Six Lectures on the Nature of the Hydrogen Bond Lecture 3 CAHB (Charge-Assisted Hydrogen Bond)

and PA/pKa Equalization Principle

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Six Lectures on the Nature of the Hydrogen Bond Gastone Gilli

The Six HB Chemical Leitmotifs (CLs)

CHARGE - ASSISTED HBs

CL # 1: (+/–)CAHB ⇒ SHB, VSHB Double Charge-Assisted HB Direct Acid-Base PA/pK_a Matching

CL # 2: (-)CAHB \Rightarrow SHB, VSHB Negative Charge-Assisted HB Acid-Base PA/pK_a Matching by Proton Loss

CL # 3: (+)CAHB \Rightarrow SHB, VSHB Positive Charge-Assisted HB Acid-Base PA/pK_a Matching by Proton Gain

Σ/Π -BOND POLARIZATION - ASSISTED HBs

CL # 4: RAHB \Rightarrow SHB, VSHB Resonance-Assisted or π -Cooperative HB PA/pK_a Matching by π -Conjugated-Bond Polarization

CL # 5: PAHB \Rightarrow **MHB Polarization-Assisted or \sigma-Cooperative HB** (*Partial*) *PA/pK_a Matching by* σ -*Bond Polarization*

NEITHER CHARGE- NOR Σ/Π-BOND POL.-ASSISTED HBs



ENOLS

CL # 6: OHB \Rightarrow W Ordinary HB No PA/pK_a Matching

⊖ D-----H∖ ⊕ A





 $\Delta pKa = -0.70$

p-TOLUIDINE

J---H-2.50_{6, Å} \

PENTACHLOROPHENOL -

CH₃

CI 1/2⊖

Cľ









Histograms of the $d(O \cdots O)$ distances derived from the CSD analysis for the **three CAHB classes** considered: (*a*) homomolecular (-)CAHBs; (*b*) homomolecular (+) CAHBs; and (*c*) from (±)CAHBs to OHBs.

CSD Classification of the O-H...O System: (-)CAHB





Very few cases of intramolecular bonds, such as the H-maleate anion, one of the most studied molecules by both diffraction and QM methods. The major part are intermolecular and occur in both organic and inorganic <u>acid salts</u>.

There are indications that (–)CAHBs become increasingly short and proton centred with the increasing acidity of the molecules involved (*cfr.* trifluoroacetates with carboxylates).

CSD Classification of the O-H...O System: (+)CAHB





(+)CAHBs can be considered H-bonds formed by two R=O bases that have captured a proton from an external strong acid.

They are essentially intermolecular and, in the majority of cases, homomolecular. This is due to the fact that, when we mix two different bases A and B in solution, the two homomolecular forms AA and BB will be inevitably more stable and easier to crystallize than the mixed form AB, for evident reasons of better pK_a matching.

CSD Classification of the O-H...O System: (+/-)CAHB



This last class of strong CAHBs can be related to the acid-base equilibrium

$$\mathbf{R}_{1} - \mathbf{O} - \mathbf{H} \cdots : \mathbf{O} - \mathbf{R}_{2} \cong \mathbf{R}_{1} - \mathbf{O} \cdots \mathbf{H}^{+} \cdots \mathbf{O}^{1/2} - \mathbf{R}_{2} \cong \mathbf{R}_{1} - \mathbf{O} \cdots \mathbf{H} - \mathbf{O}^{+} - \mathbf{R}_{2}$$

d(O…O) (Å)

between an acid, R_1 -O-H, and a base, :O- R_2 , whose conjugated acid is H-O⁺- R_2 .

This equilibrium is known to give <u>three extreme outcomes</u> (plus an infinite number of intermediate ones) modulated the value of $\Delta p K_a = p K_a (R_1 - O - H) - p K_a (H - O^+ - R_2)$.

1. When $\Delta p K_a$ is large and positive weak and neutral OHBs are formed (O-H···O).

2. When ΔpK_a is large and negative always weak but charged OHBs are obtained ($\neg O \cdots H - O^+$).

CSD Classification of the O-H...O System: PAHB



O-H…O o-Bond Cooperative or Polarization-Assisted H-bond (PAHB)

The concept of σ -bond cooperativity was firstly suggested by Jeffrey and Saenger (1991) in relation to the hypothesis that H-bonds within long chains of waters (or alcohols or phenols) should be stronger than ordinary O–H…O bonds because the increased O–H bond polarization induced by the chain has the correct sign for further strengthening the bond.

Later on, the bonds were added to the list of the Chemical Leitmotifs with the name Polarization-Assisted H-bond or PAHB (Gilli *et al.*, 2000).

The few typical cases above clearly show that the bond shortening is small, certainly because of the too small polarizability of the σ -bond, but sufficient to keep the O…O distances slightly below the shortest distances normally associated with the ordinary O–H…O bond (some 2.70 - 2.75 Å).

CSD Classification of the O-H...O System CAHB & RAHB: Symmetrization of Physical Properties



CSD Classification of the O-H...O System CAHB & RAHB: <u>Automatic</u> Equalization of VB Resonance Forms



The three ways for making iso-energetic the two VB resonance forms COV1,NCT and COV2,CT:
(*A*) by adding an electron;
(*B*) by removing an electron;
(*C*) by connecting the two oxygens by a π-conjugated chain.
(Gilli and Gilli, 2000)

CSD Classification of the O-H...O System: CAHB & RAHB: <u>Automatic</u> PA/pKa Equalization

(-)CAHB $\Delta pK_a = 0.0$ н \ 😑 2.IIa OHB Θ VERY $pK_{AH}(HO-H) = 15.7 - H^{+}$)---H---O Chemical Leitmotif # 2: **STRONG** $\mathbf{pK}_{BH}(H_2O-H^+) = -1.7$ (-)CAHB **2.II** H ~ 25-30 kcal/mol **2.IIb** $\Delta p K_a = 17.5$ [**R-D....H....A-R**]⁻ $\Delta pK_a = pK_{AH}(HO-H) - pK_{AH}(HO-H) = 15.7 - 15.7 = 0$ н $\Delta \mathbf{pK}_{a} = \mathbf{pK}_{BH}(H_{2}O-H^{+})-\mathbf{pK}_{BH}(H_{2}O-H^{+}) = -1.7 + 1.7 = 0$ Chemical Leitmotif #3: 2.VIa н н (+)CAHB H ⊕ 2.IIIa $\Delta p K_a = 0.0$ (+)CAHB WEAK ~ 4- 5 **[R-D....H....A-R]**+ VERY kcal/mol $+ \mathbf{H}^+$ **STRONG** ~ 25-31 Ĥ **2.III** н kcal/mol Ĥ. 2.IIIb н OHB **pK_{AH}**(RO-H) = 15/18 $\mathbf{pK}_{BH}(R_2C=O-H^+) = -(6/7)$ **Chemical Leitmotif #4:** EK // 2.IVa R_n-RAHB $\Delta p K_a = 0.0$ $\Delta pK_a = \sim 21-25$ **RAHB** R **R-D-H...A=R STRONG** 0-H---~ 15-22 \Leftrightarrow 2.VIb kcal/mol R=D...H-A-R **2.IV** ĭ\ KE 2.IVb **WEAK** ~ 4- 5 kcal/mol

CSD Classification of <u>Other Homonuclear</u> Systems: (-) and (+)CAHB

CHART 3.8. HOMONUCLEAR X-H...X (X = F, Cl, Br, S, N) CHARGE-ASSISTED HBs. CAHBs





The considerations made for O–H…O can be extended to other homonuclear X–H…X bonds, provided X is sufficiently electronegative and the bond formed belongs to a proper Chemical Leitmotif.

Typical (–)**CAHBs** include [Hal···H···Hal][–] (Hal = F, Cl, Br), a single case of [S···H···S][–], and a few $[N \cdots H \cdots N]^{-}$ bonds, while $[F \cdots H \cdots F]^{+}$ is the only (+)**CAHB** known to involve halogens.

Conversely, $[N \cdots H \cdots N]^+$ (+)CAHBs are quite frequent because they constitute the class of **proton sponges**, compounds with a variety of chemical applications that, for this reason, have been extensively studied from a structural point of view (Adler, 1990; Staab and Saupe, 1988; Llamas-Saiz *et al.*, 1994). **F**inally, about thirty N–H…N (±)CAHBs short enough to suggest a strict PA/p K_a matching have been retrieved, which would deserve a more detailed thermodynamic study. Only two examples are shown.

CSD Classification of <u>Heteronuclear</u> Systems: (±)CAHB



(±)CAHBs, whose strength simply depends on the more or less accidental donor/acceptor PA/pK_a matching, are not expected to be seriously affected by the heteronuclearity of the bond.

As a matter of fact, a systematic CSD study (Gilli and Gilli, 2000) carried out on **318** strong N–H···O/O–H···N bonds with $d(N \cdot \cdot \cdot O) \le$ 2.66 Å has singled out

265 (±)CAHBs (192 N-H⁺···O⁻ and 73 O-H···N),
29 (+)CAHBs (29 N-H⁺···O), and only
7 (-)CAHBs (5 N-H···O⁻ and 2 O-H···N⁻).



Summary of D...A Distances: OHBs against CAHBs (1)

D–H···A	χ _P (vdW)	dD…A) OHB	d(D··A) OHB	SHR% OHB	E _{HB}	d(D···A) CAHB	SHR% CAHB	E _{HB}
[F…H…F] [_]	394	3.59	2.49	-31	≈6	2.207	-38	42(3)
[0 …H… 0]-	3.44	3.7	2.7	-27	≈ 5	2.402	-35	27(1)
[Cl····H···Cl]-	3.16	4.22	3.69	-13	≈ 2	3.093	-30	24(3)
[N … H … N]-	3.04	3.76	3.05	-19	≈ 3	2.669	-29	12(-)
[Br…H…Br]⁻	2.96	4.46	3.91	-12	≈ 2	3.377	-24	19(3)
[S…H…S] [_]	2.58	4.24	4	-6	≈ 1	3.454	-18	13(1)
[F … H … F]+	3.98	3.59	2.49	-31	≈ 6	2.284	-36	25(2)
[<mark>O</mark> …H…O]+	3.44	3.7	2.7	-27	≈ 5	2.36	-36	32(2)
[N … H … N]+	3.04	3.76	3.05	-19	≈ 3	2.592	-31	26(2)
[O…H… <mark>O</mark>]⁺	3.44	3.7	2.7	-27	≈6	2.381	-36	<u>28.7</u>
[N····H····O] [±]	<3.24>	3.73	2.87	-23	≈ 4	2.506	-33	-
[N····H···N] [±]	3.04	3.76	3.05	-19	≈ 3	2.682	-29	<u>16.4</u>

Summary of D...A Distances: OHBs against CAHBs (2)



Full range of H-bond energies, E_{HB} , and D···A contact distances spanned by the most common OHB or CAHBs homonuclear bonds.

Horizontal lines on the bottom show the full ranges of the D…A distances from the longest (Σ, vdW) to the shortest value.

The dashed curves represent the exponential dependence of E_{HB} from $d_{\text{D...A}}$ (Gilli et al., 2008).

(Slide to be discussed below in Lecture 4)

Lecture 3. PART 2 The PA/pKa Equalization Principle Modelling the H-Bond by Thermodynamic Methods

A Proposal

Two very similar proposals come from the early thermodynamic or spectroscopic investigations on the HB and are both centered on the matching of the acid-base properties of the HB donor and acceptors moieties, what we like to call, for the sake of brevity, the PA/pK_a Equalization Principle.

With reference to any generic **D**–**H**····:**A** bond, this principle states that <u>the HB is the stronger the smaller becomes the difference of the donor-acceptor</u>

proton affinities: $\Delta PA = PA(D^{-}) - PA(A)^{*}$ or acidic constants: $\Delta pK_a = pK_{AH}(D-H) - pK_{BH+}(A-H^{+})^{*}$

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1982. Malarski, Z., M. Rospenk, and L. Sobczyk J. Phys. Chem. 86: 401–406 (1982).
1984. Meot-Ner (Mautner) M., J Am Chem Soc 106: 1257–1264 (1984).

Modelling the H-Bond by Thermodynamic Methods Thermodynamic Equations

$$AH(g) \xrightarrow{K^{\circ}_{AH}} H^{+}(g) + A^{-}(g)$$

- S || + S - S || + S
$$AH(S) + S \xrightarrow{K^{\circ}_{AH}} SH^{+}(S) + A^{-}(S)$$

$$\Delta_{AH}G^{\circ} = \Delta_{AH}H^{\circ} - T \Delta_{AH}S^{\circ} = -RT \ln K^{\circ}_{AH}$$

= gas-phase acidity of AH = gas-phase basicity of A⁻;
$$\Delta_{AH}H^{\circ} = \underline{\text{proton affinity}} \text{ of } A^{-} = PA(A^{-});$$

$$\Delta_{AH}G^{\bullet} = \Delta_{AH}H^{\bullet} - T \Delta_{AH}S^{\bullet} = -RT \ln K^{\bullet}_{AH}$$

= 2.303 RT pK_{AH} = 1.364 pK_{AH} (kcal mol⁻¹ at 25°C)

$$BH^{+}(g) \xrightarrow{K^{\circ}BH_{+}} H^{+}(g) + B(g) \qquad \Delta_{BH_{+}}G^{\circ} = \Delta_{BH_{+}}H^{\circ} - T \Delta_{BH_{+}}S^{\circ} = -RT \ln K^{\circ}_{BH_{+}}$$

= gas-phase basicity of B;
$$-S \parallel + S \qquad -S \parallel + S \qquad \Delta_{BH_{+}}H^{\circ} = proton affinity of B = PA(B);$$

$$BH^{+}(S) + S \xrightarrow{K^{\circ}BH_{+}} SH^{+}(S) + B(S) \qquad \Delta_{BH_{+}}G^{\circ} = \Delta_{BH_{+}}H^{\circ} - T \Delta_{BH_{+}}S^{\circ} = -RT \ln K^{\circ}_{BH_{+}}$$

$$= 2.303 RT pK_{BH_{+}} = 1.364 pK_{BH_{+}} (kcal mol^{-1} at 25^{\circ}C)$$

The PA/pK_a equalization (or PA/pK_a matching) principle is the hypothesis that the D-H···:A bond strength can be accounted for in terms of the differences

$$\Delta PA = PA(D^{-}) - PA(:A) \quad \text{or} \quad \Delta pK_a = pK_a(D-H) - pK_a(A-H^{+}).$$

PA and **pKa** are defined by the thermodynamic equations above, respectively in the gas phase or in a solvent (S), most frequently dimethylsulphoxide (DMSO) or water (H_2O).

Modelling the H-Bond by Thermodynamic Methods Non-Homogeneity of the Thermodynamic Indicators

Starting from the 1960', there have been numerous <u>attempts to rationalize H-bond</u> <u>strengths in terms of PA/pK_a matching</u>, with particular reference to the PA interpretation of matrix-isolation IR, ICR (ion cyclotron resonance), and PHPMS (pulsed high-pressure mass spectrometry) spectroscopies. Though many of these studies were quite successful in interpreting limited series of compounds, <u>no general rule valid for all possible types of Hbonds has never been produced.</u>

The reasons have been only recently recognized (*Gilli, Pretto, and Gilli, J Mol Struct, 2007*) in defects of the previous thermodynamic treatment, whose correction has lead to single out a number of new specific and important factors:

H-bonds are not thermodynamically homogeneous but belong to four distinct thermodynamic classes, each of which must be treated in a slightly different way:

Though PA and pK_a values are strictly correlated, they are endowed with different thermodynamic properties that hinder the application of the PA/ pK_a equalization principle to some classes of chemical leitmotifs;

The interpretation of thermodynamic data is often tricky and requires much attention to be correctly performed.

Modelling the H-Bond by Thermodynamic Methods Thermodynamic H-Bond Classes

RAHBs: RAHBs cannot be treated by PA/pK_a equalization methods because π -delocalization modifies the PA/pK_a 's of the donor and acceptor moieties.

(+/-)CAHBs, PAHBs, or OHBs: A <u>proton transfer</u> from <u>an acid</u> (HB donor) to <u>a base</u> (HB acceptor) $R-D-H\cdots:A-R' \iff R-^{1/2}-D\cdots H^+\cdots A^{1/2}-R' \iff R-^{-}D:\cdots H-A^+-R'$ $\Delta pK_a = pK_{AH}(R-D-H) - pK_{BH+}(R'-A:) \qquad \Delta PA = PA(R-D^-) - PA(R'-A:)$

(-)CAHBs: A <u>proton sharing</u> between <u>two acids</u> (HB donors) $R-D-H\cdots:D'-R' \iff [R-D\cdotsH\cdotsD'-R']^{-} \Leftrightarrow R-D:\cdotsH-D'-R'$ $\Delta pK_{a} = pK_{AH}(R-D-H) - pK_{AH}(R'-D'-H) \Delta PA = PA(R-D^{-}) - PA(R'-D'^{-})$

(+)CAHBs: A *proton sharing* between *two bases* (HB acceptors)

$$R^{+}A^{-}H^{\cdots:}A'^{-}R' \iff [R^{-}A^{\cdots}H^{\cdots}A'^{-}R']^{+} \iff R^{-}A^{\cdots}H^{-}A'^{+}-R'$$
$$\Delta pK_{a} = pK_{BH^{+}}(R^{-}A^{:}) - pK_{BH^{+}}(R'^{-}A^{:}) \qquad \Delta PA = PA(R^{-}A^{:}) - PA(R'^{-}A^{:})$$

Whenever (-) and (+)CAHBs are both homonuclear (D = D' or :A = :A') and homomolecular (R = R'), the matching condition $\Delta pK_a = 0$ will hold irrespective of the actual pK_a 's of the two interacting moieties. In this case, all HBs formed will be strong!

Modelling the H-Bond by Thermodynamic Methods Differences between PA and pK_a Quantities



The plot shows the intercorrelation between PA and pK_a values. Data are arranged in two subsets for A–H acids and :B bases with correlation equations (with removed outliers)

> PA(A-) = 329(2) + 1.61(7) pK(AH, H2O)PA(B:) = 210(1) + 2.0(1) pK(BH, H2O).

The two lines are reciprocally shifted by an amount of energy that, at the zero of pK_a , is $\Delta = \Delta(329 - 210) = 119$ kcal mol⁻¹.

What does this energy difference represent?

Considering the dissociation reactions of the acid and the base in the gas phase

 $AH(g) \Leftrightarrow \underline{H^+}(g) + \underline{A^-}(g) \qquad BH^+ \Leftrightarrow H^+(g) + B(g),$

it is evident that Δ represent the work needed to take the two charges H⁺(g) and A⁻(g) from the H-bond distance (say, *R*) to the infinity (the zero of all charged quantities).

Since $\Delta \cong 119$ kcal mol⁻¹, an approximate value of the H-bond distance can be calculated as $\langle R \rangle = 332.15/1999 = 2.23$ Å, a value that, though fairly reasonable from a physical point of view, is too inaccurate for being of practical use.

☆	HB Donors 1: Inorg
0	HB Donors 2: C-H
\triangle	HB Donors 3: Org
\Leftrightarrow	HB Acceptors 1: O
•	HB Acceptors 2: N

Modelling the H-Bond by Thermodynamic Methods Differences between PA and pK_a Indicators

(+/-)CAHB, PAHB, and OHB: A <u>proton transfer</u> from <u>an acid</u> (HB donor) to <u>a base</u> (HB acceptor) $\Delta PA = PA(R-D^{-}) - PA(R'-A:)$

(-)CAHB: A proton sharing between two acids (HB donors)

 $\Delta PA = PA(R-D^{-}) - PA(R'-D'^{-})$

(+)CAHB: A *proton sharing* between *two bases* (HB acceptors)

 $\Delta PA = PA(R-A:) - PA(R'-A:')$

The electrostatic term $\Delta \cong 119$ kcal mol-1 is

(*i*) not present in (+)CAHB (neutral A and A');

(*ii*) doubly present (charged D⁻ and D'⁻) but mutually canceling out in (–)CAHB;

(iii) present in (+/–)CAHB (charged D[–])

This leads to the conclusion that the values of ΔPA can be actually used for predicting H-bond strengths of the two proton-sharing H-bonds, but cannot be used for the much more frequent proton-transfer ones.

<u>The ΔPA use is therefore limited to (-)CAHBs and (+)CAHBs in the gas phase</u>, a type of application that can be considered of specialized concern and outside the main practical interests of chemistry and biology.

Such difficulties do not subsist in solution where no electric work is involved because the proton is exchanged within the close environment of the solvated ions and, accordingly, ΔpK_a turns out to be the correct indicator of pKa equalization for both proton-transfer and proton-sharing H-bonds.

Modelling the H-Bond by Thermodynamic Methods A Verification of the PA Equalization Principle



 ΔH° vs ΔPA plot for the gas-phase dissociation equilibrium $[-O \cdots H \cdots X]^{-} \Leftrightarrow O-H + X^{-}$ studied by *Kebarle and McMahon (1971-1984)*. X represents <u>halide ions</u> (F, Cl, Br, and I) and -O-H a series of <u>organic alcohols and acids</u>. ΔPA values are <u>positive</u> when the dissociated form is $O-H + X^{-}$ (on the right) and <u>negative</u> when this form is $O^{-} + H-F$ (blue points on the left).

The points arranged on the vertical line at $\Delta PA = 0$ show, for reference, the well known homonuclear [X··H···X]⁻(–)CAHBs.

The plot perfectly follows the **PA Equalization Principle**, larger association enthalpies being all localized on the vertical line with $\Delta PA = 0$.

Note. The form of the relationship ΔH° versus ΔPA has been investigated by many authors who mostly suggested linearity but without achieving any degree of generality. It can be said that, so far, no theoretical model of this relation has ever been proposed. Another important problem to commit to posterity!

Modelling the H-Bond by Thermodynamic Methods The pK_a Equalization Principle. Retrieving a Valid List of pK_a Values

There are two known series of acidity constants that could be a basis for H-bond-strengths prediction, one measured in water, $pK_a(H_2O)$, and the other in dimethyl-sulphoxide, $pK_a(DMSO)$, but the latter (Bordwell, 2003) is still too limited to be of practical use.

Our work makes exclusive use of $pK_a(H_2O)$ values. The main difficulty is that <u>the H-bond range of</u> <u>acidities</u> is enormous (-15< $pK_a <53$) in comparison with the autoprotolysis range of water (0< pKa <14) normally investigated. Hence, measurements must be extended to solvents considerably more acidic or basic than water and then rescaled to water by various awkward methods.

Present values were mostly taken from previous compilations (Maskill, 1985; Cookson, 1974; Smith and March, 2001; Lide, 2006; Martell et al., 2004) and complemented by a number of single values from the literature. Since all data refer to water, the short notation \mathbf{pK}_{a} instead of $\mathbf{pK}_{a}(H_{2}O)$ is used, while the symbols \mathbf{pK}_{AH} and \mathbf{pK}_{BH+} are used to distinguish between the dissociation of acids and protonated bases. Final data have been arranged in order of chemical functionality:

inorganic acids, C–H acids, and organic acids as H-bond donors; organic oxygen, nitrogen and phosphorous bases as H-bond acceptors.

Extended Tables containing more that 300 chemical classes have been compiled, together with a pictorial representation of the data in form of bar chart, called the

pK_a Slide Rule.



The pK_a slide rule summarizes in pictorial form the relationships between H-bond and acid-base properties.

D–H donors (or A–H acids) on the right and :A acceptors (or B bases) on the left define three H-bond types: left-left, right-right, and right-left, equivalent to acid-acid (–)CAHBs, base-base (+)CAHBs, and acidbase (±)CAHBs and OHBs, respectively.

For these latter, the positive and negative sign of $\Delta pK_a = pK_a(right) - pK_a(leftt) = pK_a(D-H) - pK_a(A-H^+)$ distinguishes between neutral D-H····:A and ionized ^-D :···H-A⁺ bonds. Therefore, OHBs will be neutral or doubly charged according to whether the acid (on the right side) is higher or lower than the base (on the left side) while, of course, strong (±)CAHBs remain defined as the bonds where donor and acceptor face each other on a same horizontal line.

The slide rule allows us to have, at a glance, a
 comprehensive view of the most common types of H bond donors and acceptors and of the features of the
 bonds they are going to form. It is a semiempirical
 but quite efficient solution of the H-Bond Puzzle.



C–H acids display the greatest range of values, -1≤p K_a ≤53. Unfortunately, there are very few data for very acidic C–H donors matching the p K_a of carbonyls to form strong C H…:O bonds, the only case (Bock et al., 1993) being the <u>C–H…:O< adduct</u> of trinitromethane (p K_a = 0.0) with dioxane (p $K_a \approx$ <u>2.1</u>) having C…O distance of 2.936 Å, possibly the shortest ever observed.

Other acids are in the range $-1 \le pK_a \le 40$, the strongest being trinitrophenols ($-0.7 \le pK_a \le 0.33$) and the weakest ones organic amines (pK_a around 39-40).

The order of decreasing acidity is **carboxylic acids** > **phenols** > **alcohols** and **amides** > **anilines** > **amines** and, in each class, the acidity is seen to be enhanced by halogenation or nitration.

Thioles ($6.5 \le pK_a \le 11$), enols ($8.5 \le pK_a \le 12$), and oximes ($10 \le pK_a \le 12$) are significantly more acidic than the corresponding alcohols ($15 \le pK_a \le 18$).



The acceptor range $(-12 \le pK_a \le 16)$ is shifted upwards with respect to donors, so that nitro, carbonyl, nitriles, ethers, alcohols, and sulphoxides acceptors face a few inorganic acids but no organic donors with which can only form weak H-bonds.

Likewise, the weakest donors (amines, anilines, and alcohols) do not face any known acceptor and can then form only weak and neutral $D-H\cdots$: A bonds.

The maximum donor-acceptor overlap occurs in the interval of water autoprotolysis ($0 \le pK_a \le 14$), where the greatest part of the strong H-bonds are expected.

Finally, the slide rule gives <u>a new insight</u> into the most common and familiar types of H-bonds. For instance, we speak without distinction of amine-nitro, aniline-ketone, water-water, carboxylic acid-pyridine, and perchloric acid-aniline H-bonds not considering that their ΔpK_a 's are as different as 50, 46, 17.4, -1, and -20 pK_a units corresponding to 67, 61, 23, -1.3, and -27 kcal mol⁻¹ at 25 °C and that this impressive series of numbers actually represent the energy difference between their tautomeric -D:...H-A⁺ and D-H...:A forms.



The pK_a slide rule is a tool for the graphical evaluation of the difference $\Delta pK_a = pK_{AH}(D-H) - pK_{BH+}(A-H^+)$ for the most common classes of organic compounds. HB Acceptors are on the left and HB Donors on the right. pK_a values are given for chemical class.

Results expected: $\Delta p K_a >>0$: D-H····A, weak & neutral $\Delta p K_a \approx 0$: D···H···A, strong & centered $\Delta p K_a <<0$: D····H-A+, weak & charged

pK _a ranges	of organic compounds:
C-H acids	$-11 < pK_a < 53$
Other Donors	$-1 < pK_a < 40$
Acceptors	$-12 < pK_a < 16$
All	<u>-15 < pK₂ < 53</u>
pK _a in water	$0 < pK_a < 14$

End of Lecture 3