

1,2-Dibromoethane (1,2-ジブロエタン)

Experimental Data

<u>Chemical Name:</u>	1,2-Dibromoethane	
<u>Synonym</u>	Ethylene dibromide	
	Ethane, 1,2-dibromo-	
<u>Molecular weight:</u>	187.9	
<u>Melting point:</u>	9~10°C	
<u>Boiling point:</u>	131~132°C	
<u>Chemical Structure</u>	$\text{CH}_2\text{Br}-\text{CH}_2\text{Br}$	
CAS No :	106-93-4	
MITI No :	(2)-59	
ML No :	2-(13)-38	
Source of Substance:	Tokyo Kasei Kogyo Co., Ltd.	
Lot. No.:	FA001	
Purity:	%	
Vehicle:	DMSO	

Judgement for
Chromosomal Aberration in CHL: Positive

	Treated Time (Hr)	Concen- tration (mg/ml)	No. of Metaphase	Poly- ploid (%)	Judge	Cell with Structural Chromosome Aberration (%)				Total		Judge		
						Gap	CTB	CTE	CSB	CSE	-G	+G		
DMSO	24		200	0	—	0	0.5	0	0	0	0.5	0.5	—	
	48		200	0	—	1.5	0.5	0.5	0	0	1.0	2.5	—	
Test Chemical														
	24	0.1	200	0	—	0	1.0	1.5	0	0	2.5	2.5	—	
		0.2	200	0.5	—	1.0	0.5	7.5	0	0	8.0	9.0	±	
		0.4	200	0	—	2.5	5.0	48.0	0	0	50.0	51.0	+	
		0.8				No observation for metaphase								
		1.6				No observation for metaphase								
	48	0.1	200	0.5	—	0	1.0	1.0	0	0	2.0	2.0	—	
		0.2	200	0	—	1.0	0	0.5	0	0	0.5	1.5	—	
		0.4	200	1.5	—	0	1.0	10.0	0	0.5	10.5	10.5	+	
		0.8	200	4.5	—	7.0	24.0	68.0	0	1.5	70.5	71.5	+	
		1.6				No observation for metaphase								
Positive Control														
	(MMC)	24	0.00008	200	0.5	—	6.0	21.0	61.5	0	0.5	72.0	73.0	+
		48	0.00008	200	1.5	—	9.0	25.0	79.5	0	1.5	84.0	84.5	+

IARC Evaluation : G 2A

Experimental Data

	S 9 with or without	Concen- tration (mg/ml)	No. of Meta- Phase	Poly- ploid (%)	Cell with Structural Chromosome Aberration (%)						Total		
					Judge	Gap	CTB	CTE	CSB	CSE	-G	+G	Judge
DMSO	—		200	0	—	0.5	0	0	0	0	0	0.5	—
	+		200	0	—	0.5	0	0.5	0	0	0.5	1.0	—
Test Chemical													
—	0.04	200	0.5	—	0.5	0.5	0.5	0.5	0	0	1.0	1.0	—
	0.08	200	0	—	0	0	0	0	0	0	0	0	—
	0.16	200	0	—	1.0	0	1.5	0	0	0	1.5	2.0	—
	0.24	200	0	—	1.0	0.5	1.5	0	0	0	2.0	3.0	—
	0.32	200	0	—	1.0	1.0	5.5	0	0	0	6.5	7.5	±
	0.04	200	0.5	—	0.5	0.5	1.0	0	0	0	1.5	2.0	—
	0.08	200	0.5	—	1.5	0.5	1.5	0	0	0	2.0	3.5	—
	0.16	200	1.0	—	4.5	2.5	10.0	0	0	0	11.5	15.0	+
	0.24	200	1.0	—	4.0	8.5	33.0	0	1.0	0	35.0	36.5	+
	0.32				No observation for metaphase								
Positive Control													
(B(a)P)	—	0.008	200	0	—	0.5	0	0.5	0	0	0.5	1.0	—
	+	0.008	200	0	—	3.0	4.0	16.5	0	0	19.0	20.5	+