

Mirabilis antiviral protein studies and its potential applications

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Abstract

Plants are an important source of different natural products, which have been utilised as medicine. One such natural product is protein, which has different applications like plant protection, therapeutic agent for different diseases. Ribosome inactivating proteins (RIPs) are one of such proteins which has N-glycosidase activity, due to which they can modify large mRNA and inhibit further translation. Different types of RIPs have been isolated from plants, crop plants and microorganisms. They are classified into three groups; Type I, Type II and Type III based on their physical properties. It has also been observed that more than one RIP can be isolated from one source, which indicates the presence of different isoforms in that source. One such frequently used RIP for medical purpose is Mirabilis Antiviral Protein (MAP), isolated from roots, leaves and seeds of *Mirabilis jalapa* L. MAP comes under the category of Type I RIP. MAP exhibits wide range of biological activities, such as antiviral, anticancer and antibacterial. The protein is lysin rich with pI 9.1 and molecular weight close to 24 kDa as determined by SDS-PAGE. It is highly rigid and thermostable protein, and also reported to express and maintain its antiviral activity at high temperature.

Key words: Ribosome inactivating proteins, mirabilis antiviral protein

1. Introduction

Since ancient times, different parts of plants are used as source of medicine for treatment of many diseases. They are invaluable source of phytochemicals and proteins, which have been utilized as traditional medicine in many countries. Different types of proteins have been investigated with selective toxicity and used for different purposes such as for transgenic plant protection (Logemann *et al.*, 1992; Lodge *et al.*, 1993), treatment of different types of diseases like cancer, treatment of HIV, Hepatitis B, *etc.* (Olsnes and Pihl, 1982; Pastan and Fitzgerald, 1991; Au *et al.*, 2000; Fan *et al.*, 2009). Ribosome inactivating proteins (RIPs), are one such class of proteins which are isolated not only from plants but also from certain fungi and bacteria. These proteins have N-glycosidase activity, due to which they can modify large mRNA and inhibit further translation (Stirpe *et al.*, 1992).

Most of RIP literature elucidates its isolation, characterization and different applications, since the proteins show selective toxicity (Gasperi-Campani *et al.*, 1985; Barbieri *et al.*, 1993). Also many other studies focused on enzymology, uptake of RIPs into target cells and subsequent transport to its target (Cavallaro *et al.*, 1995; Sandvig and Van Deurs, 1999). All these investigations provide a knowledge regarding biochemical and medicinal properties of RIPs. In recent years with the help of recombinant DNA technology, investigation of RIPs activities has been completely revealed. These studies not only lead to an improved understanding of RIPs expression and its N-glycosidase activity but also discovered new

enzymatic activities which suggest that RIPs biology is quite complex.

1.1 Historical view

In 1925, first RIP was extracted from pokeweed (*Phytolacca americana*), which showed the inhibition of viral infection (Irvin, 1983). Therefore, it has been linked to plant defence mechanisms. Stirpe *et al.* (1992) and Barbieri *et al.* (1993) extracted the protein from 50 plants and checked its activity. They found that most of the extracts showed *in vitro* translation inhibitory activity. Purification of these proteins led to their identification as RIPs. Further investigation showed that they are not only present in plants but also in crop plants such as wheat, maize and barley (Coleman and Roberts, 1982).

Endo *et al.* (1987) found an enzymatic activity of RIP. They discovered that RIP has a N-glycosidase activity which specifically remove adenine which is corresponding to A4324 residue in rat 28S rRNA (Endo and Tsurugi, 1987; Endo *et al.*, 1987). This adenine present in R-sarcine loop, which is a 14 nucleotide region, conserved in large rRNAs from bacteria to humans (Bailey-Serres *et al.*, 1998).

The first "A" of a GAGA sequence is the RIPs substrate, which forms the core of putative tetraloop enclosed by a short base paired stem (Correll *et al.*, 1998; Gutell *et al.*, 1993; Orita *et al.*, 1993). The translation is blocked by an irreversible modification of target A residue, by blocking EF-1 and EF-2 dependent GTPase activity. This inhibit the binding of ribosome to EF-2. Due to this translation inhibitory activity of RIPs, these proteins are toxic to cells (Nilsson *et al.*, 1986). Subsequently it has been observed that most of the RIPs remove more than one adenine from ribosome. Apart from ribosome, they also release adenine from poly-A tail and form DNA. Due to these enzymatic activities, they are also termed as polynucleotide: adenosine glycosidase (Barbieri *et al.*, 1997; Bolognesi *et al.*, 2002).

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2. Types of RIPs and their characteristics

Based on their physical properties, RIPs are classified into three groups as follows (Mundy *et al.*, 1994):

Type I: It is composed of single polypeptide chain of approximate molecular weight of 30 kDa (Barbieri *et al.*, 1993; Irvin, 1975). These are basic proteins which have not only many highly conserved active cleft residues and secondary structure within the active site region (Mlsna *et al.*, 1993; Monzingo *et al.*, 1992), but also distinctly different with respect to the overall sequence and post-translational modification (Hartley and Lord, 1993). The receptor alpha-2 macroglobulin is responsible for the binding and endocytosis of Type I RIPs. This mechanism was explained with the help of RIPs Saporin and Trichosanthin (Chan *et al.*, 2000). Study by Carvalloet *et al.* (1995) showed that there is an interaction between Type I RIPs and cell membrane mediated with the help of alpha-2-macroglobulin receptor and carried out endocytosis of Type I RIPs. Once inside cell, they disturb translation of cell.

Example: Pokeweed antiviral protein (PAP) and saporin.

Type II: These are heterodimer proteins consisting of A and B subunits, each with approximate molecular weight of 30 kDa (Stirpe *et al.*, 1978). The A chain has RIP activity and is linked to B-chain, which is endowed with galactose binding lectin through a disulphide bond (Stirpe *et al.*, 1978). B-chain binds to glycoproteins and/or glycolipids present on cell membrane of eukaryotic cells and mediate transportation of A-chain into cytoplasm (Lehar *et al.*, 1994; Olsnes and Sandvig, 1988; Sandvig *et al.*, 1976; Steeves *et al.*, 1999) then they access to rRNA and disturb the translation mechanism (Olsnes and Pihl, 1973).

Example: Ricin and Abrin

Type III: These are synthesized as inactive precursors, *i.e.* proRIPs. To form an active RIPs, proRIPs require proteolytic processing event (Peumans *et al.*, 2001). These RIPs are less common than Type I and II RIPs and are not used for therapeutic purpose. These have been characterized only from maize and barley (Walsh *et al.*, 1991; Bass *et al.*, 1992; Chaudhry *et al.*, 1994).

Example: Maize and Barley

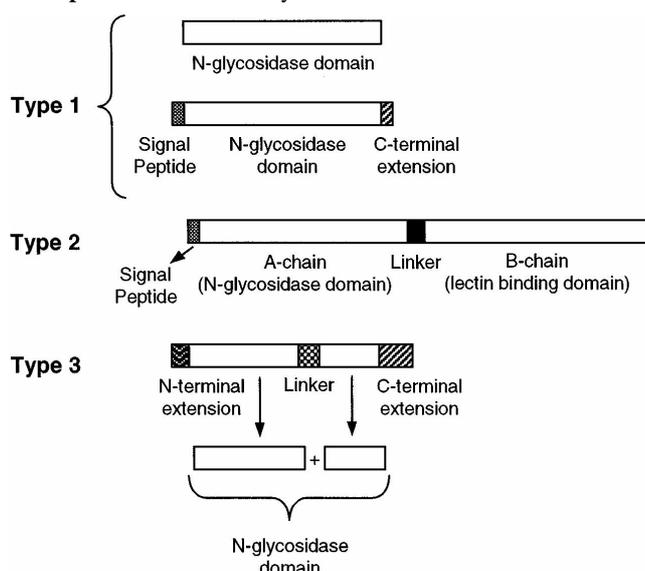


Figure 1: Classification of RIPs

3. Dispersion of RIPs

Due to their unique enzymatic activity towards eukaryotic cells, these proteins had received attention in research for their use as a therapeutic agents. To understand their mode of action and to optimize their therapeutic applications, different RIPs have been isolated from about 50 plants species covering 17 families. Some of them include many RIPs-producing species such as Cucurbitaceae, Euphorbitaceae, Nyctagaceae (Stirpe, 1992). Many plants which are from taxonomically unrelated families showed presence of RIPs which are not only structurally similar but also apparently identical in mode of action. Most of the RIPs are present in roots, leaves and seeds of the plants. It has also observed that more than one RIP can be isolated from one source, which showed similar amino acid sequence and most likely comprise a multigene family, for example, RIPs isolated from seeds of *Saponaria officinalis*, *Momordica charantia* and *Mirabilis jalapa* and leaves of *Dianthus carophyllus* and *Phytolacca americana* leaves (Stirpe *et al.*, 1992). One of such frequently used protein for medical purpose is *Mirabilis* Antiviral Protein (MAP).

Many studies have led to an improved understanding of MAP gene expression and activity against pathogens and cancerous cell also have uncovered new enzymatic activities that are suggestive of MAP biology being quite complex. This article summarizes work related to MAP activities and its different applications.

4. *Mirabilis* antiviral protein (MAP)

Mirabilis jalapa L., a native Peruvian plant, which is found to contain proteins in leaves and roots (MacBride, 1951). These proteins showed inhibition of mechanical transmission of certain plant viruses (Verma and Kumar, 1979; Kubo *et al.*, 1990; Takanami *et al.*, 1990; Vivanco, 1999). This protein is further named as *Mirabilis* Antiviral Protein (MAP), (Habuka *et al.*, 1992). This protein was purified with the help of cation exchange chromatography, by use of CM-sepharose CL6B column. The protein was revealed to be lysin rich with pI 9.1 and molecular weight close to 24 kDa as determined by SDS-PAGE (Habuka *et al.*, 1991; Takanami *et al.*, 1990). Several isoforms of MAP have been identified in the seeds of *M.jalapa*, MAP (27.788 kD), MAP-2 (30.412 kD), MAP-3 (29.771 kD), and MAP-4 (29.339 kD). Main isoform from seed, *i.e.* MAP was present in roots while the second abundant isoform from seed, *i.e.* MAP-4 was also isolated from leaf tissue (Bolognesi *et al.*, 2002).

Habuka *et al.* (1989) showed that MAP is made up of single polypeptide with 250 amino acid. They also revealed a complete amino acid sequence and it was observed that amino acid sequence has 24% homology with ricin A chain and also contain two homologous regions with ricin A and shiga like toxin. Region from Ile 163 to Ile 175, IQMVSEAAARFKYI contain glutamic acid residue (Glu 168) which was corresponding to the Glu 177 from ricin A chain and Glu 167 of shiga like toxin. The glutamic acid residue of both these proteins has been proved to be an accepted active site for ribosome inactivation (Hovde *et al.*, 1988; Schlossman *et al.*, 1989). The MAP is highly rigid and thermostable protein, this property is due to the presence of intramolecular disulphide bond (Habuka *et al.*, 1989). In *E. coli*, recombinant biological active MAP was expressed at 42°C without inactivation (Habuka *et al.*, 1990), whereas ricin A chain was expressed but inactivated at that temperature, because of absence of intramolecular disulphide bond

(O'Hare *et al.*, 1987; Piatak, 1988). In addition to this MAP also reported to maintain its antiviral activity even after the treatment at 85°C for 30 min (Takanami *et al.*, 1990).

5. Application of mirabilis antiviral proteins

A: Potential to cleave supercoiled DNA

MAP also has ability to cleave supercoiled double stranded DNA to acquire nick circular and linear forms (Zullies, 2006). This property of nucleic acid cleavage by MAP has been used to identify the presence of this protein in crude extract preparation of plant (Zullies, 2006).

B : Antibacterial activity

RIPs of *M. jalapa* leaves has bacteriostatic activity to *Staphylococcus epidermidis* with MIC value of 10 mg/ml. while it does not inhibit the growth of *Propionibacterium acnes* (Rumiyati, 2014).

C : Antiviral activity

Studies of Kubo *et al.* (1990) showed that MAP extracted from roots showed that it inhibit mechanical transmission of tobacco mosaic, cucumber green mottle mosaic, potato Y, turnip mosaic and cucumber mosaic virus. In other studies, effect of MAP extracted from roots of *Mirabilis jalapa* were studied against potato virus X, potato virus Y, potato leaf roll virus, and potato spindle tuber viroid. The roots extract containing MAP was sprayed on test followed by viral inoculation 24 h later. 100% inhibition was observed as corroborated by infectivity assays and the nucleic acid spot hybridization test. Antiviral activity of MAP extracts was observed against mechanically transmitted viruses but not against aphid-transmitted viruses. Purified MAP showed the same antiviral effect as the crude extracts (Vivanco, 1999). This antiviral activity is nonspecific regarding the characteristics of the virus and hence may have agricultural and clinical applications (Vivanco, 1999).

D: Anticancer activity

MAP was observed to be more toxic to cancerous cells than normal cells. MAP isolated from leaves showed different extent of cytotoxicity against T47D and SiHa cell lines while it was relatively less cytotoxic to mononuclear cell (Zullies, 2006). In addition, many studies had demonstrated that MAP indicated its potent anticancer activity through apoptotic pathway. Its activity was confirmed by DNA fragmentation and cell morphology change methods (Zullies, 2003, Watthanachaiyingcharoen *et al.*, 2007). In our study, we have observed that MAP isolated from roots showed more specific cytotoxic activity against cancer cell line such as MACF-7, A549, HCT 116, than the normal cell line, *i.e.* Vero (Kale and Mukundan, 2015). Recombinant RIPs can be made available on a scale which could meet the high demand of its use as a potential therapeutic agent for treating cancer cells (Salehzadeh and Arasteh, 2012).

6. Conclusion

RIPs commonly has an important enzymatic activities such as RNA N-glycosidase and adenine polynucleotide glycosidase activity. One of such RIP is *Mirabilis* antiviral protein which has vast applications in medical and therapeutic field. MAP showed inhibitory effect against plant viral infections, therefore has potential to use in agriculture. It also showed antibacterial and anticancer activity, more emphasis should be given for identifying its potential activity which will throw more insight into cytotoxic drug development and treatment of diseases.

Conflict of interest

We declare that we have no conflict of interest.

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