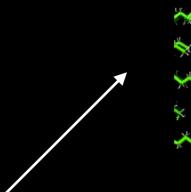
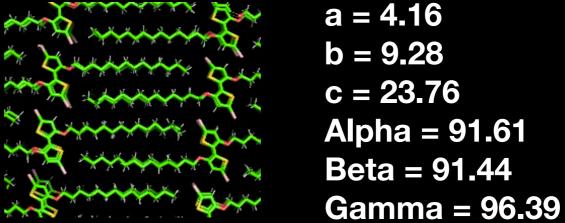
Alkylated Crystal Structure (units in angstrom)

Annealing

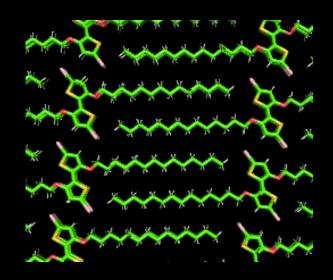
No annealing

Run times up to 60ns





Before





b = 9.298

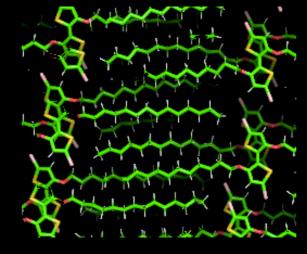
c = 23.806

Alpha = 91.86

Beta = 91.48

Gamma = 96.45





$$a = 4.23$$

b = 9.46

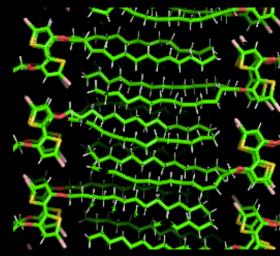
c = 24.23

Alpha = 92.90

Beta = 89.80

Gamma = 96.24

Annealed at 400K



$$a = 4.21$$

b = 9.39

c = 24.05

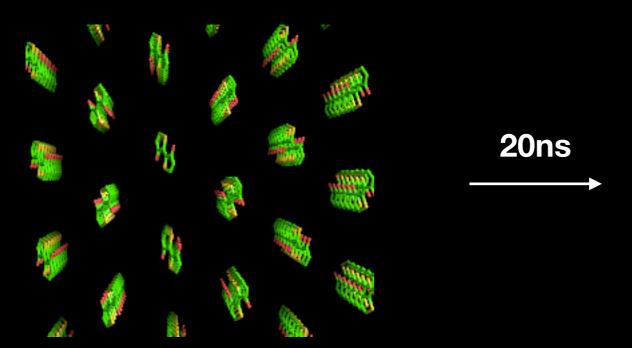
Alpha = 93.55

Beta = 89.69

Gamma = 94.87

Glycolated Crystal Structure - re-run with better optimised C-C-O Angle parameters

Before (only backbones shown)



$$a = 7.01$$

$$b = 14.02$$

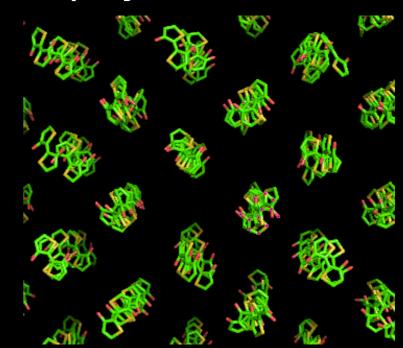
$$c = 18.01$$

$$Alpha = 90.00$$

Beta =
$$97.71$$

$$Gamma = 90.00$$

After (only backbones shown)



$$a = 7.21$$

$$b = 14.44$$

$$c = 18.55$$

Alpha =
$$90.00$$

Beta =
$$97.71$$

$$Gamma = 90.00$$

Atomic Displacements

$$dx = 0.461 + / - 2.30$$

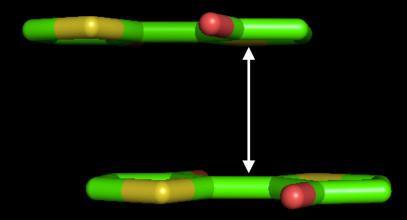
$$dy = 0.501 + /- 1.90$$

$$Dz = 0.871 + / - 1.91$$

Much more stable than last week

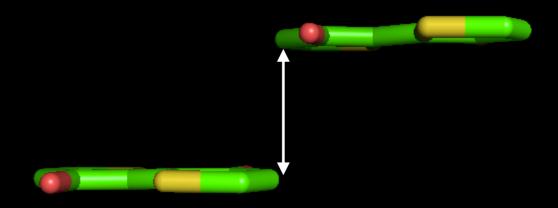
Pi-pi spacing

Alkylated structure



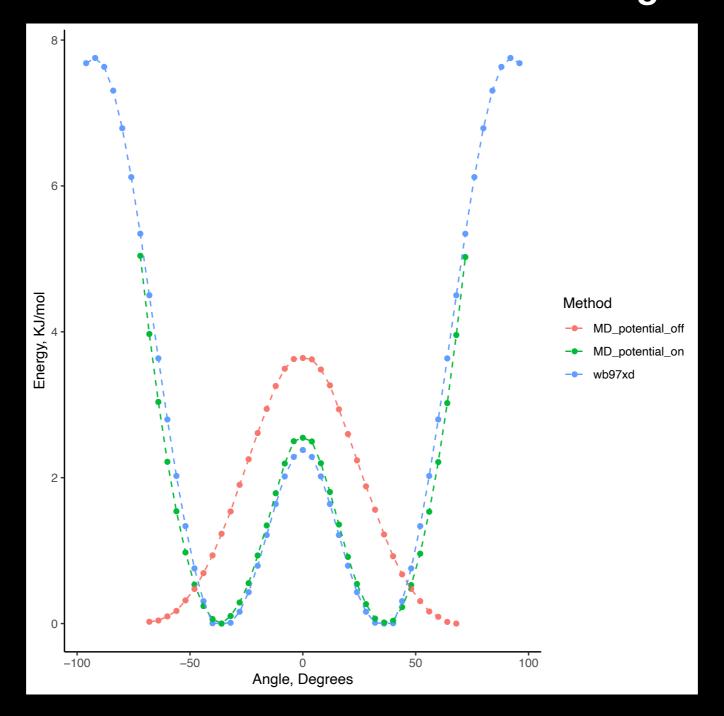
Pi-pi distance before - 3.58 Pi-pi distance after MD - 3.67

Glycolated structure



Pi-pi distance before - 3.64 Pi-pi distance after MD - 3.66 Does this even count as a pi-stack?

Outer dihedral scan now fitted also. Doing CCSD and MP2/4 to check the barrier height.



Let's do some polymer simulations!