Pregnancy-associated plasma protein A (PAPP-A) as a mortality predictor of long-term hemodialysis patients



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Patients with chronic kidney disease

- High mortality rate mainly due to cardiovascular complications
- They differ from the general population
- Classical risk factors as well as non-traditional/ uremia-related ones are involved

Basic laboratory parameters

| Parameter | HD patients | Controls | p HD vs. controls |
|----------------------------------|-------------|-----------|-------------------|
| Hemoglobin (g/L) | 106±13.2 | 141±10.1 | <0.001 |
| Creatinine (µmol/L) | 753±198 | 88±13 | <0.001 |
| Albumin (g/L) | 37.8±3.8 | 44.4±2.6 | <0.001 |
| CRP (mg/L) | 10.0±16.5 | 3.3±2.4 | 0.002 |
| Leukocytes (x10 ⁹ /L) | 6.92±1.95 | 6.41±1.70 | 0.1 |
| BMI (kg/m ²) | 25.4±4.52 | 25.5±3.42 | 0.9 |
| | | | |

Results: PAPP-A

- 27.6±15.5 mIU/L in HD vs. 9.4±2.5 mIU/L in controls, p<0.001
- Significant independent predictor
 - for overall mortality
 - HR/SD (95%CI) 1.237 (1.060-1.444), p=0.007
 - for mortality due to infection

PAPP-A pregnancy-associated plasma protein A

- metalloproteinase, cleaves IGFBP-4 → IGF-1 increase
- screening of Down syndrome in the 1st trimester
- present in ruptured atherosclerotic plaques
- biomarker of acute coronary syndrome
- increased in HD patients, related to renal function

Aim of the study

PAPP-A and related parameters

- other pregnancy protein placental growth factor -PIGF
- matrix metalloproteinases MMP-2 and MMP-9
- molecules linked to PAPP-A action IGFBP-4 and IGF-1
- established cardiac markers cTnl, BNP
- inflammatory markers CRP; retinol

Follow up of HD patients - 5 years

- + 146 patients (56%)
 - 71 cardiovascular cause
 - 42 infection
 - 14 tumour
 - 15 other cause
- 52 patients transplantation, 8 of them +
- 2 patients censored to due other reason

Laboratory methods

- PAPP-A TRACE (KRYPTOR, Brahms)
- PIGF, IGFBP-4, MMP-2 and MMP-9 ELISA (RD) Systems)
- IGF-1 IRMA
- **BNP and cTnl**-CLIA
- Retinol HPLC
- Basic nutritional and inflammatory parameters standard methods, automated analyzers

HR/SD (95%CI) 1.416 (1.115-1.798), p=0.004 - not for cardiovascular mortality

Overall mortality





→ relationship of their serum levels to prognosis of long term hemodialysis patients in 5-years followup

Study design: **Prospective observational cohort study.**

261 long-term hemodialysis patients

- follow-up for 5 years (11/2003-11/2008)
- patients from 6 HD centres in the Czech Republic
- 141 men and 120 women, mean age 64±13 years
- clinical and laboratory characteristics collected at the beginning of the study

66 healthy controls

25 men and 41 women, mean age 59±9 years

Clinical characteristics of hemodialysis patients

duration of HD treatment

median 2 years

Statistical analysis

- software SPSS v.16.0
- Survival analysis
 - Kaplan-Meier analysis
 - Cox regression univariate and multivariate analysis (forward and backward methods)
- overall mortality, cardiovascular mortality, mortality due to infection
- transplantation taken as time dependent covariate
- BCH parameters treated as continuous variables
- HR (95%CI) expressed per SD, for age per year

Other markers and mortality all increased in HD except for MMP-9

- PIGF n.s. (p=0.08-0.1)
- MMP-2 and MMP-9 n.s.
- IGFBP-4 n.s.
- IGF-1 significant in uni-variate analysis
- **cTnl** significant in both uni-variate and multi-variate analysis for overall and cardiovascular mortality
- **BNP** significant only in uni-variate analysis for overall and cardiovascular mortality
- Retinol significant in both uni-variate and multivariate analysis for overall and cardiovascular mortality (Kalousová et al. Am J Kidney Dis 2010)

Significant Independent Mortality **Predictors for Overall, Cardiovascular and** due to Infection Mortality

| | Overall mortality HR (95% CI) p | | Cardiovascular mortality HR (95% CI) p | | Mortality due to infection HR (95% CI) p | |
|----------------------|---------------------------------------|--------|--|------------|--|------------|
| PAPP-A | 1.237 (1.060-1.444) | 0.007 | | | 1.416 (1.115-1.798) | 0.004) |
| cTnl | 1.411 (1.200-1.658) | <0.001 | | <0.001 | | |
| Albumin | 0.722 (0.595-0.876) | <0.001 | 0.588 (0.447-0.774) | <0.001 | | |
| Creatinine | 0.789 (0.637-0.978) | 0.03 | | | 0.672 (0.487-0.926 | 0.02) |
| Retinol | 0.775 (0.632-0.950) | 0.01 | 0.671 (0.500-0.902 | 0.008) | | |
| Parathromone | | | | | 0.621 (0.364-1.058 | 0.08) |
| Age | 1.030 (1.011-1.048) | 0.001 | | | | |
| Diabetes mellitus | 1.625 (1.128-2.340) | 0.009 | | | 2.966 (1.474-5.966 | 0.002) |
| CVD | 1.638 (1.039-2.582) | 0.03 | 3.753 (1.699-8.289) | 0.001 | | |
| Transplantation | Transplantation | | 0.121 (0.016-0.893) | 0.04 | | |

Conclusion

This study demonstrates PAPP-A as an independent predictor of overall mortality and mortality due to infection in hemodialysis patients. Our results suggest superior relationship of PAPP-A to infection-inflammation than to cardiovascular risk in HD patients.

| duration of HD treatment | median 2 ye |
|---|-------------|
| diabetes mellitus | 33% |
| dyslipidemia | 41% |
| hypertension | 84% |
| cardiovascular disease | 61% |
| cerebrovascular disease | 24% |
| peripheral vascular disease | 25% |

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