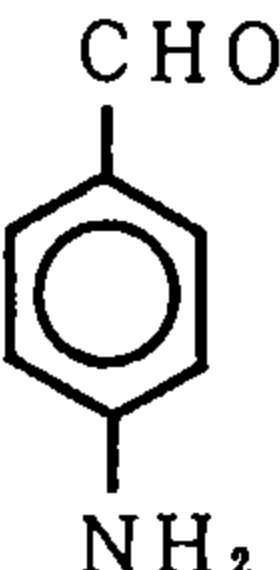


p-Aminobenzaldehyde (p-アミノベンズアルデヒド)

Experimental Data

Chemical Name: p-Aminobenzaldehyde Synonym: p-Formylaniline Formanilide Benzaldehyde, 4-amino-	Con. μg/ plate	Number of Revertants/plate									
		Base-substitution						Frame-shift			
		TA100		TA1535		WP2uvrA		TA98		TA1537	
		S9-	S9+	S9-	S9+	S9-	S9+	S9-	S9+	S9-	S9+
Molecular weight: 121.1 Boiling point: 271°C, 166°C (14mmHg) Melting point: 46.6~47.5°C Flashing point: > 110°C Chemical Structure	DMSO	(135)	(127)	(11)	(10)	(21)	(26)	(32)	(31)	(10)	(19)
		141	144	13	10	22	32	36	37	10	18
		111	138	15	11	25	31	31	37	9	13
	39	(126)	(141)	(14)	(11)	(24)	(32)	(34)	(37)	(10)	(16)
		143	136	10	8	17	32	41	39	11	15
		160	124	13	10	23	29	40	36	3	13
	78	(152)	(130)	(12)	(9)	(20)	(31)	(41)	(38)	(7)	(14)
		163	143	11	9	30	24	38	51	11	15
		157	134	11	13	22	24	40	40	13	18
	156	(160)	(139)	(11)	(11)	(26)	(24)	(39)	(46)	(12)	(17)
		152	141	7	11	26	28	41	38	7	18
		162	156	8	11	26	31	45	40	11	14
	313	(157)	(149)	(8)	(11)	(26)	(30)	(43)	(39)	(9)	(16)
		143	169	13	9	21	34	33	38	11	15
		150	170	10	6	29	31	27	37	8	10
	625	(147)	(170)	(12)	(8)	(25)	(33)	(30)	(38)	(10)	(13)
		123	171	11	9	15	28	33	32	10	11
		115	171	9	10	18	23	25	40	11	10
	1250	(119)	(171)	(10)	(10)	(17)	(26)	(29)	(36)	(11)	(11)
		121	136	5	9	23	33	30	45	9	12
		123	134	4	7	20	25	40	45	9	15
	2500	(122)	(135)	(5)	(8)	(22)	(29)	(35)	(45)	(9)	(14)
		115	147	5	8	24	23	31	50	10	18
		140	148	11	10	19	24	33	43	11	15
	5000	(128)	(148)	(8)	(9)	(22)	(24)	(32)	(47)	(11)	(17)
		96	107	11	5	11	26	33	46	10	17
		107	99	7	8	23	26	25	42	7	11
	10000	(102)	(103)	(9)	(7)	(17)	(26)	(29)	(44)	(9)	(14)
Judgement		-	-	-	-	-	-	-	-	-	-
Specific Mutagenicity											
Positive		AF2	2AA	NaN ₃	2AA	AF2	2AA 20	AF2	2AA	9AA	2AA
Control		(648)	(725)	(440)	(165)	(152)	(1452)	(410)	(363)	(543)	(144)

Mutagenicity
in Bacterial Test: Negative

IARC Evaluation: not yet cited

Experimental Data

Con. μg/ plate	Number of Revertants/plate									
	Base-substitution						Frame-shift			
	TA100		TA1535		WP2uvrA		TA98		TA1537	
	S9-	S9+	S9-	S9+	S9-	S9+	S9-	S9+	S9-	S9+
<u>DMSO</u>	(134)	(135)	(10)	(10)	(23)	(23)	(25)	(25)	(7)	(8)
	117	177	8	4	16	26	25	23	5	3
	124	179	7	7	14	28	33	30	4	3
<u>625</u>	(121)	(178)	(8)	(6)	(15)	(27)	(29)	(27)	(5)	(3)
	106	168	7	14	18	30	27	33	5	4
	136	163	8	4	15	29	24	26	8	3
<u>1250</u>	(121)	(166)	(8)	(9)	(17)	(30)	(26)	(30)	(7)	(4)
	121	141	3	7	21	25	30	36	8	2
	119	156	9	6	19	22	33	40	8	3
<u>2500</u>	(120)	(149)	(6)	(7)	(20)	(24)	(32)	(38)	(8)	(3)
	105	172	4	11	25	31	19	31	8	4
	107	162	5	14	18	30	18	27	6	4
<u>5000</u>	(106)	(167)	(5)	(13)	(22)	(31)	(19)	(29)	(7)	(4)
	105	132	8	7	24	20	25	25	7	4
	99	137	4	5	22	23	20	20	11	4
<u>10000</u>	(102)	(135)	(6)	(6)	(23)	(22)	(23)	(23)	(9)	(4)
Judgement	-	-	-	-	-	-	-	-	-	-
Specific Mutagenicity										
Positive	AF2	2AA	NaN ₃	2AA	AF2	2AA 20	AF2	2AA	9AA	2AA
Control	(696)	(763)	(308)	(145)	(200)	(1486)	(419)	(337)	(476)	(129)

Experimental Data 30% S9

Con. μ g/ plate	Number of Revertants/plate				
	Base-substitution			Frame-shift	
	TA100	TA1535	WP2uvrA	TA98	TA1537
	S9+	S9+	S9+	S9+	S9+
<u>DMSO</u>	(167)	(17)	(30)	(38)	(22)
	166	18	31	39	14
	176	18	22	24	17
<u>156</u>	(171)	(18)	(27)	(32)	(16)
	207	22	32	40	12
	181	17	23	32	15
<u>313</u>	(194)	(20)	(28)	(36)	(14)
	189	24	32	32	19
	178	18	39	36	23
<u>625</u>	(184)	(21)	(36)	(34)	(21)
	195	17	27	37	16
	174	21	29	46	6
<u>1250</u>	(185)	(19)	(28)	(42)	(11)
	185	23	39	41	17
	188	20	27	47	11
<u>2500</u>	(187)	(22)	(33)	(44)	(14)
	187	16	25	54	16
	209	21	33	34	16
<u>5000</u>	(198)	(19)	(29)	(44)	(16)
	159	12	29	57	24
	156	12	23	55	15
<u>10000</u>	(158)	(12)	(26)	(56)	(20)
Judgement	—	—	—	—	—
Specific Mutagenicity					
Positive	2AA 2.0	2AA 10	2AA 50	2AA 2.0	2AA 10
Control	(497)	(321)	(1114)	(442)	(296)