REMOVAL OF INDOXYL SULPHATE AND P-CRESOL SULPHATE AS WELL AS STIMULATION OF FORMING PROTEOLYTIC ENZYMES DURING LOW-FLUX (HDLF), HIGH-FLUX HEMODIALYSIS (HDHF) AND HEMODIAFILTRATION (HDF)

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INTRODUCTION AND AIMS:

In hemodialysis patients proteolytic enzymes and protein-bound solutes (PBS) lead to the activation of oxidative stress and inflammation causing an accelerated development of atherosclerosis. **Aims** 1. Comparison of proteolytic enzymes activity (cathepsin B, collagenase) and PBS concentrations (indoxyl sulphate and p-cresol sulphate) before and after single HDLF, HDHF and HDF procedures 2. The assessment of influence of 8-week therapy by these methods on above-mentioned markers

METHODS

METHODS: 21 patients (8F, 13M) were included and 19 completed the study. The age of the patients was 54 years \pm 14,5 (29,7-71,8), BMI- 28 \pm 6,6. The mean period of hemodialysotherapy was 20,7 \pm 15,5 (3,7 - 61) months. The study lasted for 24 weeks and consisted of 3 consecutive stages (S). Each stage lasted for 8 weeks: stage 1 (S1)-HD (LOPS low-flux Braun dialyzers), stage 2 (S2)-HD (HIPS high-flux Braun dialyzers), stage 3 (S3)- HDF with post-dilution (HIPS Braun filter). The activities of collagenase, cathepsin B (Bachem) were marked by fluorimetric method and the concentrations of indoxil sulphate and p-cresol sulphate) using HPLC method (Merck).

RESULTS

The results of the study are presented in tables 1 and 2. Lower median p-cresol values and stable values of indoxin sulphate after 8-week hemodiafiltration were observed (consecutive values for p-cresol sulphate: before the first HDLF - 45,0, after HDLF cycle before the first HDHF 40,7, after HDHF cycle before the first HDF -36,2, after HDF cycle-27,1).

			doxin sulphat	e	p-cresol sulphate				
		Ν	mean +/- SD	median [min max.]	Ν	mean +/- SD	median [min max.]		
HDLF	1	18	25,1 +/- 11,2	23,4 [2,3-45]	18	45,3 +/- 19,1	43,6 [21,2-92,9]		
	2	18	19 +/- 8,6	18,6 [1,9-34,3]	18	36,8 +/- 14,5	34,6 [16,7-63,4]		
	Δ	18	-6,1 +/- 4,5	-6 [(-13,4)-4,8]	18	-8,5 +/- 7,8	-7,4 [(-29,5)-5,4]		
	p	<0	,001		<0,001				
HDHF	1	18	23,9 +/- 12	20,2 [1,5-49,7]	18	41,9 +/- 21,5	39,9 [14,3-93,8]		
	2	18	17,3 +/- 8,6	16,8 [1,6-33,5]	18	33,4 +/- 15,5	31,4 [9,9-60,9]		
	Δ	18	-6,6 +/- 4,8	-5,6 [(-16,2)-0,1]	18	-8,6 +/- 10,4	-4,8 [(-41,8)-1,9]		
	p	<0,001				<0,001			
HDF	1	18	24,7 +/- 10,3	25,1 [2,6-42,2]	18	38,5 +/- 22	39,1 [0,1-85]		
	2	18	16,2 +/- 7,9	13,9 [2,1-31]	18	28,7 +/- 19,2	26,3 [0,3-73,6]		
	Δ	18	-8,5 +/- 4,5	-8,2 [(-16,7)-(-0,5)]	18	-9,7 +/- 6,1	-8,8 [(-26,6)-0,2]		
	p	<0	,001		<0,001				
р	HDLF vs HDHF	0,89				0,899			
	HDLF vs HDF	0,1	05		0,932				
	HDHF vs HDF	0,072				0,132			

Table 1. Changes of PBS concentrations after different dialysis types

Table 2. Changes of proteolytic enzymes (cathepsin B, collagenase) activity

			Collagenase			Cathepsin B		
		Ν	mean +/- SD	median [minmax.]	Ν	mean +/- SD	median [minmax.]	
HDLF	1	19	10,2 +/- 6,7	8,5 [5,1-32,4]	19	14,8 +/- 3,7	13,9 [9,9-24,5]	
	2	19	10,5 +/- 7,3	9,1 [3-38]	19	14,7 +/- 3,6	14,1 [10-23,3]	
	Δ	19	0,4 +/- 3,8	0,8 [(-13,8)-5,6]	19	-0,1 +/- 1,6	-0,2 [(-2,3)-4,8]	
	p	0,0	0,041			0,496		
HDHF	1	19	9,6 +/- 6,4	7,7 [4,8-30,4]	19	15,1 +/- 3,6	15 [9,8-21,8]	
	2	19	10,1 +/- 6,9	8,8 [3,8-35,2]	19	13,1 +/- 3,5	13 [4,7-21,5]	
	Δ	19	0,5 +/- 4,2	0,6 [(-15,1)-4,8]	19	-2,1 +/- 3,1	-2 [(-9,7)-3,6]	
	p	0,0	0,053			0,006		
HDF	1	19	9,6 +/- 7,4	7,7 [4,7-38,3]	19	14,7 +/- 3,6	14,2 [8,6-22,8]	
	2	19	10,7 +/- 8	9,3 [5,1-41,8]	19	13,8 +/- 4,8	14,1 [0,1-23,9]	
	Δ	19	1,1 +/- 1,8	1,2 [(-2,9)-3,5]	19	-0,9 +/- 3,2	-0,3 [(-10,7)-3,2]	
	p	0,0	0,013			0,307		
	HDLF vs HDHF	1			0,012			
p	HDLF vs HDF	0,716			0,153			
	HDHF vs HDF	0,716			0,179			

CONCLUSIONS

- 1. Single HDLF, HDHF and HDF procedures are similarly effective in removing PBS.
- 2. A tendency for more effective elimination of p-cresol sulphate during 8-week HDHF therapy was observed.
- 3. A similar stimulation of collagenase formation during all 3 types of therapies and decreased concentrations of cathepsin B after HDF (lower stimulation or more effective elimination ?) were found.

