GENOTOXICITY OF THE CYCLO-OXYGENASE-INHIBITOR SULINDAC SULFIDE IN THE FILAMENTOUS FUNGUS Aspergillus nidulans

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ABSTRACT

Sulindac sulfide is a non-steroidal anti-inflammatory drug (NSAID) with chemopreventive effect on human cancer cells. Due to the involvement of the somatic recombination in the carcinogenic process, sulindac sulfide's recombinogenic potential was evaluated by the Homozygotization Index (HI) in the filamentous fungus *Aspergillus nidulans*. The drug's recombinogenic potential was evaluated by its capacity to induce homozygosis of recessive genes from heterozygous diploid cells. Sulindac sulfide at 175 and 350 µM concentrations induced mitotic recombination in *A. nidulans* diploid cells, with HI values for genetic markers higher than 2.0, and significantly different from control HI values. The recombinogenic effect of NSAID was related to the induction of DNA strand breaks and cell cycle alterations. Sulindac sulfide's carcinogenic potential was also discussed.

Key words: antineoplastic agents, mitotic crossing-over, secondary malignances.

INTRODUCTION

Non-steroidal anti-inflammatory drugs (NSAIDs) are primarily analgesic, anti-inflammatory and antipyretic compounds which inhibit the cyclo-oxygenase-mediated production of prostaglandins (1,3). Two COX isoenzymes, COX-1 and COX-2, exist in human cells. The former (COX-1) is constitutively expressed, whereas the latter, inducible isoenzyme COX-2, is expressed only in response to tumor promoters, endotoxin, cytokines and hormones (10,27). The main product of COX-2, namely prostaglandin $E_2(PGE_2)$, is found at high levels in tumor cells. In clinical studies, PGE_2 have been associated with high metastatic potential of neoplastic cells (25,26).

Several studies have reported an inverse relationship between cancer incidence and regular use of NSAID, including aspirin (12,15). The dietary administration of NSAIDs, such as ibuprofen and celecoxib, significantly reduced the incidence, multiplicity and volume of tumors in female Spragne-Dawley rats with 7,12-dimethylbenzathracene-induced mammary carcinomas (9,24). Celecoxib also induced apoptosis in human prostate cancer cell (11).

Sulindac is a non-steroidal anti-inflammatory drug with prophylactic effects in the prevention of colon cancer. Sulindac and its metabolites, such as sulindac sulfide, inhibited the proliferation in cultured colon cancer cells and induced DNA strand breaks and cell death by apoptosis (22,23). Patients with familial adenomatous polyposis showed reduction of both the number and size of colorectal adenomas after the administration of sulindac (21).

In a variety of human tumor cells, including human lung squamous cells and human leukemia cell lines, sulindac and other NSAIDs enhance the cytotoxicity of certain anticancer drugs, such as doxorubicin, daunorubicin and epirubicin. Results suggest that active NSAIDs inhibit MRP-mediated drug efflux which confers multidrug resistance (MDR) (5,6). Classical MDR is characterized by cross-resistance to a range of chemically unrelated drugs, whereas MRP (MDR-associated protein) is a transmembrane transporter (18).

Sulindac sulfide's chemopreventive effect has been recently associated with alterations in the gene expression in gastric and ovarian cancer cells. Sulindac sulfide has been characterized as a NAG-1 gene inducer in ovarian cancer cells. Since human

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NAG-1 is linked to apoptosis, the reduced expression of NAG-1 gene may enhance tumorigenesis (12,16,17).

Since somatic recombination is mechanisma leading to carcinogenesis (7,8), current *in vivo* study shall evaluate the genotoxicity of sulindac sulfide in heterozygous cells of *Aspergillus nidulans*.

MATERIAL AND METHODS

Strains and culture media. Diploid strain was formed with UT448 and A757 haploid strains (Table 1). Minimal Medium (MM) comprised Czapek-Dox with 1% (w/v) glucose, whereas Complete Medium (CM) consisted of peptone (2,000 mg/L), yeast extract (2,000 mg/L), hydroliysed casein (1,000 mg/L), glucose (10,000 mg/L), biotin (2 mg/L), pyridoxine (50 mg/L), p-aminobenzoic acid (50 mg/L), folic acid (50 mg/L), nicotinic acid (100 mg/L), pantothenic acid (200 mg/L), choline chloride (200 mg/L), riboflavin (100 mg/L), inositol (400 mg/L) and thiamine (50 mg/L) added to MM. Selective medium (SM) comprised MM plus riboflavin, p-aminobenzoic acid, biotin, methionine and pyridoxine, while solid medium contained 1.5% agar. Incubation for strain growth occurred at 37°C.

Sulindac sulfide treatment. Drug concentration that induced the arrest of the human colon adenocarcinoma cell in G1/S phase (175 μ M) (23) and a higher concentration (350 μ M) were used in present study. They were the final ones in MM. Sulindac sulfide was dissolved in 10 μ L DMSO and added to molten MM. Solvent was per se neither visibly toxic (Fig. 1) nor recombinogenic for diploid strain (results not shown).

Evaluation of drug toxicity. Conidia of diploid strain A757//UT448 were inoculated at the center of plates with MM (control) and MM + sulindac sulfide (treatment). Ten plates were inoculated for each dose and for controls and incubated at 37°C. Measurement of colonies' diameter occurred after 48 to 144 hours incubation. Values in the presence and in the absence of the drug were compared by Student's *t* test.

Table 1. Genotype and origin of *A. nidulans* strains.

Strains	Genotype	Origin
A757	yA2 (I), $methA17$ (II),	
	pyroA4 (IV)	FGSC*
UT448	riboA1 (I), $pabaA124$ (I),	
	biA1(I), AcrA1(II), wA2(II)	Utrecht

Requirements for: riboflavin = riboA1, p-aminobenzoic acid = pabaA124, biotin = bioA1, methionine = methA17, pyridoxine = pyroA4. Conidia color: white = wA2; yellow = yA2. AcrAI, resistance to acriflavine; *FGSC = Fungal Genetic Stock Center, University of Kansas Medical Center, Kansas, USA.

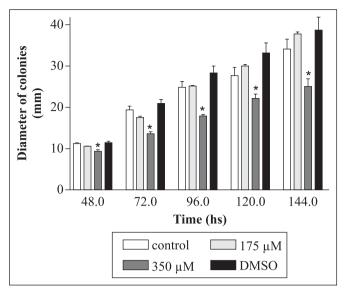


Figure 1. Growth of *A. nidulans* diploid strain A757//UT448 on plates containing Minimal Medium (control) and Minimal Medium plus sulindac sulfide (175 μ M and 350 μ M). (*) Strain growth significantly different from the control (p < 0.05).

Calculating Homozygotization Index. Conidia of diploid strain A757//UT448 were individually inoculated in plates with MM + sulindac sulfide at 175 and 350 µM concentrations. Plates were incubated for 5 days at 37°C. Treatment produced visible mitotic sectors (D1-D6), identified by their morphology which differed from the original diploid (Fig. 2A-B). Diploid sectors were purified in MM and then transferred individually to plates with CM. They were then processed by spontaneous haploidization. After haploidization, the haploid mitotic segregants, obtained from diploids D1-D6, were purified in CM and their mitotic stability evaluated. Only mitotic stable segregants at the final stage were selected for the recombinogenesis test. Conidia of each haploid segregant were individually transferred to 25 positions in CM plates. After 48 hours incubation colonies were transferred to proper selective media to determine their phenotypes. Mitotic crossing-over causes homozygotization of heterozygous-conditioned genes. If sulindac sulfide induces mitotic crossing-over in diploid strain A757//UT448, only heterozygote (+/- or -/+) or homozygote (+/-+) diploids will develop in MM. In this case, nutritional markers will segregate among the haploids at the proportion of 4+:2-. On the other hand, if NSAID does not induce crossing-over, proportion will amount to 4+:4-. This happens because the initial selection process limits the growth of diploids -/-. Homozygotization Indexes (HI) (the ratio between prototrophic segregants and auxotrophic segregants), equal to or higher than 2.0, indicate the sulindac sulfide's recombinogenic effect (13). Results were compared by Yates correct Chi-square test.

Table 2. Homozygotization Index (HI) values for markers from UT448 // A757 diploid strain after treatment with 175 μ M (D1-D3) and 350 μ M (D4-D6) of sulindac sulfide

Markers ^a	Control ^b		D1		D2		D3		D4		D5		D6	
	NS	НІ	NS	НІ	NS	НІ	NS	НІ	NS	НІ	NS	НІ	NS	НІ
ribo+	33	1.2	50	1.7	37	1.2	41	1.8	30	0.8	50	7.1*	30	1.0
ribo	27		29		30		23		36		07		29	
paba+	34	1.3	50	1.7	36	1.2	41	1.8	66	nd	50	7.1*	32	1.2
paba	26		29		31		23		0		07		27	
bi+	33	1.2	54	2.2*	44	1.9	50	3.6*	57	6.3*	50	7.1*	40	2.1
bi	27		25		23		14		09		07		19	
meth+	33	1.2	71	8.9*	61	nd	58	nd	66	nd	56	nd	51	6.4*
meth	37		08		06		06		0		01		08	
pyro+	31	1.1	49	1.6	39	1.4	41	1.8	38	1.35	32	1.3	42	2.5*
pyro	29		30		28		23		28		25		17	

^a ribo = riboflavin; paba = p-aminobenzoic acid; bi = biotin; meth = methionine and pyro = pyridoxine. ^bNot treated with sulindac sulfide. *significantly different from control (p< 0.05). nd= not determined (see text). NS = number of segregants.

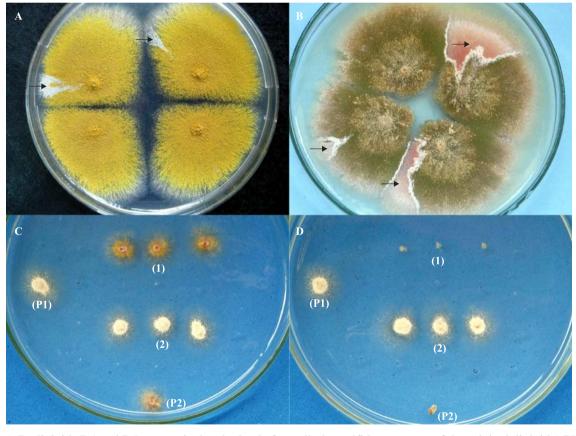


Figure 2. A-B, diploids D4 and D5, respectively, obtained after sulindac sulfide treatment of the original diploid A757//UT448 strain. Arrows indicate the mitotic segregants selected for the recombinogenesis tests. Mitotic segregants w+ meth (1) and w meth+ (2) from D2 and D3 diploids growing in the presence (C) and in the absence of methionine (D); P1 and P2 are the paternal UT448 and A757 stains, respectively.

RESULTS AND DISCUSSION

Current researchers first examined the manner sulindac sulfide affected the mycelial growth of the fungal strain. When compared to control (untreated conidia), fungal growth was significantly inhibited after cell exposition to 350 μ M sulindac sulfide (Fig. 1). Since sulindac sulfide caused strong inhibition of G1 to S phase of the cell cycle on cultured human cancer cells (14), the cytotoxic effect of NSAID in *A. nidulans* diploid cells might be explained by cell cycle delay or by cell death mechanism.

Sulindac sulfide's genotoxicity was evaluated in current study by its capacity to induce homozygosis during the mitotic propagation of the heterozygous diploid A757//UT448 strain. Results show (Table 2) that cytotoxic (350 μ M) and noncytotoxic (175 μ M) concentrations of the drug significantly increased HI values of the nutritional markers when compared to those of control.

Diploids D1 and D3, derived from A757//UT448 strain after treatment with 175 μ M sulindac sulfide, showed homozygotization indexes over 2.0 for bi and meth genes. Results indicate the occurrence of mitotic crossing-over on chromosomes I and II from the original diploid strain. HI values for meth gene from mitotic segregants derived from D2 and D3 strains could not be determined because the meth segregants obtained in the analyses were not phenotypically recombinant, but parental (w+, meth) (Fig. 2C-D). Contrastingly, all meth segregants obtained after haploidization of D1 strain were recombinants w, meth (Tables 1 and 2).

Diploid strains, obtained with 350 µM sulindac sulfide (D4 to D6), showed *ribo*, *paba*, *bi*, *meth* and *pyro* genes HI values higher than 2.0 and significantly different from respective control HI values. The absence or the reduced number of *meth* segregants derived from D4 and D5 strains, respectively, may be justified by a greater number of white (w) segregants isolated after haploidization. In the paternal UT448 strain the w gene is closely linked to the *meth*+ gene and consequently most of the w segregants have the w *meth*+ phenotype. Similar to the D2 and D3 analyses, the *meth* segregant recovered after D5 haploidization was not phenotypically recombinant (Table 2).

Although the treatment of original A757//UT448 diploid with sulindac sulfide in MM does not allow the isolation of auxotrophic diploids (-/-), recessive homozygous diploids for conidia color *y* marker may be obtained. The 175 μM sulindac sulfide treatment of A757//UT448 diploid strain only allowed the isolation of prototrophic diploids with green conidia (*y*+//*y*). On the other hand, homozygous *y*//*y* diploid strains were obtained (D4 and D6) with sulindac sulfide at 350 μM concentration. The segregation of markers from chromosome I (*ribo*, *paba* and *bi*) among the mitotic segregants of D6 strain indicates the heterozygous condition for genes: On the other hand, the absence of *paba* segregation in D4 analysis suggests

a homozygous condition for this marker in D4 diploid strain (Table 2).

As a whole, results show that sulindac sulfide is an effective somatic crossing-over inducer in diploid cells of *A. nidulans*. Sulindac sulfide's recombinogenic potential may indicate a direct effect of the drug on DNA, inducing the occurrence of DNA breaks and cell cycle alterations.

Cancer is a progressive process which results from environmental and endogenous agents' toxic effects. The disease develops from a single oncogenic mutation-bearer cell (either a proto-oncogene or a tumor suppressor gene) whose primary mutagenic event is called initiation. Since clonal expansion of an initiated cell is necessary for the fixation of the first mutation and tumor promotion, the probability is high for additional genetic alterations in decisive genes required for the tumor progression (20).

Although a normal cell may be a mutation bearer in a tumor suppressor gene (m) owing to the presence of the normal allele (m+), the loss of the heterozygous condition, caused by mitotic recombination, may be the factor which advances neoplasm (19). Actually, the loss of heterozygosity in human retinoblastoma is the most common mechanism which triggers the transformation process of a normal cell and makes it a neoplastic one (4,8). As a consequence of mitotic crossing-over, the mitotic products are rendered homozygous from all heterozygous loci distal to the point of exchange, if a recombinant and a parental chromatid segregate towards the same mitotic pole (2). Mitotic recombination, in fact, represents an important function in expressing deleterious or oncogenic mutations in heterozygous initiated cell.

Results in current study show that sulindac sulfide induces homozygosis of genes previously masked by the dominant allele. Consequently, the above anti-inflammatory drug may be characterized as a tumor promoter agent that potentially induces secondary malignancies by mitotic crossing-over.

RESUMO

Genotoxicidade de sulfeto de sulindaco em Aspergillus nidulans

Sulfeto de sulindaco é um antiinflamatório não-esteroidal com efeitos quimiopreventivos em cânceres humanos. O presente estudo teve como objetivo avaliar o potencial recombinagênico do sulfeto de sulindaco em células diplóides de *Aspergillus nidulans*. O efeito recombinagênico da droga foi demonstrado através da homozigotização de genes recessivos, previamente presentes em heterozigose. Os valores de HI (Índice de Homozigotização) para diferentes marcadores genéticos apresentaram-se maiores do que 2,0 e significativamente diferentes dos valores obtidos em sulfeto de sulindaco ausência da droga (controle). O potencial

recombinagênico do sulfeto de sulindaco foi associado à indução de quebras na molécula do DNA e a alterações no ciclo celular. O potencial carcinogênico do sulfeto de sulindaco foi discutido no presente trabalho.

Palavras-chave: agentes antineoplásicos, crossing-over mitótico, malignidade secundária, recombinação somática

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