

INTRAMOLECULAR, TWO-SUBSTITUENT, CONSECUTIVE AND SEQUENTIAL (CONSEQ)
MIGRATIONS: X-RAY STRUCTURE OF AN INDENO[1,2,3-DE]QUINOLINE PRODUCT

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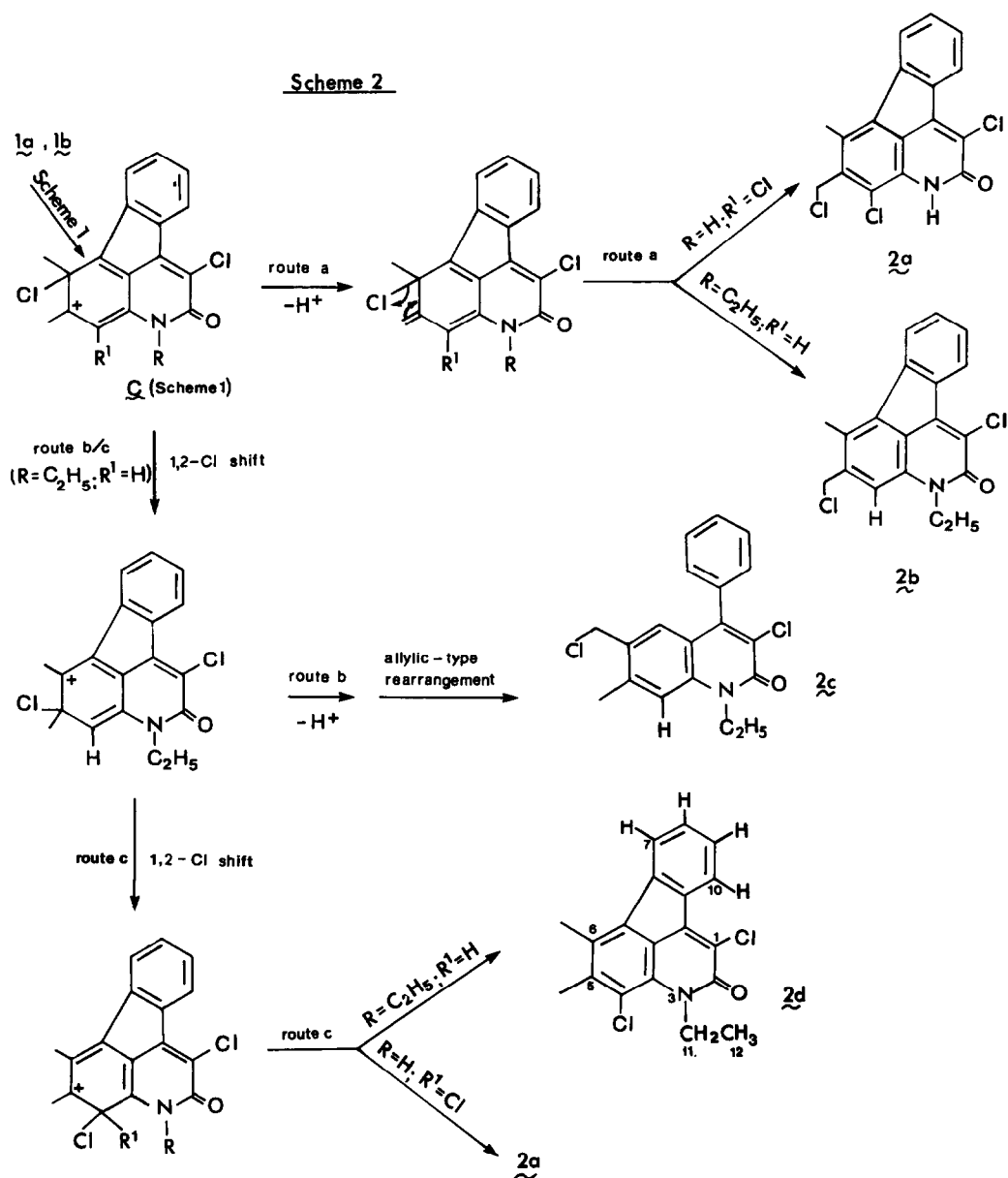
Abstract - Evidence, including the X-ray structure verification of a crucial reaction product, is presented in support of the postulation that intramolecular, two-substituent, consecutive and sequential migrations accompany the formation of certain indeno[1,2,3-de]quinolinones.

Whereas intramolecular substituent migrations accompanying chemical reactions are extensively documented, there is little concerning intramolecular, two-substituent 1,2-shifts that occur consecutively and in sequence, and here termed conseq migrations. We present supporting evidence for the latter concept which was earlier invoked¹⁻³ in an attempt to rationalise the acid-catalysed conversion of 3-keto amide (1a) to indeno[1,2,3-de]quinolin-2-one (2a). This transformation (Scheme 1) envisages the initial cyclisation of (1a) with production of an ion, (A; R = H), endowed with the propensity to undergo ring junction. There then forms an intermediate ion, (B), (the migration origin) from which issues a sequence of 1,2-substituent (conseq) migrations eventually leading to (D), (the migration terminus). Product (2a) then arises from (D) via an allylic-type rearrangement⁴ as indicated.

Such a succession of migratory events (B → D) appears to have no literature analogy and is obviously based on the validity of the structure assigned to (2a). Because of uncertainty¹ attending this latter aspect an X-ray structure verification was deemed essential. In the absence of a suitable crystal of (2a), an acceptable one of the 2-chloro derivative (3) (formed from (2a) and thionyl chloride)⁵ was grown (in CHCl₃ overlaid with hexane). The crystal structure of (3) (Fig. 1) which, incidently, is the first of the indeno[1,2,3-de]quinoline system, completely vindicates the formulation of (3), and by extension, also that of the crucial product (2a).

We next considered the feasibility of (2a) arising from (1a) via a shorter and more direct pathway (route a, Scheme 2) and which would obviate the need for the postulated consecutive and sequential two-substituent 1,2-shifts. To examine this prospect, the 3-keto amide (1b)¹ was cyclised in conc H₂SO₄ with due consideration of the reaction options available to an intervening ion (C) (R = C₂H₅; R¹ = H) (Scheme 2): One possibility (route a) envisages a proton loss followed by an allylic rearrangement and production of (2b); another option (route b) invokes a preferential⁶ 1,2-C₆ migration followed by a proton loss and an allylic rearrangement to give (2c). In essence, progression of (C) along either route a and/or route b would be expected to provide the chloromethyl derivative(s) (2b) [and/or (2c)]. A third alternative (route c) makes use of two consecutive 1,2-C₆ shifts, and should yield (ultimately), in contrast, a product (2d) void of a chloromethyl group. In the event, the ¹H-NMR spectrum (CDCl₃) of the crude, total product [TLC (CHCl₃-benzene) revealed 3 yellow constituents] exhibited no significant signal(s) attributable to

Scheme 2



accessible 3,4-dihydro-3,3,4-trichloroquinolin-2-one **4** in conc H_2SO_4 : it is contended that **4** ionises directly affording precursor ion **A** ($R = CH_3$ or C_2H_5); **A** then traverses the conseq pathway outlined in the scheme to yield the corresponding 5-chloromethylindeno[1,2,3-de]quinolin-2-one **2**. The structure of the substrate (e.g., **4a**), originally³ assigned on the basis of spectroscopic data, has now been unequivocally verified from an X-ray determination (Fig. 2). This result allows for the specification of each substituent's disposition in the species **A** ($R = C_2H_5$) immediately prior to the onset of the aforementioned cyclisation-rearrangement events.

EXPERIMENTAL

The 1H -NMR (normal) spectrum of **2d** and that of the crude reaction product from **1b** were recorded at 80MHz on a Bruker WP-80 instrument using TMS as internal standard. The NOESY experiment ($CDCl_3$, 30°C; mixing time = 0.5 sec) was done at 500 MHz using a Bruker WM-500 spectrometer. 1,4-Dichloro-5,6-dimethyl-3-ethylindeno[1,2,3-de]quinolin-2-one **2d**. A mixture of substrate **1b** (115mg; purified by preparative TLC) and conc H_2SO_4 (0.3ml) was reacted at 95°C for 3 min and diluted with water (~10ml). The acid-insoluble product was collected by filtration, washed with water, and air-dried (100mg; TLC (benzene- $CHCl_3$ 3:1) showed a mixture with **2d** as a

major component; the δ 4.85–5.5 region in the $^1\text{H-NMR}(\text{CDCl}_3)$ was free of significant absorptions). A pure sample of **2d** was obtained by chromatography: $^1\text{H-NMR}(\text{CDCl}_3)$ δ 1.49 (3H, t, $J=7\text{ Hz}$, CH_2CH_3), 2.52 (3H, s, 5- CH_3), 2.61 (3H, s, 6- CH_3), 4.76 (2H, q, $J=7\text{ Hz}$, CH_2CH_3), 7.3–7.6 (2H, m, 8-H, 9-H), 7.85 (1H, dd, $J=7, 2\text{ Hz}$, 7-H), 8.3 (1H, dd, $J=7, 2\text{ Hz}$, 10-H). The NOESY spectrum of **2d** was obtained as a 2-D contour plot. Off-diagonal peak correlations were established with assurance between 7-H and 6- CH_3 ; 8-H and 9-H; 11- CH_2 and 12- CH_3 . No significant interaction of the 11- CH_2 protons with either of the aromatic methyl groups was revealed. The expected peak correlations between 6- CH_3 and 5- CH_3 ; 7-H and 8-H; 9-H and 10-H, could not be established with a high level of confidence.

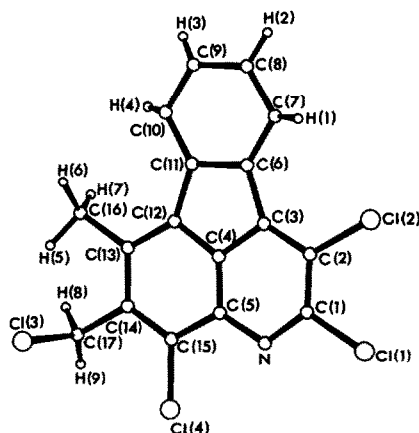


Figure 1 The structure of **3** showing the crystallographic numbering scheme

