Research Paper

A High Throughput Protein Formulation Platform: Case Study of Salmon **Calcitonin**

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Purpose. The feasibility of using high throughput spectroscopy for characterization and selection of physically stable protein formulations was studied.

Materials and Methods. A hundred aqueous formulations of salmon calcitonin (sCT) were prepared using 20 buffer compositions. The solutions had pH values between 2.5 and 10.5. The stability of the sCT formulations was analyzed over 1 week by the following assays: (1) protein concentration, (2) volume control by measuring pathlength, (3) turbidity (absorbance at 350 nm), (4) intrinsic tyrosine fluorescence, (5) 1-anilino-naphthalene-8-sulfonate (ANS) fluorescence, (6) Nile Red fluorescence. Addition of the dyes (Nile Red and ANS) was used to study protein conformational changes.

Results. After 1 day, 27 out of the 100 formulations of salmon calcitonin were stable. After 7 days, 12 stable sCT formulations remained. The best salmon calcitonin formulation was in 10 mM sodium acetate buffer with pH values between 3.5 and 5.5.

Conclusions. The findings are in accordance with the sCT formulations that were patented and used commercially. This can be considered as a proof of concept for the high throughput protein formulation platform.

KEY WORDS: high throughput screening; protein formulation; salmon calcitonin; spectroscopy; stability.

INTRODUCTION

Calcitonin is a naturally occurring polypeptide hormone involved in the regulation of calcium and bone metabolism. Calcitonin is produced in parafollicular cells of the human thyroid gland [\(1](#page-10-0)). In patients with osteoporosis, calcitonin is used to increase the bone density, prevent bone loss and fractures. In addition, calcitonin reduces acute pain due to osteolytic fractures, vertebral crush fractures, and osteolytic bone metastasis [\(2](#page-10-0)). Clinically, calcitonin is indicated for the treatment of postmenopausal osteoporosis and Paget's disease of the bone, either

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ABBREVIATIONS: ADA, N-(2-acetamido)iminodiacetic acid; ANS, 1-anilino-naphthalene-8-sulfonate; BSA, bovine serum albumin; CAPSO, 3-(cyclohexylamino)-2-hydroxy-1-propanesulfonic acid; EMEA, European Agency for the Evaluation of Medicinal Products; FDA, Food and Drug Administration; HEPES, 4-(2-hydroxyethyl) piperazine-1-ethanesulfonic acid; HTF, high throughput formulation; HTS, high throughput screening; IU, International units; MOPS, 3 morpholinopropane sulfonic acid; sCT, salmon calcitonin; TES, N-[tris (hydroxymethyl)methyl]-2-aminoethanesulfonic acid.

alone or in combination with other drugs. Calcitonin is only indicated for women who are at least 5 years beyond menopause, or for women who cannot take alternative medication [\(3](#page-10-0)).

Commercially, salmon calcitonin (sCT) is produced synthetically or by using recombinant DNA techniques. The first salmon calcitonin product (Calcimar®) was approved in 1975 by the Food and Drug Administration (FDA). A human calcitonin product (Cibacalcin®) was approved by the FDA in 1986. Salmon calcitonin consists of 32 amino acids and differs from human calcitonin by 13 residues. At present, in the USA, several salmon calcitonin formulations are approved for parenteral (50–100 IU daily, preferably by subcutaneous or intramuscular injection) or nasal (200 IU daily) administration. A list of the marketed salmon calcitonin drug products, approval date, manufacturers, route of administration and excipients is given in Table [I.](#page-1-0) The specific activity of sCT was defined as having 6,000 IU/mg in the Third International Standard by the World Health Organization ([4\)](#page-10-0). The commercial formulations contain either 200 (injection) or 2,200 IU/ml (multi-dose nasal spray), which corresponds to sCT concentrations of 0.033 and 0.37 mg/ml, respectively. sCT is formulated with an acetate buffer (sCT international standard, Calcimar®, Miacalcin® for injection, Forcaltonin®), hydrochloric acid (Miacalcin nasal spray) or citric acid (Fortical®).

By 2009, about \$10 billion worth of biopharmaceuticals will have lost patent protection [\(5\)](#page-10-0). New protein formulations that are better or bioequivalent to existing protein drugs can be marketed as 'biogenerics' after the patent has expired. An increase in 'biogenerics' and improved biopharmaceutical

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The used sCT concentration, excipients, concentration, dosage forms and storage conditions are given. The data were collected from the Food and Drug Administration (FDA), European Agency The used sCT concentration, excipients, concentration, dosage forms and storage conditions are given. The data were collected from the Food and Drug Administration (FDA), European Agency for the Evaluation of Medicinal Products (EMEA) and prescribing information of drug manufacturers. for the Evaluation of Medicinal Products (EMEA) and prescribing information of drug manufacturers.

formulations, supergenerics ([6](#page-10-0)), can be expected which would benefit from a fast drug development and formulation screening platform. In order to achieve these goals a high throughput formulation (HTF) platform was developed. The HTF platform was based on an advanced liquid handling system and a UV absorbance/fluorescence microplate reader to study the physical and chemical stability of proteins and peptides. The HTF platform can help in selecting protein stabilizing excipients for further protein formulation optimization. The use of multi-well plates allows the formulation scientist to screen for a variety of parameters, excipients and conditions using a relatively small amount of protein. As a proof of concept of the HTF platform, the sCT stability in 100 buffer solutions was characterized. The buffer solutions, prepared from 20 buffer types, have pH values between 2.5 and 10.5, and were selected based on their chemical diversity.

MATERIALS AND METHODS

Materials

The chemicals required for preparing the buffer solutions, and the dyes 1-anilino-naphthalene-8-sulfonate (ANS) and Nile Red, were provided by Sigma-Aldrich (Sigma-Aldrich Chemie GmbH, Buchs, Switzerland). The 20 10 mM buffer solutions (Tables [II](#page-2-0) and [III](#page-3-0)) consisted of: (1) glycine–hydrochloric acid, (2) citric acid–sodium citrate, (3) citric acid–sodium phosphate dibasic, (4) potassium phthalate–sodium hydroxide, (5) succinic acid–sodium hydroxide, (6) ammonium acetate–acetic acid, (7) histidine–hydrochloric acid, (8) sodium acetate–acetic acid, (9) β,β′-dimethyl glutaric acid–sodium hydroxide, (10) Tris (hydroxymethyl)aminomethane–maleate, (11) sodium phosphate monobasic–sodium phosphate dibasic, (12) N-(2-acetamido)iminodiacetic acid (=ADA)–hydrochloric acid, (13) citric acid–potassium phosphate dibasic, (14) 3-(N-morpholino)propanesulfonic acid (=MOPS)–potassium hydroxide, (15) 4-(2-hydroxyethyl)-1-piperazine-1-ethanesulfonic acid (=HEPES)–sodium hydroxide, (16) N-[tris(hydroxymethyl) methyl]-2-aminoethanesulfonic acid (=TES)–hydrochloric acid, (17) Tris(hydroxymethyl)aminomethane–hydrochloric acid, (18) glycine–sodium hydroxide, (19) 3-(cyclohexylamino)-2 hydroxy-1-propanesulfonic acid (=CAPSO)–sodium hydroxide, (20) sodium carbonate–sodium bicarbonate.

The salmon calcitonin was obtained from Novartis (Novartis AG, Basel, Switzerland). UV transparent 96-well Costar® (no. 3635) Corning (Corning Inc., New York, NY, USA) microplates were supplied by Vitaris (Vitaris SA, Baar, Switzerland). UV-Vis transparent and pressure sensitive Greiner VIEWseals™ (Greiner Bio-One GmbH, Frickenhausen, Germany) were acquired from Huber & Co. AG (Reinach, Switzerland). Sterile and single wrapped polystyrene reservoirs with 12 channels (no. 330.12.9) from Socorex (Socorex Isba S.A., Ecublens, Switzerland) contained in total 40 ml and were supplied by Milian (Milian S.A., Geneva, Switzerland). A Millipore (Millipore Corp., Bedford, Massachusetts) filtration system, comprised of a vacuum driven disposable bottle top (Stericup™) with 0.22 μm Durapore™ filter and a presterilized receiver flask (Steritop™), was used for the filtration of all non-protein solutions.

	10mM Buffers									pH								
			3.0	3.5	4.0	4.5	5.0	5.5	6.0	6.5	7.0	7.5	8.0	8.5	9.0	9.5	10.0	10.5
$\mathbf{1}$	$Glycine -$ Hydrochloric acid	$\mathbf U$ N \mathbf{A}	$\overline{\mathbf{U}}$ N $\overline{\mathbf{A}}$	\overline{U} $\mathbf N$ \mathbf{A}														
$\overline{2}$	Citric $acid$ – Sodium citrate		$\overline{\mathbf{U}}$ N $\overline{\mathbf{A}}$	$\overline{\mathbf{U}}$ $\mathbf N$ \mathbf{A}	$\overline{\mathbf{U}}$ $\overline{\mathbf{N}}$ \mathbf{A}	$\overline{\mathrm{U}}_\mathrm{N}$ $\overline{\mathbf{A}}$	$\overline{\mathrm{U}}_\mathrm{N}$ \mathbf{A}	$\overline{\mathbf{U}}_{ \mathbf{N}}$ $\overline{\mathbf{A}}$	$\overline{\mathrm{U}}_\mathrm{N}$ ${\bf A}$									
3	Citric $acid -$ Sodium phosphate dibasic		$\overline{\mathbf{U}}$ N A	\overline{U} N \mathbf{A}	$\overline{\mathbf{U}}$ $\mathbf N$ \mathbf{A}	\overline{U} A $\overline{\mathbf{N}}$	$\overline{\overline{U}}$ A $\overline{\mathbf{N}}$	$\overline{\mathbf{U}}$ $\overline{\mathbf{N}}$ $\overline{\mathbf{A}}$	$\frac{1}{\sqrt[n]{N}}$ \mathbf{A}	$\overline{\text{U}}_{\text{N}}$ $\boldsymbol{\rm A}$	$\overline{\mathrm{U}}_\mathrm{N}$ $\overline{\mathbf{A}}$	$\frac{U}{A}N$						
$\overline{\mathbf{4}}$	Citric $acid$ – Potassium phosphate dibasic		$\overline{\mathbf{U}}$ N $\overline{\mathbf{A}}$	$\overline{\mathbf{U}}$ ${\bf N}$ $\overline{\mathbf{A}}$	$\overline{\mathbf{U}}$ $\mathbf N$ $\overline{\mathbf{A}}$	$\overline{\overline{U}}$ A $\overline{\mathbf{N}}$	$\overline{\overline{U}}$ A $\overline{\mathbf{N}}$	$\overline{\mathbf{U}}$ $\overline{\mathbf{N}}$ $\overline{\mathbf{A}}$	\overline{U} $\overline{\mathbf{N}}$ $\overline{\mathbf{A}}$	$\overline{\mathbf{U}}$ $\overline{\mathbf{N}}$ $\overline{\mathbf{A}}$	$\overline{\mathbf{U}}$ ${\bf N}$ $\overline{\mathbf{A}}$							
5	β , β '-Dimethylglutaric acid – Sodium hydroxide			\overline{U} ${\bf N}$ \mathbf{A}	$\overline{\mathbf{U}}$ $\mathbf N$ \mathbf{A}	$\overline{\mathbf{U}}$ $\overline{\mathbf{N}}$ \mathbf{A}	$\overline{\mathbf{U}}$ $\overline{\mathbf{N}}$ $\overline{\mathbf{A}}$	$\overline{\text{U}}$ $\overline{\text{A}}$ $\overline{\mathbf{N}}$	$\frac{1}{\sqrt[n]{\mathbf{U}}(\mathbf{V})}$ $\overline{\mathbf{A}}$	$\overline{(\overline{U}}_{N})$ \mathbf{A}	$\overline{\mathbf{U}}$ ${\bf N}$ \mathbf{A}	$\overline{\mathrm{U}}_\mathrm{N}$ $\overline{\mathbf{A}}$						
$\overline{\mathbf{6}}$	Acetic $acid -$ Sodium acetate			$\overline{\mathbf{U}}$ $\mathbf N$ $\overline{\mathbf{A}}$	$\overline{\mathbf{U}}$ $\mathbf N$ $\overline{\mathbf{A}}$	\overline{U} \overline{A} $\overline{\mathbf{N}}$	$\overline{\mathbf{U}}$ $\overline{\mathbf{N}}$ \mathbf{A}	\overline{U} $\overline{\mathbf{N}}$ $\overline{\mathbf{A}}$										
$\overline{7}$	Potassium phthalate - Sodium hydroxide				$\overline{\mathbf{U}}$ $\overline{\mathbf{N}}$ $\overline{\mathbf{A}}$	$\overline{\mathbf{U}}$ $\frac{1}{\Delta}$ N	$\frac{1}{\overline{U}}_{N}$ $\overline{\mathbf{A}}$	$\frac{\mathbf{U}}{\mathbf{A}}$ $\overline{\mathbf{N}}$	\overline{U}_{N} $\mathbf A$									
8	Succinic acid- Sodium hydroxide				$\overline{\mathbf{U}}$ $\overline{\bf N}$ $\overline{\mathbf{A}}$	$\overline{\mathbf{U}}$ $\overline{\mathbf{N}}$ $\overline{\mathbf{A}}$	$\frac{\mathbf{U}}{\mathbf{A}}\mathbf{N}$	\overline{U} A $\overline{\mathbf{N}}$	$\frac{1}{\sqrt{10}}$ _N $\overline{\mathbf{A}}$									
$\boldsymbol{9}$	Acetic \overline{acid} – Ammonium acetate					$\overline{\mathbf{U}}$ \mathbf{A}^{N}	$\overline{\mathbf{U}}$ ${\bf N}$ \mathbf{A}	$\overline{\mathbf{U}}$ $\overline{\mathbf{N}}$ $\overline{\mathbf{A}}$	\overline{U} \overline{A}^{N}	$\mathbf U$ $\frac{3}{\Delta}N$								
10	Histidine- Hydrochloric acid							$\overline{\mathbf{U}}$ $\overline{\mathbf{N}}$ $\overline{\mathbf{A}}$	\overline{U} $\overline{\mathbf{N}}$ $\overline{\mathbf{A}}$	\overline{U} $\overline{\mathbf{N}}$ $\overline{\mathbf{A}}$	${\bf U}$ $\overline{\mathbf{N}}$ $\overline{\mathbf{A}}$	${\bf U}$ $\overline{\mathbf{N}}$ $\overline{\mathbf{A}}$						
11	$Tris -$ Maleate							$\overline{\mathbf{U}}$ $\frac{1}{\Delta}N$	$\frac{1}{\sqrt{2}}$ _N $\overline{\mathbf{A}}$	$\overline{\overline{U}}_{N}$ $\overline{\mathbf{A}}$		$\frac{\mathbf{U}}{\mathbf{A}}\mathbf{N}\frac{\mathbf{U}}{\mathbf{A}}\mathbf{N}$	$\overline{\mathrm{U}}_\mathrm{N}$ $\overline{\mathbf{A}}$					
12	Sodium phosphate monobasic - Sodium phosphate dibasic								$\overline{\mathbf{U}}$ $\overline{\underline{A}}^{N}$	$\frac{1}{ U }N$ A	$\overline{\mathbf{U}}$ $\overline{\mathbf{N}}$ $\overline{\mathbf{A}}$	$\overline{\overline{U}}_N$ A	$\overline{\mathbf{U}}$ $\overline{\mathbf{N}}$ \mathbf{A}					
13	$ADA -$ Hydrochloric acid								$\overline{\mathbf{U}}$ $\overline{\bf N}$ $\overline{\mathbf{A}}$	$\overline{\mathbf{U}}$ $\overline{\bf N}$ \mathbf{A}	$\overline{\mathbf{U}}$ ${\bf N}$ $\overline{\mathbf{A}}$							
14	$MOPS -$ Potassium hydroxide									\overline{U} $\mathbf N$ $\overline{\mathbf{A}}$	$\overline{\mathbf{U}}$ $\mathbf N$ \mathbf{A}	$\mathbf U$ $\mathbf N$ \mathbf{A}						
15	$HEPES -$ Sodiumhydroxide										$\frac{\mathbf{U}}{\mathbf{A}}$ ${\bf N}$	$\overline{\mathbf{U}}$ $\mathbf N$ $\overline{\mathbf{A}}$	$\frac{\mathbf{U}}{\mathbf{A}}\mathbf{N}$					
16	$TES -$ Hydrochloric acid										$\overline{\mathbf{U}}$ \mathbf{N} $\overline{\mathbf{A}}$	$\overline{\textbf{U}}_{\hspace{0.5mm}\textbf{N}}$ $\overline{\mathbf{A}}$	$\overline{\overline{\mathbf{U}}}$ N $\overline{\mathbf{A}}$					
17	$Tris -$ Hydrochloric acid											$\overline{\mathbf{U}}$ N $\overline{\mathbf{A}}$	$\overline{\mathbf{U}}$ $\overline{\mathbf{N}}$ \overline{A}	$\overline{\mathbf{U}}_{\mathbf{N}}$ $\overline{\mathbf{A}}$	$\frac{\mathbf{U}}{\mathbf{A}}\mathbf{N}$			
18	$Glycine -$ Sodium hydroxide														\overline{U} $\frac{1}{\Delta}N$	$\begin{array}{c c}\n\mathbf{U} & \mathbf{N} \\ \hline\n\mathbf{A} & \mathbf{N}\n\end{array}$	$\frac{\mathbf{U}}{\mathbf{A}}\mathbf{N}$	$\begin{array}{c c}\n\textbf{U} & \textbf{N} \\ \hline\n\textbf{A} & \textbf{N}\n\end{array}$
19	$CAPSO -$ Sodium hydroxide														$\overline{\mathtt{U}}$ $\overline{\mathbf{N}}$ \mathbf{A}	$\frac{1}{\left U\right _N}$	$\mathbf{U}_{\mathbf{N}}$ $\overline{\mathbf{A}}$	
$\overline{\mathbf{20}}$	Sodium carbonate- Sodium bicarbonate															$\overline{\mathbf{U}}$ $\overline{\mathbf{N}}$ $\overline{\mathbf{A}}$	\overline{U} $\overline{\mathbf{N}}$ $\overline{\mathbf{A}}$	$\begin{array}{c c}\n\mathbf{U} & \mathbf{N} \\ \hline\n\mathbf{A} & \mathbf{N}\n\end{array}$

Table II. The Table Summarizes the Results of the Turbidity, ANS and Nile Red Fluorescence Assays After 1 Day

The pH values between pH 2.5 and 10.5, with 0.5 pH unit intervals, are highlighted for the 20 buffer solutions used. Each assay was assigned a U, A, N which represents the UV turbidity assay, ANS fluorescence emission and Nile Red fluorescence emission, respectively. The samples found stable with the respective assays have a white background and unstable a grey background.

Preparation of Protein Formulations

A stock solution of 8.0 mg/ml salmon calcitonin in purified MilliQ™ Millipore water was freshly prepared. A hundred 40 mM buffer solutions with pH values between 2.5 and 10.5 (Table II) were prepared using sterile filtered 0.20 M buffer solutions. The stability of salmon calcitonin was tested at a concentration of 2.0 mg/ml. This concentration is higher than the 0.033 and 0.37 mg/ml sCT concentrations that are used in the marketed sCT drug products. Previous work showed that with an increase in calcitonin concentration a decrease in fibrillation time was determined [\(7\)](#page-10-0). It was assumed that at a 2.0 mg/ml sCT concentration during a reasonable amount of time the different aggregation properties would be distinguishable.

The sCT formulations were prepared in 96-well plates with a Sciclone ALH500 liquid handling station from Caliper Life Sciences (Hopkinton, USA). The sample preparation protocol using the liquid handling station was composed of the following steps: (1) 100 or 98 μl (if a dye was added) of water was dispensed using the bulk tool, (2) 50 μl of buffer was pipetted using the Z-8 liquid handling device with disposable tips, (3) 50 μl of sCT solution was added, similar to the dispensing of the buffer solutions, (4) 2 μ l of a dye, 0.20 mM ANS or 0.10 mM Nile Red, was simultaneously added to all 96 wells using a low-volume 96-canula array. The

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Each assay was assigned a U, A, N, T which represents the UV turbidity assay, ANS emission, Nile Red emission and tyrosine emission, respectively. The samples found stable with the respective assays have a white background and unstable a grey background. The selected stable sCT formulations complied with all four assays.

100 test formulations were composed of 2.0 mg/ml sCT in 10 mM buffer and were prepared in three variants; with 50 μM ANS, with 1 μM Nile Red and with neither. Each 96 well plate contained 20 sCT formulations (each formulation in quadruplicate), four sCT samples in water, ten buffer control solutions, one purified MilliQ water and one empty well (an example of a prepared plate is given in Fig. [2](#page-5-0)A). If 2 μl dye (ANS or Nile Red) was added to the microplate, 2 μl less water was used. The control solutions included all buffer types. The outer columns, 1 and 12, contained no sCT in buffer since in these columns evaporation is more likely to occur [\(8\)](#page-10-0). For each sCT formulation three microplates were

prepared and analyzed, making a total of 15 UV transparent Corning Costar® 96-well plates: sCT in buffers, sCT in buffers with ANS and sCT in buffers with Nile Red. After sample preparations, the plates were sealed with a Greiner VIEWseal™ and incubated at 25°C in the dark during 7 days.

High Throughput Spectroscopic Analysis

sCT is a polypeptide hormone with a molecular weight of 3,600 Da and contains 32 amino acids, one of which is the fluorophore tyrosine. The sCT fluorescence emission and absorbance was measured with a Tecan (Tecan Group Ltd.,

Männedorf, Switzerland) Safire™ microplate reader. The fluorescence was measured using the bottom optics. The fluorescence emission of tyrosine was determined between 295 and 400 nm. Excitation was at 274 nm using bandwidths for both excitation and emission of 7.5 nm. The sCT formulations plus ANS were excited at 378 nm and the emission was monitored between 460 and 520 nm (examples shown in Fig. [3A](#page-6-0) and B). The fluorescence emission of Nile Red in the presence of sCT was studied between 610 and 670 nm after excitation at 575 nm (examples shown in Fig. [3](#page-6-0)A and B). The fluorescence emission of ANS and Nile Red was analyzed with two different settings, a high and low detection sensitivity in order to characterize both strong and weak interactions between the dyes and sCT. The turbidity of sCT formulations was measured by studying the absorbance at 350 nm [\(9\)](#page-10-0). The volume or pathlength was controlled by monitoring the absorbance of water at 975 nm [\(10\)](#page-10-0). The concentration of sCT was calculated and the sample preparation controlled by measuring the UV absorbance at 274 nm [\(11\)](#page-10-0). The fluorescence and absorbance was measured at day 0, 1 and 7. The data output of the $XFluor^@$ software of the Tecan Safire™ microplate reader is in Microsoft® Excel™. Visual basic macros were developed by the authors to enable high throughput data analysis.

RESULTS

Spectroscopy

The physical stability of 2.0 mg/ml sCT was studied in 100 buffer solutions. One of the selection criteria for a stable sCT formulation is to have an absorbance value below 0.075 at 350 nm. All sCT formulations had absorbance values at 350 nm between 0.04 and 0.06 directly after sample preparation as well as after 1 day, which were similar to the absorbance values of the control solutions. Many sCT formulations at neutral pH values became turbid after 7 days, in contrast to sCT formulations at acidic and basic pH values. Some sCT formulations aggregated and formed different structures, such as gels with varying amount of transparency, white clumps or dots. In some cases a formulation contained protein aggregates that were attached to the sides of wells. Since the absorbance measurement is done in the center of the well, no increase in absorbance signal was detected. A visual inspection of all wells was performed to check for the presence of small aggregates.

The intrinsic tyrosine fluorescence decreased between day 0 and 1 from about 34,000 (relative fluorescence units) to about 24,000 RFU (see Supplementary Fig. 1). This effect was attributed to an increase in the solution temperature between day 0 and 1. The sCT formulations were prepared with buffer solutions that were stored at 4°C. The sCT formulations at pH values above 9 showed a decrease in intrinsic fluorescence intensities, caused by the lower absorbance energy of tyrosine due to deprotonation. The lower fluorescence intensities observed in potassium phthalate and Tris–maleate buffer are ascribed to the UV absorbance of these buffer solutions. The sCT formulations with a less than 6% increase or decrease in tyrosine fluorescence intensity between day 1 and 7 were considered stable. Changes in sCT stability were better distinguished, compared to the

intrinsic tyrosine assay, after addition of ANS or Nile Red. Only sCT in glycine–HCl pH 3.5 had a strong deviation in intrinsic fluorescence, with no increased ANS or Nile Red fluorescence.

The ANS fluorescence emission spectra were analyzed for each salmon calcitonin formulation. The maximum fluorescence emission of ANS was found between 480 and 490 nm. In contrast to the Nile Red fluorescence emission, the ANS maximum fluorescence emission did not shift when bound to the salmon calcitonin. Therefore, the average ANS emission at 485 nm was used for the sCT formulation evaluation (see electronic supplementary material). The measured ANS intensities differed with pH and buffer type, indicating different affinities of ANS towards the formed salmon calcitonin structures. The highest ANS fluorescence intensities were observed between pH 5 and 8.

In some formulations, binding of Nile Red to salmon calcitonin resulted in a blue shift in the fluorescence emission maxima from about 645 to about 620 nm and with a respective increase in fluorescence intensity of ~150 to 55,000 RFU (Fig. 1). Nile Red emission spectra of 20 sCT formulations measured after 1 day in a 96-well plate were recorded with a high sensitivity and are shown in Fig. [2](#page-5-0)B. The samples in the corresponding 96-well plate are shown in Fig. [2](#page-5-0)A. Measurements using high sensitivity settings are useful for fast screening of sCT formulations. However, they do not always offer detailed information on the maximum emission wavelength. Flattening of some of the fluorescence emission curves (Fig. [2B](#page-5-0)) was caused by saturation of the detector. The maximum emission intensities with corresponding wavelengths were automatically extracted from the Nile Red emission spectra recorded with low sensitivity settings. A three-dimensional representation of the emission wavelength maxima and corresponding fluorescence intensities of sCT in a 100 buffer solutions, measured 7 days after preparation, is shown in Fig. 1. The sCT formulations with Nile Red fluorescence emission maxima higher than 630 nm and average intensities lower than 300 were considered stable according to the Nile Red assay. The intensity of 300

Fig. 1. The Nile Red fluorescence after 7 days in 100 sCT formulations is shown. The black values are in three-dimensional space. Binding of Nile Red to sCT resulted in a blue shift from 645 to 620 nm and a fluorescence increase of 150 to 55,000. The stable formulations are encircled.

Fig. 2. A, B 20 sCT formulations (columns 2 to 11) and the respective controls (columns 1 and 12) are given on the top, in A. Asterisk is an example of the description of the well (sCT control in water) with the corresponding fluorescence intensity. Four replicates per formulation were prepared. Nile Red emission spectra after 7 days, using high sensitivity settings, are shown on the bottom, in **B**. The sCT in sodium citrate and sodium citrate phosphate pH 3.0 and 3.5 and in glycine pH 2.5 to 3.5 remained stable.

corresponds to twice the average value of Nile Red in presence of the control solutions. Shifts in the emission maxima of Nile Red greater than 15 nm are significant. Therefore a limit of 630 nm for the emission maximum of Nile Red was used as one of the selection criteria for obtaining stable sCT formulations.

Two buffer solutions, in which the salmon calcitonin was found stable and unstable, are sodium acetate and sodium citrate. The average of the four endpoint measurements plus the ANS and Nile Red fluorescence spectra of the four replicates per formulation are shown in Fig. [3](#page-6-0)A and B. An increase in the extrinsic fluorescence intensity is an indication of protein instability, such as aggregation. After 7 days, salmon calcitonin in sodium citrate pH 3.0 and 3.5 had an increased turbidity monitored at 350 nm (Fig. [3A](#page-6-0)). However, no change was observed in any of the three fluorescence assays (Fig. [3](#page-6-0)A). No change in ANS fluorescence was observed for sCT in sodium citrate pH 4.0, although an increased Nile Red fluorescence and increased absorbance at 350 nm were measured (Fig. [3](#page-6-0)A). All sodium acetate sCT formulations with pH values of 3.5 to 5.5 were found stable according to all four assays (Fig. [3B](#page-6-0)). The Nile Red fluorescence intensity decreased in time, which was also observed for other stable sCT formulations. This decrease can be related to the adsorption of Nile Red to the walls of the 96-well plate.

A summary of the ANS and Nile Red fluorescence and absorbance at 350 nm after 1 day of sample preparation is given in Table [II.](#page-2-0) After 1 day, an increased ANS and Nile Red fluorescence was observed for most of the sCT formulations with pH values higher than 4.5. The results after 7 days, regarding the turbidity at 350 nm, intrinsic tyrosine fluorescence, ANS fluorescence and Nile Red fluorescence assays are summarized in Table [III](#page-3-0). The absorbance spectra, tyrosine fluorescence, ANS fluorescence and Nile Red fluorescence emission spectra were analyzed for each of the sCT formulations and the average results are shown in electronically available supplementary material.

The overview Tables [II](#page-2-0) and [III](#page-3-0) and the following examples highlight the necessity of using several complemen-

Fig. 3. A The UV absorbance at 350 nm, tyrosine intrinsic fluorescence emission at 308 nm, ANS emission intensity at 485 nm and Nile Red maximum fluorescence intensity measured after 0 (filled circles), 1 (empty circles) and 7 days (x marks) are given. The ANS and Nile Red fluorescence spectra of sCT in sodium citrate after 0, 1 and 7 days are shown of the four replicates per pH value. The fluorescence intensity axes were optimized for each day. The sCT was unstable in sodium citrate with pH between 4 and 6. **B** The UV absorbance at 350 nm, tyrosine intrinsic fluorescence emission at 308 nm, ANS emission intensity at 485 nm and Nile Red maximum fluorescence intensity measured after 0 (filled circles), 1 (empty circles) and 7 days (x marks) are given. The ANS and Nile Red fluorescence emission of sCT in sodium acetate after 0, 1 and 7 days remained low, indicating no binding of the dyes to the protein occurred. A decrease in Nile Red fluorescence after 1 day was observed. The sCT was found to be stable in sodium acetate with pH values between 3.5 and 5.5 during at least 1 week.

tary methods for the selection of stable protein formulations. Salmon calcitonin in glycine–HCl buffer had after 7 days no increase in ANS and Nile Red fluorescence intensity. However, an increased turbidity (pH 2.5, 3.0 and 3.5) and variation in the intrinsic tyrosine fluorescence (pH 3.5) were measured. Salmon calcitonin was more stable at low pH values in potassium citrate phosphate than in sodium citrate phosphate. Analysis of the individual ANS fluorescence spectra at pH 3.0 revealed that the strong ANS fluorescence is caused by one out of four replicates. A strong variation in extrinsic fluorescence intensities was observed with sCT in potassium citrate phosphate, sodium citrate phosphate and sodium citrate solutions with pH values lower than 6. This is probably due to the still on-going aggregation of tree or branched-like structures that formed in time a protein gel. The sCT formulations in the same buffer solutions at pH values of 6 and higher formed, in less than a week, turbid homogeneous gels. These observations were visible with the eye.

Salmon calcitonin was found to be unstable at pH values between 4.0 and 6.0 in sodium succinate, because of an increased turbidity at 350 nm. However, the ANS and Nile Red fluorescence intensities remained low for pH 4.0 and 4.5 (Table [III](#page-3-0) and online available supplementary material). Salmon calcitonin formulations in ammonium acetate buffer pH 5.0 and higher had an increased ANS fluorescence, which was not observed for sCT in sodium acetate (Table [III](#page-3-0) and Fig. 3). The sCT aggregated in histidine solutions according to the four assays at all measured pH values. The use of sodium phosphate resulted in increased absorbance values and changes in intrinsic and extrinsic fluorescence intensities of sCT.

Sodium acetate

Fig. 3. (continued)

After 7 days, no suitable buffer solutions were found for sCT that had pH values of 6.0 or more. The salmon calcitonin turbidity at 350 nm in buffer solutions 13 to 20 (Table [III\)](#page-3-0) was low, except for ADA pH 6.0, HEPES pH 7.0 and sodium carbonate pH 10.5, where slightly increased turbidities were measured. The ANS fluorescence increased in all these solutions, where the Nile Red fluorescence intensities remained low for the following formulations: MOPS–KOH pH 6.5, HEPES pH 7.0 and 7.5, glycine–NaOH pH 9.0. Salmon calcitonin remained stable for 1 day at pH values similar to the physiological pH in the following buffer solutions: MOPS–KOH pH 6.5 and 7.5, HEPES pH 7.5, TES pH 7.0.

DISCUSSION

The proof of concept of a high throughput protein formulation platform is presented in this paper using salmon calcitonin as a model protein. The sCT physical stability was studied in 100 formulations with the following assays: (1) protein concentration, (2) volume control by measuring pathlength, (3) turbidity (absorbance at 350 nm), (4) intrinsic tyrosine fluorescence, (5) ANS fluorescence, (6) Nile Red fluorescence. The intrinsic tyrosine fluorescence assay led to the discarding of about 15 formulations that were turbid by eye. In the case of salmon calcitonin, intrinsic fluorescence is less sensitive to detect aggregation as extrinsic (1,8-ANS or Nile Red) fluorescence. Assays to control for the dispensed volume, possible evaporation and protein concentration of the samples were performed. In our experiments, no evaporation occurred during the 7 days of storage, although a reduction in pathlength was observed for sCT formulations that formed a compact and hard gel.

The described high throughput platform can be enlarged with techniques to facilitate accelerated stability testing or other complementary methods to characterize the physical (e.g. fluorescence anisotropy, fluorescence lifetime and nephelometry) and chemical (e.g. capillary electrophoresis, liquid chromatography and mass spectrometry) stability [\(12](#page-10-0)). The combination of high throughput turbidity measurements with accelerated stability tests, by means of mechanical stress, was shown to be useful in protein formulation ([13\)](#page-10-0). However, high throughput turbidity screening of protein solutions often is hindered by the requirement of high protein concentrations and the high variation in absorbance signal ([14\)](#page-10-0). This variation can be caused due to the relatively small diameter $(\emptyset \sim 0.75$ mm) of the light beam, which corresponds to less than 1% of the surface of a 96 well $(\emptyset \sim 9.0 \text{ mm})$, and the irregular growth of protein aggregates. Measurement at multiple points inside a well will reduce the variation in the turbidity signal. However, a more complicated data analysis and increased measurement time can be expected.

The fibrillation lag time of calcitonin is known to decrease with an increase in concentration ([7,](#page-10-0) [15,](#page-10-0) [16](#page-10-0)). Aggregated salmon calcitonin formed different structures as observed by eye, which is reflected in the different fluorescence and absorbance intensities. The absorbance at 350 nm was not sensitive enough to detect all forms of sCT aggregates, in particular at pH values of 7.0–10.5. If the turbidity assay would have been used as a stand-alone assay, several unstable sCT formulations would have been selected. The use of complementary fluorescence assays narrowed down the selection of stable salmon calcitonin formulations.

The fluorescence of ANS is known to increase after binding to hydrophobic regions of proteins [\(17](#page-10-0)). ANS binds primarily through electrostatic interactions of the ANS anion to the cationic groups of the protein ([18\)](#page-10-0). The binding of dyes to proteins that expose hydrophobic surfaces does not always lead to increased fluorescence intensity. No fluorescence increase was observed for up to 17 ANS anions that were bound in acidic pH to lysozyme [\(18\)](#page-10-0). BSA below pH 5.0 has as many as 100 ANS binding sites, where 95% of the protein bound ANS is non fluorescent ([18\)](#page-10-0). Similar effects may explain the low ANS or Nile Red fluorescence intensities of sCT in glycine–HCl buffer pH 2.5 to 3.5. Another explanation could be that at acidic pH, ANS is less charged and therefore it interacts with protein aggregates mainly through hydrophobic interactions rather than electrostatic interactions. Another ANS binding study, together with circular dichroism and intrinsic fluorescence analysis, characterized the different conformational states of β-lactamase between pH 2.0 and 12.0 [\(19\)](#page-10-0). The ANS fluorescence was stable in the entire pH range, except for a time-dependent increase of ANS fluorescence at pH 2.0 ([19\)](#page-10-0). In our high throughput study, no ANS fluorescence intensity increase caused by pH of buffer solutions was observed. The ANS fluorescence augmented in all sCT formulations with pH 6.0 or higher. This is attributed to the binding of ANS to the hydrophobic regions formed in sCT multimers and aggregates.

The fluorescence of Nile Red is strongly influenced by the polarity of its environment. Nile Red can interact with many proteins, is photostable and can detect the exposure or formation of hydrophobic surfaces [\(20](#page-10-0), [21\)](#page-10-0). The Nile Red fluorescence was shown in several studies to be unaffected at a variety of pH ranges: 4.5–8.5 [\(20](#page-10-0)), 6.0–10.0 ([22\)](#page-10-0). We measured a significant increase in Nile Red fluorescence emission in three salmon calcitonin formulations at pH 4.0 (sodium citrate, sodium citrate phosphate and potassium phthalate) and up to pH 10.5 compared to the control solutions. These Nile Red fluorescence intensities in the salmon calcitonin formulations were significantly higher than the Nile Red in the control buffer solutions. The Nile Red fluorescence intensity in stable sCT formulations decreased after 1 day, which is caused by slight temperature differences, as was the case for the intrinsic tyrosine, or by adsorption to walls of the microplate. Nile Red adsorbed on plastic, glass, or quartz becomes non-fluorescent ([20](#page-10-0)).

Spiking of protein formulations with the dye prior to analysis is preferred over the incorporation of the dye in the protein formulation [\(21\)](#page-10-0). Dyes might alter the protein conformation, as it has been previously reported that ANS induced folding of acid unfolded cytochrome c to the molten globule state as a result of electrostatic interactions [\(23](#page-10-0)). The fluorescence intensity of the added dyes can be influenced by changes in protein conformation or by incubation. Spiking of the formulation with the dye prior to each analysis time point can be performed. This is in particular useful when studying the stability of protein formulations over long time periods (months to a year) or when the formulations are stored at higher temperatures. However, we do not recommend spiking samples with dyes in a high throughput screening study of up to one month if no interference of the dye with the protein stability was detected. Nile Red did not change the stability of salmon calcitonin ([21\)](#page-10-0) and the dye was added once at time zero.

After 1 day of measurement, 27 from a total of 100 sCT formulations (Table [II](#page-2-0)) were found to be stable according to the turbidity, ANS and Nile Red fluorescence assays. After 7 days of sample preparation, based on the results of the turbidity, intrinsic tyrosine fluorescence, ANS fluorescence and Nile Red fluorescence assays, only 12 sCT formulations remained stable (Table [III](#page-3-0)). The formulations, with a complete white background in Table [III](#page-3-0), are: sodium citrate phosphate pH 3.0; potassium citrate phosphate pH 3.5 and 4.0; dimethyl glutaric acid pH 3.5, pH 4.0 and 4.5; sodium acetate pH 3.5, 4.0, 4.5, 5.0 and 5.5; ammonium acetate pH 4.5. The sodium acetate was the best buffer solution for a sCT drug formulation, since the protein remained stable at all measured pH values, even at pH 5.5.

The Forcaltonin®, Miacalcin® and Calcimar® commercial sCT solutions for injection contain sodium acetate (Table [I](#page-1-0)), which confirms with the results of our high throughput formulation study. The respective incorporation of hydrochloric acid and citric acid in Miacalcin® nasal spray and Fortical® nasal spray is another indication that sCT is stable in an acidic environment. However, salmon calcitonin is not generally stable at low pH values (e.g. glycine–HCl pH 2.5–3.5). The stabilizing effect of sodium acetate on salmon calcitonin is due both to sodium and acetate, since the use of sodium citrate and ammonium acetate at pH 5.0 and 5.5 resulted in an increased ANS fluorescence.

The selection of stable protein formulations depends on the set criteria per assay. These criteria need to be continuously evaluated during data analysis. Data analysis can be a time-consuming task. For example, the ANS and Nile Red emission were measured in their emission maxima over a 60 nm range and the intrinsic tyrosine over 40 nm to reduce measurement time and to facilitate data handling. With these settings and including the different absorbance assays, about 180,000 data points were acquired per day out of a total of 15

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microplates. Automated determination of average values, including standard deviations and variations of endpoint measurements (absorbance or emission value at a single wavelength) gave an indication about the reproducibility and can help in finding wells with variations in e.g. the protein concentration and volume. Analysis of absorbance and emission spectra is required before excluding samples (e.g. individual wells) from the study, as is explained further. Different graphical representation of data is possible, as can be seen in Figs. [1](#page-4-0), [2](#page-5-0), [3](#page-6-0) and the electronically available supplementary material. The fluorescence emission spectra can either be analyzed with or without the plotting of spectra. In-house build visual basic macros selected automatically the maximum intensity values with corresponding wavelengths. These values changed depending on the interaction of the dye with the protein. Line graphs or three-dimensional graphs, such as Fig. [1](#page-4-0), help to quickly understand the behavior of the protein (e.g. fluorescence intensity) in relation to the parameter tested, which was the case of the pH. Plotting of 96 fluorescence emission spectra per analyzed microplate in time showed the protein behavior per buffer and in relation to the position in the plate. Detection of a consequent error in sample pipetting and thus in concentration can be found, since in our HTF platform the protein samples are horizontally dispensed in the microplate by eight individual pipettors. Generally, evaporation starts to occur in the outer columns and rows of a well plate, which can be detected by visualizing spectra per plate.

In our high throughput study the average values of four replicates per sCT formulation were used. In several cases, one out of these four replicates could have been designated in the fluorescence assays as an outlier. However, no such statistical approach was applied and all replicates contributed to the reported data. In the ANS fluorescence emission spectra at day 0 of sCT in glycine–HCl pH 3.0, one sample showed a six times stronger fluorescence intensity than the other three replicates. After a day, this strong fluorescence decreased and became similar to the fluorescence of the replicates. Presence of air bubbles, droplets, or other technical issues could be the explanation for this increased value at day 0. Salmon calcitonin in potassium citrate phosphate buffer pH 3.0 is stable to all but the ANS emission assay. This is unexpected since sCT in potassium citrate phosphate pH 3.5 and 4.0 complied with all four assays. One out of the four replicates of sCT in potassium citrate phosphate pH 3.0 had a strong ANS fluorescence at day 0, 1 and 7. No experimental reasons could be found to eliminate this sample, which resulted in salmon calcitonin being not stable in potassium citrate phosphate buffer pH 3.0. In other formulations, one out of four sCT samples showed an increased initial fluorescence or absorbance that continued to increase in time. The remaining replicates showed, only after a prolonged time, similar spectroscopic changes caused by the irregular aggregation formation in some buffers. Automated data analysis can help pinpoint these outliers.

The assay criteria were established based on the intensities of the background values and variation in stable sCT formulations and control solutions. For each assay, the sCT formulations were awarded a score (white or grey background), either stable or not according to each assay, and presented in overview tables (see Tables [II](#page-2-0) and [III](#page-3-0)). Stable protein formulations can then be selected for further optimization, analysis or immediate use. We suggest investigating the origin of the changes if the protein formulation is considered stable with three out of four assays.

With controlled storage conditions (e.g. humidity and temperature), high throughput analysis can be performed over more than a week. There are no limitations to the length of the high throughput screening study. The formulations can be stored also outside the microplate in for example vials or deep-well storage plates. At each stability time point, the formulations can be pipetted (with or without an extrinsic dye) in a 96-well plate for characterization. A decrease in total measurement time (e.g. decrease in protein aggregation lag time) and increase in throughput can be obtained when combining accelerated stress testing, e.g. shaking or storage at elevated temperatures, with high throughput spectroscopy ([13\)](#page-10-0).

The screening of sCT formulations over a broad pH range resulted in several leads for further formulation optimization, one of which included the commercially used sodium acetate buffer. The HTF platform can be applied for fast developing stable formulations of other peptides and proteins. The HTF platform can be used with extrinsic dyes at low protein concentration $\left($ <50 μg/ml) and at high protein concentration (e.g. 100–300 mg/ml). The analytical methods have to be adapted to these extreme concentration conditions. Phenomena such as protein binding to walls at low protein concentration or high viscosity at high protein concentration have to be taken into account.

It is our opinion that, if enough protein substance is available, many parameters need to be studied to completely understand the physical and chemical properties of the drug. In this endeavor, high throughput screening platforms will provide a valuable tool to characterize protein formulations.

CONCLUSION

After 1 week, 12 out of 100 salmon calcitonin formulations were found to be stable. The physical stability study was performed in 96-well plates and the two tested parameters were buffer type and pH value. The sodium acetate buffer at pH values of 3.5 to 5.5 was the best candidate for further sCT formulation optimization. The sCT remained stable according to the turbidity, intrinsic and extrinsic fluorescence assays. It should be noted that several commercial sCT formulations contain a sodium acetate buffer. The leads found using our novel high throughput methods have already been used or patented, which validates our approach and can be considered as a proof of concept for the high throughput protein formulation platform.

The use of several complementary analytical methods permits the selection of formulations using carefully selected assay criteria. We have shown that, in some cases, an increase in turbidity was observed without an increased ANS or Nile Red fluorescence. In other formulations, an increase in fluorescence was detected without an increased turbidity. Therefore, we recommend measuring the fluorescence of at least two dyes in combination with turbidity and intrinsic fluorescence. Growth of protein aggregates at the side of wells can stay undetected if the measurement is only performed in the centre of the well. Therefore, determining turbidity at different positions in the well will result in a

Taken together, high throughput spectroscopy is a fast and versatile method for initial screening of the physical stability of protein formulations. The well-plate based platform could be enhanced with an accelerated stress testing and methods to determine chemical stability, e.g. electrophoresis, HPLC, mass spectrometry.

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