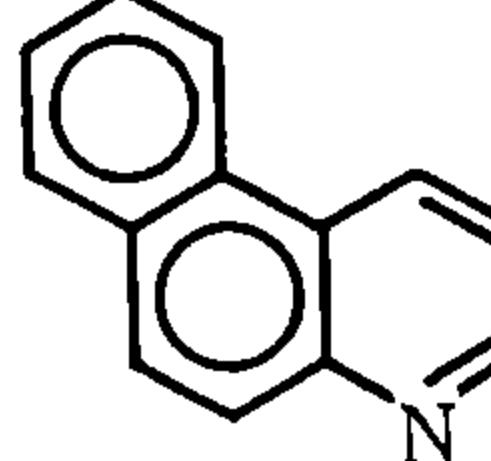


Benzo[f]quinoline (ベンゾ[f]キノリン)

Experimental Data

<u>Chemical Name:</u>	<u>Benzo[f]quinoline</u>	
<u>Synonym</u>	<u>β-Naphthoquinoline</u>	
	<u>Naphthopyridine</u>	
<u>Molecular weight:</u>	179.22	
<u>Boiling point:</u>	210-215°C	
<u>Melting point:</u>	93.5°C(93-94°C)	
<u>Chemical Structure</u>		
<u>CAS No :</u>	85-02-9	
<u>Source of Substance:</u>	Tokyo Kasei Kogyo Co., Ltd.	
<u>Lot. No.:</u>	MAL01	
<u>Purity:</u>		
<u>Vehicle:</u>	DMSO	

Judgement for
Chromosomal Aberration in CHL: Positive

	Treated Time (Hr)	Concen-ration (mg/ml)	No. of Mate-phase	Poly-ploid (%)	Judge	Cell with Structural Chromosome Aberration (%)						Total	Judge
						Gap	CTB	CTE	CSB	CSE	-G	+G	
DMSO	24		200	0	—	1.0	1.5	0.5	0	0	2.0	2.5	—
	48		200	1.0	—	0.5	0	0	0	0	0	0.5	—
Test chemical													
	24	0.05%	200	1.0	—	0.5	0	0.5	0	0	0.5	1.0	—
		0.01	200	4.5	—	2.5	2.0	1.0	0	0	3.0	5.5	±
		0.02	200	14.0	+	6.0	2.0	6.0	0.5	0	8.0	12.5	+
		0.04	200	0	—	11.0	19.0	38.5	0.5	0	49.5	52.0	+
		0.06			No observation for metaphase								
	48	0.05	200	0.5	—	0	0	0	0	0	0	0	—
		0.01	200	1.0	—	2.0	0.5	0.5	0	0	1.0	3.0	—
		0.02	200	14.0	+	3.5	3.5	4.5	0	0.5	8.5	12.0	+
		0.04			No observation for metaphase								
		0.06			No observation for metaphase								
Positive Control													
(MMC)	24	0.00008	200	0.5	—	11.5	15.0	47.5	0	0	52.5	54.5	+
	48	0.00008	200	1.0	—	8.5	28.5	83.5	0	0	86.5	87.0	+

IARC Evaluation : not yet cited

Experimental Data

S 9 with or without	Concen- tration (mg/ml)	No. of Meta- Phase	Poly- ploid (%)	Judge	Cell with Structural Chromosome Aberration (%)				Total			
					Gap	CTB	CTE	CSB	CSE	-G	+G	Judge
DMSO	—	200	0.5	—	0	0	0	0	0	0	0	—
	+	200	1.0	—	0.5	0.5	0.5	0	0	0.5	1.0	—
Test chemical												
	—	0.01%	200	1.0	—	1.0	0	0.5	0	0	0.5	1.5
		0.015	200	2.5	—	0	1.0	1.0	0	0	2.0	2.0
		0.02	200	1.5	—	0	0.5	1.0	0	0	1.5	1.5
		0.025	200	6.0	±	0.5	0	0	0	0	0.5	—
		0.03	200	8.5	±	0	0.5	0.5	0	0	1.0	1.0
	+	0.01	200	0	—	0.5	0	0.5	0	0.5	1.0	1.5
		0.015	200	1.5	—	1.0	1.5	4.0	0	0	5.5	6.5
		0.02	200	2.0	—	3.0	8.5	12.5	0	0	17.5	18.5
		0.025	189	0.5		3.7	10.1	15.3	0	0	20.6	21.2
		0.03			No observation for metaphase							
Positive Control												
(B(a)P)	—	0.008	200	2.5	—	1.5	1.5	1.0	0	0	2.5	4.0
	+	0.008	200	0	—	8.5	10.0	50.0	0	0	53.5	55.5