



# Antigenotoxic Effect of Some Lichen Metabolites

# 6

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## Abstract

Naturally occurring compounds can have protective effects towards mutagens and carcinogens, as shown by numerous studies. Several lichen species have been given much attention by researchers because their extracts and compounds have been used in traditional medicine to cure diseases such as ulcer, arthritis, tuberculosis, and cancer throughout the ages. Although a wide variety of scientific investigations on the biological activities of lichen extracts and their constituents have been performed, there is rather less research on their genotoxicity and antigenotoxic activity. To date, most results for the genotoxic/antigenotoxic activities of lichens have been obtained for lichen extracts and their constituents using the Ames/Salmonella and the *E. coli* WP2 microsome, chromosome aberration, micronucleus, sister chromatid exchange, single-cell gel electrophoresis, 8-OH-dG activity, and wing somatic mutation and recombination assays. In the present chapter, findings on antigenotoxic/genotoxic activities and its mechanisms are evaluated. By using the most common bacterial and nonbacterial assays, extracts of various lichen species have been shown to have promising antigenotoxic activity with somewhat less genotoxic activity. Lichen extracts may have a possible therapeutic potential. Therefore, these extracts must be further investigated by other multiple in vitro bioassays for development of therapeutic agents.

## Abbreviations

2-AF      2-Aminoflourene  
4-NPD    4-Nitrophenylenediamine

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8-oxo-dG	8-Oxo-2'-deoxyguanosine, 8-hydroxy-2'-deoxyguanosine
9-AA	9-Aminoacridine
AFB1	Aflatoxin B <sub>1</sub>
BrdU	Bromodeoxyuridine
CA	Chromosome aberration
CBS	Colloidal bismuth subcitrate
COMET	Single-cell gel electrophoresis
DN	Dioxidine
DXR	Doxorubicin
HPL	Human peripheral blood lymphocytes
IMA	Imazalil
MI	Mitotic index
MMC	Mitomycin C
MMS	Methyl methanesulfonate
MN	Micronucleus
MNNG	Methylnitronitrosoguanidine
SCE	Sister chromatid exchange
SCGE	Single-cell gel electrophoresis
SMART	Wing somatic mutation and recombination test

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## 6.1 Introduction

Naturally occurring organic compounds from a variety of organisms, including medicinal plants, are very important. Development of these compounds for new therapeutic agents has become a popular research objective. Investigation of the biological potential of several classes of natural agents, dietary constituents, hormones, and vitamins using *in vitro* and *in vivo* bioassays have shown that they can act as genotoxicity inhibitors as well as protectors against cytostasis or environmental carcinogens (Okai et al. 1996; Scarpato et al. 1998; Ipek et al. 2003, 2005; Mersch-Sundermann et al. 2004; Jayaprakasha et al. 2007; Zeytinoglu et al. 2008; Hoshina and Marin-Morales 2014; Delebassée et al. 2017; Makhuvele et al. 2018; Kaura et al. 2018; Kuete et al. 2018). Also, investigation of the possible genotoxicity of such agents captures the attention of researchers because of their use in folk medicine or potential applications. Most of the medicinal plants used traditionally have never been subjected to toxicological tests such as that required for modern pharmaceutical compounds. However, research has shown that quite a few plants that are used in traditional medicine or similar applications may have genotoxic or carcinogenic properties (Santos et al. 2009; Nieminen et al. 2002; Sponchiado et al. 2016; Quadros et al. 2017; Kahaliw et al. 2018; Prinsloo et al. 2018). Recently, development of an *in silico* method by using different computer models for genotoxicity and carcinogenicity prediction of secondary plant metabolites suggested that these potentials can be a first step for prioritization (Glück et al.

2018). Therefore, it becomes very important to search extracts derived from plants that contain a variety of compounds for their nontoxic, antigenotoxic, or genotoxic properties.

Lichen species have attracted much of the attention of researchers because their extracts and compounds have been used in traditional medicine in Europe, Asia, and North America (Richardson 1988; Cabrera 1996; Tilford 1997). Although lichen extracts and their compounds have been subjected to many scientific investigations for their biological activities such as immunostimulation, analgesic, anti-ulcerogenic, antipyretic, antimicrobial, antioxidative, and antitumor (Kumar and Müller 1999; Ingolfssdottir et al. 2000; Ingolfssdottir 2002; Türk et al. 2003; Tay et al. 2004; Yılmaz et al. 2004; Halici et al. 2005; Karunaratne et al. 2005; Behera et al. 2006; Zeytinoglu et al. 2008; Yeash et al. 2017; Tomović et al. 2017; reviewed in Prashith Kekuda et al. 2018), there is rather less research on their genotoxic or antigenotoxic activity. Scientific investigation of the genotoxic and antigenotoxic properties of lichens includes *in vitro* and *in vivo* studies, mostly using the aqueous, methanol, acetone, or *n*-hexane extracts. In the recent past few years, the genotoxicity and antigenotoxicity of isolated compounds from lichen species has attracted attention. In the present chapter, findings on the antigenotoxic and genotoxic activity of lichen extracts or second metabolites and the relevant mechanisms are evaluated.

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## 6.2 Bioassays for the Antigenotoxicity/Genotoxicity of Lichens

The Organization for Economic Co-operation and Development (OECD 2012) and the European Centre for the Validation of Alternative Methods (ECVAM 2012) have widely investigated the validation of mutagenicity tests. A set of assays is recommended to determine the genotoxicity of a test agent. The methods most frequently used for the assessment of genotoxic/antigenotoxic activity of lichen extracts or components based on bacterial short-term assays and mammalian test system are recommended by the OECD and the ECVAM. The Ames/*Salmonella*/microsome (Ames) and the *Escherichia coli* WP2/microsome (WP2) reverse mutation assays are the most common bacterial systems, and micronucleus (MN), chromosome aberration (CA), sister chromatid exchange (SCE), single-cell gel electrophoresis (COMET), and wing somatic mutation and recombination test (SMART) the most common nonbacterial systems, that have been used to date.

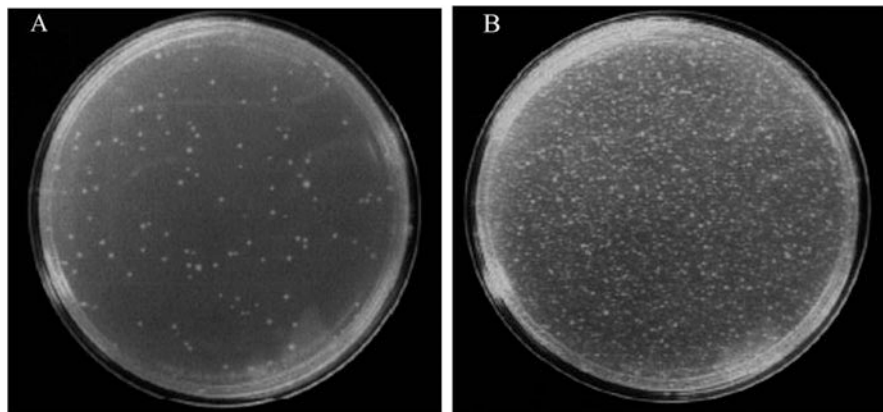
### 6.2.1 Bacterial Short-Term Assays

The Ames and the WP2 assays are short-term bacterial reverse mutation assays specifically designed to detect a wide range of chemicals or other agents which can produce genetic damage. The combination of *Salmonella typhimurium* and *Escherichia coli* strains in the tests recommended by OECD (1997) are given in Table 6.1, including their characteristics such as mutation types, plasmids, reversion

**Table 6.1** Characteristics of the most commonly used strains in the bacterial reverse mutation test, recommended in OECD guideline 471

Original bacteria/ strains	Mutation type	Plasmid	Reversion event	DNA target
<i>S. typhimurium</i> LT2				
TA97/97a	Deletion <i>hisD6610 hisO1242 rfa</i> $\Delta(\textit{gal chl bio uvrB})$	pKM101	Frameshift	-C-C-C-C-C-C-
TA98	Deletion <i>hisD3052 rfa</i> $\Delta(\textit{gal chl bio uvrB})$	pKM101	Frameshift	-C-G-C-G-C-G-C-G-
TA100	Deletion <i>hisG46 rfa</i> $\Delta(\textit{gal chl bio uvrB})$	pKM101	Base-pair substitution	-G-G-G-
TA102	Wild type <i>hisG428 rfa galE</i> <i>his</i> $\Delta(\textit{G})8476$	pKM10 pAQ1	Terminating ochre	-T-A-A-
TA1535	Deletion <i>hisG46 rfa</i> $\Delta(\textit{gal chl bio uvrB})$	None	Base-pair substitution	-G-G-G-
TA1537	Deletion <i>hisC3076 rfa</i> $\Delta(\textit{gal chl bio uvrB})$	None	Frameshift	+1 near -C-C-C- C- run of Cs)
<i>E. coli</i> B				
<i>E. coli</i> WP2 <i>uvrA</i>	<i>trpE uvrA</i>	None	Terminating ochre and DNA repair	-A-T-
<i>E. coli</i> WP2 <i>uvrA</i>	<i>trpE uvrA</i>	pKM101		-G-C-

events, and the DNA target of mutations (Maron and Ames 1983; reviewed by Mortelmans and Zeiger 2000). These assays provide a very sensitive study of potentially mutagenic pathways for the metabolism of compounds in both absence and presence of a rat liver microsomal system (S9 mix). The Ames assay employs several histidine-dependent *Salmonella* strains, each carrying different mutations in various genes in the histidine operon, pointing to different mutagenic mechanisms. When the *Salmonella* strains carrying mutations in the *his* gene are grown on a minimal media agar plate with a trace of histidine, only those bacteria that revert to histidine independence are able to form colonies (Fig. 6.1). When a mutagen is added to the plate, the number of revertant colonies per plate is increased. Base-pair substitution (A:T to G:C or G:C to A:T) and frameshift mutations (deletions) in *S. typhimurium* strains are represented to identify both types of mutation caused by a test compound. Therefore, differences in the activity of a test compound acting in these strains may yield some insight into how the compounds interact with the DNA of bacteria. Additionally, some genetic markers have been developed to make the strains more sensitive to certain types of mutagens.



**Fig. 6.1** Ames test plates of TA100 strain of *Salmonella typhimurium*. (a) Control: spontaneous revertants. (b) A mutagenic dose response to sodium azide. (From Mortelmans and Zeiger 2000)

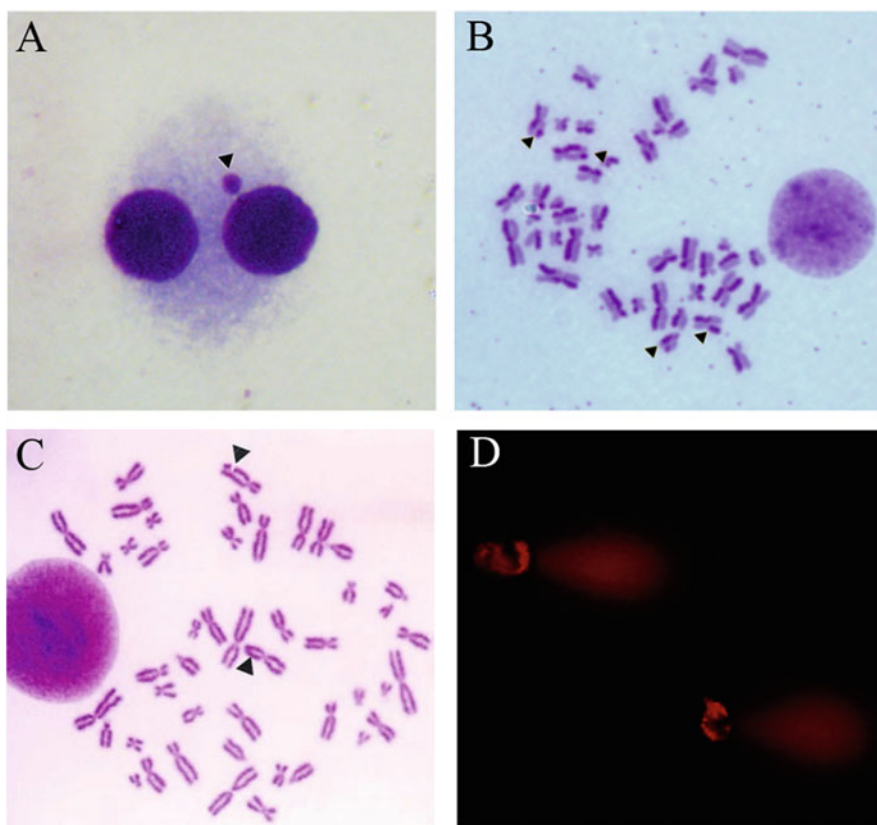
The *Escherichia coli* WP2 assay detects *trp* (–) to *trp* (+) reversion at a site blocking a step in the biosynthesis of tryptophan before the formation of anthranilic acid. The different auxotrophic WP2 strains all carry the same A:T base pair at the critical mutation site within the *trpE* gene (see Table 6.1). The most widely used *E. coli* WP2 strains, each carrying the *trpE* mutation, are WP2 (wild type for DNA repair), WP2 (pKM101), WP2 *uvrA*, and WP2 *uvrA* (pKM101) (Mortelmans and Riccio 2000; Sugiyama et al. 2016). The assay is currently used by many researchers in conjunction with the Ames assay for screening chemicals for their mutagenicity. The Ames assay procedures are the same as for the WP2 assay with the exception that limited histidine instead of limited tryptophan is used. International guidelines have been established for performing these mutagenicity assays. These assays are used worldwide as an initial screen to determine the mutagenic/antimutagenic potential of new chemicals, drugs, or natural products from plants or animals (Sponchiado et al. 2016).

Conversely, the anti-mutagenicity of a compound against a selected positive mutagen can be investigated when the two chemicals are co-administered to the bacteria in both test systems. Using known mutagenic compounds as “positive controls,” it is possible to study whether tested components can reduce DNA damage.

### 6.2.2 Nonbacterial Short-Term Assays

At present, several antigenotoxicity/genotoxicity assays, which include the chromosome aberration (CA), micronucleus (MN), somatic mutation and recombination test (SMART), sister chromatid exchange (SCE), single-cell gel electrophoresis (SCGE), or COMET assays, are available, and they are recommended to use as a set for investigations.

According to the literature, the antigenotoxic/genotoxic potential of lichens has been evaluated, commonly by MN, SCE, CA, COMET, 8-oxo-2-deoxyguanosine (8-oxo-dG) activity assays in mammalian cells and the mitotic index (MI) assay in plant cells. The purpose of the MN test is to examine the structural and numerical chromosomal damage which formed small membrane-bound DNA fragments or micronuclei in the cytoplasm of interphase cells caused by a tested agent or by clastogens and aneugens. Micronuclei can be formed by chromosome fragments lacking a centromere or whole chromosomes that are unable to migrate during cell division. The MN test can be conducted in the presence or in the absence of cytochalasin B, which is used to block cell division and generate binucleated cells (Fig. 6.2a). The cytokinesis-block micronucleus assay is a sensitive, comprehensive



**Fig. 6.2** Photomicrographs for nonbacterial genotoxicity assays. (a) A mitogen-stimulated cytokinesis-blocked lymphocyte containing one micronucleus (MN). (b) Giemsa staining of BrdU-incorporated chromosomes in human lymphocytes for sister chromatid exchange (SCE); arrowheads show chromosome breaks and sister union. (c) Chromosome aberration (CA) shows sister chromatids stained at different densities (photograph kindly provided by Dr. B. Ayaz Tuylu). (d) COMET tails of chromosomes visualized by an epifluorescence microscope (photograph kindly provided by Dr. A.T. Kopalal)

and simple methodology for measuring DNA damage, cytostasis, and cytotoxicity which can be scored easily in a variety of systems 3 and in vivo (Fenech 2007; Kirsch-Volders et al. 2011). The assay is being applied successfully for biomonitoring in vivo genotoxin exposure, for in vitro genotoxicity testing, and in diverse research fields such as nutrigenomics and pharmacogenomics.

The SCE assay is another short-term test useful for the detection of reciprocal exchanges of DNA between two sister chromatids of a duplicating chromosome in mammalian and also nonmammalian cells. Various cytomolecular protocols have been used to perform the SCE assay (Bakkali et al. 2008). SCEs result from the interchange of DNA replication products and involve DNA breakage and reunion (Wilson and Thompson 2007). Detection of SCEs requires the differential staining of sister chromatids, which usually can be achieved by the incorporation of bromodeoxyuridine (BrdU) into chromosomal DNA for two cell cycles (Fig. 6.2b). After labeling, treatment of cells with a spindle inhibitor such as colchicine is required to accumulate cells in a metaphase-like stage of mitosis (Perry and Evans 1975; Ipek et al. 2003).

The short-term in vitro mammalian cell chromosome aberration (CA) test measures the frequency of asymmetrical structural chromosome aberrations after exposure to test chemicals or mutagens. The in vitro CA test may employ cultures of established cell lines or primary cell cultures. Human peripheral blood lymphocytes (HPL) are stimulated to divide by cyclophosphamide in whole blood cultures. Cells in metaphase are analyzed for the presence of chromosomal aberrations (Fig. 6.2c) (Clare 2012).

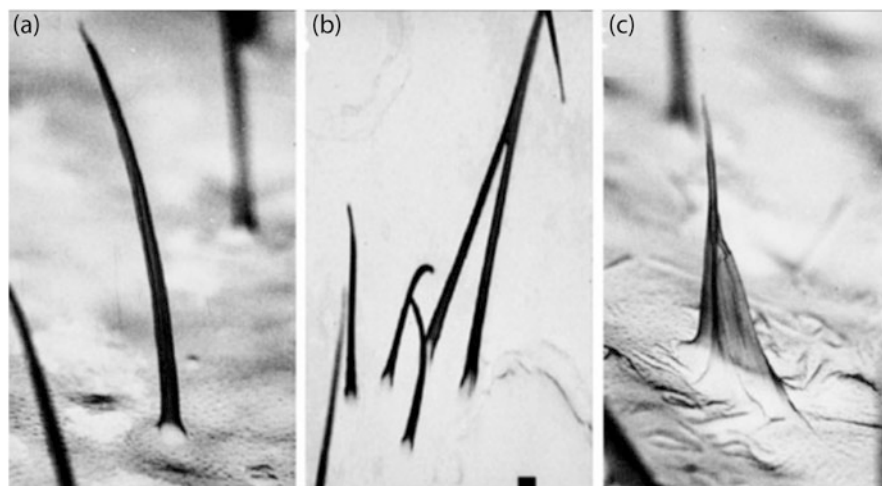
The COMET assay is used to detect the DNA-strand breaks in eukaryotic cells and named by the shape of the observed DNA distribution, which bears resemblance to a celestial comet. This well-established, highly sensitive, rapid, and simple genotoxicity test is based on the lysing of cells embedded in agarose on a microscope slide to form nucleoids containing supercoiled loops of DNA linked to the nuclear matrix. Then, electrophoresis at high pH results in structures resembling comets, as observed by epifluorescence microscopy (Fig. 6.2d). The intensity of the comet tail relative to the head reflects the number of DNA breaks (Singh et al. 1988; Collins 2004; Speit et al. 2009). Depending on experimental conditions, the migrating DNA reflects the number of single- or double-strand breaks, and alkali-labile sites, including incomplete excision repair sites, but also DNA–DNA and DNA–protein crosslinks (Santos et al. 2009; Verschaeve et al. 2010). A broad spectrum of DNA damage can then be detected either by visual classification of comet morphologies or from morphological parameters obtained by image analysis.

Level of 8-oxo-2'-deoxyguanosine (8-oxo-dG) is a frequently used biomarker of oxidative DNA damage caused by free radicals and other reactive species constantly generated in vivo. 8-Oxo-dG is later removed from DNA by the base excision repair pathway and subsequently transported into body fluids such as saliva, urine, and plasma. Such oxidative damage to DNA is probably the contributor of the age-related development of diseases such as cancer. Agents that decrease oxidative DNA damage should thus decrease the risk of cancer development. Thus, the measurement of 8-oxo-dG is the most common method of assessing DNA damage

(Halliwell 2000; Türkez et al. 2012a). An assay for the measurement of 8-oxo-dG has been developed by using a monoclonal antibody specific to 8-oxo-dG (N45.1), and the enzyme-linked immunosorbent assay (ELISA) has been well established (Toyokuni et al. 1997).

The mitotic index (MI) as a parameter for the evaluation of cytotoxic agents is the ratio of the number of cells undergoing mitosis to the number of cells not undergoing mitosis in a cell population. Mutagens can be detected cytologically by cellular inhibition, disruption in metaphase, induction of chromosomal aberrations and chromosomal fragmentation, and disorganization of the mitotic spindle and consequently of all subsequent dependent mitotic phases. MI is used as an indicator of adequate cell proliferation that can be measured by various plant test systems. Cytotoxicity tests using plant test systems *in vivo*, such as *Allium cepa* and *Zea mays*, are validated by several researchers who performed these tests jointly with other organism testing for genotoxicity (Agar et al. 2010; Gökbayrak and Sivas 2011; Aslan et al. 2012b).

Somatic mutation and recombination tests (SMART) is a fast and inexpensive mutagenicity detection test using a higher eukaryotic organism *in vivo*. Genetic damages induced by compounds that generate reactive oxygen species through different action mechanisms such as point mutations, deletions, and somatic recombination can be detected in cells of the wing imaginal disks of *Drosophila melanogaster* in this assay (Fig. 6.3). In particular, the *w/w<sup>+</sup>* eye and the *mwh/flr* wing variants of *D. melanogaster* have been extensively used to investigate the genotoxicity of a great number of chemicals, including plant extracts and pure compounds (Graf et al. 1984; Sponchiado et al. 2016; Teixeira Zafred et al. 2016).



**Fig. 6.3** Wings of *Drosophila melanogaster* illustrated by scanning electronic microscopy after a SMART mutagenicity assay: wild (a); multiple wing hair (*mwh*) mutation (b); and flare (*flr3*) mutation (c). (From Teixeira Zafred et al. 2016)



Overall these assays are those most used worldwide to determine the genotoxic/antigenotoxic potential of chemicals or natural products (as reviewed by Sponchiado et al. 2016). Thus, it is obvious that it is very important to always include bacterial and mammalian tests for genotoxicity/antigenotoxicity screening of compounds. Antigenotoxicity analysis of plant products is of valuable interest because they are a potential source of therapeutic agents, and thus safety assessments are crucial to validate their use in various areas, not only in phototherapy.

### 6.3 Antigenotoxic or Genotoxic Potential of Lichen Extracts

The first report on the genotoxicity of lichens using the Ames mutagenicity assay was on a secondary metabolite, physodic acid isolated from *Hypogymnia enteromorpha* (Ach.) Nyl. by Shibamoto and Wei (1984); the second one was on usnic acid by Al-Bekairi et al. (1991). Then, after 2007, biological activities of lichen compounds have been received more attention and several studies have been performed on the genotoxicity/antigenotoxicity of lichens in about the past 10 years. The studies up to date are summarized in two separate tables according to their activities. In Table 6.2, lichen species tested for only their antigenotoxicity or both genotoxicity and antigenotoxicity are listed. The lichen species which were tested for only their genotoxicity are listed in Table 6.3.

The first report describing the therapeutic potential of lichens against drug genotoxicity was from Geyikoglu et al. (2007) (Table 6.2). Aqueous extracts of four common lichen species collected from Giresun Province in Turkey, namely, *Dermotocarpon intestiniforme*, *Pseudevernia furfuracea*, *Parmelia pulla*, *Ramalina capitata*, and *Rhizoplaca melanophthalma* were tested for their genotoxic and antigenotoxic potentials. *Dermotocarpon intestiniforme*, *Pseudevernia furfuracea*, *Parmelia pulla*, and *Ramalina capitata* were found to be antigenotoxic at 5–10 µg/ml against colloidal bismuth subcitrate (CBS)-induced SCE and MN formation in human peripheral lymphocytes (HPL) in vitro. However, one other species, *Rhizoplaca melanophthalma*, was not antigenotoxic. The order of antigenotoxicity efficacy against CBS was *Pseudevernia furfuracea*, *Dermotocarpon intestiniforme*, *Ramalina capitata*, and *Parmelia pulla*. On the other hand, all lichen extracts tested were not genotoxic alone (Table 6.2).

After this work, a fresh aqueous extract of *Cetraria aculeatea* (Schreb.) Fr., one of the common species in Turkey, was studied for its genotoxic/antigenotoxic activities in both Ames and mammalian cell systems (Zeytinoglu et al. 2008). The extract (at 0.1–500 µg/ml) exhibited strong antigenotoxic activity against three known mutagenic agents, 4-nitrophenylenediamine (4-NPD), 2-aminoflourene (2-AF), and sodium azide (NaN<sub>3</sub>), in TA98 and TA100 strains of *Salmonella typhimurium* in the presence and absence of metabolic activation, without any mutagenic activity (Table 6.2). Preincubation of bacteria with the extract prevented the mutagenic activity of 4-NPD in the higher concentration range in both strains grown without metabolic activation than those grown with metabolic activation. It was suggested that the antigenotoxic potential of the extract was higher in the

**Table 6.2** Lichen species tested for only their antigenotoxicity and genotoxicity

Species/type of extract	Against	Assay	Cell types	Genotoxic	Anti-genotoxic	References
<i>Cetraria aculeata</i> /aqueous	4-NPD	Ames	TA98	No	Yes	Zeytinoglu et al. (2008)
	2-AF		TA100			
	NaN <sub>3</sub>					
Methanol	MMC	MN	HPL	No	No	Ceker et al. (2018)
	MNNG	Ames	TA1535	No	Yes	
	NaN <sub>3</sub>		TA1537			
	9-AA	WP2	<i>E. coli</i>		No	
	AFB1	SCE	HPL			
	AFB1	SCE, MN	HPL	NP	Yes	
<i>Cetraria islandica</i> /methanol	9-AA	Ames	TA1535	No	Yes	Kotan et al. (2011) Aslan et al. (2012b)
	NaN <sub>3</sub>		TA1537			
		MI	<i>Zea mays</i>			
<i>Cetraria olivetorum</i> /methanol	MNNG	Ames	TA1535	No	Yes	Ceker et al. (2018)
	NaN <sub>3</sub>		TA1537			
	9-AA	WP2	<i>E. coli</i>			
	AFB1	SCE	HPL			
<i>Cladonia chlorophaea</i> /methanol	MNNG	Ames	TA1535	No	Yes	Ceker et al. (2018)
	NaN <sub>3</sub>		TA1537			
	9-AA	WP2	<i>E. coli</i>			
	AFB1	SCE	HPL			
<i>Cladonia foliacea</i> /methanol		Ames	TA1535	No	Yes	Anar et al. (2013)
			TA1537			
		WP2	<i>E. coli</i>			
	AFB1	SCE	HPL	NP	Yes	

<i>Cladonia rangiformis</i> /methanol	AFB1	SCE, MN	HPL	No	Yes	Kotan et al. (2013)
<i>Dermotocarpon intestiniiforme</i> /aqueous	CBS	SCE, MN	HPL	No	Yes	Geyikoglu et al. (2007)
	CdCl <sub>2</sub>	MN				Guner et al. (2012)
	HgCl <sub>2</sub>	SCE, MN				Türkez and Dirican (2012)
<i>Evermia prunastri</i> /methanol	Imazalil	CA, MN				Türkez et al. (2012b)
	MNNG	Ames	TA1535	NP	Yes	Alpsoy et al. (2013)
			TA1537			
	Acridin	WP2	<i>E. coli</i>			
	AFB1	SCE	HPL			
<i>Lecanora muralis</i> /methanol	AFB1	SCE, MN	HPL	NP	Yes	Alpsoy et al. (2011)
<i>Parmelia pullata</i> /aqueous	CBS	SCE, MN	HPL	No	Yes	Geyikoglu et al. (2007)
<i>Peltigera rufescens</i> (Weis)/aqueous	Imazalil	CA, MN	HPL	No	Yes	Türkez et al. (2012d)
<i>Peltigera canina</i> /methanol		Ames	TA1535	No	Yes	Gomez et al. (2013)
	9-AA		TA1537			
		WP2	<i>E. coli</i>			
<i>Pseudovermia furfuracea</i> /aqueous	CBS	SCE, MN	HPL	No	Yes	Geyikoglu et al. (2007)
Methanol, acetone, hexane	AFB1	SCE, MN	HPL	No	Yes	Türkez et al. (2010)
Methanol	9-AA	Ames	TA1535	NP	Yes	Aslan et al. (2012b)
	NaN <sub>3</sub>		TA1537			
		MI	<i>Zea mays</i>			
<i>Ramalina capitata</i> /aqueous	CBS	SCE, MN	HPL	No	Yes	Geyikoglu et al. (2007)
<i>Rhizoplaca melanophthalma</i> /aqueous	CBS	SCE, MN	HPL	No	No	Geyikoglu et al. (2007)
	NaN <sub>3</sub>	MI	<i>Zea mays</i>	NP	Yes	Agar et al. (2010)
Methanol	9-AA	Ames	TA1537			
<i>Rhizoplaca chrysoleuca</i> /methanol	AFB1	SCE, MN	HPL	NP	Yes	Alpsoy et al. (2011)
	NaN <sub>3</sub>	MI	<i>Zea mays</i>			Agar et al. (2010)
	9-AA	Ames	TA1537			

(continued)

Table 6.2 (continued)

Species/type of extract	Against	Assay	Cell types	Genotoxic	Anti-genotoxic	References
<i>Usnea longissima</i> /methanol	AFB1	SCE, MN	HPL	NP	Yes	Agar et al. (2011)
<i>Umbilicaria vellea</i> /methanol	AFB1	SCE, MN	HPL	NP	Yes	Aslan et al. (2012a)
<i>Xanthoria elegans</i> /aqueous	MMC	CA, MN, SCE	HPL	No	Yes	Aydin and Türkez (2011b)
		8-oxo-dG				Türkez et al. (2012a)
<i>Xanthoparmelia somloensis</i> /methanol	AFB1	SCE, MN	HPL	NP	Yes	Aslan et al. (2012a)
Secondary metabolite						
Hypostictic acid (from <i>Pseudoparmelia sphaerospora</i> )						
Protocetraric acid (from <i>Parmotrema dilatatum</i> )	DXR	SMART	Drosophila Somatic cells	No	Yes	Guterres et al. (2017)
Psoromic acid (from <i>Usnea jamaicensis</i> )						
Salazinic acid (from <i>Parmotrema cetratum</i> )						
Usnic acid	MMS	COMET, MN	V79 cells	Yes No	Yes	Leandro et al. (2013)
(+)-Usnic acid (from <i>Cladonia arbuscula</i> )	DN	COMET	HPL	No	Yes	Prokopiev et al. (2017)
(-)-Usnic acid (from <i>Cladonia stellaris</i> )	MMS			Yes		Prokopiev et al. (2018)

CA chromosome aberration, CBS colloidal bismuth subcitrate, DN dioxidine, DXR doxorubicin, HPL human peripheral blood lymphocytes, MN micronucleus, MMS methyl methanesulfonate, MNNG methylNitrosoguanidine, NP not performed, NaN<sub>3</sub> sodium azide, SMART Somatic mutation and recombination tests, 8-OH-dG 8-Oxo-2 $\beta$ -deoxyguanosine

**Table 6.3** Lichen species tested for only their genotoxicity

Species/type of extract	Assay	Cell types	Genotoxic	References
<i>Aspicilia calcerea</i> /aqueous	CA, MN	HPL	No	Aydın and Türkez (2011a)
<i>Bryoria capillaris</i> /aqueous			No	Aydın and Türkez (2011b)
<i>Cetraria chlorophylla</i> /aqueous			No	Aydın and Türkez (2011a)
<i>Hypogymnia physodes</i> /methanol aqueous			Yes	Ari et al. (2012)
			No	Türkez et al. (2012c)
<i>Peltigera rufescens</i> /aqueous			No	Aydın and Türkez (2011b)
<i>Physcia aipolia</i> /aqueous			No	Aydın and Türkez (2011a)
<i>Ramalina polymorpha</i> /aqueous			No	Türkez et al. (2012c)
<i>Usnea florida</i> /aqueous	No	No	Türkez et al. (2012c)	
Secondary metabolite				
Physosodic acid ( <i>Hypogymnia enteromorph</i> )	Ames	TA100	No	Shibamoto and Wei (1984)
Physodalic acid ( <i>Hypogymnia enteromorph</i> )			Yes	
Physosodic acid (from <i>Pseudevernia furfuracea</i> (L) Zopf)	8-OH-dG	HPL PRCC U87MG	Yes	Emsen et al. (2016, 2018)
Psoromic acid (from <i>R. melanophthalma</i> )		PRCC U87MG		
Olivetoric acid (from <i>Pseudevernia furfuracea</i> (L) Zopf)		HPL PRCC U87MG		
Usnic acid	MNPCEs CA, MN	Mouse PCEs HPL	Yes No	Al-Bekairi et al. (1991) Polat et al. (2013)
(+)-Usnic acid (from <i>Cladonia arbuscula</i> )	MN	HPL	No	Koparal et al. (2006)
(-)-usnic acid (from <i>Cladonia stellaris</i> )				

CA chromosome aberration, HPL human peripheral blood lymphocytes, MN micronucleus, PCEs polychromatic erythrocytes, PRRC primary rat cerebral cortex cells, MNPCEs micronucleated PCEs, DN dioxidine, DXR doxorubicin, 8-OH-dG 8-Oxo-2'-deoxyguanosine, U87MG human glioblastoma cells

absence of the metabolic system and in inhibiting frameshift mutations. Result indicates a direct and specific activation of the extracts. However, in a further investigation, the extract of *Cetraria aculeata* (Schreb.) Fr. did not have antigenotoxic activity against mitomycin C (MMC) in term of MN formation in

HPL. The extract was not also genotoxic alone in the mammalian system. As reported very recently by Ceker et al. (2018), non-genotoxicity of methanol extracts of *C. aculeata* has been confirmed by Ames, WP2, and SCE assays. The extract at 5–20 µg/ml also had antigenotoxic activity, interestingly including a mammalian system. Furthermore, methanol extracts of two other species of *Cetraria*, *C. islandica* and *C. olivetorum*, showed activities similar to the activities of *C. aculeata* (Kotan et al. 2011; Aslan et al. 2012b; Ceker et al. 2018). According to the overall results, the extract of *C. aculeata* is significantly antigenotoxic in a bacterial system whereas it is not capable of inhibiting MN formation in MMC-induced human peripheral blood cells, which suggests variation in the effects of different solvent extracts and bioassay systems.

Recently, more investigation has been performed with an aqueous extract of *Dermatocarpon intestiniforme* in cultured HPL (Table 6.2). The extract at 25 and 50 ppm concentration conferred protection against cadmium chloride (CdCl<sub>2</sub>) (30 ppm)-induced MN formation despite its non-genotoxicity in the cells (Guner et al. 2012). It was also shown that the SCE and MN rates induced by mercury chloride (HgCl<sub>2</sub>) were alleviated in the cells treated with 50 µg/ml of the extract (Türkez and Dirican 2012). The extract was also antigenotoxic against imazalil (IMA)-induced CA and MN formation in cultured HPL. The lymphocytes were treated in vitro with varying concentrations of the lichen extract (25–100 µg/ml), and tested in combination with IMA (336 µg/ml). The extract alone was not genotoxic, and when combined with IMA treatment, it reduced the frequency of CAs and the rate of MNs (Türkez et al. 2012b). According to overall results of the MN, CA, and SCE assays, the extract of *Dermatocarpon intestiniforme* (5–100 µg/ml) is quite antigenotoxic against different types of clastogens or aneugens, which causes structural and numerical chromosomal damage.

One other aqueous extract of the lichen species *Peltigera rufescens* and *Xanthoria elegans* (25–100 µg/ml) has been assessed by four genotoxicity endpoints including CA, MN, SCE, and 8-oxo-dG assays in HPL (Türkez et al. 2012a, d; Aydin and Türkez 2011b). IMA- and MMC-induced frequencies of four genotoxic indices were diminished by the extract, indicating its inhibitory effect on oxidative DNA damage of reactive agents besides the structural and numerical chromosomal damages. The extract and its secondary metabolites may have potential to decrease the risk of cancer development.

The antigenotoxic effects of methanol extracts of *Rhizoplaca melanophthalma* and *Rhizoplaca chrysoleuca* against known mutagens have been evaluated in two different organisms, a plant and bacteria, using different assays (Agar et al. 2010). The extract (5–40 µg/plate) prevented NaN<sub>3</sub>-induced mitotic index partially in *Zea mays* seeds. Furthermore, it was antimutagenic against 9-aminoacridine (9-AA)-induced mutation in the TA1537 strain at all tested concentrations (0.5–5 µg/plate) in the Ames test. The inhibition rates ranged from 70.73% to 85.71%. The extract was also found to have antigenotoxic activity against AFB1-induced genetic damage (Alpsoy et al. 2011).

Several investigators have focused on the possible antigenotoxic potential of lichens against the well-known mutagen aflatoxin B<sub>1</sub> (AFB<sub>1</sub>). Türkez et al. (2010)

reported the antigenotoxic activity of another lichen species, *Pseudevernia furfuracea*, using its methanol, acetone and *n*-hexane extracts. None of these lichen extracts induced a significant number of SCEs and MN in cytokinesis-blocked HPL. Moreover, the results indicated that AFB<sub>1</sub>-induced SCEs were inhibited by the application of 50 µM methanol or acetone extracts. The positive effect of methanol, acetone, and ether extracts in decreasing the incidence of MN in comparison with an unprotected level was attained when cultures were treated simultaneously with AFB<sub>1</sub> and the extracts. Agar et al. (2011) reported that methanol extracts obtained from *Usnea longissima* suppress the mutagenic effects of AFB<sub>1</sub> in HPL examined by the SCE and MN tests. Kotan et al. (2011, 2013) also found that AFB<sub>1</sub>-induced genotoxicity has been suppressed by the methanol extract of other lichen species, *Cetraria islandica* and *Cladonia rangiformis*. The results showed that the frequencies of SCE and MN level decreased when 5 and 10 mg/ml of extract was added to AFB<sub>1</sub>-treated cultures. The methanol extracts of *Rhizoplaca chrysoleuca* and *Lecanora muralis*, at 5 and 10 µg/ml (Alpsoy et al. 2011), and *Umbilicaria vellea* and *Xanthoparmelia somloensis* (Aslan et al. 2012a) were antigenotoxic against AFB<sub>1</sub>-induced SCE and MN formation in HPL in vitro.

The methanol extract of *Evernia prunastri* (Huds.) Willd. was a strong antimutagenic on TA1537 and WP2 strains, with 37.70% and 69.70% inhibition rates against methylnitronitrosoguanidine (MNNG) and acridine-induced mutagenicity, respectively. Co-treatments of HPL with the extract and AFB<sub>1</sub> decreased the frequencies of SCE (Alpsoy et al. 2013).

The genotoxic and antigenotoxic effects of the methanol extract of *Cladonia foliacea* (Huds.) Willd., *Cladonia chlorophaea*, and *Cetraria olivetorum* were studied using WP2, Ames (TA1535 and TA1537), and SCE test systems (Anar et al. 2013; Ceker et al. 2018). According to their results, a 5 µM concentration of AFB<sub>1</sub> changed the frequencies of SCE. When 5 and 10 µg/ml concentrations of extract were added to AFB<sub>1</sub>, the frequencies of SCE were decreased. The extracts of *Cladonia chlorophaea* and *Cetrelia olivetorum* had further antigenotoxic activity against a series of known mutagens, MNNG, NaN<sub>3</sub>, and 9-AA besides AFB<sub>1</sub> (Ceker et al. 2018). On the other hand, the extracts of both species were not mutagenic in tested organisms although each has antimutagenic activity.

*Pseudevernia furfuracea* and *Cetraria islandica* were tested using their methanol extracts for both genotoxic and antigenotoxic activities. The extracts of these two species were not mutagenic in Ames and *Zea mays* mitotic index test systems. Furthermore, some extracts showed significant antimutagenic activity against 9-AA in the Ames test. Inhibition rates for 9-AA mutagenicity ranged from 25.51% (*Pseudevernia furfuracea*, 0.05 µg/plate) to 66.14% (*Cetraria islandica*, 0.05 µg/plate). In addition, all the extracts were significantly antimutagenic against NaN<sub>3</sub>, increasing the MI values of *Zea mays* (Aslan et al. 2012b). Gormez et al. (2013) showed that the methanol extract of *Peltigera canina* possesses an antigenotoxic potential in Ames and WP2 tests.

Another seven lichen species were collected from Erzurum and Artvin provinces in Turkey. *Aspicilia calcarea*, *Bryoria capillaris*, *Cetraria chlorophylla*, *Hypnogygnia physodes*, *Peltigera rufescens*, *Physcia aipolia*, *Ramalina*

*polymorpha*, and *Usnea florida* have been tested only for genotoxicity of the water extracts in cultured HPL (Table 6.3) (Aydin and Türkez 2011a, b; Türkez et al. 2012c). All tested lichen extracts up to 500 or 1000 mg/l concentration had no genotoxic effects on the cells by the application of CA and MN assays; however, antioxidant properties were shown. The methanol extract of *Hypogymnia physodes* (L.) Nyl. was also studied for its genotoxicity using CA and MN tests. Relatively higher concentrations are required for its genotoxic activity (Ari et al. 2012).

#### 6.4 Antigenotoxic/Genotoxic Potential of Lichen Secondary Metabolites

Lichen secondary metabolites exert various biological actions such as antitumor, antimicrobial, antiinflammatory, apoptotic, and cytotoxic activities (Ingolfssdottir et al. 1997; Vijayakumar et al. 2000; Huneck 2001; Tay et al. 2004; Yılmaz et al. 2004; Karunaratne et al. 2005; Mayer et al. 2005; Einarsdottir et al. 2010; Mitrovic et al. 2011; Molnar and Farkas 2010). Usnic acid is one of the most abundant lichen secondary metabolites studied for its biological activities. It has been used widely in the pharmaceutical and cosmetic industry because it has high antimicrobial activity (Ingolfssdottir 2002). Furthermore, usnic acid exhibited an antiproliferative effect on human leukemia (K562) and endometrial carcinoma (Ishikawa, HEC-50) cells (Cardarelli et al. 1997; Kristmundsdóttir et al. 2002).

A few findings present the genotoxic/antigenotoxic activities of lichen secondary metabolites. The earliest genotoxicity reports for the secondary metabolites of lichens come from Shibamoto and Wei (1984). They have tested physodic and physodalic acids isolated from *Hypogymnia enteromorpha* (Ach.) Nyl. for their mutagenicity in the Ames assay (Table 6.3). Among them only physodalic acid exhibited significant genotoxicity against *Salmonella typhimurium* strain TA 100 with or without S9 mix in both plate-incorporation and preincubation assays. Later, after several studies, usnic acid was shown to possess genotoxic activity. In a study by Al-Bekairi et al. (1991), mice were treated orally with aqueous suspensions of (+)-usnic acid enantiomers in a single dose of either 100 or 200 mg/kg. A slight increase in the micronucleated polychromatic erythrocytes (MNPCEs) without affecting DNA synthesis was reported and an effect of usnic acid on spindle apparatus was suggested. (+)-Usnic acid and (–)-usnic acid enantiomers isolated from *Ramalina farinacea* and *Cladonia foliacea*, respectively, have been found to be non-genotoxic because of the absence of MN induction in HPL (Koparal et al. 2006). Polat et al. (2013) also showed the nonmutagenicity of usnic acid by CA and MN assays.

The first findings of genotoxic and antigenotoxic potentials of (+)-usnic acid against methylmethanesulfonate (MMS)-induced chromosomal and genome damage in mammalian cells in vitro and in vivo have been shown by Leandro et al. (2013). Usnic acid alone induced DNA damage at concentrations of 60 and 120 g/ml determined by COMET assay. However, it did not induce MN formation in V79 cells at the concentrations tested, and no genotoxic effects were observed in vivo. The



combined administration of usnic acid and MMS significantly reduced the frequencies of MN and DNA damage in vitro and in vivo when compared to treatment with MMS alone (Table 6.2). Recently, more results about genotoxic/antigenotoxic activities of usnic acid enantiomers isolated from two different lichen species, *Cladonia arbuscula* and *Cladonia stellaris*, have been reported by Prokopiev et al. (2017), Prokop'ev et al. (2018). Both (+)- and (–)-usnic acids at 0.01–1.00  $\mu\text{M}$  were inhibited by genotoxicity around 37–70%, induced by DN and MMS in HPL cells, as indicated by COMET assay. On the other hand, both enantiomers exhibited genotoxic effects at concentrations of 40–300  $\mu\text{M}$ , but (–)-usnic acid induced 2.5–3.5 times more genotoxicity in COMET than (+)-usnic acid. Usnic acid triggered oxidative stress and disruption of the normal metabolic processes of breast cancer cell line MCF7 and lung cancer cell line H1299 (null for p53); however, it was not involved in DNA damage. It was suggested that the property of usnic acid as a non-genotoxic anticancer agent that works in a p53-independent manner makes it a potential candidate for novel cancer therapy (Mayer et al. 2005).

Increasing amounts information on the genotoxicity and antigenotoxicity of other secondary metabolites of lichens are appearing. Lately, different species of lichen isolated by hypostictic acid, protocetraric acid, salazinic acid, and psoromic acid have been found to act as inhibitors of DXR-induced mutagenicity in the somatic cells of *Drosophila melanogaster* (Guterres et al. 2017). Tested concentrations of the compounds at 6 mM were not genotoxic. However, psoromic acid isolated from a different species, *Rhizoplaca melanophthalma*, and other two compounds as physosodic acid and olivetoric acid were genotoxic by inducing DNA damage at different concentrations from each other in HPL, PRCC, and U87MG cells (Emsen et al. 2016, 2018). There were notable differences among the 8-OH-dG levels caused by olivetoric acid and physosodic acid (5–40 mg/l) showing the highest genotoxic effects. Therefore, it was suggested that olivetoric acid and then physosodic acid may have high potential in the treatment of glioblastoma multiform. On the other hand, physosodic acid can be used as a natural antioxidant because of its low genotoxic but high antioxidant capacity.

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## 6.5 Conclusion

The methods most frequently used for the assessment of genotoxic and antigenotoxic activities of lichen extracts and products in vitro and in vivo are described here. These methods are not meant to be comprehensive of all existing methods, but more must be in consideration for further investigation of the genotoxicity for their safety assessment or antigenotoxicity, especially of secondary metabolites alone or in combination for their synergetic activities. Positive results of in vitro/in vivo testing indicate that the tested substance is genotoxic or antigenotoxic, and negative results indicate that the test substance is not genotoxic under the conditions of the assay performed. Genotoxicity and antigenotoxicity of lichens have appeared to be evaluated using several type of assays by detecting direct or indirect base substitution and frameshift mutagenicity (Ames and WP2),

clastogenicity (chromosome breakage), and aneugenicity (chromosome lagging from dysfunction of mitotic apparatus) (MN), numerical and structural DNA damage (CA), DNA-strand breaks (COMET), oxidative DNA damage (8-oxo-dG), and somatic mutation and recombination (SMART).

Accumulated data from short-term in vitro and in vivo studies have shown that lichen extracts could possess antigenotoxic effects. There are a small number of results for extracts which do not have antigenotoxic effects and some for those having genotoxic effects. The tests generally used for this purpose were common bacterial tests such as Ames and WP2 and human lymphocytes tests such as MN and SCE. However, there are gaps in the lichen genotoxicity/antigenotoxicity data because some groups studied only mutagenicity and others antigenotoxicity without genotoxicity. Most findings are extremely promising in that lichens may have therapeutic potential, at least for cancer, because of their antigenotoxic activities without genotoxic activity. The extracts of 13 species of the 20 lichen species tested, namely, *Cetraria aculeata*, *Cetraria islandica*, *Cetraria olivetorum*, *Cladonia chlorophaea*, *Cladonia foliacea*, *Cladonia rangiformis*, *Dermatocarpon intestiniforme*, *Parmelia pulla*, *Peltigera rufescens*, *Peltigera canina*, *Pseudevernia furfuracea*, *Ramalina capitata*, and *Xanthoria elegans*, have antigenotoxic activities but they are not genotoxic (Table 6.2). The extracts of 6 species, that is, *Evernia prunastri*, *Lecanora muralis*, *Rhizoplaca chrysoleuca*, *Usnea longissima*, *Umbilicaria vellea*, and *Xanthoparmelia somloensis* are antigenotoxic, but not tested for their genotoxic activities. On the other side, the aqueous extracts of *Cetraria aculeata* and *Rhizoplaca melanophthalma* are neither genotoxic nor antigenotoxic for *E. coli* WP2 and human peripheral blood lymphocytes, respectively. The aqueous extracts of the other eight lichen species tested are not also genotoxic, except for the methanol extract of *Hypogymnia physodes* (Table 6.2).

There is minor evidence about the genotoxic and antigenotoxic activities of the secondary metabolites of lichens. Hypostictic acid, protocetraric acid, psoromic acid, and salazinic acid are antigenotoxic without genotoxic activity in *Drosophila* somatic cells. However, psoromic acid induced oxidative DNA damage in mammalian cells. Although physodic acid is nonmutagenic in the Salmonella assay, it induces oxidative DNA damage in mammalian cells.

Interestingly, secondary metabolites of the lichens tested show variation in their effects because each is either genotoxic or antigenotoxic according to the type of assays and laboratories where the work was done. Also, variation in the effective doses of the extract on different cells or test systems suggests the necessity of more in vitro and in vivo antigenotoxicity studies to know the exact potential of the extract. It may then find application for treatment. Further investigation to complete the gap and more data for other lichen species will be very useful for possible therapeutic applications.

The mechanism of antigenotoxic action of all these lichen extracts is not completely known but appears to be the antioxidative potential of their secondary metabolites, as described in Chap. 1. Because most of the extracts have been investigated for their antigenotoxicity and antioxidant activities, quite strong antioxidative activity is also indicated (Aydin and Türkez 2011a, b; Kotan et al.

2011; Polat et al. 2013; Kekuda et al. 2018). The chemoprevention potential of several lichen extracts or secondary metabolites against DNA damage induced by the known compounds such as AFB1, MNNG, MMS, IMA, and CBS strongly indicates that lichens can be a resource of new therapeutics.

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