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1 Review

Bacterial expression and re-engineering of *Gaussia princeps* luciferase and its use as a reporter protein

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ABSTRACT

Bioluminescence, the generation of visible light in a living organism, is widely observed in nature, and a large va-20 riety of bioluminescent proteins have been discovered and characterized. Luciferase is a generic term for bioluminescent enzymes that catalyze the emission of light through the oxidization of a luciferin (also a generic term). Luciferase are not necessarily evolutionary related and do not share sequence or structural similarities. Some luciferases, such as those from fireflies and *Renilla*, have been thoroughly characterized and are being used in a wide range of applications in bio-imaging. *Gaussia* luciferase (GLuc) from the marine copepod *Gaussia princeps* is the smallest known luciferase, and it is attracting much attention as a potential reporter protein. GLuc identification is relatively recent, and its structure and its biophysical properties remain to be fully characterized. Here, we review the bacterial production of natively folded GLuc with special emphasis on its disulfide bond formation and the re-engineering of its bioluminescence properties. We also compare the bioluminescent proteins with native disulfide bonds. Furthermore, we discuss and predict the domain structure and location of the catalytic core based on literature and on bioinformatics analysis. Finally, we review some examples of GLuc's emerging use in biomolecular imaging and biochemical assay systems.

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39 1. Introduction

Luciferase is a generic name for enzymes that emit light by catalyz-40ing substrates referred to as luciferin ([1], pages xix-xxi). A large variety 41 of luciferases from different biological and evolutionary origins have 42 been identified, however, few have been thoroughly characterized and 43 fewer still have been cloned and used in analytical applications. The 44 45 most commonly used luciferase in experimental imaging applications originates from the North American firefly (*Photinus pyralis*) [2]. Firefly 46luciferase (FLuc)'s molecular weight is 62 kDa (Table 1) [3]. It catalyzes 47 the substrate benzothiazole (D-luciferin) in the presence of oxygen, ATP 4849 and Mg^{+2} , and emits green light with a peak emission of 562 nm [4]. Its structure has been solved by X-ray crystallography [5]. Several FLuc var-50iants with different luminescence characteristics have been designed, 5152which has enlarged its application range [6].

Renilla luciferase (RLuc) derived from sea pansy (*Renilla reniformis*)
is also well characterized. It has a molecular weight of 36 kDa [3,7],
and it catalyzes coelenterazine to produce the excited coelenteramide
state (Table 1) emitting a blue light with a peak emission of 480 nm. Un like FLuc, RLuc does not require ATP for activity, and it is becoming an

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http://dx.doi.org/10.1016/j.bbapap.2015.05.008 1570-9639/© 2015 Elsevier B.V. All rights reserved. essential bioluminescence marker for in vivo imaging [8–10]. A physi- 58 cally and chemically stabilized RLuc variant with improved biolumines- 59 cence properties [11] and its X-ray crystal structure was reported [12]. 60 In addition to the well known FLuc and RLuc, luciferases from various 61 organisms having distinct sequences, structures, and biochemical prop- 62 erties have also been reported [13–16] (Table 1). 63

Gaussia luciferase (GLuc) is a coelenterazine-dependent luciferase 64 (Table 1) identified from the marine copepod *Gaussia princeps* [15] 65 (Fig. 1), which lives in temperate and tropical waters worldwide [17]. 66 GLuc, which catalyzes the production of a bright blue light, is secreted 67 by *Gaussia princeps* in combination with rapid swimming probably as 68 a defense mechanism [18]. GLuc's sequence (mRNA, complete cds) 69 was first reported in 1999 [15]. It is presently attracting much attention 70 for bio-imaging applications because of its small size and strong lumi-71 nescence activity, as demonstrated by over two hundred recently re-72 ported examples. Here, we review literature on GLuc, with a special 73 emphasis on the formation of disulfide bonds in bacterially expressed 74 GLuc and its functional re-designing. In addition, we discuss its structur-75 al and functional properties in the light of the literature and of bioinfor-76 matics analysis of its amino acid sequence. 77

2. Biochemical properties of GLuc

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With 185 amino acids and a molecular mass of 19.9 kDa, GLuc is 79 the smallest luciferase reported so far (Table 1, Fig. 4A). GLuc's 80

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Abbreviations: ATP, adenosine triphosphate; HPLC, high performance liquid chromatography

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1 Table 1

t1.2 Comparison of GLuc with selected luciferases.

t1.3	Species	Size (aa)	Molecular Wt (kDa)	Cofactors	Substrate	Isoelectric point	Cysteine number (frequency, %)	Reference
t1.4	Gaussia princeps (Copepod)	185	20	None	Coelenterazine	6.8	11 (5.9)	[3]
t1.5	Photynus pyralis (Firefly)	550	62	ATP, Mg ⁺²	D-Luciferin (benzothiazole)	6.59	4 (0.7)	[3]
1.6	Renilla reniformis (Sea pansy)	311	36	None	Coelenterazine	5.8	3 (1.0)	[3]
1.7	Oplophorus	196	21	None	Coelenterazine	5.0	1 (0.5)	[16]
1.8	Pleuromamma	198	22	N/A	N/A	6.6	17 (8.6)	[15]
t1.9	Vibrio harveyi (LuxA)	355	37	Long-chain aldehyde	Reduced riboflavin phosphate (FMNH2)	4.77	8 (2.2)	[13,64]
t1.10	Lingulodinium polyedrum	1241	137	None	Dinoflagellate luciferin	7.31	29 (2.3)	[14]

t1.11 FLuc contains four free cysteines. Little influence on its luminescence was observed by changing them to serine [65]. RLuc contains three free cysteines, and two of them affect its lumit1.12 nescence property when mutated to alanine [66].

bioluminescence intensity is significantly stronger (over 200 fold) than 81 that of FLuc and RLuc [3], which are commonly used as reporter pro-82 teins. Alike RLuc, GLuc catalyzes the substrate, coelenterazine, to pro-83 84 duce the excited state of coelenteramide by emitting blue light (Fig. 2) with a peak emission at 480 nm [3]. Unlike FLuc, GLuc does not require 85 ATP or other co-factors for activity. The bioluminescence activity of 86 GLuc is highly stable under a wide range of temperatures (90% and 87 88 65% of the luminescence intensity is retained after a 30-minute incuba-89 tion at 60 °C and 90 °C, respectively) [19,20] and in an acidic environment (down to pH 1.5) [21]. GLuc's bioluminescence is salt 90 concentration dependent [19,22] and activity reaches a maximum at 9150 mM NaCl [19]. GLuc's bioluminescence is strongly inhibited by 92heavy metal ions (Cu²⁺) and stimulated by monovalent ions (Cl⁻, I⁻, 93 94Br⁻) [22]. GLuc is highly specific for coelenterazine, and shows narrow substrate specificity unlike RLuc, which catalyzes coelenterazine, and 9596 also several analogs as well [23].

97 **3. Expression of recombinant GLuc**

Various expression systems for GLuc have been reported. GLuc was 98 cloned and expressed in mammalian [3,24], bacterial [21], insect [25], 99 04 and green algae [26] cells. Expression in Escherichia coli is an important step toward re-designing a protein, not only because it provides a handy 101 and inexpensive method for producing recombinant proteins, but also 102and foremost because genetic modifications are most easily introduced 103104 and assessed with this system [27]. To our knowledge, the first quantitative analysis of GLuc expression in E. coli was reported by Verhaegen 105



Fig. 1. Picture of *Gaussia princeps* provided by Professor Russel Hopcroft, University of Alaska – Fairbanks (UAF), and reproduced with the author's permission.

M and Christopoulos T.K. [18]. They designed an *E. coli* over- 106 expression system that produces a biotinylated bap (biotin acceptor 107 peptide)-GLuc fusion protein. The final yield of GLuc purified by Avidin 108 affinity chromatography from the soluble fraction of the *E. coli* culture 109 was 0.55–0.69 mg/L, but nearly 60% of the luminescence was lost during 110 affinity chromatography purification [18]. 111

4. Multiple disulfide bonds and expression of native GLuc in E. coli 112

GLuc contains eleven cysteines corresponding to an amino acid frequency of 5.95%, which is in sharp contrast to FLuc and RLuc, containing 114 respectively four and three non-disulfide bonded cysteines (Table 1). 115 Ten of the GLuc's cysteines form disulfide bonds [22,28], whereas the 116 11th cysteine is located in the pro-sequence, which presumably helps 117 GLuc's folding by reshuffling non-native disulfide bonds in vivo [29, 118 30]. Disulfide bonded proteins are usually difficult to express in *E. coli* 119 in a native form, because *E. coli* 's cytoplasm provides a reducing envipurification, often resulting into non-native disulfide bonds [27]. Even 122 if the cysteines were oxidized in the cytoplasm, *E. coli* does not have 123 the molecular machinery to reshuffle non-native disulfide bonds into 124 the correct ones, and most multi-disulfide proteins thus turn into func-125 tionally inactive aggregates [27].

To overcome the problem of non-native disulfide bonds, GLuc was 127 expressed in the *E. coli* periplasm using a prokaryotic secretion signal 128 peptide, the pelB leader sequence [31,32]. However, the yield and specific activity of the secreted GLuc were relatively low ($250 \mu g/200 \text{ mL}$ 130 culture), and the pelB cleavage during protein expression in *E. coli* was 131 inefficient [31]. Cell free systems, where the protein folding environment can be controlled, have been used in an attempt to assist native disulfide bonds formation [28,33]. In principle, some proteins can be recovered from insoluble aggregates by optimizing the refolding conditions [34]. However, with as much as five disulfide bonds, the refolding of GLuc appears to be a difficult task [19]. Finally, low expression temperature (15 °C) increases the fraction of natively folded GLuc, but 138 also decreases the final protein yield [19].

Formation of non-native disulfide bonds in recombinant proteins is 140 closely related to the low solubility of miss-folded proteins. Aggregation 141 often competes with the folding reaction [20], but GLuc, like most small 142 proteins, is intrinsically able to fold into a native structure [29]. Thus, a 143 GLuc with enhanced solubility is anticipated to increase the yield of 144 well folded protein with native-like disulfide bonds. Recently, various 145 attempts for increasing the solubility of GLuc have been reported. In 146 one of them, GLuc was fused to a highly soluble IgG-binding domain 147 of protein A (ZZ-domain, 116 amino acid residues) [35], but it remains 148 unclear whether the fused GLuc was solubilized in a native, fully func- 149 tional form with the correct disulfide bonds. On the other hand, we con-150 structed a GLuc variant with a Solubility Enhancement Peptide tag (SEP 151 tag [36-38]), consisting of nine aspartic acid tags, attached to its C ter- 152 minus, which can significantly increase protein solubility. As a result, 153 98% of the tagged GLuc was expressed in the soluble fraction at 25 °C, 154 which was fully functional and formed native-like disulfide bonds [20]. 155

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