

# Cell-free expression of human blood group A and B glycosyltransferases

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## Cell-Free Protein Synthesis

Our initial NMR studies on human blood group A and B glycosyltransferases (GTA and GTB) have shown that classical assignment procedures using e.g. 3D triple resonance spectra fail. In order to allow for combinatorial labeling approaches we had to establish a cell free expression protocol instead. Fig. 1 summarizes components required for cell-free protein synthesis (T7 RNA Polymerase, S30-extract containing more than 200 proteins and ribosomes for translation, Mg<sup>2+</sup> and K<sup>+</sup> ions, pVEX or pET-vector coding for the target protein, ATP, GTP, UTP, CTP, amino acids, tRNA, Creatine kinase/phosphocreatine for energy recycling, PEG 8000 to mimic the viscosity of cytosol and more compounds and salts [1, 2].

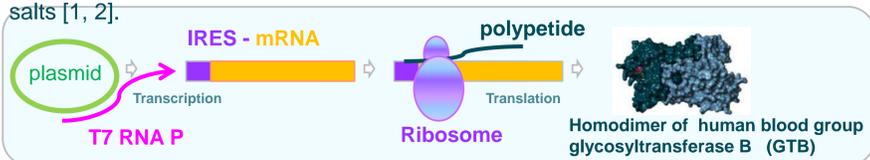


Fig. 1: Coupled system for the cell-free protein synthesis with transcription and translation

## T7 RNA Polymerase

T7 RNA P was expressed in 1 L M9-minimal medium [1] using the vector pAR1219 in BL21 (DE3). T7 RNA P has to be protected from degradation by proteases during purification. Long time storage is achieved with 100 mM NaCl and 50% glycerol at -80° C. The activity of the T7 RNA P depends strongly on Mg<sup>2+</sup>. 100 mM NaCl is an inhibitor of T7 RNA P and NaCl also inhibits the cell-free protein synthesis at >50 mM. We purified 32 mg (7 mg/ml) protein with a purity of T7 RNA P of 90%.

## S30 - Extract

An A19 strain (RNA and amino acid stabilizing mutations) was used to prepare S30-extract [1, 2] with two different media in a 10 L fermenter: 2xYTPG medium [2] and incomplete rich medium [1], a minimal medium supplemented with 0.1% NH<sub>4</sub>Cl and 4 mM MgSO<sub>4</sub>. Only 50% of our S30-extracts prepared in flask cultures were active. Only with a fermenter culture we got enough S30-extract for testing. Essential for an active S30-extract are: a short lag-phase and generation time and fast cooling before harvesting the cell pellet.

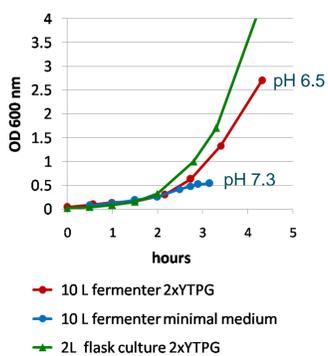


Fig. 2: Growth curve A19 in fermenter and flask culture

	10 L fermenter		1 L flask culture	
	2xYTPG	Minimal medium	2xYTPG	Minimal medium
Medium	2xYTPG	Minimal medium	2xYTPG	Minimal medium
Harvest OD <sub>600</sub>	3	0.54	1.2	0.6
pellet	66 g	13 g	2.5 g	0.6 g
S30-extract	26 ml	3.1 ml	0.7 ml	1 ml
Mg <sup>2+</sup> f.c.	10.3 mM	10.3 mM	15 mM	16 mM

	10 L fermenter		1 L flask culture	
	2xYTPG	Minimal medium	2xYTPG	Minimal medium
Quality control (25-40 mg/ml)	2xYTPG	Minimal medium	2xYTPG	Minimal medium
Protein Bradford/BSA	19 mg/ml	23 mg/ml	(250 mg/ml)	16 mg/ml

Tab. 1: Quality data S30-extracts  
The generation time of all cultures is 30-40 min. Important is a high protein concentration >25 mg/ml.

The batch-to-batch activity of S30-extract preparations also depends on the following steps: 1) Fast cell lysis; 2) Run-off to release endogenous mRNA from the ribosomes (45° C for 45 min., no preincubation mix, 400 mM NaCl); 3) Separation of the 30S and 50S subunits from the 70S ribosomes (not active in cell-free protein synthesis) by high speed centrifugation steps; 4) Cytosol matrix is exchanged against S30-buffer by dialysis (50 mM Tris/acetate, pH 8.2 (RT), 14 mM Mg(OAc)<sub>2</sub>, 60 mM KOAc, 0.5 mM DTT); 5) Endogenous amino acids have to be removed completely for stable isotope labeling of proteins for NMR measurements. Several dialysis steps with buffer exchange are sufficient.

After a last centrifugation step at 30.000xg the S30-extract is shock frozen in liquid nitrogen and stored at -80° C for max. one year.

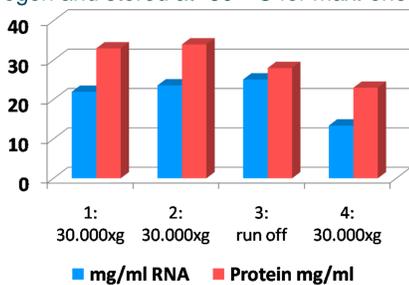


Fig. 3: Protein concentration during S30-extract preparation

From cell lysis to final S30-extract only 66 % of the total protein concentration has been retained. In any case, it is important to start with high protein concentration in the lysis.

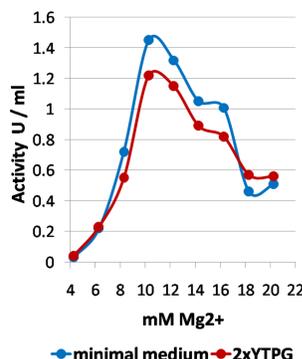


Fig. 4: Mg-Titration of S30-extract  
Usually the Mg-concentration is >15 mM. Probably this reflects the low protein concentration in both S30-extracts.

## Results: 50 µl Batch Reaction

Cell-free protein synthesis in 50 µl batch reactions: 30% S30-extract, 120 µg/ml T7 RNA P, 27 mM NH<sub>4</sub>OAc, 10.3 mM Mg(OAc)<sub>2</sub>, 100 mM KOAc, 100 mM Hepes/KOH pH 7.5, 250 µg/ml creatine kinase, 52 mM phosphocreatine di-Tris salt, 2 mM amino acids, 2 mM (Arg, Cys, Trp, Met, Asp, Glu)-mix, 2% PEG 8000, 1.2 mM ATP, 0.8 mM each CTP, GTP, UTP, 640 µM cAMP, 68 µM folic acid, 180 µg/ml tRNA, 1.7 mM DTT, 20 µg/ml plasmid, 0.1 U/ml RNasin, 1x protease inhibitor. The reaction was performed in 200 µl tubes for 20 hours at 30° C and 600 rpm (highest activity of GTB after 20 hrs) [1, 2].

	10 L Fermenter		2L flask culture	
	2xYTPG	Minimal medium	2xYTPG	Minimal medium
Medium	2xYTPG	Minimal medium	2xYTPG	Minimal medium
2 mM amino acids	1.2 U/ml	1.4 U/ml	0.2 U/ml	0.5 U/ml
1300 x10 <sup>3</sup> DPM	1300 x10 <sup>3</sup> DPM	1500 x10 <sup>3</sup> DPM	233 x10 <sup>3</sup> DPM	567 x10 <sup>3</sup> DPM
0 mM	48 DPM	65 DPM	138 x10 <sup>3</sup> DPM	150 DPM

Tab. 2: Activity of GTB with S30-extracts from fermenter and flask cultures

Intensive dialysis seems to be sufficient to remove endogenous amino acids from S30-extracts. An ultimate opinion is only possible after the first labeling and NMR experiments.

	Vector 1 pVEX-GTB	Vector 2 pIVEV-His <sub>6</sub> -TEV-GTB
Activity assay	1 U/ml	0.3 U/ml
Insoluble GTB	40%	70%

Tab. 3: Vectors for GTB cell-free synthesis

The GTB activity with vector 2 was measured after TEV cleavage.

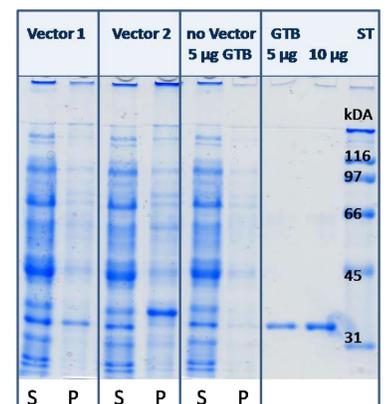


Fig. 5: SDS-PAGE of GTB batch reaction mix

The vectors are cause of different amounts of soluble S and insoluble P GTB. Reaction mix without plasmid and addition of GTB (20 hrs incubation). Reference: GTB

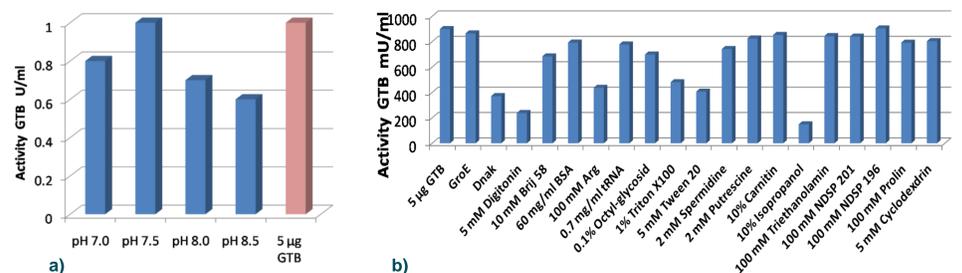


Fig. 6: Attempts to increase the amount of soluble GTB with chemical chaperons  
It seems GTB has folding problems in the cell-free expression, especially with vector 2. a) Purified GTB is stable during cell-free reaction and also the pH value of 7.5 of the reaction buffer Hepes/KOH is ideal. b) Only DnaK causes no insoluble GTB but low yield.

## Results: 50 µl Continuous Exchange Reaction

The advantage of continuous exchange cell-free protein synthesis is continuous supply of new reagents, especially ATP, and removal of side products like phosphate, which inhibit protein synthesis. Each reaction mix contains S30-extract, T7 RNA P, plasmid, tRNA and is separated from the feeding mix via a dialysis membrane (MWCO 10 kDa). The ratio of reaction mix to feeding mix is about 1:18.

	Device 1	Device 2	Device 3	Device 4	Batch
Feeding mix	950 µl	750 µl	750 µl	950 µl	-
Reaction mix	50 µl	50 µl	50 µl	50 µl	50 µl
End volume	25 µl	40 µl	45 µl	10 µl	50 µl
GTB Activity	3.9 U/ml	3.68 U/ml	1.73 U/ml	3 U/ml	0.8 U/ml

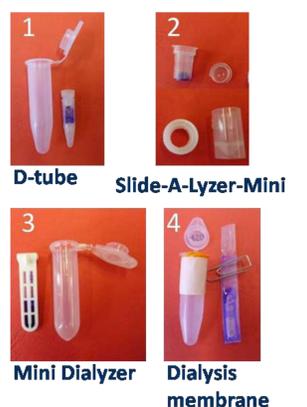


Fig. 7: Comparison of the devices with vector 1

The self made device 2 has the highest activity of GTB with a good recovery of the reaction mix. Maybe device 4 reflects only the concentration of the reaction mixture in the membrane?

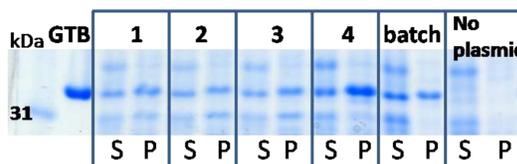


Fig. 8: SDS-PAGE of soluble and insoluble GTB in dependence of the device

Acknowledgments: We thank Monica Palcic for help and support in any respect and especially for highly stimulating discussions and many joint seminars. We thank Frank Bernhardt for many discussions. The DFG is thanked for financial support.