation status and planning measures to improve their status, including habitat restoration and identification of potential protection areas. We therefore tagged individual småblank with radio transmitters and tracked their movements over five periods (in the years 2014–2017). The aims of the study were (1) to examine the home range size and length of the river stretch used by individuals in the regulated main river and in a non-regulated tributary, and (2) to describe their habitat use in terms of water velocity and bottom substrate.

## Materials and methods

### Study area

The River Namsen (total length 174 km) originates from Lake Store Namsvatnet 450 m above sea level and enters the sea at the inner part of Namsenfjorden, Middle Norway (Fig. 1). The total catchment area is 6300 km<sup>2</sup>. The landlocked salmon, småblank, is distributed along a 90-km stretch of the upper River Namsen (from 72 to ≈ 160 km from the river mouth), in addition to several tributaries (Rikstad, 2004). Within the study area, width of the main river varies from 70 to 250 m, whilst width of the tributary varies from 30 to 50 m. This study was performed in two reaches: (a) in the upstream part of the main river, from Snåsamoen to Strompdalen (including Breidfos-sen and the mouth of River Frøyningsselva), and (b) in the major tributary (100 km<sup>2</sup> catchment area), River Mellingselva (Fig. 1), hereafter referred to as “main river” and “tributary”, respectively.



**Fig. 1** Location (red dots) of the field sites for radio tagging of småblank (*Salmo salar*) in the upper River Namsen

The main river is regulated for hydropower production and is characterized as a heavily modified water body. Nine power stations produce in total an average of 1831 GWh per year. Dams and weirs have caused major hydromorphological changes, and in total, more than half of the stretches available for småblank in the main river have changed from lotic to lentic habitat (Sandlund et al., 2014). A major impact is diversion of water from the main river, resulting in reduced catchment areas: for example, the catchment area at Bjørnstadfoss (147 km from the sea; Fig. 1) has been reduced by one-third. As a consequence, both total water discharge and mean water velocities are reduced throughout the year.

Our study site in the main river was not directly affected by dams or weirs, but by reduced water discharge due to the diversion of water. The reach has relatively uniform habitats with long slow-flowing areas combined with shallower riffles (water depths typically 0.5–1.5 m). River Mellingselva is an unregulated tributary with no anthropogenic migration

obstacles. Compared to the study site in the main river, the habitats are more diverse, with swift rapids, deep pools and waterfalls, and variable substrate categories of sand, gravel, pebble, cobble, boulder and bedrock. The water quality of the study areas in the

**Table 1** Environmental variables measured in autumn 2015

	Snåsamoen	Mellingselva
Calcium (mg/l)	< 5	< 5
Conductivity (µS/cm)	11	25
Nitrogen (µg/l)	70	85
Oxygen (%)	81	80
pH	6.8	7.0
Phosphorus (µg/l)	< 2	< 2
Temperature (°C)	12.5	13.2

Nitrogen and phosphorus were analysed by Analysesenteret in Trondheim kommune in December 2015. pH, conductivity, oxygen, calcium and temperature were measured in the field on 07.10.2015

main river and tributary is characterized by a pH over 6.0 and low contents of nitrogen and phosphorus (Table 1; nitrogen and phosphorus were analysed by “Analysesenteret Trondheim Kommune” whilst the other parameters were determined in field). Measured values were typical for these lower mountain areas.

To our knowledge, the only other fish species present in the study sites is brown trout *Salmo trutta* (Linnaeus, 1758). European minnow *Phoxinus phoxinus* (Linnaeus, 1758) has been recently introduced to the watercourse but is only found as single individuals within the study sites. The construction of fish passages in two hydro-power dams has given anadromous Atlantic salmon access to the lower part of the distribution area for småblank, hence reducing the allopatric distribution of småblank by 10 km in the main river. However, this has had no effect on our study sites further upstream.

#### Capture and tagging of fish

A total of 140 småblank (total length: mean = 169 mm, SD = 24, range 138–265 mm; body mass ( $n = 133$ ), measurements of body mass were missing for seven fish: mean 43 g; SD = 22; range 19–136 g) were captured using a backpack electrofishing apparatus (700 or 1400 Volt DC, depending on water conductivity) and tagged with radio transmitters (Advanced Telemetry Systems, ATS, USA, large model: F1440, 2.1 g in air, estimated battery life 122 days; small model: F1420, 1.3 g in air, estimated battery life 45 days). The fish were caught between 2014 and 2017 in different locations (Table 2). Individual fish were recognized by the use of different transmitter frequencies within the 142.000 to 142.700 MHz range. Before implanting the transmitters, the fish were anaesthetized in 2-phenoxy-ethanol (EEC No 204-589-7, 0.50 ml per 1 water, each fish kept for 4.0–4.5 min in the bath). During surgery (approximately 3.5 min per fish), a tube with flowing water was placed in the fish mouth, enabling oxygen uptake by the gills. Total body length and mass were measured and a small part of the adipose fin was sampled for genetic sex determination and 5–7 fish scales were sampled for age reading. To reduce the potential effects of catch and tagging on results and conclusions, tracking data from the first day after release were not included in the analyses. All the fish were caught and released in the same location. Based

on tagging area, season and year, the 140 fish were divided into eight groups, and released at six localities (Table 2).

#### Radio tracking

The study was conducted from August 2014 to August 2017, covering five tracking periods (Fig. 2). Each tracking period lasted 4–5 months. New fish were tagged for each period due to the restricted battery life of the transmitters. Tracking surveys were conducted every third week, by wading through the river using an ATS model R4500s receiver connected to a five-element Yagi antenna. During each survey, individual fish were positioned to the nearest  $\pm 1$  m. At the position of each fish, water depth was measured and water velocity and substrate were categorized (see Table 3). In situations when it was not possible to get close to the fish, due to floods or the fish being located in deep waters, the GPS-positioned location of the person tracking the fish and the direction and approximate distance to the fish were noted. During these cases, depth, velocity and substrate were not recorded. Each fish position was recorded using a Garmin GPS model 60CSx (accuracy of circa 5 m).

The tracking effort was limited by time constraints, and the tracking was therefore restricted to a certain area. In the main river (at Snåsamoen, including the confluence with the tributary Mellingselva), a total river surface area of 101,500 m<sup>2</sup> was covered during tracking. In the tributary, an area of 70,000 m<sup>2</sup> was covered (Fig. 1).

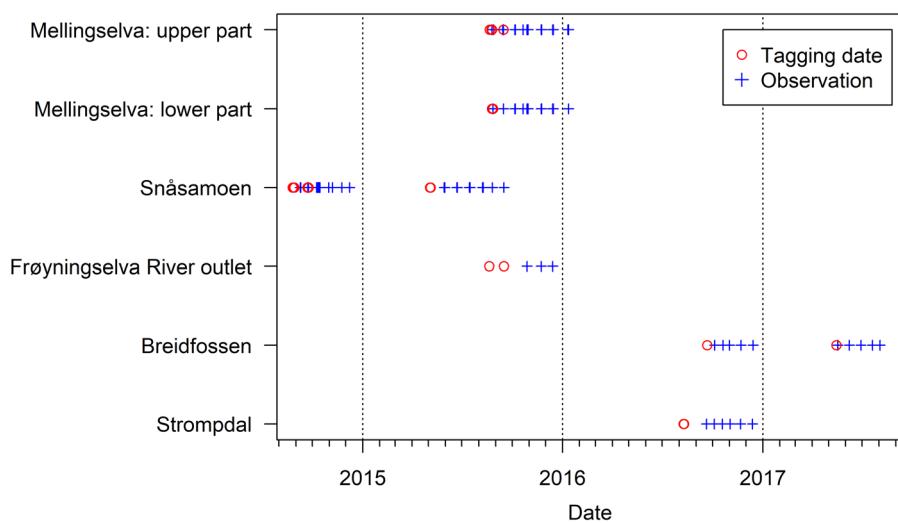
#### Determination of age, sex and body condition

The age of the fish was determined from scale analysis (Dahl, 1910; Závorka et al., 2014). Scales from 119 fish were suitable for age determination. For the remaining fish, samples only contained replacement scales that were unsuitable for age analysis.

From the 140 radio-tracked fish, genetic DNA was sampled from 132 adipose fins and analysed for sex and species determination at the Norwegian Institute for Nature Research. The sex for the eight remaining individuals was determined visually in the field during the autumn and hence prior to spawning. For the 132 individuals, the sex was determined from amplifying the male-specific sdY gene in Atlantic salmon using the marker developed by Quéméré et al. (2014),

**Table 2** Overview of the nine groups of tagged småblank

Locality	Tagging group	Date (release)	Number of days of tracked (d)	Number of fish (n)	Total body length (mm)	Mass (g)	Age	Proportion of females
Snåsamoen	1	2014-08-26	104	37	165 (140–210)	39 (23–81)	3.4 (3–4)	0.74
Snåsamoen	2	2015-05-04	135	20	163 (149–180)	29 (19–39)	4.1 (2–5)	0.85
Mellingselva: upper part	3	2015-08-21	144	25	162 (140–206)	42 (25–93)	3.2 (2–4)	0.52
Mellingselva: lower part	4	2015-08-25	140	11	153 (138–167)	34 (23–46)	3.6 (3–4)	0.45
River Frøyningsselva outlet	5	2015-08-20	116	10	158 (143–188)	34 (24–56)	3.2 (2–5)	0.89
Breidfossen	6	2016-09-21	84	8	181 (160–206)	51 (35–77)	3.5 (3–4)	0.25
Breidfossen	7	2017-05-15	80	11	176 (141–253)	47 (20–115)	3.8 (3–6)	0.29
Strompdalen	8	2016-08-09	126	18	199 (150–265)	70 (29–136)	3.9 (2–7)	0.61



together with genetic markers for differentiating Atlantic salmon and brown trout using the PCR assay developed by Karlsson et al. (2013). Because the *sdY* gene is only amplified in males, absence of an amplification was interpreted as a female, but only when the genetic markers to differentiate between species amplified well in the same PCR multiplex. DNA was extracted from fin-clips preserved in ethanol using the DNEASY tissue kit (QIAGEN). Of the 140 tagged fish, there were 87 females and 53 males. All of the tagged fish were salmon (*småblank*) and none were hybrids between salmon and trout. Fulton's body condition factor (Fulton, 1904), based on measurements of body mass and total length, was calculated for the 133 fish for which mass was measured.

#### Data analyses

The characteristics of tagged fish were first summarized using bivariate statistics. First, the relationship between body length and age was examined using Pearson's correlation and the relationship between body length and sex using a Student's *t* test. Next, the potential differences in body length, body mass and condition factor between fished tagged in the main river and those tagged in the unregulated tributary were examined using Student's *t* tests.

For fish that were unreachable by wading, bankside measurements of locations were used, which were repositioned in the GIS software Map 10.2 (desktop.arcgis.com/en/arcmap/) using field notes with estimates of the distance to the fish. Fish locations measured both within the river and from bankside measurements were used for home range estimation.

Home ranges based on GPS positions were estimated for each *småblank* when there were at least five observations using the following procedure: (1) A 50% home range polygon (excluding 50% of the positions furthest away from the home range centroid) was calculated for each *småblank* using the R function *mcp()* from the *adehabitatHR* library. A 50% limit prevented the calculated home ranges being unduly influenced by occasional long-distance movements. For comparison, 95% home ranges were also calculated. (2) The home range polygon for each *småblank* was further refined to exclude bankside areas by only retaining the intersection between the calculated home range polygon and the river channel using the R function *gIntersection()* from the *rgeos* library. Spatial

and temporal differences in 50% home ranges were tested using Mann–Whitney *U* test. Spatial differences were analysed by comparing home ranges in the main river with those in the tributary, using the home ranges estimated from observations in both parts of the river in the autumn. Temporal differences were analysed by comparing home ranges in the autumn with those in the spring, using home range data available from both seasons at Breidfossen and Snåsamoen in the main river.

Travel distances between consecutive locations for *småblank* individuals were estimated as follows: (1) For consecutive locations that did not have a boundary (river bank) between them, a simple Euclidean distance was used. (2) For consecutive locations separated by an intermediate boundary, a minimum within-river distance between the locations was estimated using the R-function *shortestPath()* from the *gdistance* library.

The effect of *småblank* characteristics on home range and maximum length of the river stretch used (defined as the maximum distance between two consecutive observations) was examined using linear regression. Both home range and maximum length of the river stretch used were log transformed to ensure a normal distribution of residuals. Length, body mass, condition factor (K), age (in years), sex and the release locality (watercourse stretch) were considered as potential predictors. Generalized variance inflation factors (GVIFs) were examined to explore potential correlation amongst predictors using the R-function *vif()* from the *car* library, following the method outlined by Zuur et al. (2009). High GVIFs were found for body length and mass when they were included in the same model, and body mass, which had the highest GVIF, was therefore excluded (reducing GVIFs to < 1.6 for body length, condition factor and age, sex and release locality). Initial models were therefore run with body length, condition factor, age, sex, and release locality, and models were simplified using a stepwise approach.

The datasets generated and analysed during the current study are available upon request in the NTNU University Museum repository NaTron, <https://natron.vm.ntnu.no/dataCollection/>.

## Results

### Body length and body condition

Body length and age were positively correlated (Pearson's correlation,  $t = 7.60$ ,  $P < 0.001$ ,  $n = 119$ ). There was no difference in body length between males (mean = 164 mm, S.D. = 25, range 138–165 mm) and females (mean = 170 mm, S.D. = 20, range 140–231 mm) (Student's  $t$  test,  $t = 1.29$ ,  $P = 0.20$ ,  $n = 132$ ). In the autumn, småblank tagged in the regulated main river had greater body length and body mass (mean length = 176 mm, mean mass = 49.2 g) than those tagged in the unregulated tributary (mean length = 159 mm, mean mass = 39.6 g) (Student's  $t$  test,  $t = 3.86$ ,  $P < 0.001$  (body length),  $t = 92.6$ ,  $P = 0.019$  (body mass),  $n = 95$ ). However, fish tagged in the regulated main river had lower condition factor (mean = 0.83, range 0.73–0.95) than those tagged in the unregulated tributary (mean = 0.95, range 0.78–1.14) (Student's  $t$  test,  $t = -7.27$ ,  $P < 0.001$ ,  $n = 95$ ). No comparison was made for the spring, because fish were then only tagged in the main river.

### Home range

The home range size could be estimated for 77 småblank. A median of eight observation locations (S.D. = 4.4, range 5–17) was available for analysis from these fish. The fish stayed within relatively small home ranges, with an average 50% home range area of 1123 m<sup>2</sup> (S.D. = 2794 m<sup>2</sup>, range 1–18 644 m<sup>2</sup>), and an average 95% home range area of 4 416 m<sup>2</sup> (S.D. = 7337 m<sup>2</sup>, range 3–44 408 m<sup>2</sup>) (Fig. 3). There was large variation in home range size amongst the fish (Fig. 3). Home ranges were larger for småblank residing in the upstream part of the study area (Snåsamoen in the main river and the tributary River Mellingselva) than in the downstream area (Strompdalen and Breidfossen) (Table 4). The 50% home range size increased with fish age (Table 4). There was no relationship between 50% home range size and fish condition, body length or sex, which were excluded from the final model through stepwise selection. Despite large differences in median values, the 50% home range size did not differ between the main river (median = 47 m<sup>2</sup>,  $n = 39$ ) and the tributary (median = 171 m<sup>2</sup>,  $n = 24$ ) for fish tracked in the autumn

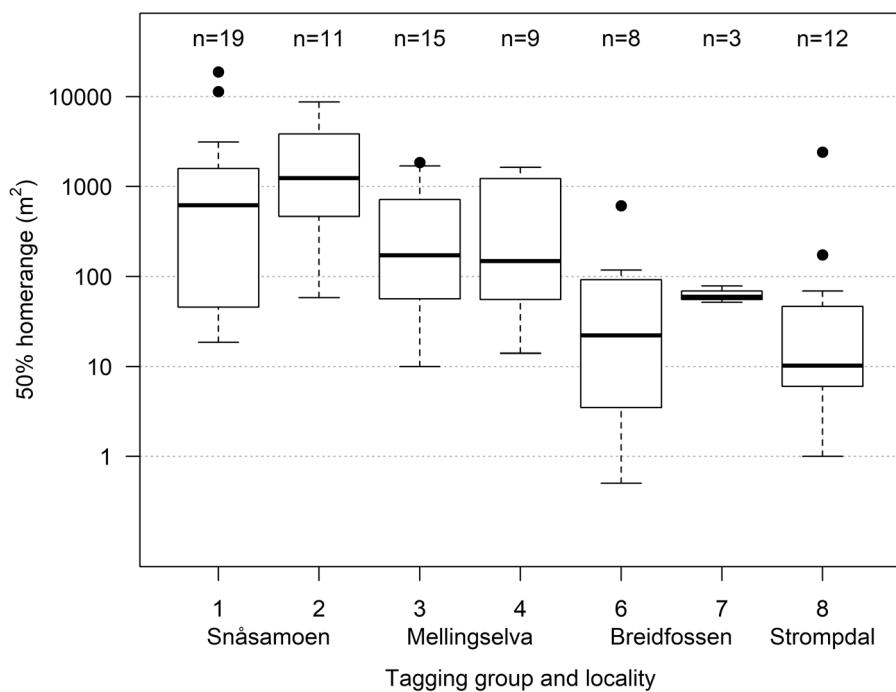
(Mann–Whitney  $U$  test,  $U = 365.5$ ,  $P = 0.15$ ). At Breidfossen and Snåsamoen in the main river, småblank were tracked both during autumn and spring. In both localities, 50% home range sizes did not differ between the spring and the autumn (Mann–Whitney  $U$  test; Breidfossen:  $U = 8$ ,  $P = 0.50$ ; Snåsamoen:  $U = 72$ ,  $P = 0.17$ ).

### Length of river stretch used

Twelve of the 140 released småblank were only detected on one occasion each, so it was not possible to determine the length of the river stretch used for these individuals. Of the remaining individuals, the median maximum length of the river stretch used (estimated as the longest distance covered between consecutive tracked locations of each individual) was 242 m (range 0–2080 m;  $n = 128$ ). The median maximum length of the river stretch used was positively correlated to body length (Table 5), but not to age or condition factor. The maximum length was also greater for males than for females. It was also dependent on locality, with småblank in the localities further upstream (Snåsamoen and Mellingselva) and Breidfossen tending to move over longer stretches (Fig. 4, Table 5).

### Habitat use

Småblank were most commonly found in areas with coarse (gravel/pebble to boulder) substrate (90%), with moderate to strong currents (79%), and at water depths shallower than 60 cm (82%) (Fig. 5, Table 3). They were, however, observed across a range of habitats including areas with fine substrate or bedrock (10%), and in either stagnant or slow waters (16%) or cascades (5%). Småblank were also observed in waters deeper than 60 cm, and all småblank in the main river at Strompdalen were found in waters deeper than 100 cm. The substrate used differed between the tributary and the main river. At Snåsamoen, småblank were mainly found in areas with coarse cobble/boulder substrates (particle size 16–35 cm), whilst småblank in the other parts of the study area were also found in habitats with finer gravel/pebble substrates (2–15 cm).



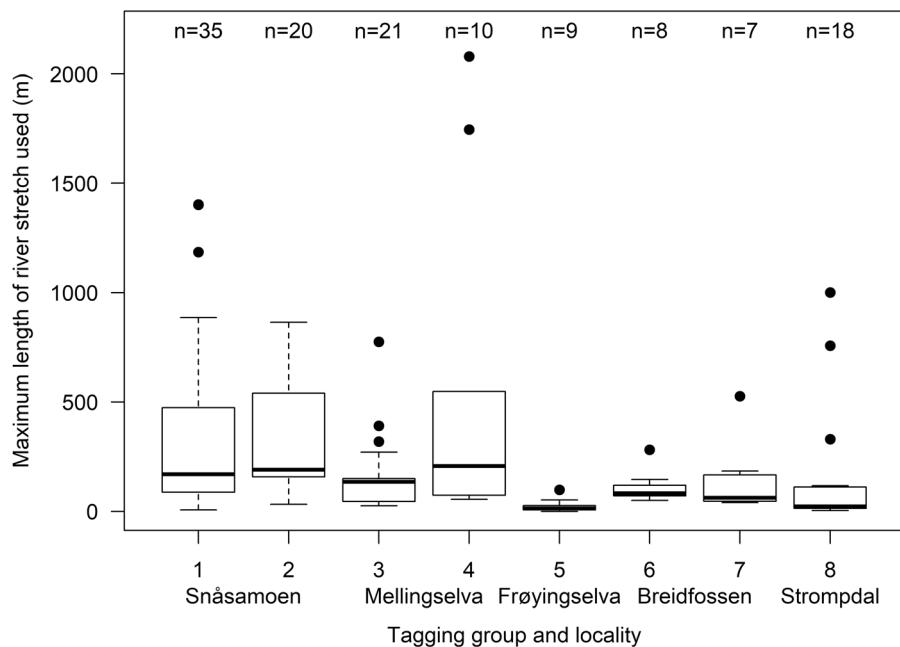
**Fig. 3** Home range estimates for radio tagged småblank (*Salmo salar*) in the upper River Namsen (50% home range size). Home ranges could not be estimated for tagging group 5 due to a low sample size of observed positions

**Table 4** Retained coefficients of stepwise linear regression model of effect of body length, condition factor, age, sex and release locality on småblank 50% home range size

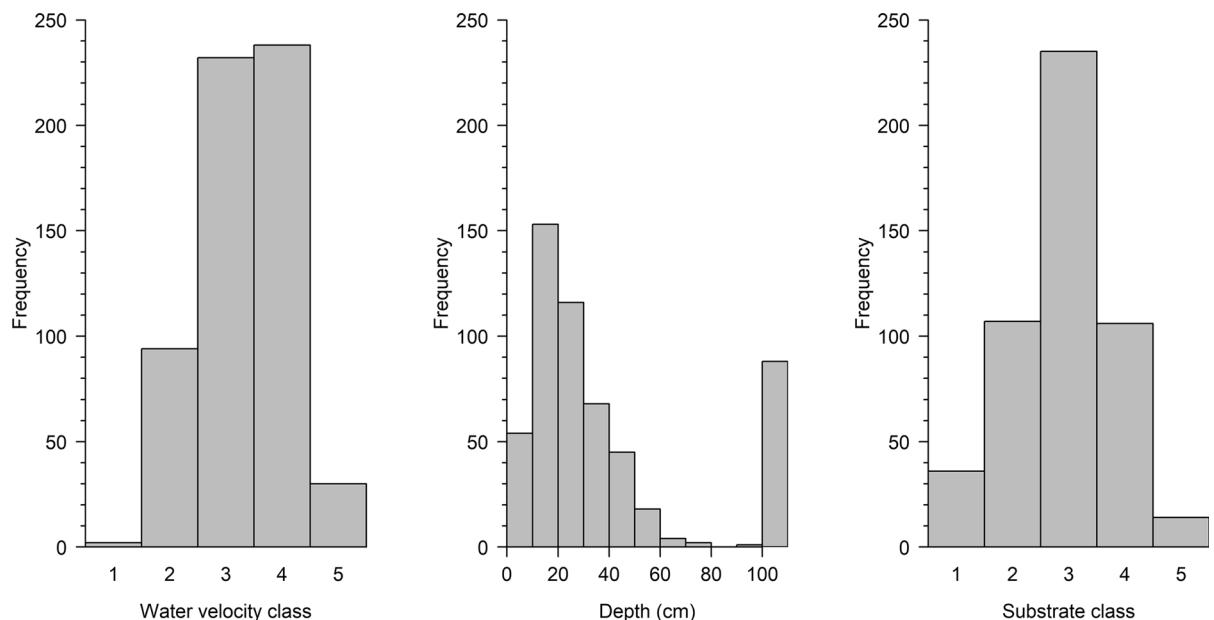
	Estimate	SE	t value	Pr(>  t )
(Intercept)	- 1.334	1.182	- 1.128	0.263
Age	1.138	0.289	3.936	< 0.001
Locality: Breidfossen	0.522	0.757	0.690	0.493
Locality: Snåsamoen	3.058	0.644	4.746	< 0.001
Locality: Mellingselva, lower part	2.250	0.799	2.817	0.006
Locality: Mellingselva, upper part	2.938	0.738	3.98	< 0.001

**Table 5** Retained coefficients of stepwise linear regression model of effect of body length, condition factor, age, sex and release locality on maximum length of the river stretch used

	Estimate	Std. Error	t value	Pr(>  t )
(Intercept)	- 0.068	1.259	- 0.054	0.957
Body length	0.018	0.006	2.876	0.005
Sex: Male	0.647	0.242	2.670	0.009
Locality: Breidfossen	1.203	0.441	2.728	0.008
Locality: River Frøyningsleve outlet	- 0.339	0.537	- 0.631	0.529
Locality: Snåsamoen	2.389	0.397	6.011	< 0.001
Locality: River Mellingselva: lower part	2.500	0.529	4.722	< 0.001
Locality: River Mellingselva: upper part	1.623	0.437	3.717	< 0.001



**Fig. 4** Maximum length of river stretch used by radio tagged småblank (*Salmo salar*) in the upper River Namsen



**Fig. 5** Habitat distribution (water velocity, depth and substrate class) of locations where radio tagged småblank were observed

## Discussion

The småblank did not display similar migratory behaviour as observed in most other landlocked populations of Atlantic salmon. Compared to the

migrations between rivers and lakes in landlocked salmon in the watercourses of Ladoga, Onega, Saimaa, Vänern and Otra (Berg, 1985; Kazakov, 1992; Anon, 2011), the småblank were relatively stationary, with small home ranges (spending 50% of their time within

an area of only 1123 m<sup>2</sup> and 95% of the time within an area of 4416 m<sup>2</sup>). They were also confined to short river stretches, as no fish moved over a longer river stretch than about 2 km, and half the fish did not use a river stretch longer than about 240 m. This shows that the tagged fish were stationary within the same river section during the entire tracking period. The short movements of småblank concur with the results in a previous study (1954–1958) from the same area, based on Carlin tags and recaptures (Berg, 1984b). Of approximately one thousand tagged fish, all recaptures were reported from the same area as the fish were tagged, except one individual recaptured 24 km downstream, and ten individuals about 10 km downstream (Berg, 1984a). Major migrations are usually associated with reproduction, seasonal preferences for different habitats (e.g. for overwintering), or as an effect of increased body size. In salmonids, smoltification stimulates downstream migration in both anadromous and non-anadromous salmon (e.g. Kendall et al., 2015). In the current study, småblank were tracked during spring, summer and autumn without any indication of seasonal or other kinds of migrations. For anadromous Atlantic salmon, spawning migrations take place during the summer and autumn and the smolt migration in the spring, so these kind of migrations should have been observed if they occurred.

Småblank residing in the upper part of the study area (Snåsamoen and Mellingselva) had larger home ranges than those residing in the downstream part (Breidfossen and Strompdalen). There might be several reasons for this difference: (1) Access to food and/or shelter may differ between the two areas, and småblank may adapt their home range accordingly. (2) The number of competitors may differ between the two areas, and the home range may be adapted to the competitive pressure. Displacement of individual salmonids by more competitive conspecifics in territorial communities of juvenile fish is well-known, and in such cases, the sub-dominant individuals may need to explore larger areas without being territorial (Grant & Kramer, 1990; Titus 1990; Grant, 1997). Experimental studies have shown that the territory size of juvenile salmonids decreases with increasing density, and increases with decreasing levels of accessible food (Kalleberg, 1958). Due to the low density of småblank in most of the studied areas, it is less likely that the populations were regulated by a high level of

territoriality. However, access to food and shelter and the level of competition may still have a major impact on the spatial distribution and habitat use of småblank, and these factors may in combination influence their home range size.

In general, home range increased with fish age, and length of the river stretch used increased with fish body length. Smaller individuals of småblank share several characteristics with juvenile anadromous salmon, which in general need access to suitable shelter (Heggenes et al., 1999; Bremset & Heggenes, 2001; Finstad et al., 2007) to reduce the predation risk (Metcalfe et al., 1987; Vehanen, 2003). In general, larger småblank may be less vulnerable to cannibalism and predation. Hence, they may be less dependent on accessible shelter and may move over larger areas than smaller individuals.

The difference in substrate use between Snåsamoen, where småblank mainly used coarse cobble/boulder substrates, and the other study areas, where fish more often were found in areas with finer gravel/pebble substrates, is probably due to differences in the bottom substrate between these localities and not to site-specific preferences. Both substrate categories provide suitable shelters for juvenile salmonids (Heggenes et al., 1999; Bremset & Heggenes, 2001; Orpwood et al., 2003), and an earlier study has indicated that this is the case for similar-sized småblank as in the current study (Norum, 2010).

Småblank mainly utilized areas with water depths shallower than 60 cm and moderate or high water velocities ( $> 50 \text{ s}^{-1}$ ). These water velocities make it possible for the fish to keep position near the river bottom and at the same time feed on drifting invertebrates. Access to drifting invertebrates depends on water velocities, since the rate of drifting material within a given river section is proportional to the speed of flow (Metcalfe, 1986; Nislow et al., 1998). The water velocities utilized by småblank in the present study (generally  $> 50 \text{ cm s}^{-1}$ ) were faster than what is normally preferred by juvenile anadromous salmonids. Degraaf & Bain (1986) and Morantz et al. (1987) found that 0+ Atlantic salmon prefer areas with water velocities slower than  $20 \text{ cm s}^{-1}$ , whilst Heggenes et al. (1995) observed that older juveniles preferred water velocities up to  $25 \text{ cm s}^{-1}$ . As småblank have a larger maximum size than juvenile salmon as well as significant larger pectoral fins (Berg, 1984a), the maximum swimming capacity of

småblank is quite high. Consequently, large småblank are able to hold position in high-velocity areas of rivers that are unsuitable for smaller individuals.

## Conclusion

In conclusion, this study shows that småblank, in contrast to other salmon populations, have relatively restricted home ranges and that they do not migrate between different parts of the river. These findings may partly explain why småblank is divided into several genetically distinct populations (Ståhl, 1987; Vuorinen & Berg, 1989; Sandlund et al., 2014). Further, the current study supports previous observations showing that småblank prefer fast-running waters, and are able to use river areas unsuited for smaller individuals such as juveniles of salmon and trout. A special adaptation to such areas with coarse river substrates implies a vulnerability to anthropogenic activities such as water diversion, dams and weirs. As a consequence of being a small population with special adaptations, further habitat degradation might inflict a decline in the småblank population and subpopulations.

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