

Fluorescent immunoliposomal nanovesicles for rapid multi-well immuno-biosensing of histamine in fish samples

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- 27 **Running head:** Immunoliposomal nanovesicles for time resolved detection of histamine
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33 ABSTRACT

Scombroid poisoning in fish-based and other food products has raised concerns due to 34 toxicity outbreaks and incidences associated with histamine, thus measuring the amount of 35 histamine toxic molecule is considered crucial quality indicator of food safety and human 36 health. In this study, liposome-based measurement of histamine was performed via rupturing 37 mechanism of sulforhodamine B dye encapsulated anti-histamine antibody conjugated 38 liposomal nanovesicles. The immunosensing ability of immuno-liposomal format was 39 40 assessed by monitoring the fluorescence at excitation/emission wavelength of 550/585 nm. Immuno-liposomal format assays were considered, one based on single wash procedure 41 (Method 1), which had a detection limit of 10 ppb and quantification limit 15-80 ppb. While 42 Method 2 based on one-by-one wash procedure had a detection limit of 2-3 ppb and 43 quantification limit 8.5 ppb–200 ppm that required 2 h 30 min to perform. In view of better 44 quantification limit. Method 2 was chosen for further tests required to validate its 45 applicability in real samples. The feasibility of Method 2 was reconfirmed in fresh mackerel 46 fish, and canned fish (tuna and salmon) with a similar detection limits but with low amplified 47 48 fluorescence signals and sufficient levels of histamine recovery from fresh mackerel (73.50-99.98%), canned tuna (79.08-103.74%) and salmon (74.56-99.02%). The specificity and 49 method accuracy were expressed as % CV in the range 5.34%-8.48%. Overall, the developed 50 multi-well sensing system (Method 2) showed satisfactory specificity, cost effectiveness, 51 rapidity, and stability for monitoring histamine toxicity as a practical food diagnostic device. 52

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Keywords: Double layered; Liposome immunosensor, Signal amplifiers, Fluorescence
quenching, Histamine toxicity

57 **1. Introduction**

58

Since 1960s optical detection systems have been used as a powerful tool to allow 59 light propagation with a minimum loss for use in sensor development strategies. Similarly, 60 nowadays, fluorescence technology has emerged as a mean of sensitivity enhancer with 61 62 reduced matrix effects, making it applicable for biosensing purposes (Chang et al., 2016). Food scientists are concerned about several issues related to the contamination of protein-rich 63 64 food products by endogenous bioactive amines as chemical messengers in biological systems (Lin et al., 2018). Nowadays, histamine contamination of food is common and considered a 65 66 serious human health and food safety issue, hence, fabrication of fluorescence assay for real 67 time monitoring of food toxicants, such as histamine, has attracted huge attention. Histamine is mainly formed in protein-rich food matrices by certain microorganisms that produce 68 histidine decarboxylase, which catalyzes the conversion of free histidine to histamine (EFSA, 69 2011). Given that fish consumption is variable, a serving size of 250 g was considered 70 reasonable to establish a maximum level of histamine in fish of 200 mg/kg (FAO, 2012), 71 72 whereas the United States Food Drug Administration (USFDA) set the histamine threshold limit at 50 mg/kg (FDA, 2011). Hence, the sensitive and selective detection of histamine is of 73 considerable importance from the safety and clinical perspectives and for studies on allergic 74 75 responses under various pathological conditions (Yan et al., 2014; Yang et al., 2015). Therefore, it is essential to develop sufficiently sensitive and rapid cost effective methods to 76 detect histamine residues for clinical and food diagnosis. 77

A number of analytical detection methods have been developed for the determination of histamine levels in food products, including reverse-phase high performance liquid chromatography (RP-HPLC), cation-exchange chromatography (CEC), gas chromatography (GC), thin layer chromatography (TLC) (Awan et al., 2008; Lapa-Guimarães and Pickova,

2004) and ELISA based assays (Luo et al., 2014). Although these analytical techniques have
provided adequate for detecting a variety of analytes, time-consuming sample processing
steps, such as, clean-up, sample derivatization, and the low optical absorbance of histamine in
the ultraviolet region are problematic.

To address these problems, biosensors have attracted interest as potential rapid 86 87 analytical sensing tools as alternatives to traditional enzyme-based detections. Several rapid, one-step electrochemical biosensors based on enzymes or nanozymes (Pérez et al., 2013; 88 89 Jiang et al., 2015; Veseli et al., 2016; Yadav et al., 2019) have been devised for the sensing of histamine. However, the specificity cannot be ensured for these biosensors as the enzymes 90 91 used can catalyze histamine and its analogues. Further, several technologies use various 92 nanomaterials in combination with electrochemical and fluorescence detection techniques as signal amplifiers (Ali et al., 2017; Pei et al., 2013; Rusling, 2012; Du et al., 2011). These 93 nanomaterials include gold nanoparticles (Du et al., 2011), quantum dots (Qian et al., 2011), 94 magnetic nanoparticles (Mani et al., 2009), silica nanoparticles (Wu et al., 2009) and carbon 95 nanomaterials (Malhotra et al., 2010), and have ability to enhance the signals while using as 96 nano carriers. Since the significant signal releasing process from various nanocarriers is 97 complicated as some of these require strong acid, base, heat and sonication treatments, which 98 might have adverse effect on biological molecules, such as enzymes, antibody and antigenic 99 100 targets (Zhao et al., 2015).

101 Thus, there is a need for more sensitive/specific signal amplifier tools, and this need 102 might be met by antibody-based immunosensing detection strategies employing fluorescent 103 nanoparticles. Although a few fluorescence-based detection devices have been used for food 104 samples safety analysis (Liu et al., 2016; Yang et al., 2017; Kaur et al., 2018; Chauhan et al., 105 2018), no fluorescence-nano-sensing-based multi-well method has yet been devised that uses 106 layer by layer arranged fluorescence dye encapsulated immuno-liposome nanovesicles for

amplifying histamine toxin or other biogenic amines detecting signals in food matrices via antigen-antibody capturing, liposome vesicle rupturing and signal release. Hence, we afforded to incorporate with a liposomal fluorescence-amplifier to determine the feasibility of a new multi-well biosensing platform for histamine toxin in food matrices.

Lumen of liposomes encapsulates any biological molecules with the help of 111 liposomal aqueous core capacity and phospholipid head groups of polar nature that can play 112 an important role for the improvement of signal amplification, and can be used as 113 114 multifunctional vesicles (Edwards and Baeumner, 2006). It has been reported that liposomes exhibited better performance for advanced biosensors as compared to other signal 115 amplification materials (Edwards and Baeumner, 2006). Also, immunosensors based on 116 117 antigen-antibody interactions exhibit superior characteristics as compared to enzyme-based immunoassays due to their high sensitivities and excellent specificities (Zhang et al., 2016). 118 Moreover, liposome-based assays where markers (antibody) and fluorescent dyes are 119 encapsulated in the liposomes as detectable molecules, provide instant signal enhancement 120 after lysis mechanism as compared to enzyme-based immunoassays which show time and 121 concentration dependent signal enhancement, thus limiting the sensitivities and the speed of 122 analysis. Specifically, the liposome-based assay proposed in this study requires comparatively 123 lesser washing and incubation steps for antigen-antibody reactions as well as eliminates the 124 125 separate reaction of secondary antibody and reaction with HRP-tagged signal generating molecules. The total assay time of the proposed method is only 2 h and 30 min as compared 126 to ELISA-based methods which require various processing steps such as coating (12 h 127 128 incubation), washing with buffer solution (5-10 min), blocking (2 h incubation) re-washing with buffer solution (5-10 min), IgG addition (1 h incubation) followed by washing with 129 buffer solution (require 5-10 min), addition of labeled enzyme (1 h incubation) followed by 130 washing with buffer solution (5-10 min), enzyme substrate reaction (30 min incubation) and 131

addition of stop solution (NaOH) followed by absorbance measurement. A few additional
factors also make the immunoliposome-based assay (after antibody conjugation) favorable as
compared to other ELISA-based histamine detection assays due to their (1) instant
fluorescence signal enhancement ability after the lysis mechanism, (2) cost effectiveness, and
(3) an alternate over other fluorescence materials.

Thus far, only very few liposome-based rapid and multi-well detection methods have 137 been developed for histamine toxin. Therefore, the current study was undertaken to construct 138 139 Sulforhodamine B dye encapsulated phospholipid bilayers with anti-histamine antibody conjugated liposomal nanovesicles as an amplifier, positioned in a 2 layered format [single 140 well one wash format (Method 1) and one by one wash format (Method 2) that unites for 141 142 strong fluorescence signal, after rupturing the constructed lipid bilayer. The developed multiwell biosensing method (Method 2) exhibited excellent properties of fluorescence excitation 143 on particular wavelength and signal amplification via tagged antibody molecule on the outer 144 surface of fluorescent liposomal nanovesicles. In order to confirm the novelty of the 145 developed method (Method 2), sensitive detection of histamine in buffer medium and fish 146 147 samples, was validated to achieve the goal of the study by developing a rapid detection method that can overcome the drawbacks associated with Method 1 and conventional 148 analytical methods, including HPLC and other liposomal sensing methods. The detection 149 sensitivity and applicability of fluorescent anti-histamine IgG conjugated liposomal 150 nanovesicles (anti-His-LNs)-based biosensing system was compared in real food matrix with 151 a conventional HPLC method to validate its simplicity, precision, rapidity, repeatability and 152 153 straightforwardness.

154

155 2. Materials and methods

157 2.1. Chemicals and reagents

158

Putrescine dihydrochloride (PUT), histamine dihydrochloride (HIS), cadaverine 159 dihydrochloride (CAD), 2-phenylethylamine (PHE), tryptamine hydrochloride (TRP), 160 spermidine trihydrochloride (SPD), spermine tetrahydrochloride (SPM), tyramine 161 hydrochloride (TYR), N-(2-hydroxyethyl) piperazine-N-2-ethanesulfonic acid (HEPES), 162 sodium azide, sodium chloride, *n*-Octyl- β -D-glucopyranoside (OG), methanol, acetone, 163 cholesterol, sucrose, Sepharose CL-4B, triethylamine, dansyl chloride, potassium phosphate 164 monobasic, ethylene diamine tetra acetic acid (EDTA), potassium phosphate dibasic, 165 dimethyl sulfoxide (DMSO) and amino acids, including histidine, glycine, alanine, lysine, 166 glutamic acid and arginine were obtained from Sigma, USA. Sulforhodamine B was 167 purchased from Molecular Probes, USA. Ammonium hydroxide, sodium hydroxide, 168 perchloric acid, and sodium hydrogen carbonate were purchased from Junsei Chemicals, 169 Japan. 1,2-Dipalmitoyl-sn-glycero-3-phosphocholine (DPPC), 1,2-dipalmitoyl-sn-glycero-3-170 [phospho-rac-(1-glycerol)] (DPPG), and 1,2-dipalmitoyl-sn-glycero-3-phosphoethanolamine 171 (DPPE) were procured from Avanti Polar Lipids, USA. N-[k-maleimidoundecanoyloxy] 172 sulfosuccinamide (sulfo-KMUS), *N*-Succinimidyl-*s*-acetylthioacetate 173 (SATA), and hydroxylamine hydrochloride were obtained from Pierce Products, USA. Monoclonal anti-174 histamine antibody (anti-histamine IgG) (cat#MBS358003; 1 mg/mL) was purchased from 175 MyBioSource, USA. Immunoplates (96 well amine binding polystyrene surface) were 176 purchased from Thermo Scientific (USA). 177

178

179 2.2. Synthesis of Sulforhodamine B-encapsulated liposome nanovesicles

181	Liposome nanovesicles were prepared using a reverse-phase evaporation method we
182	previously described with a few modifications in our earlier methodology. In brief, a lipid
183	mixture (40.3 μ mol: DPPC, 4.2 μ mol: DPPG, and 40.9 μ mol: cholesterol) was used to form a
184	phospholipid bilayer, and a fluorescent dye Sulforhodamine B at higher concentration
185	(150 mM) was selected to make fluorescent liposome nanovesicles, as we previously
186	described (Shukla et al., 2016). Non-encapsulated dye or trace of organic solvent was
187	removed from the liposome preparation by gel-filtration on a Sephadex G-50 column (1.5×18
188	cm) at room temperature. Detailed methodology for synthesizing Sulforhodamine B-
189	encapsulated liposome nanovesicles has been given in the Section 1 of the supplementary
190	information.

191

192 2.3. Surface functionalization of fluorescent liposome nanovesicles and conjugation of anti193 histamine IgG to develop immunosensor

194

195 Step I. Derivatization of anti-histamine IgG with maleimide functional group

196 Fluorescent anti-histamine conjugated liposomal nanovesicles (anti-His-LNs) as nanobiosensors were prepared as previously described with some modification [28]. Briefly, 197 0.1 mg of anti-histamine IgG was dissolved in 1 mL of 0.05 M potassium phosphate buffer 198 199 containing 1 mM of EDTA and 0.01% sodium azide (pH 7.4). A sulfo-KMUS solution was 200 prepared by dissolving 3 mg of sulfo-KMUS in 0.15 mL of solvent mixture of DMSO: MeOH (2:1, v/v). Then, a 2.25 µl of sulfo-KMUS solution was then added to 1 mL of this 201 202 anti-histamine IgG solution (0.1 mg/mL) and incubated on a shaker at room temperature for 3 h [29, 30]. Once the antibody (0.1 mg of anti-histamine IgG) derivatized with the maleimide 203 group, was dialyzed overnight with 0.02 M HEPES buffer (0.15 M NaCl, 0.01% sodium 204 205 azide), and sucrose was added to maintain its osmolarity as 427 m osmol/L. Importantly, to

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206 maintain the proper stability of Sulforhodamine B-encapsulated liposomes, all buffers used in the liposome preparation were adjusted to relatively a little higher osmolarity so as to prevent 207 208 the osmotic pressure-related swelling.

209

Step II. Removal of acetylthioacetate group from Sulforhodamine B-encapsulated liposome 210 nanovesicles 211

The N-succinimidyl-s-acetylthioacetate (SATA) was used during the first step of 212 liposome synthesis to prepare DPPE-ATA complex by mixing DPPE (7.2 µmol) and SATA 213 (14.3 µmol) with 1 mL of 0.7% trimethylamine. After incorporation of lipid constituents into 214 liposome bilayer, final obtained Sulforhodamine B-encapsulated liposome nanovesicles may 215 216 still have some amounts of acetylthioacetate group moieties from SATA, which should be removed prior to further processing reaction (Kindly refer to supplementary information 217 section for more information). 218

The total volume of Sulforhodamine B-encapsulated liposome solution was 219 measured as it may differ lot-to-lot. 0.5 M of hydroxylamine hydrochloride was prepared in 220 0.1 M HEPES solution containing 25 mM EDTA (pH 7.4) and then added into 221 Sulforhodamine B-encapsulated liposome solution in a working ratio of 1:10 (1 mL of 222 Sulforhodamine B-encapsulated liposome solution; 0.1 mL of 0.5 M hydroxylamine 223 224 hydrochloride solution) followed by flushing the flask under nitrogen for 1 min. The reaction process of deacetylation was allowed to occur at room temperature at 70 rpm shaker for at 225 least 2 h. 226

227

Step III. Conjugation of maleimide-derivatized anti-histamine IgG to Sulforhodamine B-228 encapsulated liposome nanovesicles 229

230	Conjugation of maleimide-derivatized anti-histamine IgG and fluorescent liposome
231	nanovesicles (210 μ L) was performed by adjusting the liposome solution mixture to pH 7.0
232	using 0.5 M HEPES buffer (to make same pH as required for -SH group ethylmaleimide
233	quenching, 0.02 M Tris buffer; pH 7.0), added with maleimide-derivatized anti-histamine IgG
234	solution (0.1 mg/mL of anti-histamine IgG prepared in Step I), and flushing under nitrogen
235	gas for 1 min. The reaction was allowed to proceed on a shaker at room temperature for 4 h
236	and then incubated at 4°C overnight. To quench unreacted -SH groups, 100 mM of
237	ethylmaleimide dissolved in 0.02 M Tris buffer containing 0.15 M NaCl, 0.01% NaN ₃ , and
238	0.07 M sucrose; pH 7.0 was added to the immunoliposomes reaction mixture; the osmolarity
239	of this buffer solution (421 m osmol/L) was maintained with 0.07 M sucrose using an
240	osmometer (Shukla et al., 2016). Anti-histamine IgG-tagged liposome nanovesicles were
241	separated from unreacted SH-derivatized anti-histamine IgG using a Sepharose CL-4B
242	column equilibrated with 0.02 M Tris buffer (0.15 M NaCl, 0.01% NaN ₃ , pH 7.0) containing
243	0.07 M sucrose. The desired fraction of anti-histamine IgG-tagged liposome nanovesicles
244	was collected and the solution was dialyzed (in 0.02 M Tris buffer) overnight at 4 °C in the
245	dark for achieving to improve its stability. The confirmation of free antibody released during
246	elution from Sepharose CL-4B column was analyzed in different fractions of elute buffer via
247	the Bradford method using rabbit IgG as a standard.

After this, the measured volume of anti-histamine IgG-tagged liposome nanovesicles was then treated with dropwise addition of 2% BSA solution (in 0.01 M phosphate buffer), followed by 30 min incubation at 4°C to block non-specific binding (Kim et al., 2018). This pre-blocking step was preferable in this study than our previously reported method (Shukla et al., 2016) due to lesser chances of non-specific binding that could be originated from contamination of antibody-conjugated liposome nanovesicles. As illustrated in Fig. 1, phospholipids and cholesterol moieties were used to construct a nano-sized Sulforhodamine 255 B dye encapsulated lipid bilayer (liposome) and conjugated with anti-histamine IgG to form immunoliposome nanovesicles after fractionation and purification using Sephadex and 256 Sepharose gel columns. 257 258 2.4. Characterization of fluorescent anti-histamine IgG conjugated liposomal nanovesicles 259 260 2.4.1. Morphology, size, and stability 261 Liposome nanovesicles morphologies were observed using a JEOL 2100F high-262 resolution TEM. Average diameters, polydispersity indices (PDI), and zeta potentials of 263 liposome nanovesicles were measured by dynamic light scattering (DLS) using a Zetasizer 264 265 Nano ZS particle analyzer (Malvern Instruments Ltd., Worcestershire, UK) at room temperature. Intensities of liposomal suspensions were adequately diluted with 0.02 M 266 Tris-buffered saline (TBS pH 7.0) prior to taking measurements. The PDI and zeta 267 potentials of fluorescent liposome nanovesicles were also used as measures of particle 268 homogeneity and stability. 269

270

271 2.4.2. Confirmation of Sulforhodamine B encapsulation in liposomal lipid bilayers

Lipid and phospholipid molecules can encapsulate Sulforhodamine B, a self-272 273 quenching signaling molecule at higher concentrations leading to formation of Sulforhodamine B-encapsulated liposome nanovesicles; Sulforhodamine B-encapsulation 274 efficiency was determined by measuring increases in fluorescence intensity after rupturing 275 276 lipid bilayers. In brief, 150 mM Sulforhodamine B-encapsulated liposome nanovesicles were treated with a solution of *n*-Octyl- β -D-glucopyranoside (OG) (30 mM) and few other ionic 277 and non-ionic detergent reagents such as Tween 20, Tween 80, Triton X-100, and sodium 278 279 dodecyl sulfate. After the lysis, the fluorescence intensities of released Sulforhodamine B

- from encapsulated liposome nanovesicles were recorded at 550 and 585 nm as excitation and emission wavelengths, respectively, and for control, 0.01 M HEPES buffer (pH 7.0) was used.
- 283 2.4.3. Confirmation for free anti-histamine IgG release via Sepharose CL-4B column

Anti-histamine IgG-tagged liposome nanovesicles were separated from free antihistamine IgG using a Sepharose CL-4B column. The confirmation of free antibody released during elution from Sepharose CL-4B column was analyzed by collecting different fractions of elute buffer after different time intervals and released free antibody concentration was measured via the Bradford method using rabbit IgG as a standard (Shukla et al., 2011).

- 289
- 290 2.5. Concept for detection assay format and sensitivity for histamine detection
- 291

The developed assay functions on the immuno-capturing-based fluorescence 292 controlled detection efficiency of liposome nanovesicles, which act as signal amplifiers. The 293 detection signal capacity of a liposome particle is directly proportional to its size, as 294 described previously (Shukla et al., 2016). The basic concept of the present assay is based on 295 fluorescence measurement before and after rupturing of liposomes followed by leakage of 296 encapsulated SRB dye from the liposome vesicles and finally enhanced fluorescence signals 297 298 are generated as compared to the fluorescence measurement without rupturing. In the present assay method, we simplified and specified the arrangement of liposomal vesicles for 299 generating strong signals, better sensitivity, reduced washing, single well reaction, reduced 300 301 cost, and instant signal generation by using developed immunoliposome vesicles rather than an enzyme-based assay with fluorescence detections. The overall setup for anti-His-LNs-302 based biosensing is demonstrated in Fig. 2. The liposomal nanovesicles containing 303 Sulforhodamine B amplifiers exhibit strong absorption at 500 nm and high fluorescence at 304

550 and 585 nm as excitation and emission wavelengths, respectively (data not shown),
hence these parameters were introduced into our setup to allow the detection of fluorescence
signals. Thus ability of liposome nanovesicles to act as immunosensor for histamine sensing
was assessed by monitoring the change of fluorescence at an excitation wavelength of 550
nm and emission wavelength of 585 nm.

In the present study, an arrangement strategy of fluorescent anti-His-LNs was 310 adopted to enhance the detection efficiency of the developed assay format. In brief, at first, a 311 stock solution of anti-histamine IgG-tagged liposome nanovesicles (anti-His-LNs) was 312 diluted with 0.01 M Tris buffered saline (TBS) containing 0.04 M sucrose at ratios of 1:2, 1:5, 313 1:10 or 1:50 to obtain suitable fluorescence signals in terms of intensities. The 1:10 ratio was 314 deemed optimal and used throughout the remainder of the study. Disruption of the 315 phospholipid bilayer of liposomal nanovesicles was induced by adding detergent. The release 316 of Sulforhodamine B was then detected by analyzing increases and/or decreases in 317 fluorescence intensities. Two assay formats based on two layered anti-His-LNs (anti-His-LNs 318 were mixed with histamine antigen to form immunocomplex, the procedure was done in 2 319 layers) were constructed in an effort to shorten and simplify the detection procedure. 320 Schematics of the both methods are presented in Figures 2A, B and C. 321

Method 1 involves a single well one wash procedure, in which 100 µL solution of 322 various concentrations of histamine (1, 5, 10, 20, 40, 60, 80, 100 or 200 ppb in 0.01 M 323 phosphate buffer) was added in micro-wells of 96 well amine binding polystyrene surface 324 immunoplate, followed by incubation at 37 °C for 15 min and then addition of 50 µL of anti-325 326 His-LNs and incubated for 15 min at 37 °C. These amine binding polystyrene surface 96 microwell plates allow the strong adhesion of proteins and antigenic molecules, including 327 histamine. In brief, these plates were pre-activated via maleic anhydride which allows strong 328 attachment of amine-containing molecules to microplate wells for using in binding assays. 329

Here, our target molecule is a type of biogenic amine (histamine) having -NH₂ functional
group, therefore, similar binding approach is strongly suitable in these plates.

Then, similarly the mixture of 100 μ L of histamine solution and 50 μ L anti-His-LNs 332 as a secondary layer was added to each well followed by similar incubation procedure to 333 allow the formation of immunocomplex (Fig. 2A). Wells were then washed with 0.01 M 334 phosphate buffer and 250 μ L of 30 mM OG was added to lyse the liposome nanovesicles. 335 Histamine bounded Sulforhodamine B encapsulated liposome nanovesicles were then lysed 336 and released. At final step, 200 µL of supernatant solution was transferred in to the new well, 337 and fluorescence signals of Sulforhodamine B, generated after lysis, were measured at 550 338 and 585 nm as excitation and emission wavelengths, respectively. 339

340 On the other hand, another detection assay procedure for Method 2 involved a oneby-one wash procedure (Fig. 2B). Briefly, at first 100 µL of histamine solution of various 341 concentrations (1, 5, 10, 20, 40, 60, 80, 100 or 200 ppb and higher concentrations 0.5-200 342 ppm in 0.01M phosphate buffer) was coated onto micro-wells of 96 well surface 343 immunoplate by incubating the microplate for 30 min at 37° C. Wells were then washed with 344 0.01 M phosphate buffer (3 times), and then 50 µL of anti-His-LNs were added, and wells 345 were incubated at 37 °C for 30 min to initiate the immunogenic reaction. The wells were 346 again washed with 0.01 M phosphate buffer (3 times). Similarly, a second layer of assay 347 348 format was constructed by adding 100 µL of histamine solution and 50 µL of antibody-tagged liposome nanoparticles followed by incubation at 37 °C for 30 min after each addition. Then, 349 the micro-wells were again washed three times with 0.01 M phosphate buffer and then 350 351 encapsulated Sulforhodamine B molecules (in anti-His-LNs) were released by adding 250 µL of 30 mM OG. Finally, 200 µL of lysed liposome solution (supernatant) was transferred in to 352 the new well (to avoiding false interference of any down settled aggregated immunocomplex 353 and errors due to bubbles originated after addition of OG), and signal generation in terms of 354

fluorescence intensity was measured at an excitation wavelength of 550 nm and emission wavelength of 585 nm. The fluorescence intensities of samples and blank were measured as positive and negative values, respectively. The detection results were evaluated for positive/negative (P/N) values, where P/N >2 is considered as a positive result which means histamine is present at detectable level, and P/N < 2 is considered as a negative result which means histamine is either not present or at a very low concentration.

The method for using 2 layers of liposomal vesicles in sensitive detection of 361 enterotoxic Staphylococcus aureus was originated from Yin and Wen (2017). To prove better 362 detection performance, a single layered immuno-liposomal assay format was constructed and 363 compared with current 2 layered immuno-liposomal format. Single layer for immunosensing 364 365 was formed similarly by adding 100 µL of histamine solution to coat onto micro-wells of 96 well amine binding polystyrene surface immunoplate by incubating the microplate for 30 min 366 at 37° C followed by 3 times washing (using 0.01M phosphate buffer). After which 50 µL 367 solution of anti-His-LNs was added to micro-wells followed by incubation at 37 °C for 30 368 min. Wells were then washed with 0.01 M phosphate buffer and fluorescence signals were 369 recorder after the lysis of anti-His-LNs by adding 250 µL of 30 mM OG. For measuring the 370 fluorescence signals (at excitation and emission wavelengths of 550 and 585 nm, 371 respectively), only 200 µL of lysed liposome solution was transferred to the new wells (Fig. 372 2C). 373

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375 2.6. Interference test

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The developed one-by-one assay (Method 2) was tested for its specificity against histamine. Solutions of eight other endogenous standard biogenic amines (HIS, TRP, SPD, PHE, CAD, PUT, TYR, and SPM) were prepared at 10 and 100 ppb concentrations, whereas

their commonly representing corresponding free amino acids (histidine, glycine, alanine, lysine, glutamic acid and arginine) were prepared separately at higher concentration levels (100 ppb) for interference test. The mixture of HIS was prepared at 10 ppb while other interfering amines and free amino acids were prepared at 10×excess concentrations and tested using the above-described analytical procedure. All assay sets were performed in six replicates and % CV values were calculated.

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2.7. Applicability of developed detection assay in contaminated fish, meat and ready to eat
salad products

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390 Histamine is a heat stable amine and is unaffected by high range of temperatures and imposes a great challenge in terms of public health and trade by possessing 391 scombroid fish poisoning. To confirm the applicability of the proposed sensing via Method 2, 392 the content of histamine in fresh mackerel fish, canned tuna and salmon fish, ground red meat 393 and ready to eat salad samples was detected using spiked recovery method. All the fresh 394 395 samples were transferred in hygienic conditions to the laboratory and then exposed under UV for 20-30 min to avoid any further microbial contaminations and further tested using 396 commercially available histamine detection kit (Neogen-veratox for quantification limit 2 397 398 ppm) as well as via currently developed method for the detection of lower levels of histamine (in ppb) in order to confirm the absence of histamine in each set of the sample detection 399 analysis and then spiked with histamine at different concentration levels. 400

All samples (5 g) after confirming the absence of histamine were homogenized (high speed) in 45 mL of 0.01 M phosphate buffer (pH 7.0) and homogenates were then spiked with different concentrations of histamine (1, 5, 10, 20, 40, 60, 80, 100 or 200 ppb) and final volumes made up to 50 mL. After vigorous vortexing, samples were centrifuged at 10,000

405	rpm for 10 min, 1 mL aliquots of supernatant were collected and diluted at 1:10 to reduce
406	matrix effects. In continuations of assay procedure as ascribed in earlier section for method 2,
407	collected supernatant from histamine contaminated real samples was coated onto micro-wells
408	of 96 well amine binding polystyrene surface immunoplate by incubating the microplate for
409	30 min at 37° C. These specific immunoplates have the high adhesion and easy attachment
410	capacity of amine binding polystyrene surfaces for histamine antigen present in the samples.
411	Three independent sets of spiked samples were prepared at each concentration for statistical
412	purposes.
413	
414	2.8. Assay validations
415	
416	The detection procedure was validated via calibration curve and evaluation of the
417	range of linearity, limit of detection (LOD), and limit of quantification (LOQ). The linear
418	response of histamine was determined in the concentrations, which led to the correlation
419	factor $R^2 > 0.99$. LOD and LOQ were re-calculated using the standard equations LOD = X0 +
420	3SD and $LOQ = X0 + 5SD$, respectively, where X0 was the average response of the blank
421	samples, and SD referred to the standard deviation for $n = 6$.
422	
423	3. Results and discussion
424	
425	3.1. Preparation and characterization of liposome nanovesicles
426	
427	Sulforhodamine B-encapsulated liposome nanovesicles were developed using a

428 reverse-phase method as previously described (Shukla et al., 2016) with minor modifications.

429 In brief, liposome nanovesicles were filtered through 0.8 and 0.4 μ m polycarbonate filters in

430 order to achieve uniform size distributions (Fig. 1D), which were measured by dynamic light scattering at 209.2 nm with 100% intensity and 0.965 intercept value (Fig. S1). Average 431 volume of single liposome nanoparticle calculated using average size of mixed liposome 432 vesicles is presented in Table S1. On the basis of the average size determined via dynamic 433 light scattering, it could be possible to calculate the average outer volume of single liposome 434 vesicle as 4.71×10^{-12} µL and inner volume entrapped (by assumption 4 nm lipid bilayer 435 thickness) as $4.18 \times 10^{-15} \,\mu$ L. The Sulforhodamine B content inside the liposome was assumed 436 437 as equal to Sulforhodamine B concentration used during the liposome preparation (150 mM), and other characteristics such as amount of Sulforhodamine B per liposome (6.28×10^{-13}) 438 μ mol), number of Sulforhodamine B molecules per liposome (3.78×10⁵) were calculated by 439 440 comparing the fluorescence of lysed liposomes to that of standard Sulforhodamine B solution. Amount of lipid / liposome and number of lipid molecules per liposomes were calculated 441 considering the outer surface area = $4\pi r^2$ with the assumption that only 50% of total lipid 442 molecules are present in liposome vesicle. Similarly, anti-histamine IgG molecules tagged per 443 liposomal vesicles were calculated by assuming that only 0.4% IgG was conjugated that 444 resulted in 2200 molecules of anti-histamine IgG on the surface of each liposomal vesicle. 445 Prior to antibody conjugation, the concentration of developed liposomes was calculated by 446 assuming that the Sulforhodamine B concentration inside liposomes was 150 mM. All 447 calculations were dependent on Sulforhodamine B standard curve and calculated as 448 449 encapsulated Sulforhodamine B concentrations after lysis and measured via fluorescence intensity (at an excitation wavelength of 550 nm and emission wavelength of 585 nm) for the 450 total released SRB concentration, as a result, which was calculated as 8 µmol/mL of liposome 451 solution. Based on the information given in Table S1, we calculated the µmol concentration 452 of SRB/each particle of liposome as 6.28×10^{-13} . Therefore, the concentration of liposome 453 (particles/mL) was approximately 1.27×10^{13} particles/ mL. All the calculations were 454

455 designed on the bases of our earlier reported protocols (Shin and Kim, 2008).

Previous studies have shown that fluorescence detection signal capacity of a 456 liposome nanoparticle is dependent to its particle size (Shin and Kim, 2008). Based on this 457 information, we assumed that as the size of the surface area of a liposome particle increased 458 the binding affinities of antibody molecules to its surface was enhanced. Thus, we sought to 459 prepare liposome nanovesicles with high surface areas to enhance surface antibody binding. 460 TEM observations confirmed 190-200 nm nanovesicles were spherical (Fig. 3A). Also, Shin 461 and Kim (2008) reported that as the size of the liposome increased, higher fluorescence 462 signals were obtained owing to a higher number of Sulforhodamine B molecules 463 encapsulated in the liposome. Overall, as the liposome particle size increased, the detection 464 465 signal increased.

Zeta potential and polydispersity index (PDI) are important parameters that reflect 466 the stabilities of nano-systems, and for Sulforhodamine B-encapsulated liposome 467 nanovesicles as determined by DLS their values were -42 ± 7.72 mV, and 0.191, respectively, 468 which confirmed interaction between Sulforhodamine B molecules and negatively charged 469 phospholipids (Table S1). Liposomes prepared with lower PDIs (< 1) are considered 470 monodisperse liposome vesicles, and have excellent stabilities (Shukla et al., 2011). The 471 magnitude of the zeta potential indicates the potential stability of the colloidal system. High 472 surface charges cause particles to repel each other, and thus, increase solution stability. Li et 473 al. (2016) reported similar results for liposome nanovesicles developed for targeted drug 474 delivery. 475

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Sulforhodamine B was successfully encapsulated in liposome nanovesicles and

⁴⁷⁷ *3.2. Confirmation of Sulforhodamine B encapsulation in liposomal lipid bilayers*

480 served as a strong fluorescent sensing material. To confirm the encapsulation efficiency of liposomes, results were confirmed based on the measurements of the fluorescence intensity of 481 Sulforhodamine B-encapsulated liposomes before and after lysis. In addition, we also tested 482 the abilities of a series of ionic and non-ionic detergents (Tween 20, Tween 80, Triton X-100, 483 sodium dodecyl sulfate, and OG) to rupture the membranes of Sulforhodamine B-484 encapsulated liposomes (Fig. 3B). OG at a concentration of 30 mM was found to most 485 effectively rupture lipid membrane as determined by transmission electron morphological 486 views (Fig. 3C) and Sulforhodamine B fluorescence intensities (Fig. 3D). 487

Fluorescence intensities of lysed and non-lysed liposomal nanovesicles rely on the 488 de-quenching of putatively self-quenched Sulforhodamine B (Ho et al., 2007). In order to 489 determine the concentration dependency of Sulforhodamine B release from liposomes 490 nanovesicles and generation of fluorescent signals by lysis of liposomal nanovesicles. 491 Liposome nanovesicles at different dilutions $(1:10^3, 1:10^4, 1:10^5)$ and $1:10^6)$ were treated with 492 30 mM of OG as a strong detergent, and fluorescence intensity signals were measured at 493 excitation and emission wavelengths of 550 and 585 nm, respectively. We found that 30 mM 494 OG dose-dependently induced Sulforhodamine B release, and thus, was chosen for further 495 tests (Fig. 3D). Overall our results showed Sulforhodamine B was well encapsulated by 496 liposomal vesicles, which exhibited only weak fluorescence intensity, but strong fluorescence 497 intensity after lysis, indicative of fluorescent dye leakage from vesicles (Saez et al., 1982). 498

Further, after conjugation of anti-histamine IgG to Sulforhodamine B-encapsulated liposome nanovesicles, the free antibodies were removed following the purification step via Sepharose CL-4B column. The confirmation of released free antibody was validated using different interval fractions of elute buffer by the Bradford method using rabbit IgG as a standard. During conjugation, initial concentration of anti-histamine IgG was 0.1 mg/mL while 0.021 mg/mL of free antibody release was observed during purification of anti-

505 histamine IgG from different fractions of buffer via Sepharose CL-4B column (Table S3), confirming that the free antibody was easily removed when fractioned via Sepharose CL-4B 506 507 column. The optimization of anti-histamine useful amount for the conjugation with liposomal nanovesicles was tested with various concentrations of anti-histamine IgG antibody to 508 achieve its significant bounding efficiency confirmed via Bradford test. Finally, 0.1 mg/mL 509 concentration was confirmed as the useful amount of anti-histamine IgG to form 510 immunoconjugate (Table S2). The experiment was allowed to perform in several lots during 511 the synthesis procedure. 512

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514 *3.3. Design and detection sensitivity of developed assay*

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Under optimized conditions, immunosensor assay formats (Methods 1 and 2) 516 constructed in this study based on the immunoliposome nanovesicles were applied to detect 517 different concentrations of histamine in a buffer system (Fig. 4A and 3B). Both methods 518 produced dual layer liposome-based immunocomplex with the requirement of 200 µL of 519 520 analyte (histamine) and 100 µL of anti-His-LNs. Method 1 was involved a single well one wash procedure, in which a second layer of immunocomplex was formed with anti-His-LNs 521 and was mixed together with single step washing followed by fluorescence measurements by 522 523 the release of Sulforhodamine B contents. On the other hand, Method 2 involved the multiple washing and incubation steps typical of ELISA into a one-by-one procedures as described in 524 the Experimental section in which first analyte (histamine antigen) was coated with proper 525 526 incubation time followed by 3 times washing and then the first layer of anti-His-LNs was constructed. After proper incubation period, the attached immunocomplex was washed and 527 again allowed to construct a second layer of analyte and anti-His-LNs followed by washing. 528 Method 1 and 2 resulted in LINs that exhibited concentration-dependent fluorescence 529

intensities (Fig. 4A and 4B). Lower concentration of histamine allowed the formation of antigen (histamine)–antibody (anti-His-LNs) complex only in a few amount, and large amount of anti-His-LNs remained unbounded, which can be easily washed off, and after lysis reaction, resulting in a lower range of fluorescence intensity which was observed in an increasing manner as the concentration of histamine increased (Fig. 4A and 4B).

Detection limits were defined as average fluorescence intensities of blank samples 535 plus-minus three standard deviations (Hochel and Skvor, 2009). In brief, according to the 536 537 equation: LOD = xb1 + 3sb1, where xb1 is average blank signal and sb1 is blank standard deviation. The detection limit of histamine for Method 1 was found to be 10 ppb and only 1 h 538 15 min was required to obtain signals (Fig. 4A). While, Method 2 (one by one wash) had a 539 detection limit of 2-3 ppb but 2 h 30 min was required to obtain signals (Fig. 4B). The 540 detection limit of Method 2 was preferred, presumably due to stronger analyte to anti-His-541 LNs binding due to the longer reaction time. In Method 1, the "prozone effect" was observed 542 at a histamine concentration of 80 ppb, and this may have been due to competitive reactions 543 between anti-His-LNs and higher number of histamine molecules which can be considered as 544 545 a saturation point (Fig. 4A). On the other hand, Method 2 showed minute reduction in fluorescence signals at a histamine concentration of 200 ppb (Fig. 4B); however, on an 546 increment of higher histamine concentrations (0.5 to 200 ppm), fluorescent signals again 547 increased drastically and no saturation or "prozone effect" was observed (Fig. S4), possibly 548 due to the use of multiple washes in between the arrangements of anti-His-LNs complex 549 offering a heterogeneous assay design (Vaidya et al., 1988). 550

Based on these results, although Method 1 could be implemented more easily and quickly because it did not require multiple steps of Method 2, we adopted Method 2 because it had a lower detection limit (2–3 ppb) and a quantification limits in a wider range (8.5 ppb– 200 ppm) than Method 1 (15–80 ppb). We believe that the two layered anti-His-LNs based

sensing format (Method 2) exhibited good bio-specificity and sensitivity- the emitted fluorescence was acquired from the detection of anti-histamine IgG-conjugated liposomal amplifiers that formed strong layers of immuno-complexes with the target analyte and the immuno-liposomal complex. This may be attributed to the inherited nature of the anti-His-LNs as an immunosensor which specifically captured the analyte molecule. Moreover, the appropriate blocking strategy also prevented non-specific binding between the detection surface and the target antibody molecules or liposomal amplifiers (Chang et al., 2016).

In a previous study, Shukla et al. (2012) used liposomes in a single layer format for 562 the detection of Salmonella bacterium. Here, we compared a single layered format involving 563 multiple washing stages similar with Method 2 (two layered format) using the same amount 564 565 of histamine (200 µL) and of liposome nanovesicles (100 µL) and measured detection sensitivities (Fig. 2B). We found the detection limit of histamine in single layered format was 566 60 ppb and that signals were detected in 1 h 30 min (Fig. 4C). This outcoming demonstrated 567 an immense interest in achieving better detection sensitivities by approaching a two layered 568 format of liposome nanovesicle-based diagnostic system. Finally, we compared the analytical 569 performance of Method 2 with other diagnostic methods developed for histamine detection 570 (Table 1). Method comparisons demonstrated that the sensor developed in this study has a 571 wider linear range and a lower detection limit, and does not require the complex handling 572 573 steps of previously reported sensors.

To check storage stability, fluorescence intensities of synthesized Sulforhodamine Bencapsulated liposome nanovesicles were measured after storage at 4 °C for 6 and 12 months. Significant changes in fluorescence intensity were observed after 3 months and after 12 months of storage fluorescence intensities were much reduced (Fig. S2). Therefore, in order to achieve long term stability and avoid damage of fluorescent liposome nanovesicles, we recommend the conjugation of anti-histamine IgG to fluorescent liposome nanovesicles

should be performed promptly after synthesis. Therefore, after conjugation of liposomal nanovesicles with anti-histamine IgG (immunoliposomal nanovesicle), we evaluated the stability with storage at 4 °C for 0 month to 20 months, and observed that the stability of immunoliposomal nanovesicles in terms of fluorescence intensity was not reduced until 12 months, which proves their significant stable shelf-life with sufficient encapsulation of SRB dye. However, a drastic reduction in the fluorescence intensity was noted after 12 and 14 months storage under the same conditions.

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588 *3.4. Specificity of the developed assay formats*

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In order to measure the specificity of Method 2, cross-reactivity was examined 590 versus 8 other biogenic amines (HIS, PUT, PHE, SPM, CAD, TRP, SPD, and TYR) that 591 belong to same group of amine contaminants in food products. All the examinations were 592 performed under the experimental conditions described in subsection: 2.5. Positive and 593 negative tests were conducted using fluorescence intensities and P/N values. P/N values of 594 many immunoassays provide an important means of determining positive and negative test 595 results (Saez et al., 1982). The P/N value of histamine was 7.49 ± 0.2 (>2), whereas for the 596 other biogenic amines tested, P/N values were all <2, which indicated the absence of cross 597 reactivity with other amine contaminants. The P/N value of spermidine was 2.4 ± 0.1 , 598 indicating slight interference (Fig. 5a). This might be due to close chemical interaction of 599 spermidine moieties or a few uncovered binding sites present in anti-His-LNs. The specificity 600 601 of the developed anti-His-LNs-based assay (Method 2) is probably determined by the specificity of the antibody used (Dong et al., 2017). 602

603 The selectivity was also investigated with the correspondent free amino acids found 604 mostly in fish, meat and vegetables as histidine, glycine, alanine, lysine, glutamic acid and

605 arginine, account for major proportion of the total amino acid contents. The histidine amino acid is the main interfering factor for histamine determination because the % presence of free 606 607 amino acids in meat and fish products is lesser than 1%. Fig. 5b shows the fluorescence intensity response of these free amino acids along with histamine and found the peak 608 fluorescence intensity of histamine was minimum 3 times larger at 1/10th concentration of 609 other tested free amino acids, approved for no interfering with other structural analogs. Also 610 when a mixture of all free amino acids (100 ppb) mixed with histamine at 10 ppb, 611 fluorescence intensity of histamine was slightly reduced as compare to single histamine (10 612 ppb) but was analyzed statistically different (p < 0.05), confirming acceptable specificity of 613 the proposed sensing. Few other electrocatalytic electrodes based sensing methods were 614 found to be affected by interfering analogs including histidine free amino acid (Wang et al., 615 2017; Gajjala and Palathedath, 2018). 616

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618 3.5. Applicability of the developed assay using real fish and meat samples

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In order to assess the applicability of the developed anti-His-LNs-based assay 620 (Method 2), fish food products including fresh mackerel fish and canned tuna/salmon fish 621 samples were artificially spiked with corresponding histamine concentrations. The histamine 622 623 detection limit was found to be same as in buffer medium (2–3 ppb), but fluorescence signals were slightly reduced (Fig. 6), indicating slight matrix effects, presumably due to high protein, 624 fat, or omega fatty acid contents present in fish samples (Omanovic-Miklicanin and Valzacchi, 625 626 2017). Further, average recoveries were also tested with detectable concentrations observed from fresh mackerel 73.50% to 99.98%, canned tuna 79.08% to 103.74% and 74.56% to 627 99.02% from canned salmon, respectively. The precision and accuracy of each test set were 628 expressed as % CVs ranging from 1.48% to 6.86% (Table 2). These analytical figures (% CV) 629

were consistent with the results of other reported detection approaches, indicating that themethod was reliable for the analysis of real samples (Hu et al., 2017; Taghdisi et al., 2016).

Solution obtained (after spiking of histamine into real food samples) was diluted in the ratio of 1:10 to reduce matrix effects following the USFDA guidelines and coated onto the amine group binding polystyrene 96 well immunoplates. At this stages along with histamine, a few other proteins present in food samples might also be coated onto the surfaces of immunoplate. However, the developed assay had no interfering effect on the histamine specific fluorescence signal generation, thus confirming the specificity of currently developed anti-His-LNs based assay (Method 2).

Moreover, histamine detection with good recovery was also confirmed in few other 639 640 relevant food matrices such as ground red meat and ready to eat salad. The results of an acceptable value of % histamine recoveries were found as 106.43% and 99.00% in ground 641 red meat and ready to eat salad, respectively are provided in Table S5. The total assay time of 642 the developed assay setup for all the samples was noted as 2 h 30 min, including coating, 643 blocking and washing procedures. This assay setup favors that multiple samples (96 samples 644 645 at one time) can be handled within the same time limit. These results revealed a good sensing applicability of the developed one-by-one anti-His-LNs-based assay (Method 2) in a multi-646 well biosensing system for a variety of food matrices without having any recovery loss and 647 no demand for sophisticated purification methods, such as extraction-based liquid 648 chromatography methods. 649

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651 3.6. Method comparison with conventional HPLC technique and liposome-based
652 immunomagnetic separation assay

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Although several analytical detection methods, including HPLC, GC, RP-HPLC, and

655 TLC have been reported earlier (Awan et al., 2008; Lapa-Guimarães and Pickova, 2004) for the detection of histamine, all these methods require sample pre-treatment step, derivatization 656 prior to sample injection, and able to analyze only single sample at a time. However, the 657 sensing method developed in this study does not have any requirement of pre-treatment, 658 sample derivatization and able to analyze multiple samples at the same time. Further, to 659 approve the novelty, accuracy and practical applicability of the developed anti-His-LNs-660 based assay (Method 2), fish food products, including fresh mackerel and canned tuna/salmon 661 fish samples were tested for measuring the histamine content by the proposed method and 662 high performance liquid chromatography (HPLC) method. 663

To confirm the sensitivity, rapidity, and straightforwardness of the sample pre-664 665 treatment procedure without using expensive reagents and derivatizing reagents, results were verified using a conventional HPLC method. We previously found, histamine had to be 666 derivatized with dansyl or benzovl chloride for HPLC detection (Shukla et al., 2011; Shukla 667 et al., 2014), as fish samples spiked with different biogenic amines at different concentrations, 668 including histamine, could not be analyzed without derivatization. Therefore, same fish 669 samples were extracted with 0.4 M perchloric acid and derivatized with dansyl chloride 670 followed by HPLC analyses. The results from HPLC chromatograms confirmed lower 671 histamine recovery rates than currently developed anti-His-LNs-based method (Table S4, Fig. 672 673 S3). HPLC techniques for the determination of several hazardous biogenic amines, including histamine, require sample extraction and derivatization (2–3 h), and a sample run time of 30 674 min. The assay procedure developed in the present study does not require any pre-treatment 675 676 or sample derivatization step, and was able to detect histamine in multiple samples (at least 96 samples) directly within 2 h 30 min. 677

Furthermore, histamine concentrations of < 1 ppm could not be easily detected in a
variety of food matrices by HPLC technique, which is its major limitation versus nano-based

detection techniques. Erim (2013) reported that HPLC technique can be used to quantify a single biogenic amine in a sample, and by HPLC, limit of detection of 0.02 ppm has been reported for histamine in fish products. Pradenas et al. (2016) confirmed and validated an improved HPLC technique in different food matrices for the detection of histamine with a limit of detection ranging from 0.5–20 ppm.

In addition, we have also compared the novelty, and better detection limits of anti-685 His-LNs-based method (Method 2) with other liposome-based immunomagnetic 686 concentration and separation assay (IMS) (Shukla et al., 2016). As a result, although the total 687 assay time was similar (2 h 30 min) as observed for the anti-His-LNs-based method, in 688 histamine buffer medium the detection limit was found as 10-20 ppb with quantification limit 689 690 of 10-50 ppb (Fig. 7). The practical applicability of liposome-based immunomagnetic concentration and separation assay was also compared in real fish food sample spiked with 691 different concentrations of histamine. As a result, although direct detection without sample 692 pre-treatment was confirmed, the detection and quantification limits were very poor as 20 ppb 693 and 20-50 ppb, respectively than currently developed anti-His-LNs-based assay (Method 2), 694 695 which indicates that the developed assay (Method 2) could be used efficiently for the routine analysis of histamine in real food samples than other similar detection methods. 696

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698 4. Conclusions

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A fluorescence quenching immunosensing format was developed to measure the concentration of histamine toxin in contaminated fish samples. The developed liposome amplified sensing assay (Method 2) shown preferable detection limit of 2–3 ppb with a wide range for quantification limit (8.5 ppb –200 ppm). Validation of the developed assay procedure (Method 2) was confirmed by histamine detection in artificially contaminated real

705 fish samples where a similar detection limit (2-3 ppb) was achieved within 2 h 30 min of total 706 assay time for multiple samples (96 samples at one time), including coating, blocking and 707 washing procedures without use of extraction and derivatization steps than that of conventional analytical detection methods such as HPLC. The histamine recovery rates form 708 contaminated fish samples were in the range of 73.50%-103.74% whereas repeatability 709 results as % CV were in the range of 5.34%-8.48%. In summary, the liposome-based 710 detection assay developed in this study is simple, rapid, and cost effective, and does not 711 712 require any extra pre-treatment steps. Moreover, it has great potential in a wide range of universal diagnostic applications to determine toxin analytes in real fish samples, including 713 other similar foods. 714 715 **Author contributions** 716 717 The manuscript was written through contributions of all authors. All authors have 718 given approval to the final version of the manuscript. 719 720 **Declaration of interests** 721 722 723 None. 724 Acknowledgments

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897 **Table 1**

898 Comparison of the developed liposomal vesicle-based detection assay for histamine with 899 other detection methods.

Detection base	Mechanism	Limit of detection (LOD)/ Quantification range	Reference
Silver colloid SERS substrate	Surface Enhanced Raman Spectroscopy	Linear range: 0-20 mg/kg	Janc [°] I et al., 2017
Cy5 de labeled oxido-reductases	Fluorescent based on quenching by Forster resonance energy transfer	LOD: 13 nM Linear range: 13 nM–25 nM	Gustiananda et al., 2012
Molecularly imprinted polymer film	Surface plasmon resonance sensor	Linear range: 25 µg/L – 1000 µg/L	Jiang et al., 2015
Graphene-based nano composite film with HRP (horse radish peroxidase)	Competitive electrochemical immunosensor	LOD: 0.5 pg/mL Linear range: 1 pg/mL– 1 ng/mL	Yang et al., 2015
Nanoporous alumina membranes with magnetic nanoparticles	Impedance based assay	LOD: 1 µM Linear range: 1 µM–40 nM	Ye et al., 2017
Carbon black nanoparticles	Colorimetric based chip immunoassay	0-600 µg/mL	Mattsson et al., 2017
Cu@Pd core shell nanostructures	Chrono amperometry	0.32±0.1 nM	Gajjala and Palathedath, 2018
Commercial ELISA based Kit	Enzyme linked immunosorbent assay	LOD: 2 ppm Linear range: 2.5–40 ppm	Commercial KIT (Neogen Veratox)

Gold nano-particles	Immunochromatographic test strip	LOD: 600 ng/mL (6 mg/kg)	Luo et al., 2015
Screen-printed carbon electrode and the enzyme diamine oxidase	Amperometric sensor	LOD: 0.94 mg/L	Torre et al., 2019
Superparamagnetic particle label	Magnetic immunochromatographic test	LOD: 1.2 mg/L	Moyano et al., 2019
Liposome-based immunomagnetic concentration and separation assay	Fluorescence quenching by Sulforhodamine B dye	LOD: 10–20 ppb Linear range: 15–50 ppb	Compared with current method
Immunoliposomal quenching assay	Fluorescence quenching by Sulforhodamine B dye	LOD: 2–3 ppb Linear range: 8.5 ppb– 200 ppm	This work
	JU		

901 Table 1 Continued

902

Table 2

905 Recovery test results for histamine in fish based food samples (ppb, n = 10).

Fish Food matrices	Spiked concentration (ppb)	Recovered concentration	Average recovery (%)	% CV
	2	1 47 + 1 13	73 50 %	4 98
	5	4.13 ± 1.33	82.6 %	3.44
Fresh mackerel	10	9.16 ± 1.21	91.6 %	3.09
fish	20	18.78 ± 0.80	93.9 %	3.33
	40	39.10 ± 0.54	97.75 %	2.15
	60	59.99 ± 1.81	99.98 %	2.28
	2	1.58 ± 0.10	79.08 %	6.86
	5	4.47 ± 0.07	89.45 %	1.67
	10	9.47 ± 0.37	94.76 %	4.02
Canned tuna	20	19.21 ± 0.28	96.08 %	1.48
	40	39.20 ± 1.50	98.00 %	3.83
	60	62.24 ± 2.86	103.74 %	4.31
	2	1.52 ± 0.01	76.56 %	4.40
	5	4.25 ± 0.07	85.07 %	1.80
	10	9.36 ± 0.08	93.66 %	1.67
Canned salmon	20	18.93 ± 0.72	94.67 %	3.94
	40	39.47 ± 1.05	98.69 %	2.72
	60	59.41 ± 1.20	99.02 %	2.05

908 Figure captions

909

910	Fig. 1. Preparation of the fluorescent liposomal nanovesicles as signal amplifiers. (A) DPPC,
911	DPPG and cholesterol dissolved in a chloroform/methanol mixture was used to form a thin
912	bilayer film by reverse phase evaporation; (B) Lipid bilayer hydration with aqueous
913	Sulforhodamine B at 45 °C, (C) Sulforhodamine B-encapsulated in a gel with lipid bilayered
914	vesicles; (D) Liposome nanovesicles were extruded through polycarbonate filters (0.4 and 0.8
915	μ M); (E) Fractionation using Sephadex G-50 gel for collecting Sulforhodamine B-
916	encapsulated fluorescent liposomal nanovesicles; and (F) Conjugation of anti-histamine IgG
917	into fluorescent liposomal nanovesicles with followed by fractionation using Sepharose CL-
918	4B gel for collecting immuno-liposomal nanovesicles with significant fluorescent efficiency.
919	
920	Fig. 2. Schematic representation for designed multiplexed-optic fluorescent liposomal

921 nanovesicle-based dual layered immuno-biosensing for histamine toxin.

922

Fig. 3. Characterization of synthesized liposome nanovesicles. (A) Morphological 923 examination of Sulforhodamine B-encapsulated fluorescent liposomal nanovesicles by 924 transmission electron microscopy (TEM); (B) Comparison of using ionic and non-ionic 925 detergents for rupturing the lipid membranes of Sulforhodamine B-encapsulated fluorescent 926 liposomal nanovesicles in terms of released fluorescent Sulforhodamine B dye; (C) TEM 927 images of Sulforhodamine B-encapsulated fluorescent liposomal nanovesicles before (left 928 929 side image) and after rupturing (right side image) of lipid membrane; and (D) Confirmation of the integrity of 150 mM Sulforhodamine B-encapsulated fluorescent liposomal 930 931 nanovesicles.

Fig. 4. Sensitivities of the developed double layered positioned anti-His-LNs-based biosensing formats for the detection of histamine. (A) Double layered positioned anti-His-LNs based single well one wash format; (B) Double layered positioned anti-His-LNs based one by one wash format; and (C) Single layered anti-His-LNs-based format. All experiments were conducted three times, and results are presented as means±SDs. The coefficient of variation (% CV) of fluorescence intensity (n=6) was less than 10%.

939

Fig. 5. Specificity and background interference test of the developed double layered positioned anti-His-LNs-based biosensing for the detection of histamine (A) with coexisting biogenic amines at 10 ppb and 100 ppb concentrations; and (B) with corresponding free amino acids at 100 ppb and histamine at 10 ppb. All experiments were conducted three times, and results are presented as means±SDs. The coefficient of variation (% CV) of fluorescence intensity (n=6) was less than 15%.

946

947 Fig. 6. Performance of the immunoliposomal amplified signals based sensing on spiked food 948 matrices (fresh fish, canned fish, ground meat and ready to eat salad) for the detection of 949 histamine. All experiments were conducted three times, and results are presented as 950 means±SDs. The coefficient of variation (% CV) of fluorescence intensity (n=6) was less 951 than 10%.

952

Fig. 7. Comparison of double layered positioned anti-His-LNs-based biosensing assay with
other liposome-based immunomagnetic concentration and separation assay (IMS method) for
the detection of histamine.



Fig. 1.



Fig. 2.



Jonua

Fig. 3.





Fig. 5.



Fig. 6.



Fig. 7.

Highlights

- Scombroid fish poisoning has raised concerns due to histamine related environmental toxicity
- Double layered positioned liposomal vesicles as fluorescent probe
- Anti-histamine IgG conjugated liposomal nanovesicles (anti-His-LNs)-based multiplexed biosensing system
- Anti-His-LNs-assisted rapid, sensitive and cost-effective detection of histamine toxic molecule in fish foods
- Designed sensing platform diverse the application of various ELISA-based complicated commercial detection kits

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Author contributions

The manuscript was written through contributions of all authors. All authors have given approval to the final version of the manuscript.

Journal Prevention

Declaration of interests

 \boxtimes The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

□The authors declare the following financial interests/personal relationships which may be considered as potential competing interests:

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