Monitoring of successive phosphorylations of thymidine using free and

immobilized human nucleoside/nucleotide kinases by Flow Injection

Analysis with High-Resolution Mass Spectrometry

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1 In-source fragmentation



Figure S1: Mass spectra of (a) dTTP and (b) dTDP (capillary voltage - 4 kV). In-source fragmentation of phosphate.



2 Mass spectra of dTMP conversion into dTDP

Figure S2: Mass spectra of dTMP conversion into dTDP. Analyses conducted by FIA-HRMS.

3 Calibration curves (FIA-HRMS)

Four solutions of dTMP and dTDP (from 5 to 40 μ M) and five solutions of dTTP (from 5 to 50 μ M) were prepared in ammonium acetate buffer (50 mM ionic strength, pH = 7) containing 200 μ M MgCl₂.



Figure S3: Calibration curves of dTMP (m/z 321.0482, diamonds, y = 73164x + 55415, R² = 0,999), dTDP (m/z 401.0145, squares, y = 46403x + 60855, R² = 0.999) and dTTP (m/z 480.9808, triangles, y = 21986x + 102720, R² = 0.982).



Figure S4: Monitoring of substrate phosphorylation (dT, dTMP and dTTP, 25 µM). Reaction catalyzed by free hTK1 (0.48 g/L), free hTMPK (0.04 g/L) and free hNDPK (0.04 g/L). Analyses conducted by FIA-HRMS.

Figure S4 shows product accumulation (dT, dTMP and dTTP) in function of time. hTMPK and hNDPK catalyzed reactions show fast conversion within 2 min whereras hTK1 catalyzed reaction shows linear trend. Reaction time was set at 30 min to reach plateau conversion for hTMPK and hNDPK.

4 Monitoring of product accumulation