

Antibacterial Activity of *N*-(β -Styryl)formamides Related to Tuberin¹

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A series of para-substituted *N*-(β -styryl)formamides, analogues of tuberin (4a), has been prepared and assayed for antibacterial activity. The methylthio, ethoxy, and methyl analogues 4e, 4j, and 4t were about twice as active as tuberin against *Mycobacterium phlei*. Although tuberin lacks activity against *Staphylococcus aureus*, several of the analogues described were found to inhibit this organism. The phenyl group of tuberin is not a prerequisite for activity since analogues based on naphthyl or ferrocenyl groups were also active. A quantitative structure-activity relationship further implied that an aromatic group need not be present, suggesting the synthesis of the cyclohexyl and *n*-amyl analogues which were found to possess high activity.

The growth of various species of mycobacteria is inhibited by the weak antibiotic tuberin although other bacteria are unaffected.² We have synthesized a series of analogues, some of which show activity greater than that of tuberin against mycobacteria, while others are also active against *Staphylococcus aureus*.

Chemistry. The required *N*-(β -styryl)formamides 4a-t were prepared from the cinnamic acids 1a-t as shown in Scheme I.³ The same methods were also applicable to the preparation of the furyl, thienyl, 2-naphthyl, 1-naphthyl, ferrocenyl, cyclohexyl, and *n*-amyl analogues 5a-g.

Results and Discussion

A group of six analogues, 4a-f, covering a wide range of substituent π , π^2 , \mathfrak{F} , and \mathcal{R} , was selected from a list of candidates, using the determinant ($X'X$) criteria,⁴ for initial synthesis and screening against *Mycobacterium phlei*. The resulting minimum inhibitory concentrations (MIC), when plotted in the π - σ biological activity coordinate system as described by Darvas,⁵ showed an apparent correlation. Further compounds were then prepared leading to the data displayed in Figure 1 and Table I. Maximum activity about two times that of tuberin (4a, R = MeO) was obtained in a small region enclosing the EtO, Me, and MeS substituents. Values of the MIC against *M. phlei* and *S. aureus* are listed in Table I. Significant activity against *S. aureus* was observed for several of the compounds, in contrast to the inactivity of tuberin.

From the graphical structure-activity correlation it was clear that compounds with more than two to three times the activity of tuberin were unlikely to be obtained from para-substituted *N*-(β -styryl)formamides. This was confirmed by a quantitative structure-activity (QSAR) relation (see below). Compounds with major modifications of the aromatic system were therefore prepared. The most active of these, the 2-naphthyl analogue 5c, had about four times the activity of tuberin against *M. phlei*. The ferrocenyl analogue 5e had an MIC equal to that of tuberin (see Table II).

The observation that large changes in the aromatic system did not necessarily impair activity, and the results of a QSAR study, suggested that the aromatic portion of tuberin was not a requirement. This was confirmed by the preparation of the cyclohexyl and *n*-amyl analogues 5f,g which were found to be about one-half and two times as potent as tuberin.

QSAR Studies. Table III gives the relevant data for QSAR studies. Lipophilicities were determined by three procedures (the interrelations are more fully discussed in ref 6): (a) classical shake-flask, (b) LC on C18 Corasil using 25 wt % MeOH/pH 7.0 buffer, and (c) LC on C18 Corasil coated with 1-octanol using pH 7.0 buffer saturated with 1-octanol. Shake-flask and the new C18 Corasil/octanol procedures gave excellent agreement (correlation coefficient 0.99); however, C18 Corasil/25 wt % MeOH showed

systematic deviations due to hydrogen-bonding substituents. If $D_{\text{ha}} = 1$ for hydrogen-bonding acceptors and $D_{\text{ha}} = 0$ for non-hydrogen-bonding substituents, then we obtain eq 1 relating $\log P$ and $\log k'$. All terms are significant at

$$\log P = 1.7 (\pm 0.2) + 1.2 (\pm 0.2) \log k' - 0.3 (\pm 0.2) D_{\text{ha}} \quad (1)$$

$n = 19; r = 0.973; s = 0.161; F_{2,16} = 140$

the 95% confidence limits or better; n = number of points, s = standard deviation of points from the line, r = correlation coefficient, F is the overall significance parameter, and 95% confidence limits are given in parentheses. As shown in eq 2, the $\log P$ values are not those of substituted

$$\log P = 1.9 (\pm 0.2) + 0.98 (\pm 0.13) \pi_{\text{benzene}} + 0.3 (\pm 0.4) \mathfrak{F} \quad (2)$$

$n = 19; r = 0.977; s = 0.143; F = 166$

benzenes but deviate⁵ because of the electronegative formamide group. (The \mathfrak{F} term is significant at the 90% confidence limits.) Note that π_{benzene} was used in the initial planning, but this should have only a minor effect on the overall design, the main purpose of which was to obtain a spread in substituent types.

The resulting QSAR using $\log k'$ and $\log P$ are given in eq 3-5 where $pA = \log 1/\text{MIC}_M$. The unsubstituted

$$pA = -0.3 (\pm 1.4) + 2.9 (\pm 1.1) \log P - 0.5 (\pm 0.2) (\log P)^2 - 2.6 (\pm 0.9) \mathfrak{F}^2 - 1.7 (\pm 0.5) \mathcal{R}^2 + 0.4 (\pm 0.3) D_{\text{ha}} \quad (3a)$$

$n = 19; r = 0.952; s = 0.161; F_{5,13} = 25$

$$pA = -1.1 (\pm 2.4) + 3.1 (\pm 1.9) \log P - 0.5 (\pm 0.4) (\log P)^2 - 1.8 (\pm 1.5) \mathfrak{F}^2 - 1.6 (\pm 0.9) \mathcal{R}^2 + 0.5 (\pm 0.5) D_{\text{ha}} \quad (3b)$$

$n = 20; r = 0.856; s = 0.283; F_{5,14} = 8$

$$pA = 3.1 (\pm 0.5) + 1.2 (\pm 1.0) \log k' - 0.6 (\pm 0.5) (\log k')^2 - 2.1 (\pm 1.4) \mathfrak{F}^2 - 1.1 (\pm 0.7) \mathcal{R}^2 \quad (4)$$

$n = 18; r = 0.867; s = 0.257; F_{4,13} = 10$

$$pA = 3.4 (\pm 0.3) + 0.8 (\pm 0.3) \pi_{\text{benzene}} - 0.4 (\pm 0.2) \pi_{\text{benzene}}^2 - 2.2 (\pm 1.0) \mathfrak{F}^2 - 1.6 (\pm 0.6) \mathcal{R}^2 + 0.4 (\pm 0.3) D_{\text{ha}} \quad (5)$$

$n = 19; r = 0.943; s = 0.175; F_{5,13} = 21$

compound 4c was omitted because of very large deviations of the observed activity from calculated values in all equations examined. (See, for example, eq 3b where 4c deviates by 2.6 standard deviations and the quality of the correlation is considerably reduced by including this point.) Since 4c is the only para-unsubstituted compound,

Table I. Physical Data and Antibacterial Activity of Tuberin Analogues

No.	R	Mp or bp (mm), °C	Recrystn solvent	Formula ^a	MIC, µg/mL	
					<i>M. phlei</i>	<i>S. aureus</i>
4a	CH ₃ O	135-136	Me ₂ CO-hexane	(Tuberin)	80	<i>b</i>
4b	NO ₂	128-130	EtOAc-hexane	C ₉ H ₈ N ₂ O ₃	320	3
4c	H	100-101	C ₆ H ₆	C ₉ H ₉ NO	640	50
4d	<i>t</i> -Bu	98-99	C ₆ H ₆	C ₁₃ H ₁₇ NO	80	200
4e	MeS	130-131	EtOAc	C ₁₀ H ₁₁ NOS	40	100
4f	Me ₂ N	158-160	EtOAc	C ₁₁ H ₁₄ N ₂ O	640	<i>b</i>
4g	Et ₂ N	150 (0.05)	EtOAc	C ₁₃ H ₁₈ N ₂ O	320	<i>b</i>
4h	HO	197-198	Me ₂ CO	C ₉ H ₉ NO ₂	<i>c</i>	<i>b</i>
4i	AcO	112	EtOAc	C ₁₁ H ₁₁ NO ₃	640	<i>b</i>
4j	EtO	114-115	EtOAc	C ₁₁ H ₁₃ NO ₂	40	<i>b</i>
4k	<i>i</i> -PrO	92-93	EtOAc-hexane	C ₁₂ H ₁₅ NO ₂	320	<i>b</i>
4l	<i>n</i> -BuO	110-111	EtOAc-hexane	C ₁₃ H ₁₇ NO ₂	80	<i>b</i>
4m	F	75-76	EtOAc	C ₉ H ₈ FNO	160	<i>b</i>
4n	Cl	90-92	EtOAc	C ₉ H ₈ ClNO	80	200
4o	Br	95-96	EtOAc-hexane	C ₉ H ₈ BrNO	160	100
4p	I	128-129	EtOAc-hexane	C ₉ H ₈ I ₂ NO	160	200
4q	F ₃ C	103-105	Me ₂ CO-hexane	C ₁₀ H ₈ F ₃ NO	80	100
4r	CN	127, 141 ^d	Me ₂ CO-hexane	C ₁₀ H ₈ N ₂ O	640	<i>b</i>
4s	PhSO ₂	138-140	EtOAc	C ₁₅ H ₁₃ NO ₃ S	320	<i>b</i>
4t	Me	124-125	C ₆ H ₆	C ₁₀ H ₁₁ NO	40	<i>b</i>

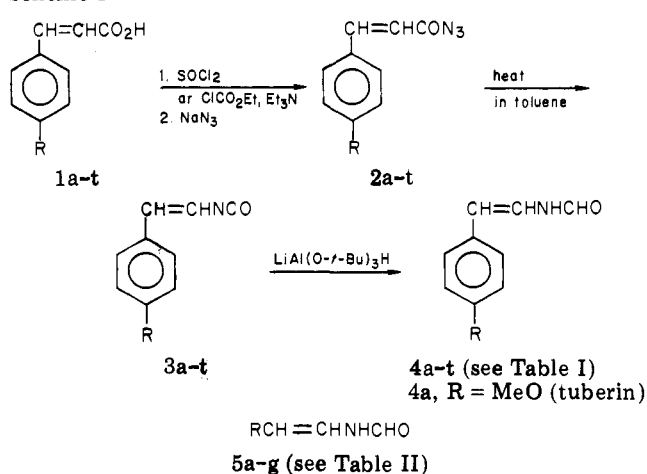
^a Analyzed for C, H, and N. ^b Inactive at 200 µg/mL. ^c Inactive at 640 µg/mL. ^d Double melting point.

Table II. Physical Data and Antibacterial Activity of Tuberin Analogues

No.	R	Mp, °C	Recrystn solvent	Formula ^a	MIC, µg/mL	
					<i>M. phlei</i>	<i>S. aureus</i>
5a	2-Furyl	109-110	EtOAc-hexane	C ₇ H ₇ NO ₂	<i>c</i>	<i>b</i>
5b	2-Thienyl	85-86	C ₆ H ₆	C ₇ H ₇ NOS	320	25
5c	2-Naphthyl	157	Me ₂ CO-hexane	C ₁₃ H ₁₁ NO	20	<i>b</i>
5d	1-Naphthyl	137-138	EtOAc-hexane	C ₁₃ H ₁₁ NO	80	<i>b</i>
5e	Ferrocenyl	128-130	CHCl ₃ -hexane	C ₁₃ H ₁₃ FeNO	80	200
5f	Cyclohexyl	<i>d</i>		C ₉ H ₁₅ NO	160	<i>b</i>
5g	<i>n</i> -Amyl	<i>d</i>		C ₈ H ₁₅ NO	40	<i>b</i>

^{a-c} See corresponding footnotes in Table I. ^d Gum.

Scheme I



metabolic deactivation is a possible cause. Equations 3-5 were obtained by searching all regressions of $\log k'$ (or $\log P$), $(\log k')^2$ [or $(\log P)^2$], \mathcal{F} , \mathcal{F}^2 , \mathcal{R} , \mathcal{R}^2 , MR, D_{ha} , and the cross product $(\log k')D_{ha}$ [or $(\log P)D_{ha}$]. Note that there are only 18 $\log k'$ values (Table III), since 4c is omitted and 4g could not be measured due to lack of compound. All terms are significant at the 95% confidence limits by *t* tests, and no terms with statistical significance can be added to these equations. $\log P$ and $(\log P)^2$ terms [and $\log k'$ and $(\log k')^2$] are highly correlated because of the

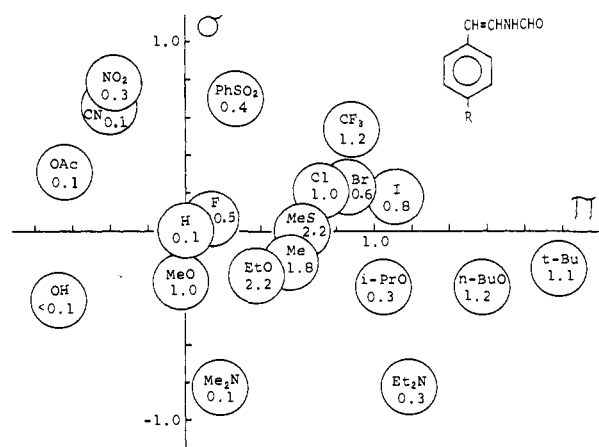


Figure 1. Antibacterial activities of tuberin analogues in the π - σ coordinate system. Substituent R and activity (molar basis) against *Mycobacterium phlei*, relative to tuberin, are shown.

small range; using RIDGE regression,⁹ which can deconvolute this interrelation, we find the same optimal $\log P_0$ and $\log k_0$, however.

The following conclusions can be drawn. (1) The optimum electronic effect occurs with electronically neutral substituents since there is no linear term in \mathcal{F} or \mathcal{R} (the same is true if σ_p is used in place of \mathcal{F} and \mathcal{R}). This observation suggested the preparation of cyclohexyl and *n*-amyl analogues. (2) $\log P_0 = 2.87$ (2.7-3.13, 95%

Table III. Data for QSAR Study on 4-X-PhCH=CHNHCHO Tuberin Analogues

X	pA_{obsd}^a	pA_{calcd}^b	$\text{Log } k'^c$	$\text{Log } P^d$	π_{benzene}^e	σ_p^e	\mathfrak{F}^e	\bar{r}^e	D_{ha}^f
3,4-(CH) ₄	3.99	3.80	1.16 ^g	3.22	1.32	0.04	0.03	0.01	0
OEt	3.68	3.61	0.65	2.31	0.38	-0.24	0.22	-0.44	1
SMe	3.68	3.65	0.77	2.55	0.61	0.00	0.20	-0.18	0
Me	3.60	3.74	0.66	2.45	0.56	-0.17	-0.04	-0.13	0
O- <i>n</i> -Bu	3.44	3.45	1.62 ^g	3.32 ^h	1.33 ^{i,j}	-0.32	0.25	-0.55	1
CF ₃	3.43	3.41	1.09	3.06	0.88	0.54	0.38	0.19	0
<i>t</i> -Bu	3.40	3.36	1.71 ^g	3.82 ^h	1.98	-0.20	-0.07	-0.13	0
Cl	3.36	3.37	0.79	2.70	0.71	0.23	0.41	-0.15	0
OMe	3.34	3.16	0.28	1.94 ^k	-0.02	-0.27	0.26	-0.51	1
I	3.23	3.33	1.14	3.18	1.12	0.18	0.40	-0.19	0
Br	3.15	3.31	0.96	2.87	0.86	0.23	0.44	-0.17	0
F	3.01	2.89	0.33	2.11	0.14	0.06	0.43	-0.34	0
SO ₂ Ph	2.95	2.92	0.78	1.95	0.27	0.70	0.56	0.18	1
NEt ₂	2.83	2.81	^j	2.90	1.18	-0.90	0.01	-0.91	1
O- <i>i</i> -Pr	2.81	3.07	0.93 ^g	2.61	0.36	-0.45	0.30	-0.72	1
NO ₂	2.78	2.57	0.24	1.94	-0.28	0.78	0.67	0.16	1
OAc	2.51	2.55	0.16	1.32 ^k	-0.64	0.31	0.41	-0.07	1
NMe ₂	2.47	2.39	0.48	2.03	0.18	-0.83	0.10	-0.92	1
CN	2.43	2.69	0.04	1.62 ^k	-0.57	0.66	0.51	0.19	1
H	2.36	3.42 ^j	0.21	1.95 ^k	0.00	0.00	0.00	0.00	0

^a $\text{Log } 1/\text{MIC}_{\text{molar}}$ vs. *M. phlei*. ^b From eq 3a. ^c C18 Corasil/25 wt % MeOH/pH 7.00 phosphate buffer (0.001 M) or as noted. $k' = (t_x - t_0)/t_0$ where t_x is the retention time of compound and t_0 is for the unretained compound. ^d C18 Corasil/octanol/pH 7.00 phosphate buffer (0.01 M); see ref 6. ^e Pomona College Medicinal Chemistry Project or see ref 7. ^f $D_{\text{ha}} = 1$ if H-bond acceptor. ^g Determined in 40 wt % MeOH and extrapolated to 25 wt % by $\text{log } k'_{25} = 0.825 + 1.169 \text{ log } k'_{40}$ ($n = 4$; $r = 0.990$) which was established for $D_{\text{ha}} = 0$ substituents. OEt was found to be +0.05 higher than calculated and this increment was added to O-*n*-Bu and O-*i*-Pr (i.e., assume parallel lines with displacement due to H bonding). ^h For $D_{\text{ha}} = 0$, $\text{log } P = 1.68 + 1.25 \text{ log } k'$ ($n = 9$; $r = 0.994$), which is used for *t*-Bu, while for OR substituents $\text{log } P = 1.65 + 1.03 \text{ log } k'$ ($n = 3$; $r = 1.0$), which is used for O-*n*-Bu. ⁱ From $\pi = -1.79 + 0.968 \text{ log } P - 0.382 \mathfrak{F}$ ($n = 19$; $r = 0.979$; $s = 0.143$; reverse of eq 2). ^j H omitted from eq 3-5; NEt₂ omitted from eq 1 and 4 (not determined); O-*n*-Bu omitted from eq 2, unreliable literature value. ^k $\text{Log } P$ by shake-flask: X = H, 1.93 (± 0.01); X = OMe, 1.92 (± 0.02); X = CN, 1.58 (± 0.01); X = OAc, 1.40 (± 0.01). The estimated standard deviation by interpolation using standards is 0.03.

confidence limits). Note that $\text{log } P$ for the cyclohexyl and *n*-amyl analogues 5f and 5g is 2.69 and 2.65, respectively. (3) The origin of the D_{ha} term is interesting. If it were present in eq 3 because of an activity enhancing effect, the term would be expected to appear in eq 4, but it cannot be added with statistical significance. Substitution of eq 1 in eq 3 should give rise to a cross-product term ($\text{log } k' D_{\text{ha}}$); however, this term also cannot be added to eq 4. This suggests that alkane ($\text{log } k'$) may be a better model for the waxy¹⁰ mycobacteria than octanol ($\text{log } P$). The D_{ha} term in eq 3 and 5 "corrects" octanol back toward a purer alkane phase. The high activity of the cyclohexyl and *n*-amyl analogues tends to support the view that D_{ha} represents a correction of the lipophilicity scale rather than a special activating effect. (4) Finally, eq 2 and 3a can be used to calculate a MIC of 1133 $\mu\text{g}/\text{mL}$ for the inactive 4-OH compound 4h which was excluded from the QSAR. This further confirms the above QSAR.

A QSAR study of substituted phenyl 4-aminosalicylates vs. *M. tuberculosis* and *M. bovis* has appeared;¹¹ however, while electronic effects were found to be important, the mode of action of these compounds probably differs from that of the tuberins. The same may also be true of a study of 3-benzoylacrylic acids vs. *M. tuberculosis* H37Rv where an electronic term improved the correlation.¹² The Hammett ρ value for this term was negative, in contrast to a positive ρ value found in the case of 2,6-dialkoxy-phenylpenicillins,¹³ which, it was argued,¹² showed potential structural similarities in part. Optimal electronic effects have been found, for example, in a study of acaricidal hydrazones.¹⁴

The optimal $\text{log } P_0 = 2.87$ for tuberins vs. *M. phlei* is far from the general $\text{log } P_0$ found for relatively nonspecific agents (ca. 4 for gram-negative and 6 for gram-positive bacteria¹⁵); however, it is close to the $\text{log } P$ found for some of the more lipophilic penicillins, such as dicloxacillin ($\text{log } P = 2.83$ and 2.45).

Experimental Section

Melting points were determined with a Mel-Temp apparatus and are corrected. Standard agar dilution methods (in duplicate) were used for the determination of MIC values vs. *M. phlei* (estimated standard deviation less than 0.15 log units) and microdilution methods (in triplicate) for *S. aureus* (estimated standard deviation less than 0.3 log units). QSAR regressions were performed on the Proprietary Computer Systems, Inc., APL system. Literature $\text{log } P$ values were from Hansch et al.⁷ or the Pomona College Medicinal Chemistry Project.

Synthesis of *N*-(β -Styryl)formamides. Compounds 4a-e and 4i-t were synthesized via the acid chloride according to Scheme I, following experimental procedures previously described.³ Compounds 4f and 4g were prepared via the mixed anhydride according to Scheme I.

***N*-(p -Hydroxy- β -styryl)formamide (4h).** A mixture of *N*-(p -acetoxy- β -styryl)formamide (4i) (200 mg, 1.03 mmol) in 3 mL of Me₂SO was added to a stirred suspension of Fleischmann's yeast (2 g) in 55 mL of 10% phosphate buffer at 42 °C. After 18 h the mixture was extracted with ether and the ethereal solution dried (Na₂SO₄). Evaporation of the solvent and crystallization of the residue from acetone gave 4h (149 mg, 95%).

Partition Coefficients. Procedures are described in ref 6. For the shake-flask procedure, both phases were analyzed and quadruplicates were run. $\text{Log } k'$ values were determined with reference to 25 wt % MeOH by linear regression from 40 wt % MeOH for the more lipophilic compounds ($r = 0.990$) with non-hydrogen-bonding substituents. Hydrogen-bonding substituents showed different sensitivities to solvent change and were placed on the 25 wt % scale, if they could not be measured directly, by appropriate corrections using overlapping series.

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References and Notes

- (1) Publication No. 499 from the Syntex Institute of Organic Chemistry.
- (2) K. Ohkuma, K. Anzai, and S. Suzuki, *J. Antibiot., Ser. A*, 15, 115 (1962).

