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Fluorescent RNA tags: current and future applications

Sven Hennig^{*,1}  & Saskia Neubacher^{*,1}

¹Department of Chemistry & Pharmaceutical Sciences, Vrije Universiteit Amsterdam, De Boelelaan 1108, Amsterdam 1081 HZ, The Netherlands

*Authors for correspondence: s.a.neubacher@vu.nl

“The past 8 years of research on FLAPs have broadened the panel for *in vitro* and *in vivo* applications and facilitated a general implementation of the technique”

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The rising interest in the field of RNA research, especially regarding diverse (short and long) noncoding RNAs, demands for novel techniques to identify, functionally characterize and also chemically target these molecules [1,2]. Fluorescent light-up aptamers (FLAPs) are RNA sequences that bind to small molecules and thereby enhance their intrinsic fluorescence significantly [3]. They can be encoded on the DNA level and are then either transcribed in the cell and functionalized as FLAPs on the RNA level, or they can be *in vitro* transcribed and purified for further applications. FLAPs have been known since the discovery of a fluorescence-enhancing RNA aptamer for malachite green in 2003 [4]. However, it was only in 2011 that they gained importance when Paige and colleagues published an RNA aptamer that binds the cell-permeable and nontoxic derivative (Z)-4-(3,5-difluoro-4-hydroxybenzylidene)-1,2-dimethyl-1H-imidazol-5(4H)-one (DFHBI) of the chromophore 4-hydroxybenzylidene imidazolinone (HBI) found in GFP [5]. This combination of RNA and small molecule, called Spinach, caused a plethora of follow-up publications focusing on either the variation or the optimization of different physicochemical features, resulting in RNA–fluorogen pairs named Baby Spinach, Broccoli, Corn or Mango, and others [6]. Currently, there are various small-molecule chromophores each in combination with a specific RNA aptamer sequence available. The most important variations appear within their spectral characteristics, the chemical nature of the chromophore, the size and stability of the formed RNA aptamer and their stoichiometry [3].

The past 8 years of research on FLAPs have broadened the panel for *in vitro* and *in vivo* applications and facilitated a general implementation of the technique. In the specific case of Mango, where the fluorogen carries a biotin moiety, the tag can be used to affinity pull down and purify fused RNA of interest, but also associated proteins (or RNAs), facilitating the identification of novel endogenous RNA–protein complexes. Tagging RNAs of interest to FLAPs has been used in early *in vivo* applications for highly abundant RNAs, such as the 5S ribosomal RNA (rRNA) or U6 snRNA [5]. Detection and localization of the tagged RNA of interest and quantitative measurements of polymerase III transcription in live mammalian cells were shown [7]. More recently, improved FLAPs were successfully used to detect low-abundance RNAs *in vitro* (FASTmiR), thereby substituting labor intensive methods (e.g. northern blotting, DNA–nanosilver clusters or *in situ* hybridization) and also in live cell imaging [8,9]. This allowed for quantitative and dynamic studies of mRNA transcription in cells. To measure mRNA and protein levels simultaneously, a FLAP can also be fused to a protein-encoding mRNA. A study where red fluorescent protein (mRFP1) was analyzed in *Escherichia coli* indicates that RNAs have a much higher turnover in cells than proteins [10]. FLAPs can also be fused to other aptamers that can then selectively bind to a target molecule (such as EGFR) to label cell surfaces [11].

Attached to so-called sensor RNAs, FLAPs enable the detection of biomolecules, such as metabolites, proteins or RNAs. Upon binding of the analyte to the sensor unit, a structural rearrangement causes the FLAP unit to fold and bind its fluorogen, which then results in fluorescence increase. These bifunctional RNAs have been used to

detect many metabolites *in vitro* or *in vivo*, including adenosine triphosphate (ATP), adenosine 5'-diphosphate (ADP), flavin mononucleotide (FMN), cyclic-di-nucleotide monophosphates (e.g. c-di-GMP, c-di-AMP, c-AMP-GMP), S-adenosyl-L-homocysteine (SAH) and S-adenosylmethionine (SAM), theophylline (TH) and thiamine 5'-pyrophosphate (TPP) [12]. Similar approaches were used to detect and image proteins (such as streptavidin and MS2 coat protein) [13] and RNAs (mRNAs and micro RNAs) [8]. FLAPs are also used as tools in high-throughput screening, whereas their utilization is currently rather limited. In one example, the activity of an RNA-modifying enzyme was screened using a chemically modified fluorogen that simply causes the FLAP to light-up upon enzyme activity. For sure, great challenges lie ahead of high-throughput screening utilization of FLAPs, but the first steps have been made and clearly a great potential is hidden in this underexplored area [14].

Interactions between biomacromolecules are of great importance both for the understanding of cellular functions and also for the screening of potential interaction modulators [15,16]. A split version of the aptamer dBroccoli was shown to be able to reassemble and generate fluorescence [17]. Using this system cellular RNA–RNA interactions could be detected directly, and further downstream using a reporter gene as readout. In the Corn aptamer, homodimerization of the aptamer is a prerequisite for fluorogen binding and hence fluorescence [18]. This particular feature could potentially be used to build a system for the detection of RNA–RNA interactions. However, it is still unclear whether the homodimerization could prove to be too strong for this application in its current state. While most applications use fluorescence intensity as readout, in the context of intermolecular interactions, proximity-based Förster resonance energy transfer (FRET) is also a possible readout that can offer advantages. In the so-called 'apta-FRET', Spinach and Mango were fused together and used for small-molecule sensing and target RNA sensing [19]. This demonstrates the potential of FLAPs, especially to study bimolecular interactions *in vivo*. In principle, FRET can also occur between FLAPs and other fluorophores. In combination with protein fluorophores, the detection of RNA–protein interactions becomes possible. Spinach as a green fluorophore was used together with mCherry as a red protein fluorophore in a FRET-based system to detect and quantify the interaction between pp7-RNA and the PP7-coat-protein [20]. This is the first example that shows in principle a genetically encodable protein–RNA detection system.

FLAPs require the external addition of fluorogens, which is one of the major restrictions of their applications. *In vivo*, FLAPs are therefore currently limited to cultured cells. However, with growing knowledge in the field of RNA and its tertiary structures, it might be possible to generate intrinsic fluorescence directly from RNA. Until then, further research will be needed to improve properties of FLAPs, such as stability and fluorescence brightness. Being a genetically encodable fluorescent tag, FLAPs already offer many areas of research a tool to monitor RNA transcription, processing and trafficking. The first examples now have shown that this tool can be expanded to investigate interactions between RNA and RNA or RNA and proteins for basic and applied research.

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