# Application of the Particle Filter to COHERENS V2 model of Lake Säkylän Pyhäjärvi

Akiko Mano\* and Takayuki Shuku†

# Abstract:

The particle filter (PF, Gordon et al, 1993; Kitagawa, 1996) was applied to determine the optimal parameters for the sediment transport for the lake Säkylän Pyhäjärvi model with COHERENS Version 2. After having obtained the optimal parameters, time series of the TSM (Total Suspended Matters) concentrations at the automatic station were compared among the model result of (1) before- (2) after applying PF and (3) the monitored observation data. The model result with using the optimal parameters after applying PF was well-reproduced the observation data. The application of PF can be considered as a quite usable way to improve the model reproducibility.

# 1. Introduction

Our research was focused on suspended solid simulation in Lake Säkylän Pyhäjärvi (Fig. 1), the largest lake in southwestern Finland, with the COHERENS V2 model (COHERENS version 2(Luyten, 2011)). The aim of this study is to apply PF to the model to determine the optimal parameters related to sediment transport to improve the model reproducibility.

The lake has two inlets from Ylaneenjoki and Pyhäjoki rivers and one outlet from Eurajoki river. These inflow and outflow river water and meteorological conditions such as wind are considered to be the main driving forces of water circulation in the lake.

# 2. Data and materials

To put it plainly, the model itself and all data such as the boundary conditions, the initial conditions and the observation data were based on ones used in satellite data assimilation (Mano, 2013) as described below.

The boundary conditions at river mouths were used the following data. Time series of river discharge and TSM loading at each river mouth were extracted from Hertta-database of Finnish

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Environmental Administration. Those of temperature were done from VEMALA, the water quality component of the Watershed Simulation and Forecasting System (Vehviläinen B, et al., 2005) of the Finnish Environment Institute. Meteorological data was obtained from Finnish Meteorological Institute which contains wind speed and direction, air temperature, humidity, cloud coverage and air pressure. The initial condition for the TSM concentration on the lake surface was converted from Turbidity of the satellite data on June 26 in 2009 constructed in Finnish Environment Institute. The observation data was monitored on the surface of the lake at the automatic station at the main basin of the lake by Finnish Environment Institute.

The parameters related to the sediment transport were used as shown in Table 1. PF was applied for these four parameters.

**Table 1** Four parameters related to the sediment transport. Original value means one used in the model for satellite data assimilation.

| Parameter name    | Notes                                       | Original value |  |  |
|-------------------|---|----------------|--|--|
| bio_sinkrate      | settling velocity [m/s]                     | -0.000 000 03  |  |  |
| sedimentationrate | sedimentation rate (ex. 0.1 means that 10 % | 0.1            |  |  |
|                   | of sedimentation occurs par day.)           |                |  |  |
| alphas            | resuspension rate parameter [g/m2/s]        | 0.0025         |  |  |
| rnsed             | resuspension exponential factor             | 3.0            |  |  |

# 3. Methods

PF is a method for estimating probability density functions (PDFs) of state variables (Gordon, 1993; Kitagawa, 1996), and it can be applied to parameter identification of input parameters in numerical simulation. The considered sets of parameters called "particles" in PF. The simulation code of the PF was developed in (Shuku et al., 2012). Here, four parameters shown in Table 1 were regarded as "particles", and the observation data was regarded as the reference data. Preparing a number of sets of four parameters and performing a number of model calculations from COHERENS with using the parameters ("particles") were required for PF.

The PF usually requires about 1000 sets of parameters (particles) by a random function to find out the optimal ones. Nevertheless, by using not a random function but LDS (Low-discrepancy sequence (Tezuka, 1995)), the required sets will be reduced to 200 sets. This is because LDS generates more efficient numbers for PF to converge the results than a random function does. The images of the numbers using random function and LDS are shown in Fig. 1. Because of the time limitation, LDS was used for generating 200 particles. An application developed by one of the authors (Shuku) for generating numbers with LDS was used in this study, and the mean and the deviation for it are shown in Table 2.

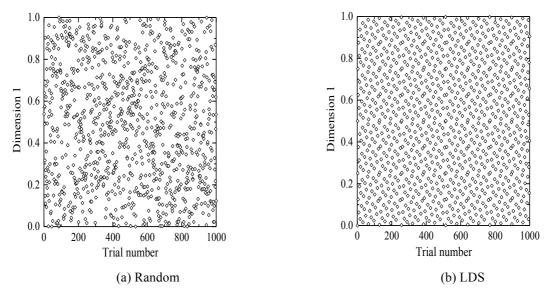


Fig. 1 Image of the numbers using a random function(a) and LDS(b).

**Table 2** Mean and deviation for an application with LDS for four parameters for generating 200 sets of parameters.

| Parameter name    | Mean         | Deviation  | Considered range            |
|-------------------|--------------|------------|-----------------------------|
| bio_sinkrate      | -0.000 002 2 | -0.000 001 | -0.000 003 2 ~ -0.000 001 2 |
| sedimentationrate | 0.5          | 0.5        | 0.0 ~ 1.0                   |
| alphas            | 0.001 501    | 0.001 499  | 2.00E-06 ~ 0.003            |
| rnsed             | 11. 5        | 10.505     | 1 ~ 22                      |

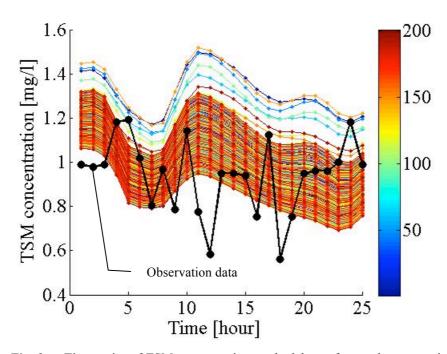
Procedures were as follows. Some test calculations were performed with the parameters to guess the optimal parameters approximately in advance in order to narrow the range of values to be considered as particles. Here, only "bio\_sinkrate" was focused because of the time limitation and that its sensitivity was the largest of the four as mentioned in the report (Mano, 2013). The tested values were in the hundredfold to hundredth range of the original value, -0.000 000 03. As the results of the test calculations, -0.000 002 20 was assumed to be close in value to the optimal one according to time series of the TSM concentration at the lake surface at the automatic station with the monitored observation data. An application for generating sets of parameters with LDS in an Excel file format was made by one of the authors (Shuku) for this study. 200 particles were generated with the setting shown in Table 1. The generated particles are shown in Fig. A-1 and listed in Table A-1. The parameters for each case were read from the file named "param.txt "(See Table A-1) automatically. This is because a directory's name of a case was made to be equivalent to a line number in "param.txt". Extracted time series of TSM concentration at the automatic station were gathered from 200 cases. With the data and observation data provide the optimal parameters with PF.

The calculation term was 13:00, June  $26 \sim 12:00$ , July 6, the period when the CHL concentrations seem to be low according to satellite image and monitored the CHL

concentration at the automatic station. For the PF application, calculation term is June 26, 13:00(close to the time when satellite data was obtained, 12h59m14s)  $\sim 12:00$ , June 29. Only the results from June 28  $12:00 \sim$  June 29 12:00 were used for running the PF application. For comparing the before-after results, the ending time was extended until 12:00, July 6. The calculation term started from June 26 for all cases, the satellite data of the day was used as the initial value of the TSM concentration on the lake surface.

# 4. Results

Fig. 2 shows time series of the TSM concentration extracted from 200 cases and the observation data for 24 hours. The observation data is TSM concentration with no biomass converted from Turbidity monitored at the automatic station.



**Fig. 2** Time series of TSM concentration on the lake surface at the automatic station. 200 cases of model result and an observation data.

PF was applied to the data of 200 cases and observation data. Fig. 3 shows the PF results for four parameters. The converged values show the optimal parameters. Thus the optimal values of parameters are shown in Table 3.

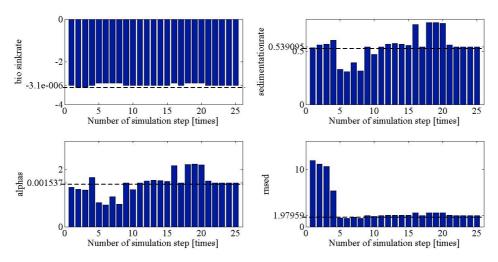


Fig. 3 PF result for four parameters

 Table 3
 Original values and optimal values for four parameters

| Parameter name    | Original value | Optimal value |
|-------------------|----------------|---------------|
| bio_sinkrate      | -0.000 000 03  | -0.000 003 1  |
| sedimentationrate | 0.1            | 0.539 095     |
| alphas            | 0.0025         | 0.001 537     |
| rnsed             | 3.0            | 1.979 59      |

# 5. Discussions and conclusions

Comparisons with the results before-after applying the optimal parameters were performed. Fig. 4 shows the time series of TSM concentration at the automatic station. "Before" and "After" are the results from calculations with using the original value and the optimal value in Table 3, respectively. As PF was applied to the results of "Before" from day 3 to 4 in Fig. 4 to obtain the optimal values for the four parameters, the results after day 4 shows the pure predictions. According to Fig. 4, the results of "After" are dramatically well-reproduced the observations. Thus it can be regarded that applying PF improved the model prediction in this case.

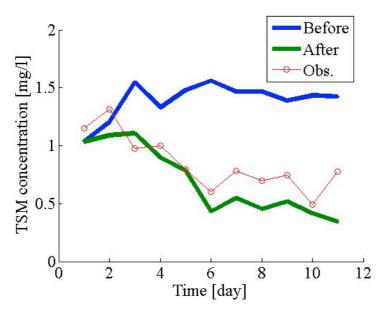
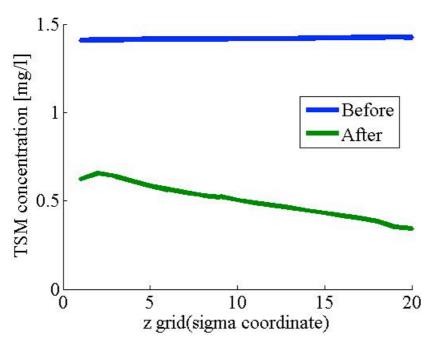


Fig. 4 Time series of TSM concentration at the automatic station.

Fig. 5 shows the comparison of vertical profile at the automatic station on 11th day of the calculation term. In the case of after applying optimal parameters, the trend, the deeper part has the larger TSM concentration, was reproduced as monitored at the automatic station (Mano, 2013).



**Fig. 5** Comparison of vertical profile of before-after applying the optimal parameters determined from PF on 11th day of the calculation term at the automatic station. z=20 and z=1 mean the lake surface and the bottom, respectively.

According to these results, to determine the optimal parameters with using PF seems to be very usable. This is because, not only time series at the surface but also the vertical profile, to apply PF increased the reproducibility of the model dramatically. However, to make sure the usability of PF, more tests would be required.

Two ideas for future works to make sure the usability of PF would be considered. One would be to apply PF to other calculation term, for example, not for 24 hours from June 28 12:00 but for 24 hours from on June 27 12:00. The other would be to apply PF from June 26 13:00  $\sim$  July 6 12:00 (11 days) with using a coarser grid model. To apply PF for longer term could provide us more reliable clues for consider the usability of PF.

# Acknowledgment

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# Assimilation of satellite data to 3D hydrodynamic model of Lake Säkylän Pyhäjärvi

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# **Abstract:**

To analyze the applicability of occasional initialization (OI) of total suspended matter (TSM) concentration field based on turbidity derived from satellite data to numerical simulation, dispersion studies of suspended matter in Lake Säkylän Pyhäjärvi (lake area 154 km²; mean depth 5.4 m) were conducted using the 3D COHERENS simulation model. To evaluate the practicality of OI, five cases with different initialization frequencies were conducted: 1) every time, when satellite data were available; 2) every 10 d; 3) 20 d; and 4) 30 days; 5) control run without repeated initialization. To determine the effectiveness of initialization frequency, three methods of comparison were used: simple spatial differences of TSM concentration without biomass in the lake surface layer; averaged spatial differences between initialization data and model results; and time series of TSM concentration and observation data at 1 m depth at the deepest point of the lake. Results showed that OI improves the prediction significantly when applied often, i.e. at 10 day intervals. Even if it is applied less often, OI improves the prediction.

# **Key words**

COHERENS; model prediction; suspended particulate matter; suspended solids; turbidity; water quality

This paper has already been submitted to water science and technology

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# WP 4

Statistical analysis for habitat modelling and relationship between land use and river water quality

# Juvenile salmon patch identification and comparison using Echelon analysis

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# **Abstract:**

We studied a habitat patch identification technique for juvenile Atlantic salmon utilizing the suitability index and Echelon analysis. After identification of patches, we evaluated each identified patch using the likelihood ratio statistic so that the best suitability areas for salmon were determined. Laser and sonar are interferometric measuring systems we used for our salmon data. As a result of comparison between these two data sets, we found that sonar was more optimistic than laser for measuring suitability index.

# 1. Introduction

Various patch identification methods for habitats have been published (Fortin et al., 1995; Plotkin et al., 2002). A patch is defined as a spatially homogeneous area where at least one variable has similar attributes either of category or quantitative value (Fortin et al., 2005). Therefore, a patch was adopted in studies for both plants and animals. Patches can be identified using various variables such as tree or animal abundance and percentage coverage of trees. However, a solid method of identifying patches has not been established yet.

In the present study, we identified juvenile salmon patches using Echelon analysis. Oda et al. (2012) suggested the technique for identification of patches in a forest using Echelon analysis. We presented that the technique can also be used for an animal, juvenile salmon, and evaluated the identification patches.

In this paper, we first explained the patch identification method we used, and identified juvenile salmon patches based on the suitability index. We then assessed these identified

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patches using the two different patch evaluation methods called laser and sonar, and finally compared the data derived from these two patch evaluation methods for measuring suitability index.

# 2. Survey site and data

The subarctic river Utsjoki is a tributary of a highly productive Atlantic salmon (*Salmo salar*, L.) river, Tana River. We modeled a stream stretch, which is known to have a strong Atlantic salmon juvenile population. The modeling procedure was undertaken for two different datasets with different accuracy: laser and sonar.

We applied life stage-specific habitat preference criteria (HPC) for depth, velocity and substrate preference by Atlantic salmon juvenile (Mäki-Petäys et al., 2002, 2004, unpublished data at http://www.rktl.fi/www/uploads/pdf/raportti284.pdf) to acquire a combined suitability index (CSI) value (product of habitat preference values). For this analysis we used only one discharge situation (20 m³s⁻¹) recognized as a normal summer flow.

# 3. Echelon analysis

Echelon analysis (Myers et al., 1997; Kurihara et al., 2000; Ishioka et al., 2007) is a method to show phase structure in Echelon dendrogram based on response variables and neighboring information.

# 3.1. Echelon analysis procedure

A phase structure of data can be shown using Echelon dendrogram (Figure 3.1): The surface value is shown such as contour lines in the left picture in Figure 3.1. The middle and right pictures show a lateral view. The right picture is Echelon dendrogram.

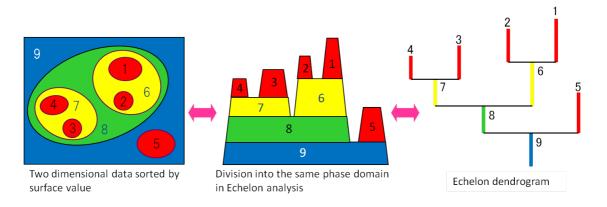


Fig. 3.1 The diagram of Echelon analysis.

We call the Echelon numbers 1, 2, 3, 4 and 5 as a "peak", the Echelon numbers 6, 7 and 8 as a "foundation" and the Echelon number 9 as a "root".

Each cell has a value as presented in Figure 3.2, and each cell has neighboring information. For example, cell [C3] is located at the center of its adjacent cells of [B2], [C2], [D2], [B3], [D3], [B4], [C4] and [D4].

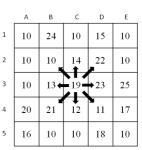


Fig. 3.2  $5 \times 5$  array data.

# 1. Establish peaks

At first, the cell [E3], with the maximum value in the 5×5 array data, is identified (Figure 3.3). Among adjacent cells of the cell [E3], the largest value, the cell [D3] is identified. The third ([D2]) and fourth ([C3]) values are identified in the same manner. The cell [C3] has the adjacent cell [B4], which is larger than the cell [C3]. Therefore, the first peak consists of three cells ([D2], [D3], [E3]). Other peaks {G(2), G(3), G(4)} are found in same manner (Figure 3.4).

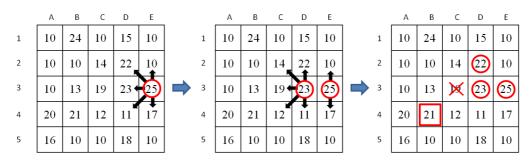


Fig. 3.3 Established peaks procedure.

# 2. Establish foundations

Excluding the four peaks identified earlier, the cell [C3] is the maximum value in  $5 \times 5$  array data based on its spatial location relationship (Figure 3.5), followed by the cell [E4]. Next, the cell [C2] becomes foundation between the G(2) and foundation [E4], [A5] and [D1]. The rest of cells are included in same [C2] foundation, since these are inevitably adjacent to peaks and the foundations.

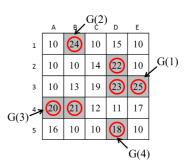


Fig. 3.4 Established peaks.

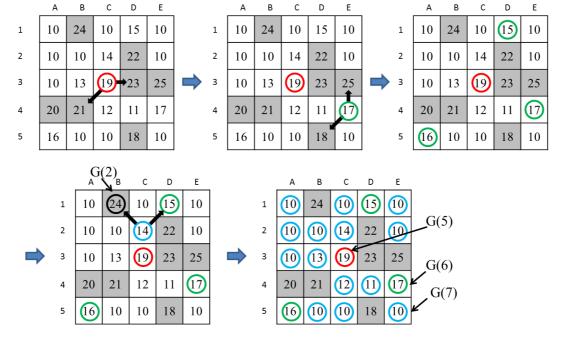


Fig. 3.5 Established foundations.

# 3. Establish Echelon dendrogram

Echelon dendrogrm can be established when peaks and foundations are identified in  $5\times 5$  array data. Four identified peaks: G(1) - G(4) are linked to foundations in order to draw a dendrogram (Figure 3.6). The foundation G(5) links the peaks G(1) and G(3), and the cell E[4] links the peak G(4) and foundation G(5). The cells [A5] and [D1] belong to the foundation of cell [E4]. These three cells establishes the foundation G(6). The rest of cells are the root G(7).

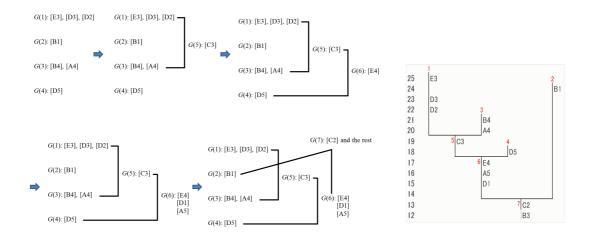


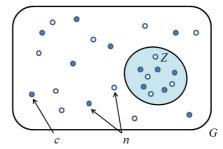
Fig. 3.6 Echelon dendrogram procedure.

# 2.2. Hotspot Detection

Establishing an Echelon dendrogram makes it easy to detect statistical significance among groups like G(1). This is an advantage of Echelon analysis. We call this significant area "hotspot". There is a subset area Z in the area G.  $p_1$  is the population probability with an attribute within the area Z, and  $p_2$  is the population probability with an attribute outside the area Z. The probabilities of all individuals with attributes are mutually independent. The hypothesis is as follows:

$$H_0: p_1 = p_2 = p$$
 v.s.  $H_1: p_1 > p_2$ 

Where n(G) is the total population in the area G, n(Z) is the population within the area Z, c(G) is the number of attributes in the area G, and c(Z) is the number of the attributes within the area Z (Figure 3.7). Poisson model is used.



**Fig. 3.7** Population n and the number of the attribute c in area G. Comparing  $p_1 = c(Z)/n(Z)$  and  $p_2 = c(G)/n(G)$ .

The probability of the number of points is c(G) in the area G is

$$\frac{\exp[-p_1 n(Z) - p_2 n(\bar{Z})][p_1 n(Z) + p_2 n(\bar{Z})]^{c(G)}}{c(G)!}$$
(3.1)

The density at location x in the area G is

$$\frac{p_1 n(x)}{p_1 n(Z) + p_2 n(\overline{Z})} \quad \text{if} \quad x \in Z$$
 (3.2)

$$\frac{p_2 n(x)}{p_1 n(Z) + p_2 n(\overline{Z})} \quad \text{if} \quad x \in \overline{Z}$$
 (3.3)

The likelihood function of Poisson model is

$$L(Z, p_{1}, p_{2}) = \frac{\exp[-p_{1}n(Z) - p_{2}n(\overline{Z})][p_{1}n(Z) + p_{2}n(\overline{Z})]^{c(G)}}{c(G)!}$$

$$\times \prod_{x_{i} \in Z}^{n_{i} \in Z} \frac{p_{1}n(x)}{p_{1}n(Z) + p_{2}n(\overline{Z})} \prod_{x_{i} \in Z^{c}}^{n_{i} \in Z^{c}} \frac{p_{2}n(x)}{p_{1}n(Z) + p_{2}n(\overline{Z})}$$

$$= \frac{\exp[-p_{1}n(Z) - p_{2}n(\overline{Z})}{c(G)!} p_{1}^{c(Z)} p_{2}^{c(\overline{Z})} \prod_{x_{i}}^{n} n(x_{i})$$
(3.4)

Let x(Z) be a random variable for the number of attributes within the area Z. Anywhere under the area Z,

$$x(A) \sim Poisson(p_1 n(A \cap Z) + p_2 n(A \cap \overline{Z}))$$
 (3.5)

Under the null hypothesis,

$$x(A) \sim Poisson(pn(A))$$
 (3.6)

To maximize the likelihood function, the maximum likelihood function over the area Z is calculated. The maximum likelihood estimators  $\hat{p}_1 = c(Z)/n(Z)$  and  $\hat{p}_2 = c(\overline{Z})/n(\overline{Z})$  are estimated.

$$L(Z) = \frac{\exp[-c(G)]}{c(G)!} \left(\frac{c(Z)}{n(Z)}\right)^{c(Z)} \left(\frac{c(\overline{Z})}{n(\overline{Z})}\right)^{c(\overline{Z})} \prod_{x_i}^{n} n(x_i)$$
 (3.7)

The likelihood ratio  $\lambda(Z)$  is the maximum value in the subset area within the area G to detect hotspots.

$$\lambda(Z) = \frac{\max_{Z} L(Z)}{L_0} = \frac{\left(\frac{c(Z)}{n(Z)}\right)^{c(Z)} \left(\frac{c(\overline{Z})}{n(\overline{Z})}\right)^{c(\overline{Z})}}{\left(\frac{c(G)}{n(G)}\right)^{c(G)}}$$
(3.8)

 $L_0$  is the following likelihood function under the null hypothesis,

$$L_0 := \sup_{p} \frac{\exp[-pn(G)]}{c(G)!} p^{c(G)} \prod_{x_i}^{n} n(x_i) = \frac{\exp[-c(G)]}{c(G)!} (\frac{c(G)}{n(G)})^{c(G)} \prod_{x_i}^{n} n(x_i)$$
(3.9)

The test statistic  $\lambda(Z)$  can be

$$\lambda(Z) = \left(\frac{c(Z)}{e(Z)}\right)^{c(Z)} \left(\frac{c(\overline{Z})}{e(\overline{Z})}\right)^{c(\overline{Z})} \tag{3.10}$$

where e(Z) is the expectation of the attribute within the area Z, and e(G) is equal to c(G).

# 2.3 Hotspot detection procedure using Echelon dendrogram

The hotspot detection of Echelon analysis is as follow:

- 1. Draw the Echelon dendrogram for target data.
- 2. Scan the areas from upper Echelon to the bottom, based on the hierarchical structure determined in Step 1.
- 3. Detect the hotspot, which takes the maximum log likelihood ratio;  $\log \lambda(Z)$ .

The significance of the hotspot candidate is evaluated using Monte Carlo simulation.

A p-value based on Monte Carlo simulation can be obtained as follow:

- 1. Generate a random data set under the null hypothesis when we condition with conditions on the total number of attribute c(G).
- 2. Calculate the  $\log \lambda(Z)$  from the simulated data.
- 3. Repeat a process of Step 1 and 2 multiple times.

# 4. Define as p = R/(#SIM + 1)

where R is the rank of the test statistics from the real data set among all data sets and #SIM is the number of simulated data sets that have been generated.

Echelon analysis detects two hotspot candidates: most likely cluster and secondary cluster. The most likely cluster is the highest  $\log \lambda(Z)$ , and the secondary cluster  $\log \lambda(Z)$  is the second highest.

# 4. Patch identification method

# 4.1. Patch identification method

First, an area with significantly high suitability was detected based on the Echelon analysis hotspot detection method (Figure 4.1 and 4.2, Table 4.1) and salmon suitability index. The vertical line of Echelon dendrogram is the suitability index. If the number of areas is too large, we can set the maximum number of hotspot areas in advance. This maximum number is K. K is the maximum number of areas in most likely cluster. The secondary cluster can also be detected in the same manner. Here, K=500 (approximately 10% of the total) was adapted (Figure 4.3). Detected areas included not good suitability (<0.5). Therefore, we proposed another patch identifying method (Figure 4.4). The top ten patches of maximum suitability within each patch area were shown (Figure 4.5 and 4.6).

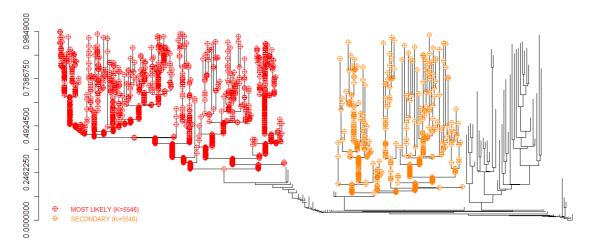


Fig. 4.1 Echelon dendrogram (laser) marked significantly high areas.

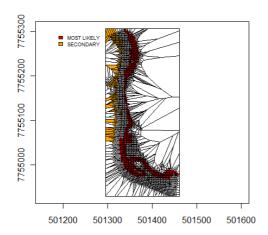
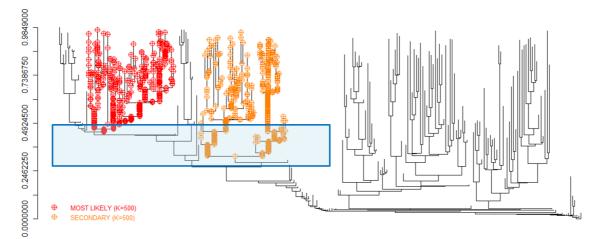


Table 4.1 Significantly high areas (laser).

|                    | Most likely | Secondary |
|--------------------|-------------|-----------|
| The number of area | 1707        | 489       |
| $\log \lambda$     | 383.07      | 19.49     |
| <i>p</i> -value    | 0.001       | 0.001     |

Fig. 4.2 Significantly high areas (laser).



**Fig. 4.3** Echelon dendrogram (laser) marked significantly high areas (*K*=500).

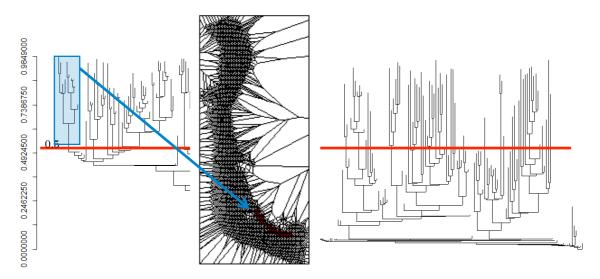


Fig. 4.4 Proposed patch identifying method.

# 4.2. Patch evaluation

Some identified patches are too small, therefore a patch size criterion or patch evaluation were needed. Eq. (3.10) can be useful as patch evaluation. The result is shown in Table 4.2 and 4.3. Obviously,  $\log \lambda$  of the too small patch was small, therefore these  $\log \lambda$  were utilized to evaluate patches.

**Table 4.2** The number of cell and  $log\lambda$  (laser).

| ID                 | 1     | 2     | 3      | 4    | 5     | 6    | 7    | 8     | 9    | 10   |
|--------------------|-------|-------|--------|------|-------|------|------|-------|------|------|
| The number of cell | 10    | 102   | 770    | 4    | 79    | 2    | 12   | 81    | 22   | 10   |
| logλ               | 42.61 | 22.92 | 208.08 | 0.78 | 17.51 | 0.51 | 2.91 | 13.22 | 5.58 | 1.92 |

**Table 4.3** The number of cell and  $log\lambda$  (sonar).

| ID                 | 1     | 2     | 3     | 4    | 5    | 6     | 7     | 8    | 9    | 10   |
|--------------------|-------|-------|-------|------|------|-------|-------|------|------|------|
| The number of cell | 158   | 80    | 61    | 2    | 9    | 267   | 140   | 24   | 2    | 19   |
| logλ               | 44.79 | 23.18 | 15.56 | 0.64 | 2.84 | 75.44 | 33.18 | 6.81 | 1.02 | 6.13 |

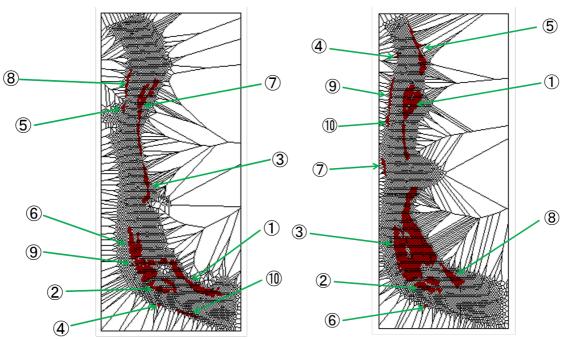
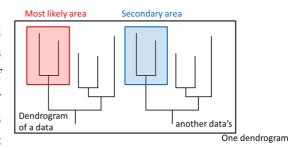


Fig. 4.5 Identified ten patches (laser).

Fig. 4.6 Identified ten patches (sonar).

# 5. Comparing two kind of data: laser and sonar

logλ are indexes evaluated only in same area. Laser data and sonar cannot be compared using logλ. An advantage of Echelon analysis is easy scan. We can compare two kinds of data by making a dendrogram from both laser and sonar. A conventional method is detecting significance area in each area, it cannot compare some areas. New method can assess same area as one area and one dendrogram (Figure 5.1). The hypothesis is as follows:

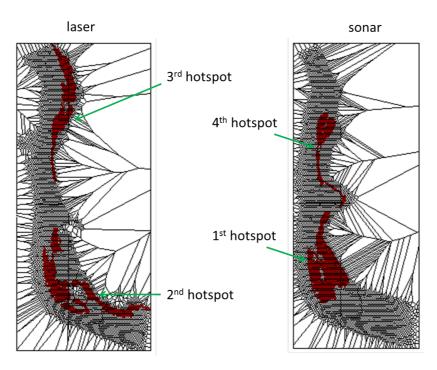


**Fig. 5.1** The Echelon dendrogram with two kind of data method.

H<sub>0</sub>: All area's suitability indexes are equal.

 $H_1$ : not  $H_0$ 

The result is Figure 5.2. The first and forth hotspot were sonar data, second and forth were laser data. This result might have been related that the laser measurement accuracy was higher than the sonar.



**Fig. 5.2** Detected hotpot areas (*p*-value<0.5).

# 6. Discussion

In this study, we confirmed that the method of patch identification was effective in salmon habitat suitability. The identification of patches through Echelon dendrogram and salmon suitability index was more useful than detecting significant areas from spatial scan statistic. It is possible to show the best habitat for salmon by evaluating patches. However, in previous finding (Vadas and Orth, 2001), suitability index  $\geq 0.75$  were defined as optimal. We should further examine the critical suitability index. We also presented comparison method with two data (sonar and laser) using Echelon analysis. We want to establish the comparing method because it has much room to study.

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# Water quality affects diatom diversity in Finnish rivers

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# Abstract:

Land use has a great impact on water quality, which can affect diatom diversity and ecology. We assessed statistically these relationships using a spatial method. As the results, using classification and regression trees, we demonstrated low pH having a great impact on diatom diversity.

# 1. Introduction

Technical development has made a study based on spatial information much easier. This type of study technique is utilized in various disciplines such as Population Biology (Bellier et al., 2013) and Epidemiology (Takahashi et al., 2008). A study about water quality also uses spatial information. LaBeau et al. (2014) presented relationship between land use and phosphorus (P) and estimated the amount of P loading.

We studied the impact of water quality on diatoms using land use data. We found the spatial relationship between water quality and diatom utilizing Echelon analysis. Then, we assessed the impact of water quality on diatoms using classification and regression trees (CART).

In this paper, we explained the survey area, data and analysis methods followed by the results of Echelon analysis and CART. We also identified the substances that impact on diatoms.

# 2. Materials and methods

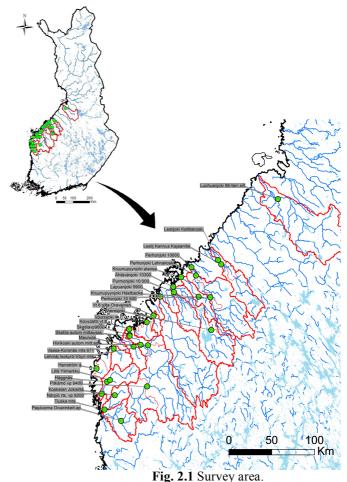
Figure 2.2 shows our study procedure. We analyzed land use data, water quality data and diatoms data utilizing Echelon analysis and CART.

# 2.1. Study area

study area was mid-western part of Finland (Figure 2.1). The water quality data were collected from the 13 catchment areas (rivers) with land use data and had altogether 29 sampling sites.

# 2.2. Land use data

Land use data was measured in 2006. Land use of each catchment area was classified



in 43 types, and they were grouped into ten classes based on soil characteristics (Table 2.1) because of too detailed to estimate a missing value.

Table 2.1 Ten classes of land use grouped based on soil.

| Class of land use | Original land use  |
|-------------------|--|
| Artificial area   | Density constructed urban areas, sparsely constructed urban areas, industrial and service areas, traffic areas, harbor areas, airport areas, area            |
|                   | for land taking, disposal sites, under construction, parks and green areas   |
|                   | in urban areas, summer cottages, sport areas   |
| Agricultural area | Fields, fruit tree and berry production areas, stock feeding areas, small agricultural mosaic  |
| Mineral soil area | Deciduous forests on mineral soil, evergreen forests on mineral soil, mixed forest on mineral soil, rare tree areas, cc <10-30% on mineral ground            |
| Peat area         | Deciduous forests of peat land, evergreen forests on peat land, mixed forest on peat land, rare tree areas, cc <10-30% on peat ground, peat production areas |
| Rocks area        | Evergreen forests on rocks, mixed forest on rocks, rare tree areas, cc <10-30% on rock ground, rock ground   |

| Wet area | Moor land with rare growing, inland wetlands on the land, inland          |
|----------|---|
|          | wetlands in the water, open swamps, wetlands on the land area close to    |
|          | the sea, wetlands in the water close to the sea                           |
| Others   | Natural grasslands, thin tree groups and moors, rare tree areas, cc <10%, |
|          | rare tree areas over boreal forest, rare tree areas, cc <10-30% on below  |
|          | electricity line, sandy beaches   |
| River    | River   |
| Lake     | Lake  |
| Sea      | Sea   |

# 2.3. Water quality data

Water quality was surveyed from 2006 to 2012, and the number of sampling sites was 29. However, none of the sites had complete data (i.e. without missing variables). The Table 2.2 shows the percentages of missing variables pertaining to the water quality sampling sites.

**Table 2.2** The percentages of missing variables.

| Substance              |      | Alkalinity | Al                   | Cd  | COD    | Particu<br>matt |      | Total S | Cr    |      |
|------------------------|------|------------|----------------------|-----|--------|-----------------|------|---------|-------|------|
| Missing value rate (%) |      | 51.3       | 76.9                 | 79. | 9 59.0 | 70.             | 7    | 100     | 80.2  |      |
|                        |      |            |                      |     |        |                 |      |         |       |      |
| Cu                     | Pb   | Ni         | Total Orga<br>Carbon |     | Fe     | Turbidity       | Zn   | SO4     | Color | рН   |
| 80.2                   | 80.2 | 79.9       | 92.3                 |     | 94.1   | 63.0            | 80.2 | 95.6    | 51.3  | 49.1 |

# 2.4. Diatom data

Diatoms state was evaluated using two methods: number of type-specific species ( $TT_{40}$ ) and percent model affinity (PMA). Both  $TT_{40}$  and PMA are between 0 and 1: close to 0 means the bad state, close to 1 means the best.

 $TT_{40}$  (Aroviita et al., 2008) is a taxon type specific index, which tells about how many taxa are found in the studied site compared to the reference site. The higher the index value, the higher the number of the same taxa in the sites. A low index value shows that there is a difference in the diatom community at the study site compared to a reference community.

PMA (Novak and Bode, 1992) describes the similarity of the reference and studied diatom community. If the index value is high, the communities in the studied and reference environment area are the same. If the value is low, the community in the studied site differs from the one in the reference site.

These data were surveyed in summer from 2007 to 2012, and the number of survey areas

varied from year to year. 2010 was the highest number of survey with 17 areas, and the lowest was 2008 with no survey taken place.

# 2.5. Models for land use data and water quality

Water quality data had multiple missing values; therefore, we made the complete data using two kinds of method; near sampling site data and estimating generalized linear model (GLM). We preferably used the sampling site data from the same catchment area, near sites and the same river. Nevertheless, if data had a missing value, we estimated them using GLM according to equation 2.1.

$$Y_j = \beta_0 + \sum_i \beta_i X_i + \varepsilon \tag{2.1}$$

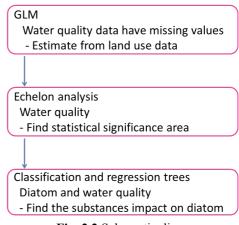


Fig. 2.2 Schematic diagram.

where  $Y_i$  is a variable like alkalinity and  $X_i$  are year, season and each land area like artificial area or agriculture area of Table 2.1.

However, the estimation the substance like metals was not successful due to too many missing values. Therefore, we estimated the missing values of alkalinity, COD, turbidity and pH.

# 2.6. Spatial model for water quality

Establishing an Echelon dendrogram makes it easy to detect statistically significant areas. For additional information, see a section 3 of "Juvenile salmon patch identification and comparison using Echelon analysis".

We detected hotspot areas from 3D data based on time series data, therefore time (season and year) and areas information were included in significantly high areas. Statements of significance are based on  $p \le 0.05$ .

# 2.7. Models for water quality and diatoms

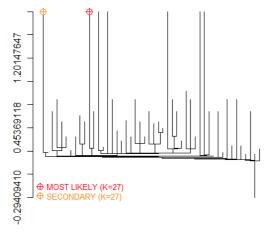
Classification and regression trees (CART) were obtained by recursively partitioning the data space and fitting a simple prediction model within each partition. As a result, the partitioning can be represented graphically as a decision tree (Loh, 2011).

A variable was divided based on impact on the response variable. In this case, response variables were the diatom indexes and explanatory variables were the water quality data.

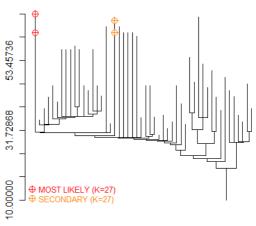
# 3. Results

# 3.1 Hotspot areas of alkalinity, COD, pH and turbidity

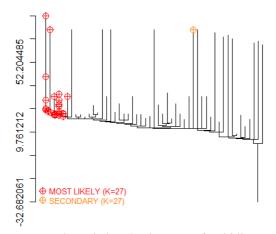
We detected hotspot areas of alkalinity, COD, pH and turbidity using an Echelon analysis. Each Echelon dendrogram is shown in Figures 3.1, 3.2, 3.3 and 3.4, and Tables 3.1, 3.2, 3.3 and 3.4 explain the hotspot areas. The red symbols represent the most likely significance areas and the orange symbols represent the secondary likely areas in each figure. The p-values of all the most likely hotspot areas and secondary areas were less than 0.5; therefore, the detected areas were the statistically significant areas. K in figures is the maximum number of areas in the most likely cluster. The secondary cluster can also be detected in the same manner. Here, K=27 (approximately 10% of the total) was adapted. However, the number of diatom samples was 47.



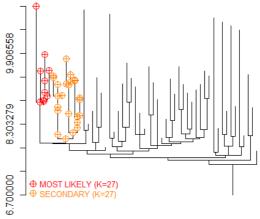
**Fig. 3.1** The Echelon dendrogram of alkalinity (2006-2012).



**Fig. 3.2** The Echelon dendrogram of COD (2006-2012).



**Fig. 3.3** The Echelon dendrogram of turbidity (2006-2012).



**Fig. 3.4** The Echelon dendrogram of low pH (2006-2012).

**Table 3.1** The Hotspot areas of alkalinity (2006-2012).

|           | <i>p</i> -value | season | year | sampling site |
|-----------|-----------------|--------|------|---------------|
| hotspot   | 0.001           | summer | 2006 | Purmoj.10k    |
| secondary | 0.001           | summer | 2012 | Purmoj.10k    |

**Table 3.2** The Hotspot areas of COD (2006-2012).

|           | <i>p</i> -value | season         | year         | sampling site            |
|-----------|-----------------|----------------|--------------|--------------------------|
| hotspot   | 0.001           | summer<br>fall | 2012<br>2012 | Purmoj.10k<br>Purmoj.10k |
| secondary | 0.001           | summer<br>fall | 2011<br>2011 | Purmoj.10k<br>Purmoj.10k |

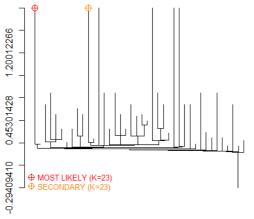
**Table 3.3** The Hotspot areas of turbidity (2006-2012).

|           | <i>p</i> -value | season | year | sampling site  |
|-----------|-----------------|--------|------|----------------|
|           |                 | spring | 2006 | Purmoj.10k     |
|           |                 | spring | 2006 | Lapua 9900     |
|           |                 | spring | 2006 | Vt8Oravainen   |
|           |                 | spring | 2006 | Koivulahti vt8 |
|           |                 | spring | 2006 | Maunula        |
|           |                 | spring | 2006 | VaasaKorsnäs   |
|           |                 | spring | 2006 | Harrström a    |
|           |                 | spring | 2006 | Närpiö9200     |
|           |                 | spring | 2006 | Tiukka         |
|           |                 | summer | 2006 | Purmoj.10k     |
|           |                 | summer | 2006 | Vt8Oravainen   |
|           |                 | summer | 2006 | Harrström a    |
| hotspot   | 0.001           | fall   | 2006 | Lapua 9900     |
|           |                 | fall   | 2006 | Vt8Oravainen   |
|           |                 | fall   | 2006 | Koivulahti vt8 |
|           |                 | fall   | 2006 | VaasaKorsnäs   |
|           |                 | fall   | 2006 | Närpiö9200     |
|           |                 | spring | 2007 | Vt8Oravainen   |
|           |                 | spring | 2007 | Maunula        |
|           |                 | spring | 2007 | Harrström a    |
|           |                 | spring | 2007 | Närpiö9200     |
|           |                 | spring | 2007 | Tiukka         |
|           |                 | summer | 2007 | Vt8Oravainen   |
|           |                 | summer | 2007 | Harrström a    |
|           |                 | fall   | 2007 | Vt8Oravainen   |
| secondary | 0.001           | summer | 2012 | Purmoj.10k     |

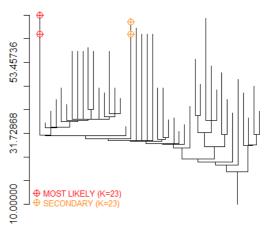
**Table 3.4** The Hotspot areas of low pH (2006-2012).

|           |                 | •      |      | ` /               |
|-----------|-----------------|--------|------|-------------------|
|           | <i>p</i> -value | season | year | sampling site     |
|           |                 | fall   | 2006 | Lestj.Kattilakosk |
|           |                 | fall   | 2006 | Perhonj10600      |
|           |                 | fall   | 2006 | Kruununpyyn. ala  |
|           |                 | fall   | 2006 | Ähtävänj10300     |
|           |                 | fall   | 2006 | Purmoj.10k        |
| hotspot   | 0.001           | fall   | 2006 | Lapua 9900        |
|           |                 | fall   | 2006 | Vt8Oravainen      |
|           |                 | spring | 2007 | Kruununpyyn. ala  |
|           |                 | spring | 2007 | Purmoj.10k        |
|           |                 | spring | 2007 | Lapua 9900        |
|           |                 | spring | 2007 | Vt8Oravainen      |
|           |                 | fall   | 2006 | Maunula           |
|           |                 | fall   | 2006 | VaasaKorsnäs      |
|           |                 | fall   | 2006 | Harrström a       |
|           | 0.001           | fall   | 2006 | Närpiö9200        |
|           |                 | fall   | 2006 | Tiukka            |
|           |                 | spring | 2007 | Maunula           |
|           |                 | spring | 2007 | Harrström a       |
|           |                 | spring | 2007 | Närpiö9200        |
|           |                 | summer | 2007 | Närpiö9200        |
|           |                 | fall   | 2007 | Purmoj.10k        |
|           |                 | fall   | 2007 | Lapua 9900        |
| secondary |                 | fall   | 2007 | Vt8Oravainen      |
|           |                 | fall   | 2007 | Maunula           |
|           |                 | fall   | 2007 | VaasaKorsnäs      |
|           |                 | fall   | 2007 | Harrström a       |
|           |                 | fall   | 2007 | Närpiö9200        |
|           |                 | fall   | 2007 | Tiukka            |
|           |                 | spring | 2008 | Purmoj.10k        |
|           |                 | spring | 2008 | Lapua 9900        |
|           |                 | spring | 2008 | Vt8Oravainen      |
|           |                 | spring | 2008 | Koivulahti vt8    |
|           |                 | spring | 2008 | Maunula           |
|           |                 | spring | 2008 | Harrström a       |
|           |                 | spring | 2008 | Närpiö9200        |

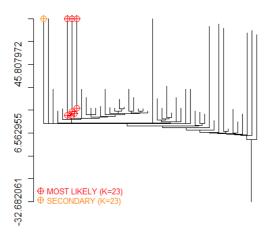
Many hotspot areas included the year 2006, however diatoms were not surveyed in this year. Therefore, we tried Echelon analysis again excluding year 2006. Each Echelon dendrogram is shown in Figures 3.5, 3.6, 3.7 and 3.8, and Tables 3.5, 3.6, 3.7 and 3.8 explain the hotspot areas.



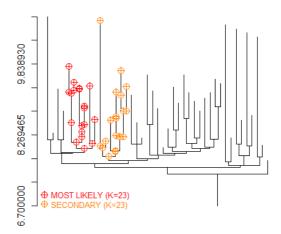
**Fig. 3.5** The Echelon dendrogram of alkalinity (2007-2012).



**Fig. 3.6** The Echelon dendrogram of COD (2007-2012).



**Fig. 3.7** The Echelon dendrogram of turbidity (2007-2012).



**Fig. 3.8** The Echelon dendrogram of low pH (2007-2012).

**Table 3.5** The Hotspot areas of alkalinity (2007-2012).

|           | <i>p</i> -value | season | year | sampling site |
|-----------|-----------------|--------|------|---------------|
| hotspot   | 0.001           | summer | 2012 | Purmoj.10k    |
| secondary | 0.001           | summer | 2011 | Purmoj.10k    |

**Table 3.6** The Hotspot areas of COD (2007-2012).

|                   |                 | _      |      |               |
|-------------------|-----------------|--------|------|---------------|
|                   | <i>p</i> -value | season | year | sampling site |
| hatanat           | 0.001           | summer | 2012 | Purmoj.10k    |
| hotspot           | 0.001           | fall   | 2012 | Purmoj.10k    |
| a a a a m d a m v | 0.001           | summer | 2011 | Purmoj.10k    |
| secondary         | 0.001           | fall   | 2011 | Purmoj. 10k   |

**Table 3.7** The Hotspot areas of turbidity (2007-2012).

|           | <i>p</i> -value | season | year | sampling site |
|-----------|-----------------|--------|------|---------------|
|           |                 | spring | 2007 | Purmoj.10k    |
|           |                 | summer | 2007 | Purmoj.10k    |
|           |                 | fall   | 2007 | Purmoj.10k    |
| hotanot   | 0.001           | spring | 2008 | Purmoj.10k    |
| hotspot   | 0.001           | summer | 2008 | Purmoj.10k    |
|           |                 | fall   | 2008 | Purmoj.10k    |
|           |                 | spring | 2009 | Purmoj.10k    |
|           |                 | summer | 2009 | Purmoj.10k    |
| secondary | 0.001           | summer | 2012 | Purmoj.10k    |

**Table 3.8** The Hotspot areas of low pH (2007-2012).

|           |                 |        | 1    | ,                 |
|-----------|-----------------|--------|------|-------------------|
|           | <i>p</i> -value | season | year | sampling site     |
|           |                 | spring | 2007 | Maunula           |
|           |                 | spring | 2007 | Harrström a       |
|           |                 | spring | 2007 | Närpiö9200        |
|           |                 | summer | 2007 | Närpiö9200        |
|           |                 | fall   | 2007 | Purmoj.10k        |
|           |                 | fall   | 2007 | Lapua 9900        |
|           |                 | fall   | 2007 | Vt8Oravainen      |
|           |                 | fall   | 2007 | Maunula           |
|           |                 | fall   | 2007 | VaasaKorsnäs      |
| hotspot   | 0.001           | fall   | 2007 | Harrström a       |
| •         |                 | fall   | 2007 | Närpiö9200        |
|           |                 | fall   | 2007 | Tiukka            |
|           |                 | spring | 2008 | Purmoj.10k        |
|           |                 | spring | 2008 | Lapua 9900        |
|           |                 | spring | 2008 | Vt8Oravainen      |
|           |                 | spring | 2008 | Koivulahti vt8    |
|           |                 | spring | 2008 | Maunula           |
|           |                 | spring | 2008 | Harrström a       |
|           |                 | spring | 2008 | Närpiö9200        |
|           |                 | fall   | 2008 | Lestj.Kattilakosk |
|           |                 | fall   | 2008 | Perhonj10600      |
|           |                 | fall   | 2008 | Purmoj.10k        |
|           |                 | fall   | 2008 | Lapua 9900        |
|           |                 | fall   | 2008 | Vt8Oravainen      |
|           |                 | fall   | 2008 | Koivulahti vt8    |
|           |                 | fall   | 2008 | Maunula           |
|           |                 | fall   | 2008 | VaasaKorsnäs      |
| secondary | 0.001           | fall   | 2008 | Harrström a       |
| ·         |                 | fall   | 2008 | Närpiö9200        |
|           |                 | fall   | 2008 | Tiukka            |
|           |                 | spring | 2009 | Perhonj10600      |
|           |                 | spring | 2009 | Kruununpyyn. ala  |
|           |                 | spring | 2009 | Ähtävänj10300     |
|           |                 | spring | 2009 | Purmoj.10k        |
|           |                 | spring | 2009 | Lapua 9900        |
|           |                 | spring | 2009 | Vt8Oravainen      |

| spring | g 2009  | Koivulahti vt8   |
|--------|---------|------------------|
| spring | g 2009  | Maunula          |
| spring | g 2009  | Harrström a      |
| spring | g 2009  | Närpiö9200       |
| summ   | er 2009 | Kruununpyyn. ala |

# 3.2 Water quality impact on diatom diversity

We assessed the impact of water quality on diatom diversity utilizing CART (Figure 3.9 and 3.10). We got different results on diatom indexes. In case of  $TT_{40}$ , the important variables were turbidity, alkalinity and pH. On the other hand, the important substances for PMA were pH and COD.

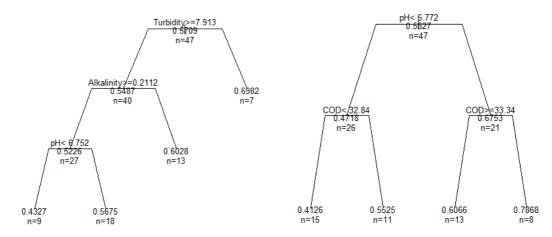


Fig. 3.9 Response variable: TT<sub>40</sub>.

Fig. 3.10 Response variable: PMA.

# 4. Conclusions

In this study utilizing CART, it was found that the diatom diversity was impacted by multiple variables. On the other hand, we detected statistically significant areas of these variables using Echelon analysis. Water alkalinity and COD did not have a significant effect on the diatom indices. Instead, pH and turbidity had an effect on some sampling sites. However these areas did not match diatom survey area because of the difference of survey time. If the diatom had been surveyed at the same time and the same site of detected hotspot areas, we might have obtained more reliable results. However, we can also assume that the circumstances have not changed between the years. Further studies are needed to explore the role of the catchment area use on the diatom responses.

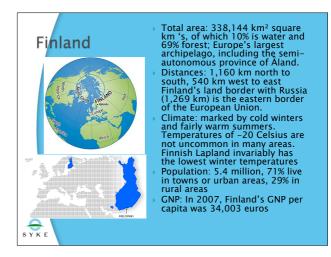
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WP 5

Dissemination of the project results







# Why water quality models New legislation in late 1960's forced point loaders (like industry and municipalities) to apply permission to lead purified waste waters to recipient waters (lakes, rivers, coastal areas etc..) from the Water Courts and During the Assesment Procedure ("Katselmustoimitus") the effects on the water quality was assessed and also the compensations were ordered Models were applied for Determining the optimal loads Determining the optimal loads Determining the optimal locations for water water outlets Model applications from 1970's by Virtanen, Koponen and Sarkkula

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# Later needs for modelling

- Problems with eutrophication and related diffuse loads increased the need of WQ assessment and modelling
- Effects of global change need to be assessed
- European wide approach: Water Framework Directive

S Y KETE15

- Fate and transport of harmfull substances (heavy metals, organic compounds etc..) is to be predicted since many recipient waters are also used for water supply
- Hydrological (water resources) modelling was fist needed for optimal regulation of our large lakes.
   Modelling started in early 1980's by Vehviläinen



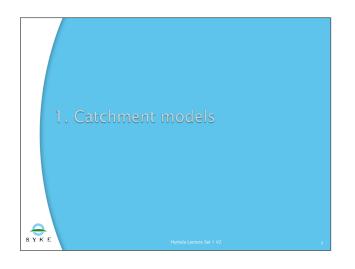
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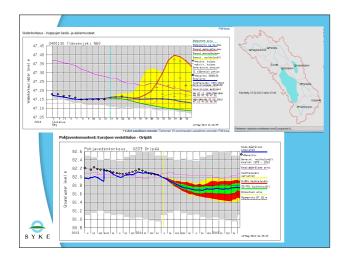
# Presently many new needs

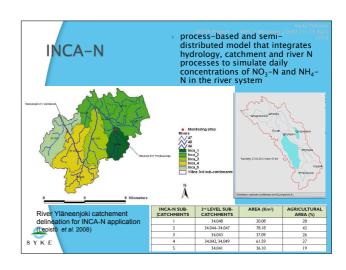
- Oil spill models
  - Operative models for oil combatting
- Habitat modelling for river restoration
- Socio-economical modelling
  - See CONPAT-project

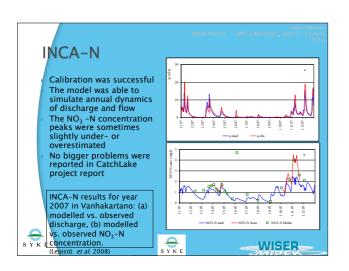
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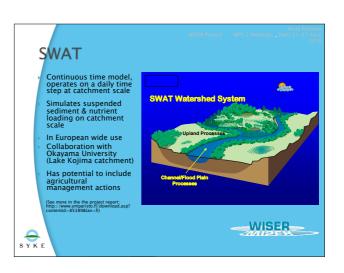


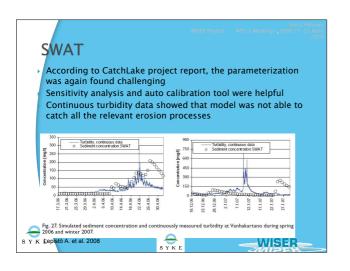
# Hydrological simulation and forecast A country wide system (WSFS): http://www.ymparisto.fl/en-US/Waters and sea/ Hydrological-situation and forecasts/ Hydrological-situation and forecasts/ Hydrological-forecasts and maps/Hydrological forecasts and maps \$2826174829 Used for operational forecasts Used also for research Based on water balance computation on watersheds (3rd division level) Dynamic links Meteorological data forecasts (FMI) Hydrological observations (water levels, river discharges, water temperature, ground water level) Links to satellite automated imaging products (snow coverage, snow water equivalent) Automatic and tailored products











Lake mass balance  $\frac{dm}{dt} = I - O - S$ Where, m= total amount of phosphorus in a lake, I=total load of phosphorus entering to the lake, O=amount of phosphorus leaving the lake, S=retention (the amount of phosphorus (P) sediment to the lake) First applications by Piontelli and Tonolli (1969) In Vollenweiders (1969) solution the lake was considered as a continuously stirred tank (CSTR) and sedimentation of P was taken as following the first order kinetics with settling coefficient  $\sigma$  and phosphorus concentration (c) at the outlet was taken as mean concentration in the lake

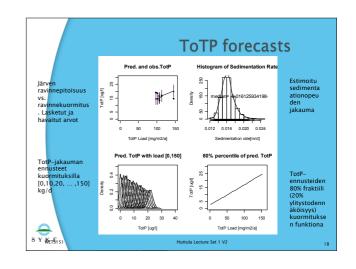
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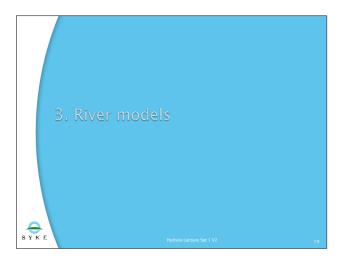
S Y KETE15

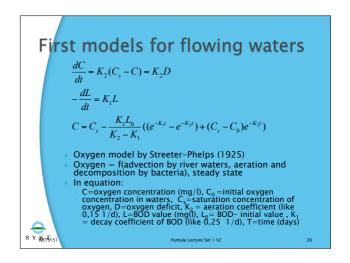
Vollenweiders equation  $\frac{dm}{dt} = I - Qc - \sigma m$  $\rho = \frac{Q}{V} \Rightarrow \frac{dc}{dt} = \frac{I}{V} - \rho c - \sigma c$ Where, Q= water discharge at the outlet and  $\rho=$ exchange coefficient,  $\sigma$ =settling coefficient It is expected here that the lake volume is constant (eq. an annual mean) and introduce In steady state we mark phosphorus concentration as  $c_{ss}$  it can be solved for time t S Y KETS15

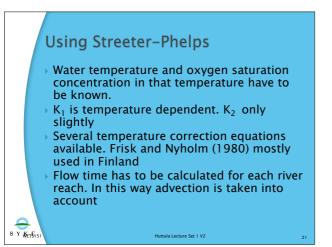
Huttula Lecture Set 1 V2

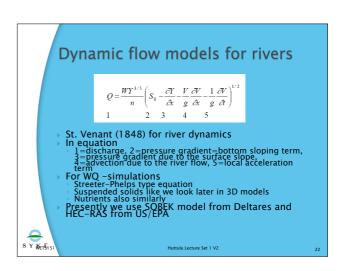
Solution for Vollenweider-equation  $c = c_{ss} - (c_{ss} - c_0)e^{-(\rho + \sigma)(t - t_0)}$ Where c<sub>0</sub> is phosphorus concentration at time Comments Settling description with first order kinetics is too Model can be used for studying the effects of loading options S Y KETE15

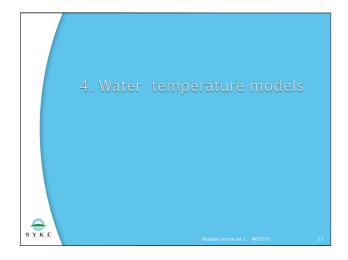


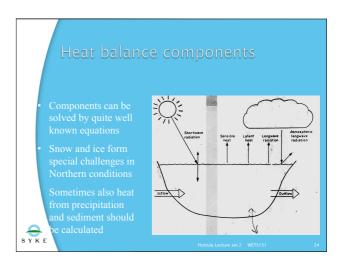


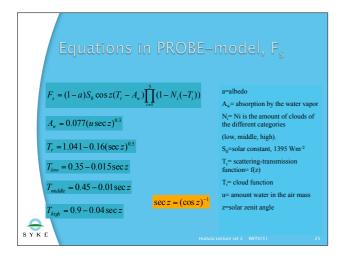


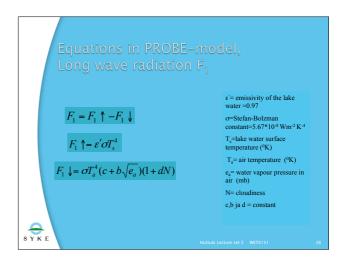


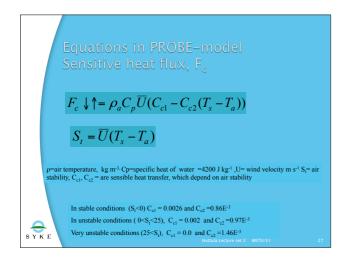


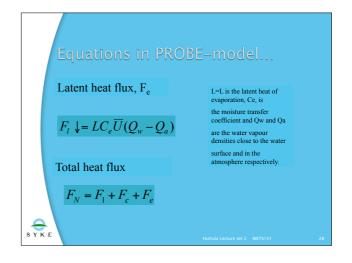


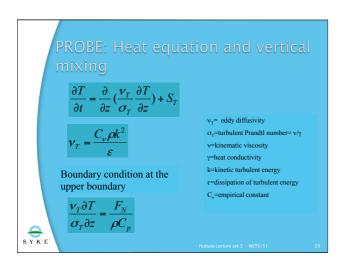


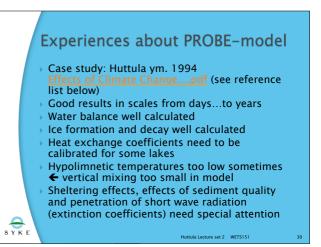






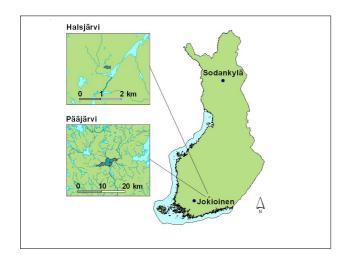


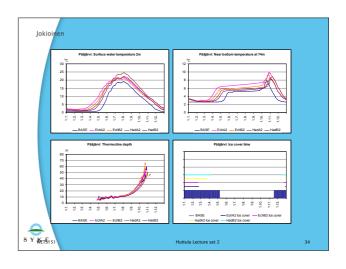


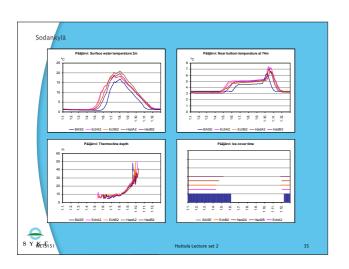


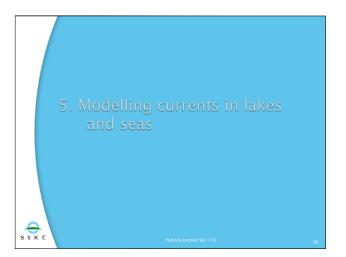
# MyLake • One dimensional vertical lake model • Andersson and Saloranta 2000 • Ice model by Leppäranta (1991) and Saloranta (2000) • The vertical diffusion coefficient from the stability frequency N² (Hondzo and Stefan, 1993) • Utilises the MATLAB Air-Sea Toolbox (http://sea-mat.whol.edu/air\_sea-ntml/) for calculation of radiative and turbulent heat fluxes, surface wind stress and astronomical variables • Vertical mixing is based on the energy calculation between kinetic energy from wind and potential energy of layer(s) to be mixed • Model package with documentation available from TH

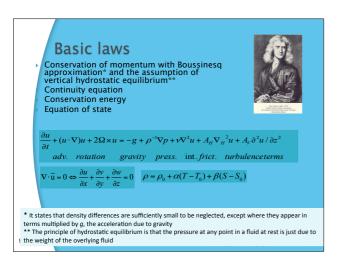
# FINESSI-project Web tool for assessing the effects of global chage in Finland <a href="https://www.finessi.info/finessi">www.finessi.info/finessi</a> Lake Pääjärvi (area= 13.5 km², max depth= 87 m) and Halsjärvi (area= 0.5 km², max depth= 6 m) Meteorological data from Jokioinen Calibration for Pääjärvi Link FINESSI uses 6 GCM-models Link We Echam and Hadley Center

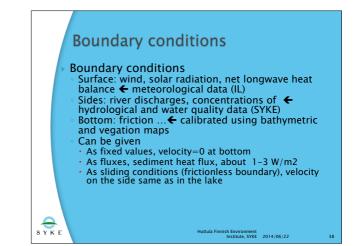


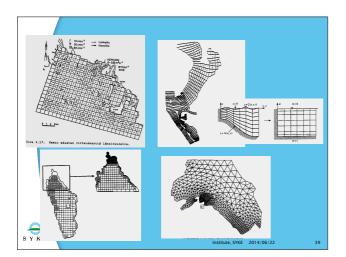


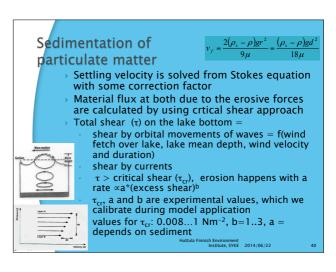




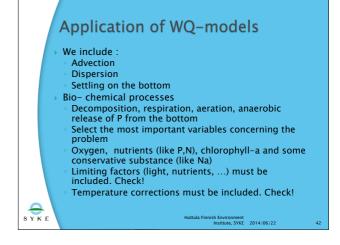


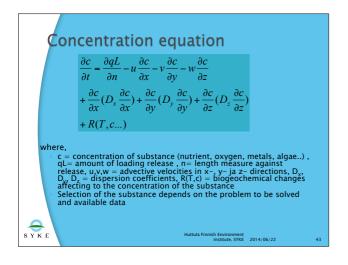


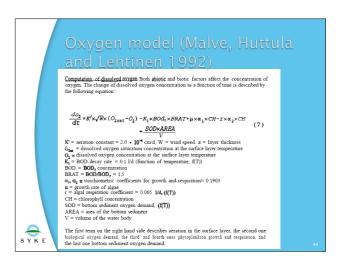


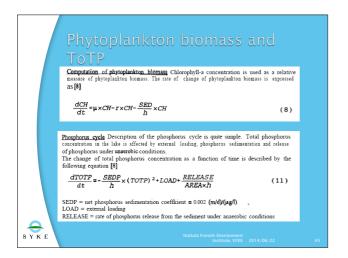


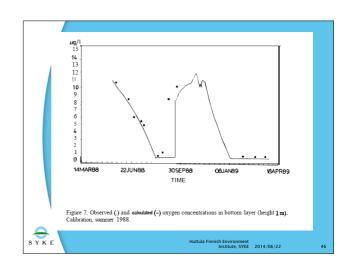
6. Modelling water quality or hydrochemical and hydrobiological modelling

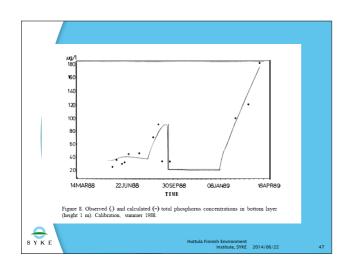


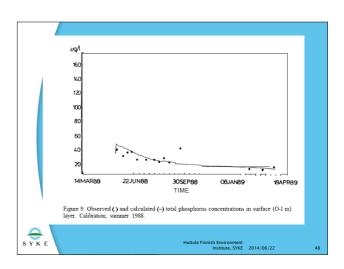


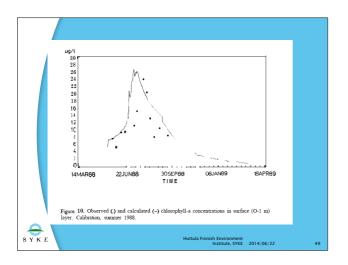


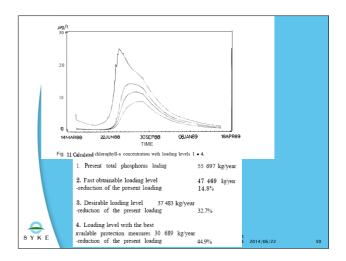


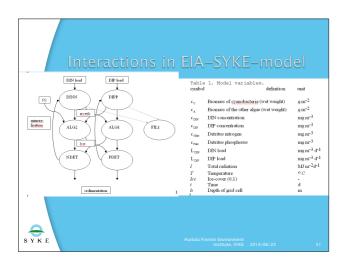








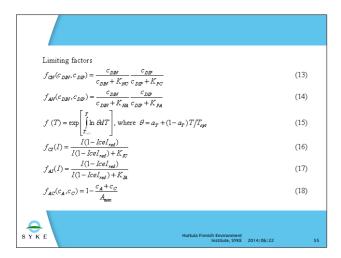




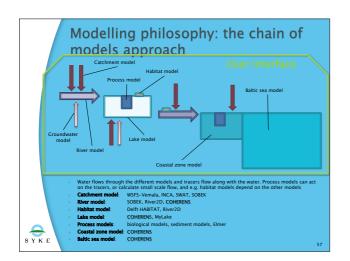
| Table<br>Symbol  | 2. Model parameters<br>definition                            | reference                  | value   | unit                               |
|------------------|--|----------------------------|---------|------------------------------------|
| -,               |  |                            |         |                                    |
| $N_{inC}$        | Nitrogen in cyanobacteria                                    | Redfield, 1958             | 0.0193  |                                    |
| $P_{i\pi C}$     | Phosphorus in cyanobacteria                                  | Redfield, 1958             | 0.00268 |                                    |
| $N_{isA}$        | Nitrogen in the other algae                                  | Redfield, 1958             | 0.0193  | -                                  |
| $P_{inA}$        | Phosphorus in the other algae                                | Redfield, 1958             | 0.00268 |                                    |
| $\mu_{c_{\max}}$ | Maximal growth rate of cyanobacteria                         | calibration                | 0.5     | d-1                                |
| $\mu_{Ano}$      | Maximal growth rate of the other algae                       | Olli et al., 1996          | 0.7     | d-1                                |
| R case           | Maximum loss rate of cyanobacteria                           | calibration                | 0.1     | d-1                                |
| R same           | Maximum loss rate of the other algae                         | calibration                | 0.15    | d-1                                |
| K <sub>NC</sub>  | Half-saturation coefficient of DIN for cyanobacteria         | Tyrrell, 1999              | 0       | mg m-3                             |
| Kpc              | Half-saturation coefficient of DIP for cyanobacteria         | Kononen & Leppanen, 1997   | 2       | mg m-3                             |
| K <sub>NA</sub>  | Half-saturation coefficient of DIN for the other algae       | calibration                | 7       | mg m-3                             |
| KpA              | Half-saturation coefficient of DIP for the other algae       | calibration                | 1       | mg m-3                             |
| K <sub>IC</sub>  | Half saturation coefficient of radiation for cyanobacteria   | calibration                | 20      | MJ m <sup>-2</sup> d <sup>-1</sup> |
| K <sub>IA</sub>  | Half saturation coefficient of radiation for the other algae | calibration                | 15      | MJ m <sup>-2</sup> d <sup>-1</sup> |
| $C_{\min}$       | Minimum biomass of cyanobacteria                             | calibration                | 0.5     | g m <sup>-2</sup>                  |
| $A_{nin}$        | Minimum biomass of the other algae                           | calibration                | 0.01    | g m <sup>-2</sup>                  |
| $A_{\rm nac}$    | Maximum total biomass of algae                               | calibration                | 300     | g m <sup>-2</sup>                  |
| $\beta_0$        | Maximal detritus nitrogen mineralisation rate                | Garber, 1984               | 0.018   | d <sup>-1</sup>                    |
| $\gamma_0$       | Maximal detritus phosphorus mineralisation rate              | Garber, 1984               | 0.043   | d-1                                |
| $\nu_{N  m det}$ | Settling rate of detritus nitrogen                           | Heiskanen & Tallberg, 1999 | 1       | m d <sup>-1</sup>                  |

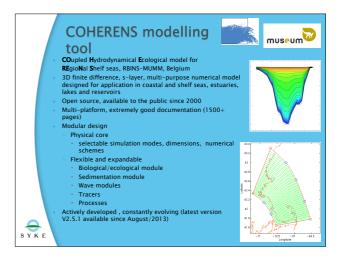
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Settling rate of detritus nitrogen
                                                                                                       Heiskanen & Tallberg, 1999 1
                                                                                                                                                                   m d-1
\nu_{N 
m det}
                 Settling rate of detritus phosphorus
                                                                                                       Heiskanen & Tallberg, 1999 1
\nu_{\it Pdet}
                 Sedimentation rate of detritus nitrogen
                                                                                                                                                                   m d-1
\mathbb{S}_{\mathrm{Hdet}}
                 Sedimentation rate of detritus phosphorus
                                                                                                       Lehtoranta, 1998
                                                                                                                                                   0.00
                                                                                                                                                                   m d<sup>-1</sup>
                Optimal temperature
                                                                                                                                                                   °C
°C
°C
                 for the growth of cyanobacteria
                                                                                                       Kononen & Leppänen, 1997 25
                for the growth of cyanobactena
for the growth of the other algae
for losses
for destitus nitrogen mineralisation
for destitus phosphorus mineralisation
Coefficient for temperature limiting factor
                                                                                                        calibration
                                                                                                       calibration
Garber, 1984
Garber, 1984
                                                                                                                                                    25
18
18
                 for the growth of cyanobacteria
                                                                                                        calibration
                                                                                                                                                    1.14
                  for the growth of the other alga-
for losses
                                                                                                                                                    1.001
1.05
1.31
1.60
0.5
                 for detritus nitrogen mineralisation
for detritus phosphorus mineralisation
Radiation attenuation by ice
                                                                                                        Garber, 1984
Garber, 1984
                Depth of mixing layer
                                                                                                        calibration
                                                                                                                                                    20
                                                                                          Huttula Finnish Environment
Institute, SYKE 2014/06/22
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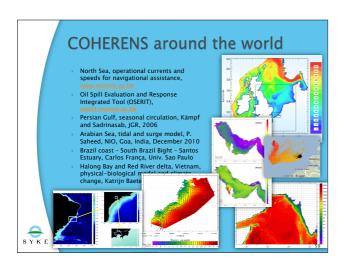
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Table 3. Model equations, rates and limiting factors.
Equations
  \frac{\partial c_c}{\partial t} = (\mu_c - \mathbb{R}_c) c_c
                                                                                                                                                                                                                                             (1)
  \frac{\partial c_A}{\partial t} = (\mu_A - R_A)c_A
                                                                                                                                                                                                                                             (2)
  \frac{\partial \mathcal{L}_{\mathit{DBN}}}{\partial t} = \beta \; c_{\mathit{NDet}} - \mu_{\mathit{A}} N_{\mathit{insF}} \,_{\mathit{A}} h_{\mathit{mix}}^{-1} - \mu_{\mathit{C}} N_{\mathit{insC}} c_{\mathit{C}} h_{\mathit{mix}}^{-1} + L_{\mathit{DBN}}
  \frac{\partial \mathcal{L}_{DB}}{\partial t} = \gamma \ c_{Pdat} - \mu_A P_{ind} c_A h_{mix}^{-1} - \mu_C P_{inC} c_C h_{mix}^{-1} + L_{DB}
                                                                                                                                                                                                                                             (4)
  \frac{\partial \mathcal{C}_{N \, \mathrm{det}}}{\Delta} = N_{\mathrm{ind.}} \ R_A c_A h_{\mathrm{mir}}^{-1} + N_{\mathrm{inc.}} \ R_C c_C h_{\mathrm{mir}}^{-1} - \beta c_{N \, \mathrm{det.}} - \nu_{\mathrm{N \, det.}} c_{N \, \mathrm{det.}} h^{-1} - S_{N \, \mathrm{det.}} c_{N \, \mathrm{det.}} h^{-1}
                                                                                                                                                                                                                                             (5)
  \frac{\partial \mathcal{E}_{\mathrm{Pdet}}}{\partial t} = P_{\mathrm{IMA}} \ R_{\mathrm{A}} \mathcal{C}_{\mathrm{A}} h_{\mathrm{min}}^{-1} + P_{\mathrm{InC}} \ R_{\mathrm{C}} \mathcal{C}_{\mathrm{C}} h_{\mathrm{min}}^{-1} - \chi_{\mathrm{Pdet}} - \nu_{\mathrm{Pdet}} \mathcal{E}_{\mathrm{Pdet}} h^{-1} - S_{\mathrm{Pdet}} \mathcal{C}_{\mathrm{Pdet}} h^{-1}
                                                                                                                                                                                                                                             (6)
Rates
  \mu_{\mathrm{C}} = \mu_{\mathrm{Casec}} \, f_{\mathrm{CM}}(c_{\mathrm{DBM}}, c_{\mathrm{DBM}}) \ f_{\mathrm{CM}}(I) \ f(T) f_{\mathrm{AC}}(c_{\mathrm{A}}, c_{\mathrm{C}})
 \begin{split} \mu_{A} &= \mu_{\text{base}} \, f_{AN}(c_{DR}, c_{DR}) \, \, f_{AI}(I) \, \, f(T) f_{AC}(c_{A}, c_{C}) \\ R_{C} &= \mathbb{R}_{\text{Chook}} \, f(T) (c_{C} - C_{\min}) / c_{C} \end{split}
                                                                                                                                                                                                                                             (8)
(9)
  R_{A} = \mathbb{R}_{A \max} f(T) \left( c_{A} - A_{\min} \right) / c_{A}
                                                                                                                                                                                                                                              (10)
  \beta = \beta_0 f(T)
                                                                                                                                                                                                                                             (11)
 y = y_0 f(T)
```

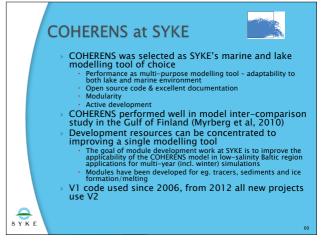


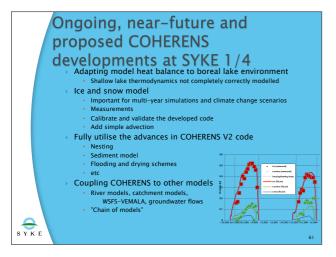


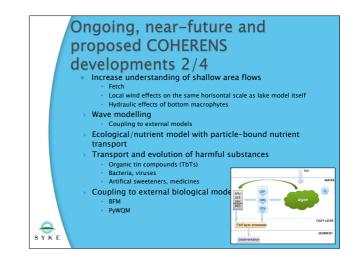


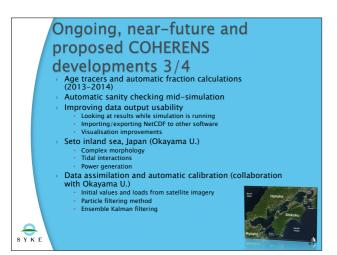


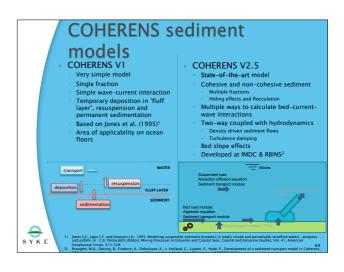




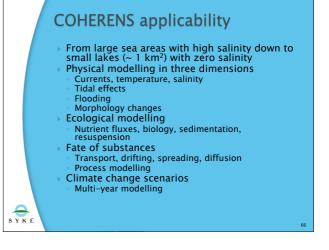












# **Basic lessons**

- Solution orientation >
- Model to be applied has to solve the problem at hand
- As simple model as possible
- We should have good understanding what happens in
- Model (=Tool) known and at hand
- Well documented
- Open source code
- Possibility to modify the model
  Collective experience and team work
- Data collection
  - Calibration
- Validation



# "We have the skills, ambition and courage to open new avenues for improving the environment."

# Related literature

Huttula T., 1994: Suspended sediment transport in Lake Säkylän Pyhäjärvi. Aqua Fennica, Vol. 24,2: 171–185

Huttula T., 1994: Suspended sediment transport in Lake Sākylān Pyhājārvi. Aqua Fennica, Vol. 24, 2. 171-185

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# Future prospects

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As presented in this report, various topics related to aquatic problems have been addressed by researchers of OU and SYKE in this project. During this project period, our collaboration has been related mainly to scientific approaches, but it has not remained restricted to them. We have enjoyed mutual understanding through the scientific discussion and social events that have taken place during our reciprocal visits. We have sought mutual understanding in terms of culture and history, which has deepened our collaborations and has made them much more fruitful.

This project officially ends in June 2014, but we regard that event as a starting point for our next step forward. Now we have come to understand that there are many not only important but also interesting problems that we share. We are poised to proceed to wider and deeper research collaborative efforts.

Furthermore, several young scientists and graduate students have participated in this project. Especially for them, this project has provided an invaluable experience. We believe they will extend international collaborative efforts of this kind throughout their scientific careers.

