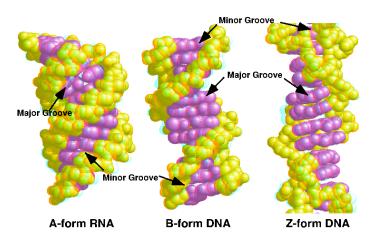
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A-form

- · Most RNA and RNA-DNA duplex in this form
- Shorter, wider helix than B.
- Deep, narrow major groove not easily accessible to proteins
- Wide, shallow minor groove accessible to proteins, but lower information content than major groove.
- Favored conformation at low water concentrations
- Base pairs tilted to helix axis and displaced from axis
- Sugar pucker C3'-endo (in RNA 2'-OH inhibits C2'-endo conformation)

B-form

- Most common DNA conformation in vivo
- Narrower, more elongated helix than A.
- Wide major groove easily accessible to proteins
- Narrow minor groove
- Favored conformation at high water concentrations (hydration of minor groove seems to favor B-form)
- Base pairs nearly perpendicular to helix axis
- Sugar pucker C2'-endo

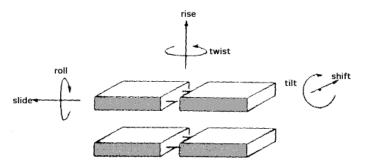
Z-form

- Helix has left-handed sense
- Can be formed in vivo, given proper sequence and superhelical tension, but function remains obscure.
- Narrower, more elongated helix than A or B.
- Major "groove" not really groove
- Narrow minor groove
- Conformation favored by high salt concentrations, some base substitutions, but requires alternating purine-pyrimidine sequence.
- N2-amino of G H-bonds to 5' PO: explains slow exchange of proton, need for G purine.
- Base pairs nearly perpendicular to helix axis
- GpC repeat, not single base-pair
 - o P-P distances: vary for GpC and CpG
 - o GpC stack: good base overlap
 - o CpG: less overlap.
- Zigzag backbone due to C sugar conformation compensating for G glycosidic bond conformation
- Conformations:
 - o C; anti, C2'-endo
 - o G; syn, C3'-endo

The geometry of the DNA forms can be used to describe the differences seen.

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| | Α | В | z |
|---------------------------|--------------|------------------|-----------------------------|
| Helix sense | Right handed | Right- handed | Left handed |
| Repeating unit | 1 bp | 1bp | 2 bp |
| Rotation/bp | 33.6° | 35.9° | 60°/2 |
| Mean bp/turn | 10.7 | 10.0 | 12 |
| Inclination of bp to axis | +19° | -1.2° | -9° |
| Rise/bp along axis | 2.3Å | 3.32Å | 3.8Å |
| Pitch/turn of helix | 24.6Å | 33.2Å | 45.6Å |
| Mean propeller twist | +18° | +16° | 0° |
| Glycosyl angle | anti | anti | C: anti, G: syn |
| Sugar pucker | C3'-endo | C2'-endo | C: C2'-endo, G: C3'-endo |
| Diameter | 26Å | 20Å | 18Å |

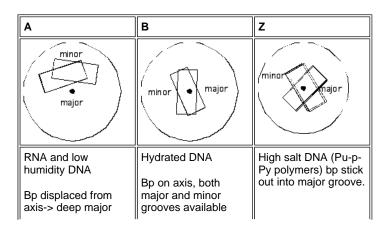


Grooves and stacking of bases:

Major and minor groove definition:



Sugars are on minor groove side of the base pair



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| groove | | |
|--------|--|--|
|--------|--|--|

Sugar pucker:

Differences --> differences in forms of helices

| A | В | Z |
|--|-------------------|----------------------|
| A-form RNA | B-form DNA | αβε |
| | | Z-form DNA |
| | | |
| C3'-endo (favored in RNA due to steric problems with 2'OH) | C2'-endo | C: C2'-endo |
| , | | G: C3'-endo |
| Glycosyl (c) anti | Glycosyl (c) anti | C: Glycosyl (c) anti |
| | | G: Glycosyl (c) syn |