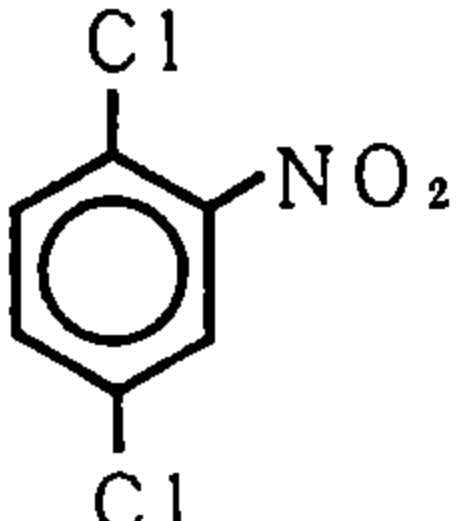


1,4-Dichloro-2-nitrobenzene (1,4-ジクロロ-2-ニトロベンゼン)

Experimental Data

Chemical Name: Synonym	1,4-Dichloro-2-nitrobenzene		Treated Time (Hr)	Concentration (mg/ml)	No. of Meta-phase	Poly-ploid (%)	Cell with Structural Chromosome Aberration (%)							Total	Judge			
	p-Dichloronitrobenzene	2,5-Dichloronitrobenzene					Judge	Gap	CTB	CTE	CSB	CSE	-G	+G				
Molecular weight:	192.0		CMC	24	200	0.5	—	0	0	0	0	0	0	0	—			
Melting point:	54-57°C			48	200	0	—	0.5	0	0.5	0	0	0.5	1.0	—			
Boiling point:	266-269°C		Test Chemical															
Flashing point:	> 110°C			24	0.2	200	0.5	—	1.0	0.5	0	0	0	0.5	1.5	—		
Chemical Structure					0.3	200	0.5	—	0.5	0	0	0	0	0	0.5	—		
CAS No :	89-61-2				0.4	200	0.5	—	1.0	0	0.5	0	0	0.5	1.5	—		
MITI No :	(3)-261				0.5	200	0.5	—	2.0	0	0	0	0	0	2.0	—		
Source of Substance:	Wako Pure Chem. Ind., Ltd.				0.6	200	0	—	1.0	0.5	0	0	0	0.5	1.5	—		
Lot. No.:	LAR4761			48	0.2	200	0	—	0	0	0	0	0	0	0	—		
Purity:	%				0.3	200	0.5	—	0.5	0	0	0	0	0	0.5	—		
Vehicle:	1%CMC				0.4	200	1.5	—	0.5	0	0	0	0	0	0.5	—		
Judgement for Chromosomal Aberration in CHL: Positive					0.5	200	0.5	—	0	0	0	0	0	0	0	—		
					0.6	200	5.5	±	1.0	0	1.0	0	0	1.0	2.0	—		
Positive Control																		
Judgement for Chromosomal Aberration in CHL: Positive	(MMC)	24	0.00008	200	0	—		3.5	7.5	25.0	0.5	0	29.0	31.5	+			
		48	0.00008	200	0	—		2.0	7.5	35.0	0	0.5	39.0	39.5	+			

IARC Evaluation : not yet cited

Metaphase was not observed at the concentration of 0.7mg/ml.

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
CMC	—	200	0.5	—	0	0	0	0	0	0	0	—
	+	200	0	—	1.0	0	0	0	0	0	1.0	—
<b>Test Chemical</b>												
—	0.038	200	0	—	0	0.5	0.5	0	0	0.5	0.5	—
	0.075	200	0	—	0	0	2.0	0	0	2.0	2.0	—
	0.15	200	0	—	0	0	0	0	0	0	0	—
	0.3	200	0.5	—	0.5	0	0	0	0	0	0.5	—
	0.45	200	0	—	1.0	1.5	0.5	0.5	0	2.5	3.5	—
+	0.038	200	1.0	—	0.5	0	0.5	0	0	0.5	1.0	—
	0.075	200	0	—	1.0	0	1.5	0	0	1.5	2.5	—
	0.15	200	6.0	±	1.5	1.0	5.5	0	0	6.0	7.0	±
	0.3	200	4.5	—	3.5	6.0	10.0	0	1.0	15.0	17.0	+
	0.45	No observation for metaphase										—
<b>Positive Control</b>												
(B(a)P)—		200	0.5	—	0	0	0.5	0	0	0.5	0.5	—
+		200	0	—	4.5	0	22.5	0	0	24.5	26.0	+