

2, 4, 6-Trimethylaniline (2, 4, 6-トリメチルアニリン)

Chemical Name: 2, 4, 6-Trimethylaniline
 Synonym: 2, 4, 6-Trimethylbenzenamine
 Mesidine

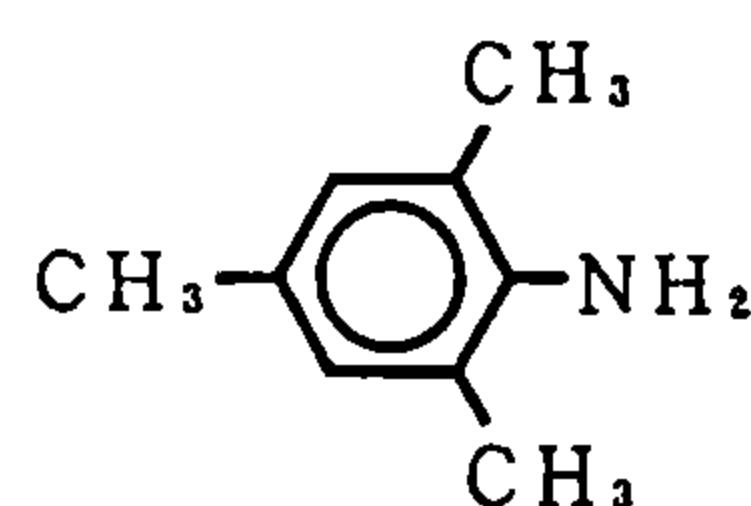
Molecular weight: 135.2

Melting point: -5°C

Boiling point: 232-233°C

Flashing point 96°C

Chemical Structure



CAS No: 88-05-1

MITI No: (3)-132

Source of Substance: Wako Pure Chem. Ind. Ltd.

Lot. No: LAQ5139

Purity: %

Vehicle: DMSO

Judgement for

Chromosomal Aberration in CHL: Positive

Experimental Data

	Treated Time (Hr)	Concen- ration (mg/ml)	No. of Meta- phase	Poly- ploid (%)	Judge	Cell with Structural Chromosome Aberration (%)					Total		
						Gap	CTB	CTE	CSB	CSE	-G	+G	
DMSO	24		200	0	-	0.5	0	0	0	0	0	0.5	
	48		200	0.5	-	0	0	0.5	0	0	0.5	0.5	
Test Chemical													
	24	0.1	200	0.5	-	0.5	0.5	0	0	0	0.5	1.0	
		0.2	200	0.5	-	0.5	0.5	1.0	0	0	1.5	2.0	
		0.4	200	1.0	-	1.5	0.5	0.5	0	0	1.0	2.5	
		0.6				No observation for metaphase							
		0.8				No observation for metaphase							
	48	0.1	200	0	-	0.5	0	1.0	0	0	1.0	1.5	
		0.2	200	1.0	-	0.5	0	0	0	0	0	0.5	
		0.4	200	0	-	1.0	0	0.5	0	0	0.5	1.5	
		0.6				No observation for metaphase							
		0.8				No observation for metaphase							
Positive Control													
(MMC)	24	0.00008	200	0	-	6.0	14.5	40.0	0	0	49.0	50.0	
	48	0.00008	200	2.0	-	6.5	12.0	46.5	0	0	54.5	55.0	

IARC Evaluation: G 3

Experimental Data

S 9 with or without	Concen- tration (mg/ml)	No. of Meta- phase (%)	Poly- ploid	Judge	Cell with Structural Chromosome Aberration (%)						Total	Judge	
					Gap	CTB	CTE	CSB	CSE	-G	+G		
DMSO	—	200	0.5	—	0	0	0	0	0	0	0	—	
	+	200	0.5	—	1.0	0.5	0	0	0	0.5	1.5	—	
Test Chemical													
—	0.05	200	0	—	0.5	0	0	0	0	0	0.5	—	
	0.10	200	0.5	—	1.0	0	0	0	0	0.5	1.5	—	
	0.15	200	0.5	—	1.0	0	0	0	0	0	1.0	—	
	0.20	200	0.5	—	0.5	0	0	0	0	1.5	2.0	—	
	0.25	200	0	—	0	0	0	0	0	1.5	1.5	—	
	0.05	200	1.0	—	1.5	0	0.5	0	0	0.5	2.0	—	
	0.10	200	0.5	—	3.0	1.5	10.0	0	0	10.5	12.0	+	
	0.15	200	0	—	5.0	7.5	38.0	0	0	39.0	40.5	+	
	0.20				No observation for metaphase								
	0.25				No observation for metaphase								
Positive Control													
(B(a)P)				—	0.016	200	1.0	—	0.5	0	1.5	0	1.5
				+	0.016	200	0.5	—	4.5	10.0	45.5	0	50.0
									0	0	50.5	+	