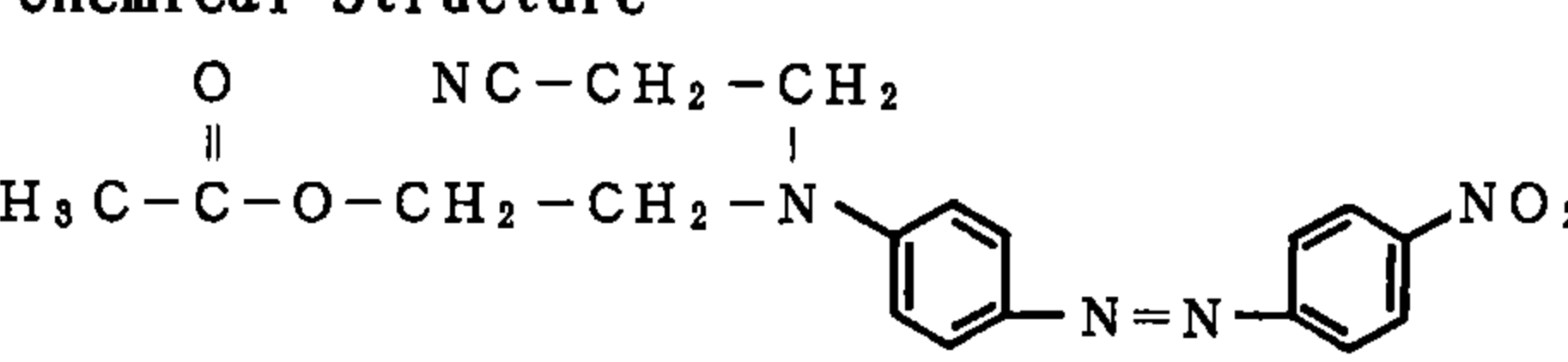


4-(4-Nitrophenylazo)-N-(2-cyanoethyl)-N-(2-acetoxyethyl)aniline

4-(4-ニトロフェニルアゾ)-N-(2-シアノエチル)-N-(2-アセトキシエチル)アニリン

Experimental Data

Chemical Name: 4-(4-Nitrophenylazo)-N-(2-cyanoethyl)-N-(2-acetoxyethyl)aniline	Treated time (Hr)	Concentration (mg/ml)	No. of Meta-phase	Polyploid (%)	Judge	Cell with Structural Chromosome Aberration (%)						Judge		
						Total		CTB	CTE	CSB	CSE			
						-G	+G							
Synonym						Gap	CTB	CTE	CSB	CSE	-G	+G		
Molecular weight: 381.0	Q													
Melting point: 115 °C	C M C		200	0	-	0	0	0.5	0	0	0.5	0.5	-	
Boiling point: °C			200	0.5	-	0.5	0	0.5	0	0	0.5	1.0	-	
Chemical Structure	Test Chemical													
	24	0.5	200	1.0	-	1.5	0.5	0	0	0	0.5	2.0	-	
			1.0	200	0	-	1.5	1.5	0.5	0	0	1.0	3.5	-
			2.0	200	0	-	2.0	2.5	1.0	0	0	3.5	5.5	-
			3.0	189	0	-	4.3	2.1	2.1	0	0	4.2	8.0	±
			4.0				No observation for metaphase							
CAS No : 6657-36-9	48	0.5	200	0	-	0.5	0.5	0.5	0	0	1.5	2.0	-	
Source of Substance: Tokyo Kasei Kogyo Co., Ltd.		1.0	200	0	-	1.5	0.5	0.5	0	0	0.5	2.0	-	
Lot. No. :		2.0				No observation for metaphase								
Purity:		3.0				No observation for metaphase								
Vehicle: 1% CMC	Positive Control (MMC)													
Judgement for Chromosomal Aberration in CHL: Positive	24	0.0001	200	0	-	3.5	24.0	24.0	0	0	29.0	31.0	+	
IARC Evaluation : not yet cited	48	0.0001	200	0.5	-	3.0	41.5	41.5	0	1.0	48.5	50.0	+	

Experimental Data

S 9 with or without	Concent- ration (mg/ml)	No. of Meta- phase	Poly- ploid (%)	Judge	Cell with Structural Chromosome Aberration (%)							Judge	
										Total			
					Gap	CTB	CTE	CSB	CSE	-G	+G		
C M C	-	200	0.5	-	0	0	0	0	0	0	0	-	
	+	200	0	-	0	0	0	0	0	0	0	-	
Test Chemical													
	-	2.0	200	2.0	-	0.5	0.5	0	0	0	0.5	1.0	-
		4.0	200	4.0	-	0	0.5	0	0	0	0.5	0.5	-
		8.0	200	0	-	4.0	3.5	9.0	0	0	11.5	14.0	+
		16.0	146	1.4		8.9	18.5	28.8			39.1	42.5	+
	+	2.0	200	1.0	-	1.0	0	0	0	0	0	1.0	-
		4.0	200	0	-	1.5	0.5	0	0	0	0.5	2.0	-
		8.0	200	0	-	0.5	0	1.0	0	0	1.0	1.5	-
		16.0	200	0.5	-	1.0	0.5	4.0	0	0	4.5	5.5	±
Positive Control													
(B(a)p)	-	0.016	200	0	-	0.5	0.5	0	0	0	1.0	1.5	-
	+	0.016	200	0.5	-	3.0	6.5	39.5	0	0	42.0	43.0	+