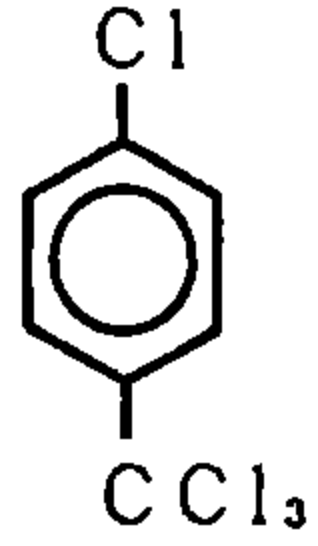


p-Chlorobenzotrichloride (p-クロロベンゾトリクロリド)

Chemical Name:	p-Chlorobenzotrichloride
Synonym	1-Chloro-4-(trichloromethyl)-benzene
	Benzene, 1-chloro-4-(trichloromethyl)-
Molecular weight:	229.92
Melting point:	0.6 °C
Boiling point:	24.5 °C
Flashing point:	>110 °C
Chemical Structure	
CAS No :	5216-25-1
NITI No :	(3)-89, (3)-93
Source of Substance:	Tokyo Kasei Kogyo Co., Ltd.
Lot.No. :	FBNO1
Purity:	%
Vehicle:	1%CMC

Experimental Data

	Treated Time (Hr)	Concentration (mg/ml)	No. of Meta-phase	Poly-ploid (%)	Judge	Cell with Structural Chromosome Aberration (%)							
						Total		CSB	CSE	-G	+G	Judge	
						Gap	CTB						CTE
CMC	24		200	0.5	—	0.5	0	0	0	0	0	0.5	—
	48		200	0	—	0.5	0	0	0	0	0	0.5	—
Test Chemical													
	24	0.025	200	0	—	1.0	1.5	0.5	0	0	2.0	3.0	—
		0.05	200	0	—	0.5	1.5	3.5	0	0	4.5	5.0	±
		0.1	200	0.5	—	0.5	3.5	7.0	0	0	10.5	10.5	+
		0.2	200	2.0	—	2.0	5.0	13.5	0	0	16.0	17.0	+
		0.4	200	1.0	—	4.0	11.5	29.0	0	0	35.0	36.0	+
	48	0.025	200	0.5	—	0.5	0	0.5	0	0	0.5	1.0	—
		0.05	200	0	—	0.5	0	1.5	0	0	1.5	2.0	—
		0.1	200	1.0	—	0	1.0	2.5	0	0	3.0	3.0	—
		0.2	200	0.5	—	2.5	2.0	4.5	0	1.0	7.0	8.5	±
		0.4	200	0	—	4.5	3.5	12.0	0	0	13.5	14.5	+
Positive Control													
(MMC)	24	0.00008	200	0	—	7.0	13.0	41.5	0	0	46.0	48.0	+
	48	0.00008	200	2.0	—	11.5	12.5	46.0	0	0	53.0	53.5	+

Judgement for Chromosomal Aberration in CHL: Positive

IARC Evaluation : not yet cited

Experimental Data

S 9 with or without	Concent- ration (mg/ml)	No. of Meta- phase	Poly- ploid (%)	Judge	Cell with Structural Chromosome Aberration (%)							Judge
					Gap	CTB	CTE	Total		-G	+G	
								CSB	CSE			
CMC	-	200	2.0	-	0	0	1.0	0	0	1.0	1.0	-
	+	200	0.5	-	0	1.5	1.0	0	0	2.5	2.5	-
Test Chemical												
-	0.026	200	0.5	-	0.5	0	1.5	0	0	1.5	2.0	-
	0.064	200	0	-	0.5	1.5	3.5	0	0	4.5	4.5	-
	0.16	200	0.5	-	1.0	3.5	7.5	0	0	9.5	10.5	+
	0.4	200	0.5	-	5.5	6.5	46.5	0	0	48.5	50.0	+
	1.0				No observation for metaphase							
+	0.026	200	1.0	-	0	1.5	1.0	0	0	2.5	2.5	-
	0.064	200	0	-	1.5	0.5	8.0	0	0	8.0	9.0	±
	0.16	200	0	-	3.0	3.0	18.0	0	0	21.0	22.5	+
	0.4	200	0	-	8.5	18.5	61.5	0	0.5	66.5	67.5	+
	1.0				No observation for metaphase							
Positive Control												
(B(a)P)	-	200	0	-	1.0	0	0.5	0	0	0.5	1.5	-
	+	200	1.0	-	3.0	5.5	25.0	0	0	27.5	29.0	+