

# **Molecules**

ID, group, name, Summenformel, molecular formula (image/ChemDraw), SMILES-String

## **Single Molecules**

ID, *further information*

## **Triplet-Doublet pairs**

ID, Doublet, Linker, Chromophore

(Linker includes possibility to define bimolecular systems and supramolecular systems)

## **Radical pairs**

ID, Radical 1, Radical 2

## **Triplet-Triplet pairs**

ID, Triplet 1, Triplet 2

## **Measurement**

ID, Molecule, Method, Temperature, Solvent, *Concentration*, Date, *Location*, *Device*, *Series*, Path to the dataset, data corrected? Data evaluated?

## **TREPR**

ID, Frequency (XQW), Excitation wavelength, *Excitation energy*, Attenuation, *Number of scans*, *Repetition rate*, *Mode*

## **CWEPR**

ID, Frequenz (XQW), Attenuation

## **UVVIS**

ID, Dimension of the cuvette

## **Fluorescence**

ID, Excitation or emission?, Wavelength, OD (Absorbance)

## **FS-TA**

ID, Excitation wavelength, Excitation energy

## **NS-TA**

ID, Excitation wavelength, Excitation energy

## **NMR**

ID, Frequency

## **QY**

ID, Excitation wavelength

## **SPT**

ID, Excitation wavelength, Filter

## **PULSE EPR (TN, PEANUT, PELDOR, saturation recovery...)**

ID, Type (e.g. PEANUT, PELDOR...), Path to DSC-File