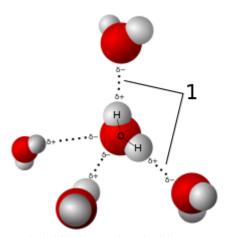
Hydrogen bond

A **hydrogen bond** (often informally abbreviated **H-bond**) is a primarily <u>electrostatic</u> force of attraction between a <u>hydrogen</u> (H) atom which is <u>covalently bound</u> to a more <u>electronegative</u> atom or group, particularly the second-row elements <u>nitrogen</u> (N), <u>oxygen</u> (O), or <u>fluorine</u> (F)—the hydrogen bond donor (Dn)—and another electronegative atom bearing a <u>lone pair</u> of electrons—the hydrogen bond acceptor (Ac). Such an interacting system is generally denoted Dn–H···Ac, where the solid line denotes a polar <u>covalent bond</u>, and the dotted or dashed line indicates the hydrogen bond. The use of three centered dots for the hydrogen bond is specifically recommended by the <u>IUPAC</u>. [4] While hydrogen bonding has both covalent and electrostatic contributions, and the degrees to which they contribute are currently debated, the present evidence strongly implies that the primary contribution is covalent. [5]



Model of hydrogen bonds (1) between molecules of water

Hydrogen bonds can be <u>intermolecular</u> (occurring between separate molecules) or <u>intramolecular</u> (occurring among parts of the same molecule). Depending on the nature of the donor and acceptor atoms which constitute the bond, their geometry, and environment, the energy of a hydrogen bond can vary between 1 and 40 kcal/mol. This makes them somewhat stronger than a <u>van der Waals interaction</u>, and weaker than fully <u>covalent</u> or <u>ionic bonds</u>. This type of bond can occur in inorganic molecules such as water and in <u>organic molecules</u> like <u>DNA</u> and proteins.

The hydrogen bond is responsible for many of the anomalous physical and chemical properties of compounds of N, O, and F. In particular, intermolecular hydrogen bonding is responsible for the high boiling point of water (100 °C) compared to the other group 16 hydrides that have much weaker hydrogen bonds. [11] Intramolecular hydrogen bonding is partly responsible for the secondary and tertiary structures of proteins and nucleic acids. It also plays an important role in the structure of polymers, both synthetic and natural.

Weaker hydrogen bonds^[12] are known for hydrogen atoms bound to elements such as sulfur (S) or chlorine (Cl); even carbon (C) can serve as a donor, particularly when the carbon or one of its neighbors is electronegative (e.g., in chloroform, aldehydes and terminal acetylenes). Gradually, it was recognized that there are many examples of weaker hydrogen bonding involving donor other than N, O, or F and/or acceptor Ac with electronegativity approaching that of hydrogen (rather than being much more electronegative). Though these "non-traditional" hydrogen bonding interactions are often quite weak (\sim 1 kcal/mol), they are also ubiquitous and are increasingly recognized as important control elements in receptor-ligand interactions in medicinal chemistry or intra-/intermolecular interactions in materials sciences. The definition of hydrogen bonding has gradually broadened over time to include these weaker attractive interactions. In 2011, an IUPAC Task Group recommended a modern evidence-based definition of hydrogen bonding, which was published in the IUPAC journal *Pure and Applied Chemistry*. This definition specifies:

The hydrogen bond is an attractive interaction between a hydrogen atom from a molecule or a molecular fragment X–H in which X is more electronegative than H, and an atom or a group of atoms in the same or a different molecule, in which there is evidence of bond formation. [15]

As part of a more detailed list of criteria, the IUPAC publication acknowledges that the attractive interaction can arise from some combination of electrostatics (multipole-multipole and multipole-induced multipole interactions), covalency (charge transfer by orbital overlap), and dispersion (London forces), and states that the relative importance of each will vary depending on the system. However, a footnote to the criterion recommends the exclusion of interactions in which dispersion is the primary contributor, specifically giving Ar—CH₄ and CH₄—CH₄ as examples of such interactions to be excluded from the definition. [4]

Nevertheless, most introductory textbooks still restrict the definition of hydrogen bond to the "classical" type of hydrogen bond characterized in the opening paragraph.

Contents

Bonding

Definitions and general characteristics

Bond strength

Structural details

Spectroscopy

Theoretical considerations

History

Hydrogen bonds in small molecules

Water

Bifurcated and over-coordinated hydrogen bonds in water

Other liquids

Further manifestations of solvent hydrogen bonding

Hydrogen bonds in polymers

DNA

Proteins

Cellulose

Synthetic polymers

Symmetric hydrogen bond

Dihydrogen bond

Dynamics probed by spectroscopic means

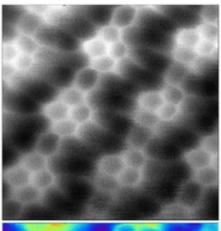
Application to drugs

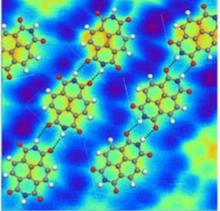
Hydrogen bonding phenomena

References

Further reading

External links





AFM image of napthalenetetracarboxylic diimide molecules on silver-terminated silicon, interacting via hydrogen bonding, taken at 77 K.[1] ("Hydrogen bonds" in the top image are exaggerated by artifacts of the imaging technique.[2][3])

Definitions and general characteristics

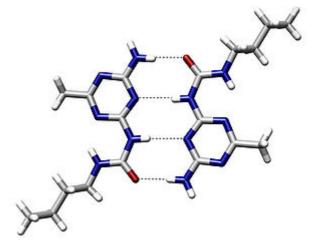
A hydrogen atom attached to a relatively electronegative atom is the hydrogen bond *donor*. [17] C-H bonds only participate in hydrogen bonding when the carbon atom is bound to electronegative substituents, as is the case in chloroform, CHCl₃.[18] In a hydrogen bond, the electronegative atom not covalently attached to the hydrogen is named proton acceptor, whereas the one covalently bound to the hydrogen is named the proton donor. While this nomenclature is recommended by the IUPAC, [4] it can be misleading, since in other donoracceptor bonds, the donor/acceptor assignment is based on the source of the electron pair (such nomenclature is also used for hydrogen bonds by some authors [19]). In the hydrogen bond donor, the H center is protic. The donor is a Lewis acid. Hydrogen bonds are represented as H···Y system, where the dots represent the hydrogen bond. Liquids that display hydrogen bonding (such as water) are called **associated liquids**.

The hydrogen bond is often described as an <u>electrostatic</u> <u>dipole-dipole interaction</u>. However, it also has some features of <u>covalent bonding</u>: it is directional and strong, produces interatomic distances shorter than the sum of the van der Waals radii, and usually involves a limited number of interaction partners, which can be interpreted as a type of <u>valence</u>. These covalent features are more substantial when acceptors bind hydrogens from more electronegative donors.

Bond strength

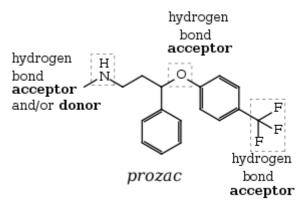
Hydrogen bonds can vary in strength from weak (1–2 kJ mol⁻¹) to strong (161.5 kJ mol⁻¹ in the ion $\overline{\text{HF}}_2^-$). [20][21] Typical enthalpies in vapor include:

- F-H···:F (161.5 kJ/mol or 38.6 kcal/mol), illustrated uniquely by HF₂⁻, bifluoride
- O-H···:N (29 kJ/mol or 6.9 kcal/mol), illustrated water-ammonia
- O-H···:O (21 kJ/mol or 5.0 kcal/mol), illustrated water-water, alcohol-alcohol
- N-H···:N (13 kJ/mol or 3.1 kcal/mol), illustrated by ammonia-ammonia
- N-H···:O (8 kJ/mol or 1.9 kcal/mol), illustrated water-amide
- OH₃+···:OH₂ (18 kJ/mol^[23] or 4.3 kcal/mol)



An example of <u>intermolecular</u> hydrogen bonding in a <u>self-assembled</u> dimer complex. [16] The hydrogen bonds are represented by dotted lines.

<u>Intramolecular</u> hydrogen bonding in <u>acetylacetone</u> helps stabilize the enol tautomer.



Examples of hydrogen bond donating (donors) and hydrogen bond accepting groups (acceptors)

The strength of intermolecular hydrogen bonds is most often evaluated by measurements of equilibria between molecules containing donor and/or acceptor units, most often in solution. The strength of intramolecular hydrogen bonds can be studied with equilibria between conformers with and without hydrogen bonds. The most important method for the identification of hydrogen bonds also in complicated molecules is crystallography, sometimes also NMR-spectroscopy. Structural details, in particular distances between donor and acceptor which

Cyclic dimer of acetic acid; dashed **green** lines represent hydrogen bonds

are smaller than the sum of the van der Waals radii can be taken as indication of the hydrogen bond strength.

One scheme gives the following somewhat arbitrary classification: those that are 15 to 40 kcal/mol, 5 to 15 kcal/mol, and >0 to 5 kcal/mol are considered strong, moderate, and weak, respectively.

Structural details

The X–H distance is typically ≈ 110 pm, whereas the H···Y distance is ≈ 160 to 200 pm. The typical length of a hydrogen bond in water is 197 pm. The ideal bond angle depends on the nature of the hydrogen bond donor. The following hydrogen bond angles between a hydrofluoric acid donor and various acceptors have been determined experimentally: [25]

| Acceptordonor | VSEPR geometry | Angle (°) |
|------------------------|-----------------|-----------|
| HCNHF | linear | 180 |
| H ₂ CO···HF | trigonal planar | 120 |
| H ₂ O···HF | pyramidal | 46 |
| H ₂ S····HF | pyramidal | 89 |
| SO ₂ ···HF | trigonal | 142 |

Spectroscopy

Strong hydrogen bonds are revealed by downfield shifts in the ${}^{1}H$ NMR spectrum. For example, the acidic proton in the enol tautomer of <u>acetylacetone</u> appears at δ_{H} 15.5, which is about 10 ppm downfield of a conventional alcohol. [26]

In the IR spectrum, hydrogen bonding shifts the X-H stretching frequency to lower energy (i.e. the vibration frequency decreases). This shift reflects a weakening of the X-H bond. Certain hydrogen bonds - improper hydrogen bonds - show a blue shift of the X-H stretching frequency and a decrease in the bond length. Honds can also be measured by IR vibrational mode shifts of the acceptor. The amide I mode of backbone carbonyls in α -helices shifts to lower frequencies when they form H-bonds with side-chain hydroxyl groups. [28]

Theoretical considerations

Hydrogen bonding is of continuing theoretical interest. According to a modern description O:H-O integrates both the intermolecular O:H lone pair ":" nonbond and the intramolecular H-O polar-covalent bond associated with O-O repulsive coupling. [29]

Quantum chemical calculations of the relevant interresidue potential constants (compliance constants) revealed large differences between individual H bonds of the same type. For example, the central interresidue $N-H\cdots N$ hydrogen bond between guanine and cytosine is much stronger in comparison to the $N-H\cdots N$ bond between the adenine-thymine pair. [30]

Theoretically, the bond strength of the hydrogen bonds can be assessed using NCI index, <u>non-covalent</u> <u>interactions index</u>, which allows a visualization of these <u>non-covalent interactions</u>, as its name indicates, using the electron density of the system.

From interpretations of the <u>anisotropies</u> in the <u>Compton profile</u> of ordinary ice that the hydrogen bond is partly covalent. [31] However, this interpretation was challenged. [32]

Most generally, the hydrogen bond can be viewed as a <u>metric</u>-dependent <u>electrostatic</u> <u>scalar field</u> between two or more intermolecular bonds. This is slightly different from the <u>intramolecular bound states</u> of, for example, <u>covalent or ionic bonds</u>; however, hydrogen bonding is generally still a <u>bound state</u> phenomenon, since the <u>interaction energy</u> has a net negative sum. The initial theory of hydrogen bonding proposed by <u>Linus Pauling</u> suggested that the hydrogen bonds had a partial covalent nature. This interpretation remained controversial until <u>NMR techniques</u> demonstrated information transfer between hydrogen-bonded nuclei, a feat that would only be possible if the hydrogen bond contained some covalent character. [33]

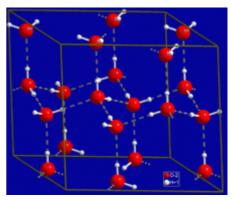
History

The concept of hydrogen bonding once was challenging. Linus Pauling credits T. S. Moore and T. F. Winmill with the first mention of the hydrogen bond, in 1912. Moore and Winmill used the hydrogen bond to account for the fact that trimethylammonium hydroxide is a weaker base than tetramethylammonium hydroxide. The description of hydrogen bonding in its better-known setting, water, came some years later, in 1920, from Latimer and Rodebush. In that paper, Latimer and Rodebush cite work by a fellow scientist at their laboratory, Maurice Loyal Huggins, saying, "Mr. Huggins of this laboratory in some work as yet unpublished, has used the idea of a hydrogen kernel held between two atoms as a theory in regard to certain organic compounds."

Hydrogen bonds in small molecules

Water

A ubiquitous example of a hydrogen bond is found between <u>water</u> molecules. In a discrete water molecule, there are two hydrogen atoms and one oxygen atom. Two molecules of <u>water</u> can form a hydrogen bond between them that is to say <u>oxygen-hydrogen bonding</u>; the simplest case, when only two molecules are present, is called the <u>water dimer</u> and is often used as a model system. When more molecules are present, as is the case with liquid water, more bonds are possible because the oxygen of one water molecule has two lone pairs of electrons, each of which can form a hydrogen bond with a hydrogen on another water molecule. This can repeat such that every water molecule is H-bonded with up to four other molecules, as shown in the figure (two through its two lone pairs, and two through

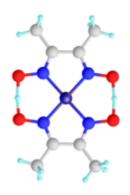


Crystal structure of hexagonal ice. Gray dashed lines indicate hydrogen bonds

its two hydrogen atoms). Hydrogen bonding strongly affects the <u>crystal structure</u> of <u>ice</u>, helping to create an open hexagonal lattice. The density of ice is less than the density of water at the same temperature; thus, the solid phase of water floats on the liquid, unlike most other substances.

<u>Liquid</u> water's high <u>boiling point</u> is due to the high number of hydrogen bonds each molecule can form, relative to its low <u>molecular mass</u>. Owing to the difficulty of breaking these bonds, water has a very high boiling point, melting point, and viscosity compared to otherwise similar liquids not conjoined by hydrogen bonds. Water is unique because its oxygen atom has two lone pairs and two hydrogen atoms, meaning that the total number of bonds of a water molecule is up to four.

The number of hydrogen bonds formed by a molecule of liquid water fluctuates with time and temperature. From <u>TIP4P</u> liquid water simulations at 25 °C, it was estimated that each water molecule participates in an average of 3.59 hydrogen bonds. At 100 °C, this number decreases to 3.24 due to the increased molecular motion and decreased density, while at 0 °C, the average number of hydrogen bonds increases to 3.69. A more recent study found a much smaller number of hydrogen bonds: 2.357 at 25 °C. The differences may be due to the use of a different method for defining and counting the hydrogen bonds.



Structure of <u>nickel</u> <u>bis(dimethylglyoximat</u> which features two linear hydrogen-bonds.

Where the bond strengths are more equivalent, one might instead find the atoms of two interacting water molecules partitioned into two <u>polyatomic ions</u> of opposite charge, specifically <u>hydroxide</u> (OH^-) and <u>hydronium</u> (H_3O^+). (Hydronium ions are also known as "hydroxonium" ions.)

Indeed, in pure water under conditions of standard temperature and pressure, this latter formulation is applicable only rarely; on average about one in every 5.5×10^8 molecules gives up a proton to another water molecule, in accordance with the value of the <u>dissociation constant</u> for water under such conditions. It is a crucial part of the uniqueness of water.

Because water may form hydrogen bonds with solute proton donors and acceptors, it may competitively inhibit the formation of solute intermolecular or intramolecular hydrogen bonds. Consequently, hydrogen bonds between or within solute molecules dissolved in water are almost always unfavorable relative to hydrogen bonds between water and the donors and acceptors for hydrogen bonds on those solutes. [40] Hydrogen bonds between water molecules have an average lifetime of 10^{-11} seconds, or 10 picoseconds.

Bifurcated and over-coordinated hydrogen bonds in water

A single hydrogen atom can participate in two hydrogen bonds, rather than one. This type of bonding is called "bifurcated" (split in two or "two-forked"). It can exist, for instance, in complex natural or synthetic organic molecules. [42] It has been suggested that a bifurcated hydrogen atom is an essential step in water reorientation. [43]

Acceptor-type hydrogen bonds (terminating on an oxygen's lone pairs) are more likely to form bifurcation (it is called overcoordinated oxygen, OCO) than are donor-type hydrogen bonds, beginning on the same oxygen's hydrogens. [44]

Other liquids

For example, <u>hydrogen fluoride</u>—which has three lone pairs on the F atom but only one H atom—can form only two bonds; (ammonia has the opposite problem: three hydrogen atoms but only one lone pair).

Further manifestations of solvent hydrogen bonding

- Increase in the melting point, boiling point, solubility, and viscosity of many compounds can be explained by the concept of hydrogen bonding.
- Negative azeotropy of mixtures of HF and water
- The fact that ice is less dense than liquid water is due to a crystal structure stabilized by hydrogen bonds.
- Dramatically higher boiling points of NH₃, H₂O, and HF compared to the heavier analogues PH₃, H₂S, and HCl, where hydrogen-bonding is absent.
- Viscosity of anhydrous phosphoric acid and of glycerol
- Dimer formation in <u>carboxylic acids</u> and hexamer formation in <u>hydrogen fluoride</u>, which occur even in the gas phase, resulting in gross deviations from the ideal gas law.
- Pentamer formation of water and alcohols in apolar solvents.

Hydrogen bonds in polymers

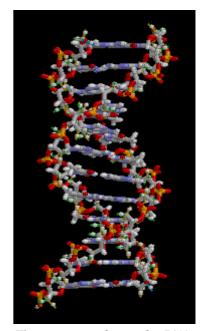
Hydrogen bonding plays an important role in determining the three-dimensional structures and the properties adopted by many synthetic and natural proteins. Compared to the C-C, C-O, and C-N bonds that comprise most polymers, hydrogen bonds are far weaker, perhaps 5%. Thus, hydrogen bonds can be broken by chemical or mechanical means while retaining the basic structure of the polymer backbone. This hierarchy of bond strengths (covalent bonds being stronger than hydrogen-bonds being stronger than van der Waals forces) is key to understanding the properties of many materials. [45]

DNA

In these macromolecules, bonding between parts of the same macromolecule cause it to fold into a specific shape, which helps determine the molecule's physiological or biochemical role. For example, the double helical structure of <u>DNA</u> is due largely to hydrogen bonding between its <u>base pairs</u> (as well as <u>pi stacking</u> interactions), which link one complementary strand to the other and enable replication.

Proteins

In the secondary structure of proteins, hydrogen bonds form between the backbone oxygens and amide hydrogens. When the spacing of the amino acid residues participating in a hydrogen bond occurs regularly between positions i and i+4, an alpha helix is formed. When the spacing is less, between positions i and i+3, then a 3_{10} helix is formed. When two strands are joined by hydrogen bonds involving alternating residues on each participating strand, a beta sheet is formed. Hydrogen bonds also play a part in forming the tertiary structure of protein through interaction of R-groups. (See also protein folding).



The structure of part of a DNA double helix

<u>Bifurcated H-bond</u> systems are common in alpha-helical <u>transmembrane</u> <u>proteins</u> between the backbone amide C=O of residue i as the H-bond acceptor and two H-bond donors from residue i+4: the backbone amide N-H and a side-chain hydroxyl or thiol H⁺. The energy preference of the

bifurcated H-bond hydroxyl or thiol system is -3.4 kcal/mol or -2.6 kcal/mol, respectively. This type of bifurcated H-bond provides an intrahelical H-bonding partner for polar side-chains, such as <u>serine</u>, threonine, and <u>cysteine</u> within the hydrophobic membrane environments. [46]

The role of hydrogen bonds in protein folding has also been linked to osmolyte-induced protein stabilization. Protective osmolytes, such as <u>trehalose</u> and <u>sorbitol</u>, shift the protein folding equilibrium toward the folded state, in a concentration dependent manner. While the prevalent explanation for osmolyte action relies on excluded volume effects that are entropic in nature, recent <u>circular dichroism</u> (CD) experiments have shown osmolyte to act through an enthalpic effect. [47] The

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Hydrogen bonding between <u>guanine</u> and <u>cytosine</u>, one of two types of base pairs in DNA

molecular mechanism for their role in protein stabilization is still not well established, though several mechanisms have been proposed. Recently, computer <u>molecular dynamics</u> simulations suggested that osmolytes stabilize proteins by modifying the hydrogen bonds in the protein hydration layer. [48]

Several studies have shown that hydrogen bonds play an important role for the stability between subunits in multimeric proteins. For example, a study of sorbitol dehydrogenase displayed an important hydrogen bonding network which stabilizes the tetrameric quaternary structure within the mammalian sorbitol dehydrogenase protein family. [49]

A protein backbone hydrogen bond incompletely shielded from water attack is a <u>dehydron</u>. Dehydrons promote the removal of water through proteins or <u>ligand binding</u>. The exogenous dehydration enhances the <u>electrostatic</u> interaction between the <u>amide</u> and <u>carbonyl</u> groups by de-shielding their <u>partial charges</u>. Furthermore, the dehydration stabilizes the hydrogen bond by destabilizing the <u>nonbonded state</u> consisting of dehydrated isolated charges. [50]

<u>Wool</u>, being a protein fibre, is held together by hydrogen bonds, causing wool to recoil when stretched. However, washing at high temperatures can permanently break the hydrogen bonds and a garment may permanently lose its shape.

Cellulose

Hydrogen bonds are important in the structure of $\underline{\text{cellulose}}$ and derived polymers in its many different forms in nature, such as cotton and flax.

Synthetic polymers

Many polymers are strengthened by hydrogen bonds within and between the chains. Among the synthetic polymers, a well characterized example is nylon, where hydrogen bonds occur in the repeat unit and play a major role in crystallization of the material. The bonds occur between carbonyl and amine groups in the amide repeat unit. They effectively link adjacent chains, which help reinforce the material. The effect is

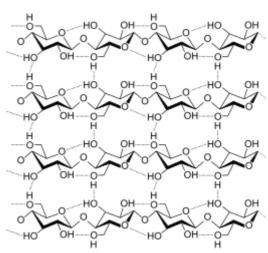
Para-aramid structure

great in <u>aramid</u> <u>fibre</u>, where hydrogen bonds stabilize the linear chains laterally. The chain axes are aligned along the fibre axis, making the fibres extremely stiff and strong.

The hydrogen-bond networks make both natural and synthetic polymers sensitive to <u>humidity</u> levels in the atmosphere because water molecules can diffuse into the surface and disrupt the network. Some polymers are more sensitive than others. Thus <u>nylons</u> are more sensitive than <u>aramids</u>, and <u>nylon 6</u> more sensitive than nylon-11.

Symmetric hydrogen bond

A <u>symmetric hydrogen bond</u> is a special type of hydrogen bond in which the proton is spaced exactly halfway between two identical atoms. The strength of the bond to each of those atoms is equal. It is an example of a <u>three-center four-electron bond</u>. This type of bond is much stronger than a "normal" hydrogen



A strand of cellulose (conformation I_{α}), showing the hydrogen bonds (dashed) within and between cellulose molecules

bond. The effective bond order is 0.5, so its strength is comparable to a covalent bond. It is seen in ice at high pressure, and also in the solid phase of many anhydrous acids such as <u>hydrofluoric acid</u> and <u>formic acid</u> at high pressure. It is also seen in the <u>bifluoride</u> ion $[F-H-F]^-$. Due to severe steric constraint, the protonated form of Proton Sponge (1,8-bis(dimethylamino)naphthalene) and its derivatives also have symmetric hydrogen bonds $([N-H-N]^+)$, although in the case of protonated Proton Sponge, the assembly is bent. [52]

Symmetric hydrogen bonds have been observed recently spectroscopically in <u>formic acid</u> at high pressure (>GPa). Each hydrogen atom forms a partial covalent bond with two atoms rather than one. Symmetric hydrogen bonds have been postulated in ice at high pressure (Ice X). <u>Low-barrier hydrogen bonds</u> form when the distance between two heteroatoms is very small.

Dihydrogen bond

The hydrogen bond can be compared with the closely related <u>dihydrogen bond</u>, which is also an <u>intermolecular</u> bonding interaction involving hydrogen atoms. These structures have been known for some time, and well characterized by <u>crystallography</u>; <u>[53]</u> however, an understanding of their relationship to the conventional hydrogen bond, <u>ionic bond</u>, and <u>covalent bond</u> remains unclear. Generally, the hydrogen bond is characterized by a proton acceptor that is a lone pair of electrons in nonmetallic atoms (most notably in the <u>nitrogen</u>, and <u>chalcogen</u> groups). In some cases, these proton acceptors may be <u>pi-bonds</u> or <u>metal complexes</u>. In the dihydrogen bond, however, a metal hydride serves as a proton acceptor, thus forming a hydrogen-hydrogen interaction. <u>Neutron diffraction</u> has shown that the <u>molecular geometry</u> of these complexes is similar to hydrogen bonds, in that the bond length is very adaptable to the metal complex/hydrogen donor system. <u>[53]</u>

Dynamics probed by spectroscopic means

The dynamics of hydrogen bond structures in water can be probed by the <u>IR spectrum</u> of OH stretching vibration. In the hydrogen bonding network in protic organic ionic plastic crystals (POIPCs), which are a type of phase change material exhibiting solid-solid phase transitions prior to melting, variable-temperature infrared spectroscopy can reveal the temperature dependence of hydrogen bonds and the dynamics of both the anions and the cations. The sudden weakening of hydrogen bonds during the solid-solid phase transition seems to be coupled with the onset of orientational or rotational disorder of the ions.

Application to drugs

Hydrogen bonding is a key to the design of drugs. According to <u>Lipinski's rule of five</u> the majority of orally active drugs tend to have between five and ten hydrogen bonds. These interactions exist between <u>nitrogen-hydrogen</u> and <u>oxygen-hydrogen</u> centers. [56] As with many other rules of thumb, many exceptions exist.

Hydrogen bonding phenomena

- Occurrence of <u>proton tunneling</u> during DNA replication is believed to be responsible for cell mutations.
- High water solubility of many compounds such as ammonia is explained by hydrogen bonding with water molecules.
- Deliquescence of NaOH is caused in part by reaction of OH⁻ with moisture to form hydrogen-bonded H₃O₂⁻ species. An analogous process happens between NaNH₂ and NH₃, and between NaF and HF.
- The presence of hydrogen bonds can cause an anomaly in the normal succession of <u>states of matter</u> for certain mixtures of <u>chemical compounds</u> as temperature increases or decreases. These compounds can be liquid until a certain temperature, then solid even as the temperature increases, and finally liquid again as the temperature rises over the "anomaly interval" [58]
- Smart rubber utilizes hydrogen bonding as its sole means of bonding, so that it can "heal" when torn, because hydrogen bonding can occur on the fly between two surfaces of the same polymer.

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■ George A. Jeffrey. *An Introduction to Hydrogen Bonding (Topics in Physical Chemistry)*. Oxford University Press, USA (March 13, 1997). ISBN 0-19-509549-9

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- The Bubble Wall (https://web.archive.org/web/20141203035841/http://www.magnet.fsu.edu/edu cation/tutorials/slideshows/bubblewall/index.html) (Audio slideshow from the National High Magnetic Field Laboratory explaining cohesion, surface tension and hydrogen bonds)
- isotopic effect on bond dynamics (http://scitation.aip.org/content/aip/journal/jcp/129/19/10.1063/ 1.3006032)

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