Chemical and microbiological changes of salted Caspian Kutum
(Rutilus frisii kutum) roe

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Abstract
Salted Kutum roe was prepared by soaking in brine to a traditional procedure. Proximate
composition (moisture, protein, lipid, pH, ash and salt content), total volatile basic nitrogen
(TVB-N), fatty acid profiles and microbiological characteristics of raw and salted roe of
Caspian white fish (Rutilus frisii kutum) were determined. The results indicated that protein
and moisture contents were significantly higher in raw roe compared to salted one (28.81 and
61.07% to 23.99 and 51.57%, respectively). Significant differences in ash and salt contents
were obtained among different treatments. TVB-N in all samples was less than 25mg/100g.
The gas chromatography (GC) analysis showed that the major saturated fatty,
monounsaturated fatty acids and poly unsaturated fatty acids were 16:0, 18:1n-9, and 22:6n-3,
respectively. Aerobic plate count (APC) and Total Coliform Count (TC) in all samples were
significantly higher (p<0.05) in raw roe (5.33 logCFU/g and 210 MPN/g) compared to salted
roe (1.23 log CFU/g and 1.2 MPN/g). None of the samples contained Escherichia coli,
Salmonella, Clostridium perfringens or yeast.

Keywords: Rutilus frisii kutum, Roe, Salting, Microbial population

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Introduction
Roe of many fishes are consumed as food worldwide, either as a whole sac or as loose eggs (Scano et al., 2009). Roe, as referred to the oocytes when they are gathered in skeins, and incorrectly named as eggs, the non-fertilized oocytes are processed as a by-product collected from the female fish captured before spawning or during the slaughtering of farmed fish (Mahmoud et al., 2008). Caviar, ikura and tobiko are the well-known roe products in some countries. Salted-dried (karasumi) and salted–fermented (karashi mentaiko) are the most popular form in many middle and far eastern countries and a few European countries (e.g. Greece, Spain, Italy, and Norway) (Bledsoe et al., 2003; Bekhit et al., 2008). Recently, these products have grown in the world and currently enjoy expanding international and domestic markets (Mahmoud et al., 2008).

Many procedures are employed to process and preserve this highly perishable foodstuff rich in health-beneficial n-3 polyunsaturated fatty acids (PUFA) (Scano et al., 2009). Salting and drying are two methods used by many ancient civilizations for food preservation, while nowadays they are used not only to preserve food but also to give them a special flavor (Rodrigo et al., 1998; Katselis et al., 2005; Gallart-Jornet et al., 2007; Bekhit et al., 2008;).

The potential health benefits related to fish and fish-derived products are due to the presence of nutrients such as proteins, minerals, vitamins, and n-3 polyunsaturated fatty acids (n-3 PUFA) (Bledsoe et al., 2003; Rosa et al., 2009), that which are important for human health (Rosa et al., 2009). Furthermore, fish roe has a high percentage of phosphatidylcholine (PC) that is effective for the improvement of learning ability and the lowering of plasma lipids (Shirai et al., 2006).

The Kutum (R. frisii kutum) (also known as Caspian White Fish or Mahi-Sefid in Persian) is the most popular and highly valued fish among the Iranian seafood products particularly in the Guilan and Mazandaran provinces in north of Iran. Several types of roe called as Ashpal or Ashbal, are used in these areas, which are consumed as grilled, cured, salted, or mixed with other ingredients. Carp roe is less common and barbel roe is also occasionally used. In this study we investigated the proximate composition and effects of salting on chemical and microbiological characteristics of Kutum roe.

Materials and Methods
All analytical grades of chemicals and reagents were purchased from Merck (Darmstadt, Germany) or Sigma (St Louis, MO). Kutum was caught in the southwestern coast of Caspian Sea (Anzali port) by beach seine. The average weight of fish specimens and ovaries were 1800±20g and 185±5g respectively. All ovaries were in final stage of maturity. Dried Rose madder (Rubia tinctorum) plant, whose root is a source of red dye, used in food processing, was purchased from the local shops.

Preparation of salted roes
Roes were collected from 24 freshly killed fishes. The whole ovaries were divided in two parts; the first part was packed in aseptic bags, placed in ice (ice/roе ratio 2/1) and immediately transferred to the laboratory within 1h for further analyzes. The other
parts, were used for salting; salted roes were prepared according to the traditional recipe. In this method, a mixture of 70 g salt and 5 g madder for each sample were prepared and covered each ovary wrapped in paper and completely immersed in brine saturated with salt. The experiments were carried out in triplicate and samples were stored at 20°C for 45 days.

**Chemical composition**

**Moisture, protein and lipid contents**

The moisture content of roe samples was determined by drying samples to constant weight at 102-105°C for 20 to 24 h according to the AOAC standard method (AOAC, 1995). The crude protein content of roe samples was measured using the Kjeldahl method. Approximately 2 g of roe sample was used for protein content determination by using a factor 6.25 following the AOAC standard method (AOAC, 1995). The roe lipids were extracted using a chloroform and methanol mixture (ratio 1:1) according to the protocol of Bligh and Dyer (Bligh and Dyer, 1959).

**pH value, salt content and ash determination**

Samples of raw and salted fish roes (10 g) were homogenized in sterile blenders with 10 mL of distilled water to make a thick slurry. The pH of this slurry was measured by a calibrated portable digital pH meter (Beckman Φ40, Krefeld, Germany) (Kung et al., 2008). NaCl content was determined by the volumetric method of Volhard. The homogenized roe samples was titrated with 0.1 M AgNO₃, using 10 % w/v K₂CrO₄ solution as an indicator. The salt content was calculated as percentage of the sample. Ash was determined after heating the samples overnight at 550°C (AOAC, 1995).

**Determination of total volatile basic nitrogen (TVB-N)**

The TVB-N content of the Kutum roe samples were measured by the method of Conway’s dish. Each ground sample (10 g) of roes was extracted with 20 mL of 6% trichloroacetic acid (TCA) and filtered. The TCA extracts of the roe samples were absorbed by boric acid and then titrated with 0.02 N HCl. The TVB-N content was expressed in mg/100 g of fish roes (Kung et al., 2009).

**Fatty acid analysis**

The total lipids (TL) were extracted from each sample by the method of Bligh and Dyer. Fatty acid methyl esters were prepared by transmethylatation using 2M KOH in methanol and n-heptane. Extracted lipids (10 mg) were dissolved in 2 mL heptane followed by 4 mL of 2M methanolic KOH. The tube vortex and centrifuged at 4000 rpm xg unit for 10 minutes (Shirai et al., 2006). The heptane layer was taken for GC analyses. Fatty acids were measured by the gas chromatograph (DANI 1000, DANI Instruments SpA, Italy) equipped with a fused silica capillary SGE column (30mx0.25 mm ID). The carrier gas was helium with a split injection of 40:1. The temperature profiles were as follows: initial temperature, 175°C; heating rate, 1° C/min; final temperature, 220 °C (final time, 20 min); injector temperature, 250 °C, and detector temperature, 270 °C. The fatty acids were identified by comparison of the retention times with those of standard purified fatty acids (Shirai et al., 2006).
Microbiological analysis

Aerobic Plate Count (APC)
A 25 g portion of each test sample was homogenized at high speed for 2 minutes in a sterile blender with 225 mL of sterile potassium phosphate buffer (pH 7.0). The sterile blender was prepared by autoclaving for 15 minutes at 121 °C. The homogenates were serially diluted with a sterile phosphate buffer, and 1.0 mL aliquots of the diluted homogenates were inoculated into an aerobic plate count (APC) agar (Difco, Detroit, MI, USA) containing 0.5% NaCl. Bacterial colonies were counted after incubating the plates at 35 °C for 48 hours. The bacterial numbers in the roe samples were expressed as log_{10} colony-forming units (CFU/g) (Huang et al., 2010).

Salmonella spp.
A 25 g of sample and 225 mL buffered peptone-water were homogenized using a filter Stomacher. The analysis depends on the principles of pre-culturing for 18-24 hours in Buffered Peptone-Water at 35-37 °C, the selective culturing for 24 hours in Selenite Cystine Broth at a temperature of 35 °C, and the identification of the suspected colonies through the current biochemical and serological tests (Valcheva et al., 2011).

C. perfringens
A 10 g of sample and 100 mL buffered peptone-water were homogenized and further 10-fold dilutions' were made. The analysis was on incubation for 24 hours in Sulphite polimyxin sulphadiazine agar at a temperature of 35-37 °C and the identification of the Clostridia through the biochemical test (Huang et al., 2010).

Total Coliform and E. coli
Analyses of total coliform and E. coli in roe samples were conducted by using three tubes according to the most probable number (MPN) method (Kung et al., 2009). Lauryl sulphate tryptose broth (LST broth) and brilliant green lactose bile (2 %) broth (BGLB broth) were used for presumptive and confirmed tests for total coliform, respectively. E. coli was determined by the LST broth and E. coli (EC) broth. Cultures that showed positive production of gas were then confirmed by eosin methylene blue agar (EMBA) (Huang et al., 2010).

Yeast
A 10 g of sample and 100ml peptone-water were homogenized and further 10-fold dilutions' were made. The analysis depends on incubation for 5 days in Dichloran Rose Bengal Chloramphenicol agar at a temperature of 25 °C; yeasts colonies were counted (Altug and Bayrak, 2003).

Microbiological data were transformed into logarithms of the number of colony-forming units (cfu/g). All plates were visually examined for the typical colony types and morphological characteristics associated with each growth medium. All counts were performed in triplicate.

Statistical analyses
Experiments were performed in triplicates and they were expressed as mean ± standard deviation. Data were evaluated by software program SPSS version 12.0. Data were subjected to T-test. The level of significance was set at p<0.05.

Results
Chemical composition
Table 1 shows the results of proximate composition (moisture, crude protein, fat, ash and pH) of raw and salted Kutum roes. The moisture contents varied significantly from 61.1 to 51.5% (raw and salted roes, respectively). The mean protein contents of raw and salted roes were 28.8 and 24.0 % respectively which was in good agreement with 28.5% (trout roe) (Mahmoud et al., 2008).

<table>
<thead>
<tr>
<th>Sample</th>
<th>Protein</th>
<th>Lipid</th>
<th>Moisture</th>
<th>Ash</th>
<th>Salt</th>
<th>pH</th>
</tr>
</thead>
<tbody>
<tr>
<td>Raw Roe</td>
<td>28.81±1.1a</td>
<td>6.84±2.2a</td>
<td>61.07±0.3a</td>
<td>1.3±1.8a</td>
<td>0.35±0.08a</td>
<td>7.49±0.05a</td>
</tr>
<tr>
<td>Salted Roe</td>
<td>23.99±0.5b</td>
<td>6.4±0.7a</td>
<td>51.57±0.6b</td>
<td>17.74±1.1b</td>
<td>5.13±0.3b</td>
<td>5.1±0.09b</td>
</tr>
</tbody>
</table>

Mean ± Standard deviation and data have expressed as % of wet matter.
* a–b Values in the same column with different letters are statistically different (p<0.05).

There was no significant difference between lipid values. Lipid contents of raw and salted samples were 6.84% and 6.4%, respectively (p>0.05). Ash content of brined roe (Table 1) was much higher than that of raw samples. Water losses associated with brining resulted in increased ash content. Marked differences were found for pH of the samples. Significant decrease was observed in the pH value of salted roe, indicating the effect of salting process. Salt content in salted roe (5.1%) was significantly higher than raw sample (0.3%), which was the result of salting process. The average of TVB-N value of the raw samples (10.9 mg/100gr) was significantly higher than salted roe (5.1 mg/100gr) (Figure 1); both levels were under determined levels (30 mg/100gr).

**Figure 2:** Comparison of TVB-N[^1] measured in raw and salted roe of Kutum

[^1]: Mean values from three independent determinations. Standard deviations are expressed by bars.
**Fatty acid composition**

Fatty acid composition of Kutum roe are presented in Table 2. C16:0, C16:1, C18:0, C18:1 and C22:6 were the main fatty acids of roe. Monounsaturated fatty acids (MUFA) represented the most dominant class of total fatty acids (49.58%) followed by saturated fatty acids (27.93%).

<table>
<thead>
<tr>
<th>Fatty acid composition of total lipid (TL) in raw and salted Kutum roe (%)</th>
<th>Raw Roe</th>
<th>Salted Roe</th>
</tr>
</thead>
<tbody>
<tr>
<td>aSFA</td>
<td></td>
<td></td>
</tr>
<tr>
<td>14:0</td>
<td>0.68±0.00</td>
<td>0.62±0.1</td>
</tr>
<tr>
<td>15:0</td>
<td>18.26±0.3</td>
<td>18.93±0.6</td>
</tr>
<tr>
<td>16:0</td>
<td>0.85±0.05</td>
<td></td>
</tr>
<tr>
<td>18:0</td>
<td>6.05±0.3</td>
<td>7.48±0.1</td>
</tr>
<tr>
<td>20:0</td>
<td>0.00</td>
<td>0.00</td>
</tr>
<tr>
<td>∑</td>
<td>27.93</td>
<td>27.3</td>
</tr>
<tr>
<td>bMUFA</td>
<td></td>
<td></td>
</tr>
<tr>
<td>14:1n-5</td>
<td>0.00</td>
<td>0.14±0.02</td>
</tr>
<tr>
<td>15:1</td>
<td>0.37±0.01</td>
<td></td>
</tr>
<tr>
<td>16:1n-7</td>
<td>12.02±0.5</td>
<td>4.52±0.3</td>
</tr>
<tr>
<td>17:1n-7</td>
<td>1.48±0.03</td>
<td></td>
</tr>
<tr>
<td>18:1n-9</td>
<td>27.15±1.3</td>
<td>12.67±1.3</td>
</tr>
<tr>
<td>18:1n-7</td>
<td>7.39±0.5</td>
<td>6.18±0.5</td>
</tr>
<tr>
<td>20:1n-9</td>
<td>1.18±0.07</td>
<td>0.39±0.08</td>
</tr>
<tr>
<td>∑</td>
<td>49.58</td>
<td>23.9</td>
</tr>
<tr>
<td>cPUFA</td>
<td></td>
<td></td>
</tr>
<tr>
<td>18:2n-6</td>
<td>1.33±0.02</td>
<td>2.11±0.4</td>
</tr>
<tr>
<td>18:3n-3</td>
<td>0.63±0.08</td>
<td>0.35±0.05</td>
</tr>
<tr>
<td>20:2n-6</td>
<td>0.46±0.04</td>
<td>0.33±0.02</td>
</tr>
<tr>
<td>20:3n-3</td>
<td>0.00</td>
<td></td>
</tr>
<tr>
<td>20:4n-6</td>
<td>1.63±0.1</td>
<td>4.74±0.6</td>
</tr>
<tr>
<td>20:5n-3</td>
<td>4.37±0.3</td>
<td>7.6±0.5</td>
</tr>
<tr>
<td>22:6n-3</td>
<td>5.17±0.9</td>
<td>21.62±1.4</td>
</tr>
<tr>
<td>∑</td>
<td>13.59</td>
<td>36.75</td>
</tr>
<tr>
<td>PUFA/SFA</td>
<td>0.49</td>
<td>1.34</td>
</tr>
</tbody>
</table>

a SFA, saturated fatty acids.
b MUFA, monounsaturated fatty acids.
c PUFA, polyunsaturated fatty acids.
d Samples were analyzed in triplicate and results are expressed as mean value ±SD (p<0.05).

According to the results of the analyses of fatty acids, there was significant difference between SFA and MUFA values (p<0.05). The percentage of MUFA differed from 49.58 to 23.9 % in raw and salted samples. The predominant MUFA in both samples was oleic acid (C18:1n9, 27.15 and 12.67%). A significant difference was found between the processing and the MUFA; showing a decreasing trend (Fig. 2). The samples had < 40% PUFA (13.6 and 36.7%, raw and salted roe), eicosapentaenoic acid (EPA) (C20:5n3) had the highest percentage in raw and salted samples.
Kutum roe (5.1 and 21.6%, respectively). Results showed that salted roe had higher percentage of DHA and EPA compared to raw samples.

![Graph showing SFA, MUFA, and PUFA levels in raw and salted Kutum roe](image)

**Figure 2**: Values of the total Saturated Fatty acid (SFA), Monosaturated Fatty Acid (MUFA), and Polyunsaturated Fatty acid (PUFA) measured in raw and salted roe of Kutum.  

1 Expressed as percentage of total fat.

**Microbiological analysis**

The content of aerobic plate count (APC), total coliform (TC) and *E. coli*, *Salmonella*, *C. perfringens*, fungi and yeast counts which were obtained in this study are summarized in Table 3.

**Table 3**: The results of microbiological analysis of raw and salted Kutum roe.

<table>
<thead>
<tr>
<th>Sample</th>
<th>APC (log cfu/g)</th>
<th>TC (MPN/g)</th>
<th><em>E. coli</em> (MPN/g)</th>
<th><em>Salmonella</em> (cfu/g)</th>
<th><em>C. perfringens</em> (cfu/g)</th>
<th>Fungi (cfu/g)</th>
<th>Yeast</th>
</tr>
</thead>
<tbody>
<tr>
<td>Raw Roe</td>
<td>(5.33 ±1.5) a</td>
<td>(210±0.0) a</td>
<td>0</td>
<td>ND</td>
<td>ND</td>
<td>0</td>
<td>4×10²</td>
</tr>
<tr>
<td>Salted Roe</td>
<td>(1.23 ±0.05) b</td>
<td>(1.2 ±2.07) b</td>
<td>0</td>
<td>ND</td>
<td>ND</td>
<td>0</td>
<td>ND</td>
</tr>
</tbody>
</table>

1 Means ± SD. Values are average of triplicate (n=3) analyses ± standard deviation. Means within rows followed by different letters denote significant differences (p<0.05). ND: Not detected

None of the raw and salted roe samples contained *E. coli*, *Salmonella* and *C. perfringens*. The mean value of APC level in raw roe (5.33×10³ cfu/g) was significantly higher than salted roe (1.23×10³ cfu/g). The average content of
Coliforms in raw roe was 210 mpn/g which was significantly higher than salted samples (1.2 mpn/g). The mean content of yeasts in raw roe was $4 \times 10^2$ cfu/g, but not detected in salted roe.

**Discussion**

Salting leads to a reduction in moisture that has been attributed to the denaturing effect of salt on proteins. High brine concentration caused dehydration of the samples, due to the difference in solute concentration between the brine solution and roe water. Jittinandana and others (2002) indicated that denaturation of muscle protein facilitated diffusion of water from fish. Loss of the protein content is one of the main biochemical changes during salting (Aristoy and Toldra, 1991). Sodium chloride increased protein solubility (Martinez-Alvarez and Gomez-Guillen, 2005); According to Jittinandana and others (2002) soaking in high brine solution may have caused salting-out of proteins that led to precipitation and dehydration of proteins and, in turn, exclusion of water molecules; that is dependent on salting time and salt concentration (Aristoy and Toldra, 1991; Andersen et al., 2007).

Lipid content in raw and proceeded Kutum roe was about 6% and it ranged from 3% in pacific herring (*Clupea pallasi*) roe (Rodrigo et al.,1998), 12% in *Acipenser huso* (Altug & Bayrak, 2003), 14.5% in Ikura (salted salmon roe), 3.7% in Tarako (salted pollock roe), 3.2% in Tobiko (salted flyingfish roe), 3% in Kazunoko (salted herring roe) (Shirai et al.,2006), 25.7% in salted mullet roe (Lu et al., 2006) and 9.8% white fish (*Coregonus albula*) roe (Kaitaranta,1980).

In the present study, the ash content of salted Kutum roe was higher than salted hake and ling roe (*Molva molva*) (6.9%) and mullet roe (2.6%) (Rodrigo et al., 1998). This is mainly affected by salting process, which remains a high amount of minerals to the roe (Rodrigo et al., 1998). High brine concentration increased water phase salt content of samples, and it corresponded with increased product ash content (Jittinandana et al., 2002).

Marked differences were found for pH of the samples. Significant decrease was observed in the pH value of salted roe, indicating the effect of salting process. In other studies pH of salted roes ranged between 5.5-5.86; as the result of salting process (Tsai et al., 2005; Celik et al., 2012).

The properties of fish and fish products vary as a result of alterations in water and salt concentration. Diffusion has been known as the most important mass transfer mechanism of sodium and chloride. Differences in osmotic pressures between cells and salting agent cause the abruption of salting agent into fish and excretion of water out of fish (Gallart-Jornet et al., 2007).

TVB-N is one of the most widely indicators of fish quality and spoilage (Kung et al., 2008). According to the quality categorization of fish and fish products proposed by Lang (1983), TVB-N value should be lower than 25 mg/100 g, for “high quality”, less than 30 mg/100g for “good quality”, lower than 35 mg/100 g for “limit of acceptability” and more than 35 mg/100 g for “spoilt” (Özyurt et al., 2009). In both group samples were under determined levels (30 mg/100gr). Based on
this category, salted and raw samples may be placed in high quality group. The TVB-N content of salted roe was nearly half of the level of raw samples, indicating that the addition of salt could apparently have some inhibitory effects on spoilage of salted roe samples; it was in agreement with Kung et al (2008).

The main fatty acids (SFA) of roe were found similar pattern in Kutum roe by Ghomi and Nikoo (2010). Most of the studies have shown that saturated fatty acids of fish and fish products are between 20 and 30%, which it was in agreement with these studies (Rincón-Cervera et al., 2009; Rosa et al., 2009; Shin et al., 2010). Palmitic acid had highest proportions. In 15 previous studied species, the predominant SFA was palmitic acid (Rincón-Cervera et al., 2009) and this fatty acid is required for fish growth and formation of roe in females (Huynh et al., 2007). Rincón-Cervera et al. (2009) reported oleic acid as the main component of MUFA in 13 species. Brined roe had less lipid content which is possibly due to decreasing of the MUFA through processing, PUFA percentage of salted roe increased. Scano et al. (2009) suggested that the fatty acid profiles of salted and dried mullet roe were not affected by the processing.

Some reports have indicated that the fatty acid composition of fish roe is impacted by diet but does not depend on the level of maturation (Shirai et al., 2006). Analysis of fatty acid composition of salted roe of salmon, pollock, flying fish, and herring represented that DHA value was higher than EPA (17.4, 22.2, 27.9 and 22.6%, respectively). Mahmoud et al (2008) studied fatty acid profile of rainbow trout (Oncorhynchus mykiss) roe and they indicated that the PUFA of rainbow trout roe was about 40-50% of total fatty acid content which was higher than that of raw and salted Kutum roe. Ghomi and Nikoo (2010) suggested the Kutum roe as a good source of PUFA can be used in food products.

Tsai et al. (2005) reported that higher salt content (>5%) had some inhibitory effects on bacterial growth which results in agreement. Besides, the positive effect of salt on microbial growth (Wheaton and Lawson, 1985), rose madder (Rubia tinctorum) have antimicrobial effects on some bacteria (Mehrabian et al., 2000). APC is a criteria for determination of the general microbiological quality of the product (FAO, 1992). Coliforms in raw roe was higher than salted samples. The existence of coliform in processed food products indicates that the processing have been conducted under inefficient hygienic conditions (ICMSF, 1987). These samples revealed that salt had some inhibitory effect on coliform bacterial growth. Some studies proved that most bacteria responsible for fish meat spoilage are stressed by salt content above 1% and would die or at least would be stopped further development, as the salt content increase from 6% to 8% (Wheaton and Lawson, 1985; Kung et al., 2008). During the process of excerption of the roe from the fish, the bacteria on the surface of the fish may be transferred to the roe.

The yeast was not detected in salted roe. Among the micro-organisms, the yeasts are known to be causing the food spoilage and organoleptic spoilage that were visually noticed depending on the time and
production intensity. This can affect the commercial quality of the product in a negative manner (Altug and Bayrak, 2003).

This study represents a report on the proximate composition and quality changes of raw and salted Kutum roe. Despite the increasing levels of TVN in raw samples, a decrease occurred for the salted samples. Although TVB-N values seems to be a good spoilage factor for both samples, but to draw a better conclusion these values should be interpreted in combination with quality changes (lipid oxidation,…) and shelf storage life of the products are required. The total count and APC of raw samples significantly were higher than that of salted samples representing the preservative effect of salting. Overall most of the results obtained in this study indicate that salting might be a suitable procedure for preserving quality of Kutum roe. It had some inhibitory effects on bacterial growth. Therefore, further research is suggested to investigate the effects of salting on quality and sensory changes during storage.

References


