

[ベンジルクロロド]

Chemical Name; Benzyl chloride

Synonym ; α -Chlorotoluene

(Chloromethyl)benzene

 α -クロロトルエン

(クロロメチル)ベンゼン

塩化ベンジル

Molecular Weight ; 126.59

Melting Point ; -43 °C [CHCD, Aldrich]

-48 °C [CHCD]

-48 ~ -43 °C [Merck]

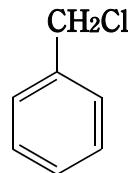
Boiling Point ; 179 °C [CHCD, Merck]

177 ~ 181 °C [Aldrich]

Flashing Point ; 73 °C [Aldrich]

Molecular Formula; C₇H₇Cl

Chemical Structure



CAS No. ; 100-44-7

MITI No. ; (3)-39, (3)-102

ML No. ; -

Specified Chemical Substances; -

Source of Substance; Wako Junyaku Kogyo Co., Ltd.

Lot No. ; APQ5976

Purity ; 100.0 %

Vehicle ; DMSO

Experimental Data without Metabolic Activation

Substance	Time (h)	Concen- tration (mg/ml)	No. of Metaphase	Polyploid (%)	Judge- ment	Cell with Structural Chromosome Aberration (%)						Judge- ment
						Gap	Chromatid CTB	Chromatid CTE	Chromosome CSB	Chromosome CSE	Total -G	
Test Chemical	24	0.01	200	0.5	-	0.0	0.0	1.0	0.0	0.0	1.0	1.0
						0.0	0.0	0.0	0.0	0.0	0.0	0.0
	48	0.02	200	0.5	-	0.0	2.0	2.0	0.0	0.0	3.5	3.5
						0.0	2.5	5.5	0.0	0.0	7.0	7.0
		0.03	200	0.5	-	1.5	7.0	17.5	0.0	0.0	19.5	19.5
						2.5	12.0	42.0	0.0	0.0	46.0	46.5
		0.04	200	4.0	-	0.0	0.0	0.0	0.0	0.0	0.0	0.0
						0.0	0.5	2.5	0.0	0.0	3.0	3.0
		0.05	200	4.0	-	0.0	0.5	3.0	0.0	0.0	3.0	3.5
						0.5	0.5	3.0	0.0	0.0	3.0	3.5
Positive Control	24	0.00004	200	0.5	-	0.5	11.0	55.5	0.0	0.0	59.5	59.5
	48	0.00004	200	0.0	-	1.0	9.0	69.5	0.0	0.0	70.5	70.5

※ There was no observation for metaphase with both treatment of 24Hr and 48Hr at 0.06mg/ml.

Judgement for

Chromosomal Aberration in CHL ; Positive

IARC Evaluation

; not yet cited

Experimental Data with Metabolic Activation

Substance	Treatment		No. of Metaphase	Polyploid (%)	Judge- ment	Cell with Structural Chromosome Aberration (%)						Judge- ment		
	S9 mix	Concen- tration (mg/ml)				Gap	Chromatid CTB	CTE	Chromosome CSB	CSE	Total -G			
DMSO	—		200	0.5	—	0.0	0.5	0.5	0.0	0.0	1.0	1.0	—	
	+		200	0.0	—	0.0	0.0	0.5	0.0	0.0	0.5	0.5	—	
Test Chemical	—	0.025	200	0.0	—	0.0	0.0	0.0	0.0	0.0	0.0	0.0	—	
	—	0.05	200	1.0	—	0.5	1.5	4.5	0.0	0.0	5.5	6.0	±	
	—	0.1				No observation for metaphase								
	—	0.2				No observation for metaphase								
	—	0.3				No observation for metaphase								
	+	0.025	200	0.5	—	0.0	0.0	0.0	0.0	0.0	0.0	0.0	—	
Positive Control [B(a)P]	+	0.05	200	0.5	—	0.0	0.0	1.0	0.0	0.0	1.0	1.0	—	
	+	0.1	200	1.0	—	0.0	2.0	1.5	0.0	0.0	3.5	3.5	—	
	+	0.2	200	1.0	—	0.0	0.5	12.5	0.0	0.0	12.5	12.5	+	
	+	0.3				No observation for metaphase								
	—	0.01	200	2.5	—	0.0	0.5	0.0	0.0	0.0	0.5	0.5	—	
Positive Control [B(a)P]	+	0.01	200	0.5	—	0.5	6.0	28.0	0.0	0.0	31.0	31.0	+	