Toxicological mechanisms of current flame retardants

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1. Introduction

There is an ever-increasing exposure to complex mixtures of chemicals in our daily lives. One such family of compounds that are produced at high volume are flame retardants (FRs). FRs are introduced into materials with the purpose of preventing the initiation and spread of fire. They are present in nearly all manufactured items and are able to diffuse out of materials and contaminate surrounding environments [1]. Accordingly, many FRs have been detected in household dust [2], and have been detected in human breast milk and serum as a consequence of daily exposure [3]. More worryingly, FRs are able to be transferred from mother to infant during pregnancy and breast-feeding stages [4]. In addition, previously-used FRs have been shown to elicit a wide range of toxicological effects and have thus been banned from use. Considering the structural similarity to their toxic predecessors, their persistence, bioaccumulation, and lack of insight of toxicological and molecular mechanisms, currently-used FRs pose a significant risk.

We employed a bacterial gene profiling assay to investigate *in vitro* effects of 12 currently-used FRs on a selection of general bacterial stress responses. Such responses included responses to oxidative stress, DNA damage, membrane damage, and general cell lesions such as protein degradation and growth arrest [5]. Bacterial biosensors are frequently used to assess ecotoxicological impacts of compounds since they are particularly useful in compound screening and classification according to mode of action [6].

2. Results and discussion

2.1. Gene inductions

A majority of FRs significantly affect multiple stress genes (Table 2). However, only a few genes were induced at any significant level (>2-fold) (data not shown). These genes included *ClpB*, *RecA*, and *MicF*, indicating possible effects on protein, DNA, and membrane integrity associated with these compounds. Additionally, these effects could be a result of reactive oxygen species (ROS) production attributed to these compounds, as evidenced by the significant induction of *KatG*, *Zwf*, *Soi28*, and *Nfo*.

					Gene	ral Cell						Membrane	
r	Oxidative Stress				Lesions		DNA Damage					Damage	
Flame Retardant	KatG	Zwf	Soi28	Nfo	ClpB	UspA	RecA	UmuDC	Ada	SfiA	Nfo	MicF	OsmY
тсрр	**	**		***		**	*	**	**	*	***	**	*
ТРР	**				**	*	****					**	**
ТВЕР	***	*	*		*	***			**			**	****
TDCPP	**	*			**	**			**			**	*
TCEP	**	**		**	**		**	**	****		**	***	**
TnBP	*	*	*		*	*						**	****
ТЕР			**	**	*			**	**	**	**		
DOPO		****	**	****		**	**		***	**	****		***
HBCD		**	*		**	*	*		*	***			
ТВВРА	**	**			**		****	*				**	
тврн	*		*		**	**	*						**
ТВС	***	***	**	**		****	***		**		**	**	***

Table 1: Significance of inductions of stress genes by FRs at highest concentration. *p<0.05 **p<0.01 ***p<0.001 ***p<0.0001.

2.2. Clustering

Clustering analysis identified two additional clusters involving *OsmY*, and *UspA* stress genes, responsive to membrane damage and growth arrest respectively (Figure 1). These FRs therefore affect multiple toxicological modes of action which include protein, membrane, and DNA damage, along with growth arrest. Additionally, ROS production could be the underlying mechanism resulting in these effects, in agreement with previous studies showing ROS induction in response to FRs [7].

3. Conclusions

FRs effect several toxicological modes of action on prokaryotic cells. Given that many of these bacterial genes have eukaryotic homologues, such results are relevant to higher biological systems [8]. Finally, the lack of any notable gene induction following DOPO treatment along with its excellent fire-retardation supports its increasing interest as an alternative to halogenated FRs [9]. Full details of this work have recently been published [10].

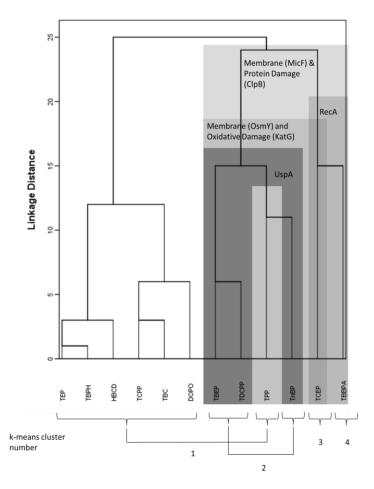


Figure 1: Clustering using hierarchical and k-means algorithms

4. References

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