



detected *in vivo* together with the newly found hydroxylated metabolic product.

The last work of the first part was focused on identification of *in vitro* phase I/II metabolites of newly developed thiosemicarbazone analog - di(2-pyridyl)ketone-4-methyl-4-cyclohexyl-3-thiosemicarbazone (DpC) using UPLC-QTOF. Initially, the chemical structures of the metabolites were proposed based on the high resolution analysis with accurate mass measurement. These were subsequently confirmed by the fragmentation study. Ten phase I metabolites were detected, among which the products of the thiocarbonyl group oxidation, *N*-demethylation, hydroxylation, cyclohexyl ring opening coupled with water addition as well as the various combinations of these reactions were identified. In the case of phase II conjugates, only two glucuronides were detected, while the sulphates and glutathione conjugates were not found.

The second part of the thesis was intended to investigate the feasibility of the HILIC mode for the simultaneous analysis of dexrazoxane and its polar metabolites (B, C, ADR-925). The particular attention was also pay on description of the retention mechanism under MS compatible chromatographic conditions. The separation of all compounds in one analytical run was achieved only on the zwitterion-based stationary phase. This systematic study of the retention behavior confirmed the participation of mixed mode retention mechanism, among which the partitioning, adsorption and ion-exchange could be involved.