

[3'-アミノ-4'-メトキシアセトアニリド]

Experimental Data without Metabolic Activation

Chemical Name; 3'-Amino-4'-methoxyacetanilide

Synonym ; 5-Acetamino-2-methoxyaniline

5-Acetamido-2-anisidine

N-(3-Amino-4-methoxyphenyl)-

acetamide

5-アセトアミノ-2-メトキシアニリン

5-アセトアミド-2-アニシジン

N-(3-アミノ-4-メトキシフェニル)アセトアミド

Molecular Weight ; 180.21

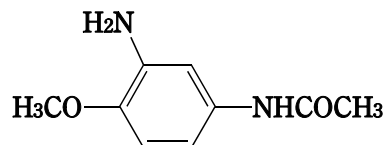
Melting Point ; 106 - 109 °C

Boiling Point ; — °C

Flashing Point ; — °C

Molecular Formula; C₉H₁₁N₂O₂

Chemical Structure



CAS No. ; 6375-47-9

MITI No. ; (3)-731 , (3)-2797

ML No. ; —

Specified Chemical Substances; —

Source of Substance; Tokyo Kasei Kogyo Co., Ltd.

Lot No. ; A001

Purity ; 96.0 %

Vehicle ; DMSO

| Substance | Treatment | | No. of Metaphase | Polyploid (%) | Judge-ment | Cell with Structural Chromosome Aberration (%) | | | | | | | |
|------------------|-----------|-----------------------|------------------|---------------|------------|--|------|------|-----|-----|-------|------|------------|
| | Time (h) | Concentration (mg/ml) | | | | Chromatid | | | | | Total | | Judge-ment |
| | | | | | | Gap | CTB | CTE | CSB | CSE | -G | +G | |
| DMSO | 24 | 1.0 % | 200 | 2.0 | — | 0.0 | 0.5 | 0.0 | 0.0 | 0.0 | 0.5 | 0.5 | — |
| | 48 | 1.0 % | 200 | 1.5 | — | 0.0 | 1.0 | 1.0 | 0.0 | 0.0 | 2.0 | 2.0 | — |
| Test Chemical | 24 | 0.25 | 200 | 1.0 | — | 0.0 | 1.5 | 0.0 | 0.0 | 0.0 | 1.5 | 1.5 | — |
| | | 0.5 | 200 | 1.5 | — | 0.5 | 1.0 | 1.0 | 0.0 | 0.0 | 2.0 | 2.0 | — |
| | | 1.0 | 200 | 0.0 | — | 0.5 | 1.0 | 0.5 | 0.0 | 0.0 | 1.5 | 2.0 | — |
| | | 2.0 | 200 | 0.5 | — | 1.0 | 2.5 | 4.5 | 0.0 | 0.0 | 6.5 | 7.5 | ± |
| | | 3.0 | 200 | 0.5 | — | 3.0 | 15.5 | 12.5 | 0.0 | 0.0 | 23.5 | 25.0 | + |
| | 48 | 0.25 | 200 | 0.0 | — | 0.5 | 1.0 | 1.0 | 0.0 | 0.0 | 2.0 | 2.0 | — |
| | | 0.5 | 200 | 0.0 | — | 0.0 | 0.5 | 0.5 | 0.0 | 0.0 | 1.0 | 1.0 | — |
| | | 1.0 | 200 | 2.0 | — | 1.0 | 2.0 | 4.0 | 0.0 | 0.0 | 5.0 | 6.0 | ± |
| | | 2.0 | 200 | 1.0 | — | 0.5 | 3.5 | 4.5 | 0.0 | 0.0 | 7.5 | 8.0 | ± |
| | | 3.0 | 200 | | | No observation for metaphase | | | | | | | |
| Positive Control | 24 | 0.00004 | 200 | 3.0 | — | 2.0 | 11.0 | 15.0 | 0.0 | 0.0 | 23.0 | 23.0 | + |
| [MMC] | 48 | 0.00004 | 200 | 2.5 | — | 1.5 | 15.0 | 30.0 | 0.0 | 0.0 | 40.0 | 40.5 | + |

※ There was no observation for metaphase with both treatment of 24Hr and 48Hr at 4.0 mg/ml.

Judgement for

Chromosomal Aberration in CHL ; **Positive**

IARC Evaluation ; not yet cited

Experimental Data with Metabolic Activation

| Treatment | | | No. of Metaphase | Polyploid (%) | Judge- ment | Cell with Structural Chromosome Aberration (%) | | | | | | | Judge- ment |
|--------------------------------|-----------|-------------------------------|---------------------|------------------|----------------|---|------|------------|-----|-------|------|------|----------------|
| Substance | S9 mix | Concent- ration (mg/ml) | | | | Chromatid | | Chromosome | | Total | | | |
| | | | | | | Gap | CTB | CTE | CSB | CSE | -G | +G | |
| DMSO | - | 1.0 % | 200 | 0.5 | - | 0.5 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.5 | - |
| | + | 1.0 % | 200 | 1.5 | - | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | - |
| Test Chemical | - | 0.019 | 200 | 1.0 | - | 1.0 | 0.0 | 0.5 | 0.0 | 0.0 | 0.5 | 1.5 | - |
| | | 0.078 | 200 | 2.5 | - | 0.0 | 0.5 | 0.5 | 0.0 | 0.0 | 1.0 | 1.0 | - |
| | | 0.31 | 200 | 1.0 | - | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | - |
| | | 1.3 | 200 | 4.0 | - | 0.0 | 1.0 | 0.5 | 0.0 | 0.0 | 1.5 | 1.5 | - |
| | | 5.0 | 200 | 3.5 | - | 0.0 | 0.5 | 2.0 | 0.0 | 0.0 | 2.0 | 2.0 | - |
| | + | 0.019 | 200 | 1.0 | - | 0.0 | 1.0 | 4.5 | 0.0 | 0.0 | 4.5 | 4.5 | - |
| | | 0.078 | 200 | 3.0 | - | 2.0 | 7.5 | 19.5 | 0.0 | 0.0 | 21.0 | 21.5 | + |
| | | 0.31 | 200 | 6.0 | ± | 2.5 | 9.5 | 22.0 | 0.0 | 0.0 | 24.5 | 24.5 | + |
| | | 1.3 | 200 | 2.0 | - | 1.5 | 12.5 | 20.5 | 0.0 | 0.0 | 23.5 | 23.5 | + |
| | | 5.0 | 200 | 2.5 | - | 2.5 | 19.0 | 30.0 | 0.0 | 0.0 | 34.5 | 34.5 | + |
| Positive Control [B(a)P] | - | 0.01 | 200 | 0.0 | - | 0.5 | 0.5 | 0.0 | 0.0 | 0.0 | 0.5 | 1.0 | - |
| | + | 0.01 | 200 | 2.0 | - | 1.5 | 6.0 | 34.0 | 0.0 | 0.0 | 36.5 | 36.5 | + |