Notes to the Excercise

- 1. Install NMRFAM-Sparky or alternative program for NMR.
- 2. Open *ucsf files and adjust spectral properties (S/N ration, colors, levels). Create *save and *proj files. (SPARKY)
- 3. Inspect the skeleton of solved compound (attached figure).
- 4. Assign max. of ¹H and ¹³C resonances, determine position of 2 fragments. Specify stereochemistry of all ¹H signals/substituents.

Few comments

- · HSQC spectrum is not fully evolved for all crosspeaks :-(
- Two bond couplings in HMBC are in some cases smaller than three-bond.
- · For crosspeaks in t1-noise check their presence in opposite dimension.
- · Always double check the individual assignments steps!
- · Solvent chloroform with water residuals, t = 5°C.

Requested outputs

- 1. Complete assignment of crosspeaks → send *save files.
- 2. Build final structure of the molecule.
- 3. Prepare a comprehensive report with comments describing logically the procedure of your work. Wrap your report together with *save files and deposit it via IS.