

評価方法はまだ不足しているが、⑤、⑥で示した方法は、今後、他の合併症発症率、あるいは、保存期 CKD 患者での合併症発症率を検討する上で応用できると考えられた。

以上の結果をふまえて、今後我々は本研究事業で下記の 3 つの研究を行う (図 3)。以下に説明する。

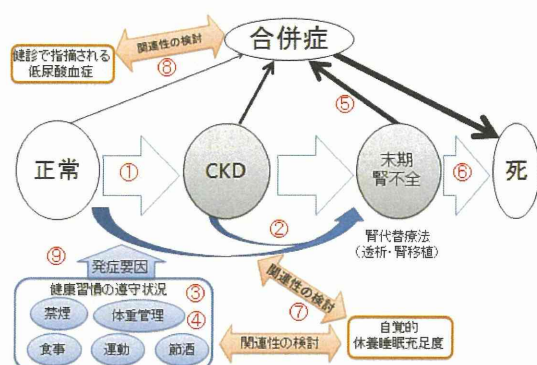


図 3. 今後行う予定の研究

⑦自覚的休養睡眠充足度と生活習慣・CKD との関係

睡眠障害の改善は重要視されている。事実、21 世紀における第二次国民健康づくり運動 (健康日本 21 (第二次)) の目標の一つに、睡眠による休養を十分とれていない者の割合の減少があげられている。しかし、具体的に何をどうすれば、睡眠による休養を十分とれていない者の割合を減らせるのかは明らかではない。睡眠と生活習慣病には相互に密接な関連があることから、睡眠障害の改善は生活習慣病対策にも重要である可能性が考えられる。そこで、特定健診の標準的な問診票で得られる自覚的睡眠休養充足度がどのような生活習慣と関連するかを明らかにする。これが明らかになれば、特定健診での保健指導時に、生活習慣のみならず睡眠障害にもあわせて介入することが可能になり、さらに睡眠障害の改善が健診受診者の生活習慣の行動変容へのモチベーションに繋

がる可能性が考えられる。

⑧低尿酸血症の疫学

健診時に発見された低尿酸血症に対してどのような対応をすべきかについての明確な答えはない。尿酸は活性酸素のスカベンジャーであり、生体内の全抗酸化作用の 3 分の 2 を担うため、低尿酸血症患者で動脈硬化性疾患が多い、あるいは生命予後が不良である可能性があるが、それを支持する疫学データはない。そもそも低尿酸血症は頻度が低く、疫学データが不足しているのが現状である。本研究事業で作成される全国コホート群のような大規模の調査で検討する必要がある。

⑨ 5 つの健康習慣と CKD の詳細な検討

③で行った検討に加えて、より多数例で、より長期の観察期間で、詳細な検討を行う。さらに保健指導等による 5 つの健康習慣の改善による CKD 新規発症への効果もあわせて検討を行う予定である。

これらの研究により、より包括的な CKD 評価が可能になり、ひいては地域の実情に即した効果的な CKD 地域医療連携システムの制度設計に繋がると考える。

E. 結論

現時点で可能な、地域における CKD の包括的評価方法をまとめた。健診時の血清クレアチニン測定は、個人の腎機能評価のみならず、集団としての評価も可能にする。集団としての評価は、地域の実情に即した効果的な CKD 地域医療連携システムの制度設計を行う上で、極めて有用な情報になると考える。より包括的な評価を行うために、本研究事業でさらなる臨床疫学研究を開始した。

G. 研究発表

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H. 知的財産権の出願・登録状況

1. 特許取得 なし。
2. 実用新案登録 なし。
3. その他 なし。

分担研究報告書

「自治体の特定健診データからみた CKD の実態調査
～血清クレアチニンを測定しない場合の CKD 見逃し率の推定等～」

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研究要旨

特定健診は CKD を早期発見できる絶好の機会であるが血清クレアチニンが必須項目となっていないため、CKD を見逃す可能性が指摘されている。本研究では血清クレアチニンを自主的に測定している 20 の自治体の 578,965 人の特定健診のデータを用いて検証した。対象住民における CKD の頻度は、18.6%であった。eGFR < 60 ml/min/1.73m² の 81,572 人のうち尿蛋白陰性者は 72,956 人 (89.4%)であった。eGFR < 60 ml/min/1.73m² および/または尿蛋白 1+以上で定義された CKD 102,061 人の住民のうち、尿蛋白陰性者は 71.5%であった。すなわち、特定健診では血清クレアチニンを測定しなければ、CKD の 70~90%を見逃す可能性があった。特定健診では、血清クレアチニンの測定を必須化することが必要である。

A. 研究目的

特定健診は CKD を早期発見できる絶好の機会であるが血清クレアチニンが必須項目となっていない。そのため、CKD を見逃す可能性が指摘されている。本研究では血清クレアチニンを自主的に測定している自治体の特定健診のデータを用いて、もし、血清クレアチニンが測定されなかったらどの程度 CKD を見逃す可能性があるかを検証することを目的とした。

B. 研究方法

対象者: 表 1 に示す 20 府県の 2008 年の健診受診者のうち血清クレアチニンが自主的に測定されていた 578,965 人のデータを対象とした。男性 42%、年齢 61 ± 10 (平均 ± SD) 歳であった。

解析

SPSS version 17.0 (IBM, Chicago, IL)を用いた。

表 1. 解析に用いた住民

	度数	パーセント	有効パーセント	累積パーセント
有効				
茨城県	15708	2.7	2.7	2.7
沖縄県	137996	23.8	23.8	26.5
宮崎県	46234	8.0	8.0	34.5
宮城県	15772	2.7	2.7	37.3
熊本県	10913	1.9	1.9	39.1
高知県	29	.0	.0	39.1
佐賀県	2834	.5	.5	39.6
埼玉県	3433	.6	.6	40.2
新潟県	35579	6.1	6.1	46.4
神奈川県	50054	8.6	8.6	55.0
石川県	6005	1.0	1.0	56.1
大阪府	18707	3.2	3.2	59.3
長崎県	6708	1.2	1.2	60.4
長野県	11718	2.0	2.0	62.5
東京都	26161	4.5	4.5	67.0
徳島県	4328	.7	.7	67.7
栃木県	5838	1.0	1.0	68.7
福岡県	145426	25.1	25.1	93.9
福島県	9221	1.6	1.6	95.5
北海道	26301	4.5	4.5	100.0
合計	578965	100.0	100.0	

(倫理面への配慮)

匿名化された健診データを用いる後ろ向き解析であるため、倫理的な問題は生じない。個人情報情報は取り扱わない。

C. 研究結果

1. CKD の頻度

eGFR<60 ml/min/1.73m²の頻度は14.6%、尿蛋白1+以上の頻度は5.2%で、どちらかまたは両者を有するCKDの頻度は18.6%であった。

2. eGFR による CKD のステージの頻度

G1 4.1%, G2 15.4%, G3a 71.2%, G3b 7.7%, G4 1.1%, G5 0.5% で、ステージ別の頻度には男女で大きな差異は見られなかった。

3. 血清クレアチンを測定しない場合の CKD の見逃し率 (表 2, 表 3, 表 4, 表 5)

血清クレアチンと尿蛋白を測定した住民 560,758 人のうち eGFR < 60 ml/min/1.73m² の CKD は 81,572 人であった。そのうち尿蛋白陰性者は 72,956 人 (89.4%) であった。すなわち、血清クレアチンを測定しなければ、eGFR < 60 ml/min/1.73m² の住民の 89.4% が CKD と認識さ

れないという結果であった。また、尿蛋白陰性群での CKD の頻度は 13.7% であった。

表 2. eGFR(1: eGR < 60 ml/min/1.73m²) と尿蛋白(0: 陰性)のクロス表

		UP01		合計	
		0	1		
GFR01	0	度数	458697	20489	479186
		GFR01 の %	95.7%	4.3%	100.0%
		UP01 の %	86.3%	70.4%	85.5%
1		度数	72956	8616	81572
		GFR01 の %	89.4%	10.6%	100.0%
		UP01 の %	13.7%	29.6%	14.5%
合計		度数	531653	29105	560758
		GFR01 の %	94.8%	5.2%	100.0%
		UP01 の %	100.0%	100.0%	100.0%

eGFR < 60 ml/min/1.73m² and/or 尿蛋白 1+ 以上で定義された CKD 102,061 人の住民のうち、尿蛋白を有するのはわずか 28.5% であった。すなわち、尿蛋白のみの測定では CKD の 71.5% を見逃す可能性があることが示された。eGFR によるステージ別の見逃し率は、G3a 91.8%, G3b 76.9%, G4 41.6%, G5 42.6% で、特に G3 で見逃し率が高いことが分かった (CKD の定義から G1 と G2 における CKD では尿蛋白は 100% 陽性である)。

表 3. CKD (1: eGR < 60 ml/min/1.73m² and/or 尿蛋白 1+以上) と尿蛋白(0:陰性)のクロス表

		UP01		合計	
		0	1		
GFRgrade	G1	度数	0	4355	4355
		%	.0%	100.0%	100.0%
	G2	度数	0	16134	16134
		%	.0%	100.0%	100.0%
	G3a	度数	66276	5917	72193
		%	91.8%	8.2%	100.0%
	G3b	度数	6041	1812	7853
		%	76.9%	23.1%	100.0%
	G4	度数	450	632	1082
		%	41.6%	58.4%	100.0%
	G5	度数	189	255	444
		%	42.6%	57.4%	100.0%
合計		度数	72956	29105	102061
		%	71.5%	28.5%	100.0%

年齢別にみると、年齢が上がるにつれて、CKDが増加するとともに、尿蛋白陰性のCKDの割合も増加することも示された。すなわち、CKDのうち、尿蛋白陰性の割合は40歳台では54.1%、50歳台では65.5%、60歳台では73.4%、70歳以上では74.2%であった。

表4. 全体での年代別CKDの割合

		CKD01		合計
		0	1	
年齢10歳刻み	40歳未満	度数 11145	2114	13259
		% 84.1%	15.9%	100.0%
40歳代	度数	53395	6608	60003
	%	89.0%	11.0%	100.0%
50歳代	度数	85605	13922	99527
	%	86.0%	14.0%	100.0%
60歳代	度数	198978	45513	244491
	%	81.4%	18.6%	100.0%
70歳以上	度数	109578	36786	146364
	%	74.9%	25.1%	100.0%
合計	度数	458701	104943	563644
	%	81.4%	18.6%	100.0%

表5. CKD群での年代別尿蛋白の有無

		UP01		合計
		0	1	
年齢10歳刻み	40歳未満	度数 1110	549	1659
		% 66.9%	33.1%	100.0%
40歳代	度数	3089	2624	5713
	%	54.1%	45.9%	100.0%
50歳代	度数	8423	4438	12861
	%	65.5%	34.5%	100.0%
60歳代	度数	33118	12032	45150
	%	73.4%	26.6%	100.0%
70歳以上	度数	27216	9462	36678
	%	74.2%	25.8%	100.0%
合計	度数	72956	29105	102061
	%	71.5%	28.5%	100.0%

4. リスク因子の有無による解析

(1) 高血圧

高血圧の有無とCKDの有無が同時に判断出来た住民492,118人のうち、高血圧患者は229,751人、46.7%であった。高血圧患者でのCKDは55,086人、24%と、この住民群におけるCKD95,063人、19.3%に比して高頻度であった。CKDのうち尿蛋白陰性はこの住民群では71.5%であったが、高血圧患者では66.8%であった。尿蛋白の陽性率はこの住民全体では

5.2%であったが、高血圧を有する住民では7.8%と高かった。

(2) 糖尿病

糖尿病の有無とCKDの有無が同時に判断出来た住民319,661人のうち糖尿病患者は54,790人、17.1%であった。糖尿病患者でのCKDは14,377人、26.2%、この住民におけるCKD61,116人、19.1%に比して高頻度であった。CKDのうち尿蛋白陰性はこの住民では71.5%、糖尿病患者では49.9%であった。尿蛋白の陽性率は全住民では5.2%であったが、糖尿病を有する住民では13.0%と高かった。

(3) 脂質異常症

脂質異常症の有無とCKDの有無が同時に判断出来た住民513,556人のうち脂質異常症患者294,266人、57.3%であった。脂質異常症でのCKDは58,953人、20.0%で、この住民でのCKD93,083人、18.1%に比して高頻度であった。この住民群におけるCKDのうち71.5%は尿蛋白陰性であったが、脂質異常症患者では70.2%であった。CKDのうち尿蛋白陰性はこの住民では71.5%、脂質異常症では70.2%であった。尿蛋白の陽性率は全住民では5.2%であったが、脂質異常症を有する住民では6.0%とやや高かった。

(4) 肥満

BMIが増えるとCKDの頻度が上昇することが示された(表5)。

表5. BMI と CKD のクロス表

		CKD01		合計
		0	1	
18.5未満 やせ	度数	27238	4573	31811
	%	85.6%	14.4%	100.0%
18.5~25.0 普通	度数	312684	65581	378265
	%	82.7%	17.3%	100.0%
25.0~35.0 軽度肥満	度数	102971	29543	132514
	%	77.7%	22.3%	100.0%
35.0~40.0 中等度肥満	度数	12622	4151	16773
	%	75.3%	24.7%	100.0%
35.0~40.0 高度肥満	度数	1437	513	1950
	%	73.7%	26.3%	100.0%
40.0以上 超肥満	度数	258	113	371
	%	69.5%	30.5%	100.0%
合計	度数	457210	104474	561684
	%	81.4%	18.6%	100.0%

また、今回の横断データの解析では、高齢になるにしたがい尿蛋白陰性の CKD が増加していることも示された。いわゆる腎硬化症の頻度が増加していることを反映している可能性がある。腎硬化症は末期腎不全で透析導入になる原因疾患として増加しつつあることを考えると、高齢化社会を迎えて重要な知見と思われる。また、高血圧、糖尿病、脂質異常症、肥満では CKD の頻度が高くなることも示された。

D. 考察

今回の約 58 万人の特定健診データからは、血清クレアチンを測定しない場合の CKD の見逃しが、CKD 全体の 71.5%にもなることが示されたことは意義が大きい。特に、eGFR < 60 ml/min/1.73m²の CKD に限ると実に 89.4%が見逃されることになる。

特定健診は CKD のスクリーニングとして計画されているわけではないが、CKD を早期に見つけて対策をたてることのできる絶好のチャンスである。CKD は末期腎不全のみならず心血管疾患の高危険群であるから、特定健診を活かして CKD 対策を立てることは国民の健康を維持するためには喫緊の課題である。そのためには尿蛋白のみでは全く不十分であり、血清クレアチンを測定することが必須であることが示された。

E. 結論

特定健診では血清クレアチンを測定しなければ、CKD の 70~90%を見逃す可能性が示された。特定健診で血清クレアチニンの測定を必須化することが必要である。

G. 研究発表

1. 論文発表 なし。
2. 学会発表 なし。

H. 知的財産権の出願・登録状況

1. 特許取得 なし。
2. 実用新案登録 なし。
3. その他 なし。

「特定健康診査による慢性腎臓病早期発見早期治療の財源影響に関する研究」

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研究要旨

特定健康診査による個人リスク評価に基づく、保健指導と連結した効果的な慢性腎臓病（CKD）地域連携システムの制度設計の、医療経済面の基礎研究として、特定健康診査による腎機能検査項目に着目して CKD 早期発見早期治療の財源影響に関する研究を行った。現行の尿蛋白のみを必須項目としている政策では医療費の削減が生じていることが明らかになった。

A. 研究目的

特定健康診査による個人リスク評価に基づく、保健指導と連結した効果的な慢性腎臓病（CKD）地域連携システムの制度設計の医療経済面の基礎研究として、特定健康診査による CKD 早期発見早期治療の医療財源への影響を明らかにすることを目的とした。特に、健康診査項目としての尿蛋白と血清クレアチニンの影響を明らかにすることを目的とした。本研究の結果は、腎臓病対策の観点からの特定健康診査における腎機能検査項目を検討するに際し有用な経済エビデンスのひとつとなる。

B. 研究方法

特定健康診査の腎機能検査項目に関する費用効果分析として発表している経済モデル (Kondo, Yamagata, et al. Clin Exp Nephrol.

2012;16(2):279-91.) を用いて、この経済モデルが 15 年間有効であると仮定して、国立社会保障・人口問題研究所の全国将来推計人口を当てはめて、財源影響分析を行った。分析対象とした財源の範囲は保険者が負担する特定健康診査での腎機能検査費と、慢性腎臓病とその続発症としての慢性腎不全及び心血管疾患にかかる診療報酬である。なお、使用した経済モデルによる費用効果分析の結果としては、尿蛋白のみが必須項目とされているものの約 60% の保険者が血清クレアチニンも健診項目に含めている現状から血清クレアチニンも必須項目化する場合や、尿蛋白のみを必須項目とする現状維持の場合の、いずれも費用対効果に優れる政策であることが明らかにされている。

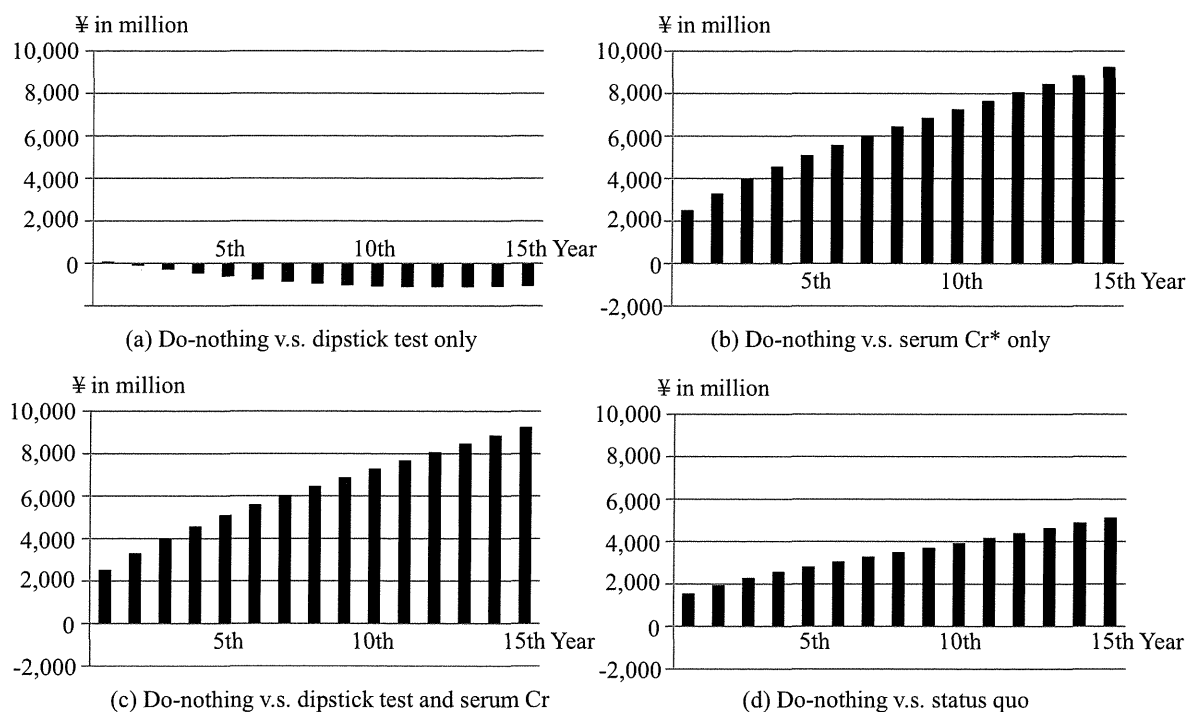
(倫理面への配慮)

経済モデル研究のため倫理面の問題はない。

C. 研究結果

Figure 1 が特定健康診査で腎機能検査を行わない (Do-nothing) 場合と (a) 尿蛋白のみを行う場合、(b) 血清クレアチンのみを行う場合、(c) 尿蛋白と血清クレアチンの両方を行う場合、(d) 現状である尿蛋白を全対象者で行い血清クレアチンを 60%の対象者で行う場合、を比較した財源影響分析の結果である。

(a) では、2 年目から財源影響が負となっており、いわゆる医療費の削減が生じることが明らかになった。9 年目以降では年間 10 億円程度の削減が生じる。しかし、(b)、(c)、(d) では財源影響は正となっており、10 年目にはそれぞれ、72 億円、72 億円、39 億円の医療費の増大が生じる。



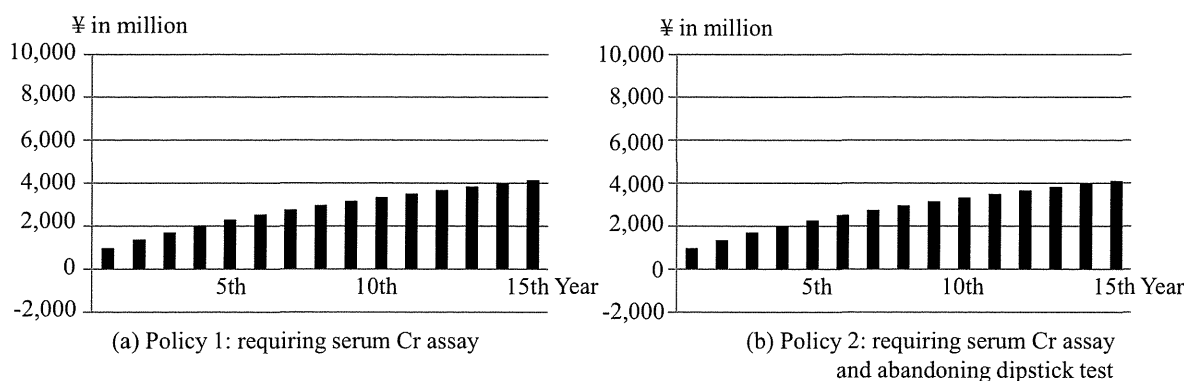
* Cr stands for creatinine

Figure 1 Budget impact model estimators

Figure 2 が特定健康診査で尿蛋白を全対象者で行い血清クレアチンを 60%の対象者で行う現状の場合と、(a) 加えて血清クレアチンを必須項目化する場合、(b) 血清クレアチンのみ必須項目化して尿蛋白を必須項目から除外し全保険者が尿蛋白を行わない場合、を比較した財源影響分析の結果である。(b) で全保

険者が尿蛋白を行わないことを仮定したのは特定健康診査での糖尿病検査で尿糖が必須項目から除外され採血による HbA1c のみが必須項目化される場合を考えてのことである。

(a) と (b) とともに財源影響は正となっており、10 年目にはそれぞれ、33 億円、33 億円の医療費の増大が生じる。



* Cr stands for creatinine

Figure 2 Budget impacts

D. 考察

本研究の結果としてもっとも着目すべきことは、特定腎機能検査を行わない場合と尿蛋白のみを行う場合の比較において、医療費の削減が生じていることが明らかにされたことである。この比較は、現行の尿蛋白のみを必須項目としている政策を反映しているものと捉えることができる。この意味では、現行の政策は医療費適正化へ貢献していると示唆される。

一方で、血清クレアチニンの必須項目化にもなっては、医療費の増大が見込まれた。ただし、この結果に基づいて直ちに特定健康診査で血清クレアチニンを行うべきではないとは言えないことに注意しなければならない。既述のように特定健康診査で血清クレアチニンを行うことについては費用効果分析によって「支払いに見合う価値」があることが示されている。また、本研究班が取り組んでいる保健指導と連結した効果的な慢性腎臓病（CKD）対策のような効果の大きな早期治療法の開発が進めば、血清クレアチニンについても尿蛋白のように財源影響が負に転じていく可能性は小さくない。さらに現在検討が進められている「かかりつけ医/非腎臓専門医と腎臓専門医の協力を促進する

慢性腎臓病患者の重症化予防のための診療システムの有用性を検討する研究」(FROM-J) や腎専門医での大規模な慢性腎臓病の疫学研究である日本CKDコホート研究(CKD-JAC)の結果が明らかとなることにより、CKDの長期的予後改善可能な治療法が確立されれば、健診項目の財源影響にとどまらず、特定健診全体に対する医療経済的に良好な効果が期待される。

本研究の最大の限界は、経済モデルに依存していることである。しかし、健康診断による早期発見早期治療の財源影響を検討するにあたってフィールドで前向き研究を行うことは著しく困難であるうえに、数(十)年後に結果が得られたとしても、時機を得た政策決定に資することは不可能である。こうした観点から経済モデルに依存したアプローチは正当化できるだろう。

E. 結論

特定健康診査による腎機能検査項目に着目してCKD早期発見早期治療の財源影響に関する研究を行ったところ、現行の尿蛋白のみを必須項目としている政策では医療費の削減が生じ

ていることが明らかになった。

本研究は3年計画の1年目であるので次年度以降の研究計画にふれておく。2年目では、保健指導と連結した効果的な慢性腎臓病（CKD）対策の医療経済面を研究する。この研究に関しては、平成24年3月に終了した厚生労働科学研究費補助金（腎疾患対策研究事業）「かかりつけ医/非腎臓専門医と腎臓専門医の協力を促進する慢性腎臓病患者の重症化予防のための診療システムの有用性を検討する研究」（FROM-J）から報告される臨床エビデンスなどを利用する予定である。3年目では、本研究班の先行研究である平成20-22年度厚生労働科学省科学研究費補助金（循環器疾患等生活習慣病総合研究事業）「今後の特定健康診査・保健指導における慢性腎臓病（CKD）の位置付けに関する検討」で構築した全国の約58万人のコホート群の追跡データから報告される臨床エビデンスなどを利用して、地域連携システムの制度設計の医療経済面の研究を行う予定であ

る。

G. 研究発表

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2. 学会発表

なし。

H. 知的財産権の出願・登録状況

1. 特許取得

なし。

2. 実用新案登録

なし。

3. その他

なし。

研究成果の刊行に関する一覧表

研究成果の刊行に関する一覧表

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Association of High Pulse Pressure With Proteinuria in Subjects With Diabetes, Prediabetes, or Normal Glucose Tolerance in a Large Japanese General Population Sample

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OBJECTIVE—To examine whether there is a difference in the association between high pulse pressure and proteinuria, independent of other blood pressure (BP) indices, such as systolic or diastolic BP, among subjects with diabetes, prediabetes, or normal glucose tolerance.

RESEARCH DESIGN AND METHODS—Using a nationwide health checkup database of 228,778 Japanese aged ≥ 20 years (mean 63.2 years; 39.3% men; none had pre-existing cardiovascular disease), we examined the association between high pulse pressure, defined as the highest quintile of pulse pressure (≥ 63 mmHg, $n = 40,511$), and proteinuria ($\geq 1+$ on dipstick, $n = 12,090$) separately in subjects with diabetes ($n = 27,913$), prediabetes ($n = 100,214$), and normal glucose tolerance ($n = 100,651$).

RESULTS—The prevalence of proteinuria was different among subjects with diabetes, prediabetes, and normal glucose tolerance (11.3 vs. 5.0 vs. 3.9%, respectively; $P < 0.001$). In subjects with diabetes, but not those with prediabetes or normal glucose tolerance, high pulse pressure was associated with proteinuria independently of significant covariates, including systolic BP (odds ratio 1.15 [95% CI 1.04–1.28]) or diastolic or mean BP (all $P < 0.01$). In patients with diabetes, a +1 SD increase of pulse pressure (+13 mmHg) was associated with proteinuria, even after adjustment for systolic BP (1.07 [1.00–1.13]) or diastolic or mean BP (all $P < 0.05$).

CONCLUSIONS—Among the Japanese general population, there was a significant difference in the association between high pulse pressure and proteinuria among subjects with diabetes, prediabetes, and normal glucose tolerance. Only in diabetes was high pulse pressure associated with proteinuria independent of systolic, diastolic, or mean BP levels.

In the systemic circulation, the kidney has unique features: vascular resistance in the glomerular afferent arterioles is low, and the myogenic response of the glomerular arterioles is insensitive to changes in the other BP indices of systolic blood pressure (BP), including pulse pressure (1–3). These characteristics suggest that pressure pulsatility may contribute to barotrauma-induced renal microvascular injury, and in turn causes glomerular ultrastructural changes (e.g., podocyte loss and glomerular basement membrane thickness) (1–6).

In fact, several cross-sectional studies performed in general or hypertensive populations have demonstrated a significant association between pulse pressure and albuminuria (7,8), and some longitudinal studies have underscored the importance of pulse pressure as a risk factor for increased albuminuria in general or hypertensive populations (9,10); however, few studies have directly examined the impact of high pulse pressure on albuminuria with adjustment for other BP components, such as systolic BP, diastolic BP, and/or mean BP levels. Since renal autoregulation is particularly impaired in patients with diabetes (1–3,11–13), we hypothesized that the association between high pulse pressure and albuminuria would be more prominent in patients with diabetes than in subjects without diabetes (14–16); as of yet, however, there have been no studies examining this hypothesis directly in a large database. Furthermore, the association of pulse pressure with albuminuria has never been explored in prediabetics, who are classified as being at an intermediate stage between normal glucose tolerance and diabetes (17), but prediabetics have been shown to have a significantly increased risk of developing not only diabetes but also cardiovascular disease (18).

In the current study, therefore, we examined the association of high pulse pressure with proteinuria separately in each of subjects with diabetes, prediabetes, and normal glucose tolerance, using a large nationwide

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Received 18 November 2011 and accepted 14 February 2012.

DOI: 10.2337/dc11-2245

This article contains Supplementary Data online at <http://care.diabetesjournals.org/lookup/suppl/doi:10.2337/dc11-2245/-/DC1>.

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database of subjects recruited from the national health checkup system in Japan.

RESEARCH DESIGN AND METHODS

Study population

This study was performed as a part of the prospective ongoing "Research on the Positioning of Chronic Kidney Disease in Specific Health Check and Guidance in Japan" project. A new annual health check program, "The Specific Health Check and Guidance in Japan", was started by the Japanese government in 2008, targeting early diagnosis and intervention for metabolic syndrome. The target population comprises Japanese citizens between the ages of 40 and 74 years. In Japan, there are 47 administrative divisions (prefectures), and 13 of these prefectures (Yamagata, Miyagi, Fukushima, and Niigata from the Tohoku region in northeastern Japan; Tokyo, Kanagawa, and Ibaraki from the Kanto region in central Japan; Osaka, Okayama, and Kochi from the Kansai, Tyugoku, or Shikoku region in western Japan; and Fukuoka, Miyazaki, and Okinawa from the Kyushu region in southern Japan), which were randomly distributed across Japan, agreed with the aims of this study and performed data collection prospectively from 2008 to 2009. Data were sent to an independent data center, the non-profit organization Japan Clinical Research Support Unit after anonymization in a linkable fashion, and verified by trained staff (K.I. and Y.O.). After that, the database was locked with a security password, which contained the participant's information managed by a research ID number but did not contain the participant's name, and was sent to each investigator on a recordable compact disc.

There were a total of 346,942 subjects (mean age, 63.4 years; 41% [$n = 141,938$] men) for whom information on age, sex, BP, BMI, habitual smoking or drinking, use of antihypertensive drugs, and previous history of cardiovascular diseases (i.e., stroke and cardiac diseases such as angina and myocardial infarction) were available, as well as data on the serum creatinine level and dipstick urine test for proteinuria (19). Some of the regions participating in our project (i.e., Okinawa and Osaka) concomitantly performed regular health checkups for employees as legally mandated in Japan; as a result, the database used in the present analysis also included subjects aged 20–39 years ($n = 2,025$). Among the 346,942 subjects, 29,820 subjects with a previous

history of cardiovascular disease, 243 subjects with chronic kidney disease stage 5 (estimated glomerular filtration rate [eGFR] < 15 mL/min/1.73m²), and 47 subjects with both were excluded from the present analysis. Moreover, 88,101 subjects with insufficient blood sampling data of glucose and lipid parameters were excluded. Supplementary Table 1 shows the differences in clinical characteristics between subjects who were included in the present analysis ($n = 228,778$) and those who had missing data ($n = 88,101$).

The study was conducted according to the guidelines of the Declaration of Helsinki and Ethical Guidelines for Epidemiological Research (1 November 2007, Ministry of Education, Culture, Sports, Science, and Technology and Ministry of Health, Labor, and Welfare of Japan). Ethical approval from the respective institutional review boards was also granted.

Baseline measurement

All subjects completed a self-administered questionnaire to document their medical history, current medications, smoking habits (current smoker or not), and alcohol intake (daily drinker or not). The study physicians performed a physical examination of each subject and rechecked their medical history to improve the precision of the information. Body height and weight were measured in light clothing without shoes, and the BMI was calculated (kg/m²). BP measurement and blood and urine sampling were performed at each local medical institution to cooperate with the nationwide medical checkup. According to the recommendations of the Japanese Ministry of Health, Labor, and Welfare (<http://www.mhlw.go.jp/bunya/shakaihoshou/iryouseido01/info03a.html>), BP was measured by medical staff using a standard sphygmomanometer or an automated device on the right arm after the subject had rested for 5 min in a seated position with the legs not crossed. Conversation as well as alcohol/caffeine consumption was also avoided before measurement. Pulse pressure was calculated as systolic BP – diastolic BP, and mean BP was calculated as diastolic BP + (pulse pressure/3).

Blood samples were collected after an overnight fast and were assayed within 24 h with an automatic clinical chemical analyzer. All measurements were conducted locally rather than at a central laboratory without calibration among different laboratories, despite the fact that beginning several years ago, standardized methods to measure laboratory data were recommended

and widely adopted by the activity of the Japan Society of Clinical Chemistry.

The value for hemoglobin A_{1c} (HbA_{1c}) was estimated as a National Glycohemoglobin Standardization Program equivalent value calculated with the following equation (20): HbA_{1c} (%) = HbA_{1c} (Japan Diabetes Society) (%) + 0.4%.

Diabetes was defined in accordance with American Diabetes Association guidelines (17) as a fasting glucose concentration of 126 mg/dL or higher, HbA_{1c} 6.5% or higher, or self-reported use of antihyperglycemic drugs. Diagnosis of prediabetes was based on the new American Diabetes Association criterion of impaired fasting glucose (fasting plasma glucose 100–125 mg/dL) or HbA_{1c} 5.7–6.4%, or both (17).

Urinalysis by the dipstick method was performed on a single spot urine specimen collected in the early morning after overnight fasting. Urine dipstick results are interpreted by the medical staff in each local medical institution and recorded as –, ±, 1+, 2+, and 3+. In Japan, it is recommended and widely adopted by the activity of the Japanese Committee for Clinical Laboratory Standards (<http://jccls.org/>) that all urine dipstick tests be manufactured so that a urine dipstick result of 1+ will correspond to a urinary protein level of 30 mg/dL. In the current study, proteinuria was defined as 1+ or more. eGFR was derived using the following equation (21): eGFR (mL/min/1.73 m²) = 194 × age (years)^{-0.287} × serum creatinine (mg/dL)^{-1.094} (if women × 0.739).

Statistical analysis

All statistical analyses were performed with SPSS version 18.0 J software (SPSS, Chicago, IL). Data were expressed as the means ± SD (age, BMI, eGFR, and BP values) or median and interquartile range (glucose and lipid parameters). Clinical parameters and BP or metabolic values according to the presence of diabetes or prediabetes were compared using ANOVA, and categorical parameters were compared with the χ^2 test. We subdivided the study population according to the quintiles of pulse pressure, and the prevalence of proteinuria ($\geq 1+$) was compared by χ^2 test among each group of the quintiles of pulse pressure separately in subjects with diabetes, prediabetes, or normal glucose tolerance, respectively. The highest quintile of pulse pressure (≥ 63 mmHg, $n = 40,511$) was defined as the high pulse pressure group in the present analysis.

Next, we used a multivariable logistic regression analysis to examine the independent

association of high pulse pressure with proteinuria ($\geq 1+$) separately in subjects with diabetes, prediabetes, or normal glucose tolerance, respectively. In the initial model (Model 1), these associations were assessed with adjustment for age, sex, BMI, current smoking and daily drinking, the presence of antihypertensive medications, and eGFR. Extended models were used to assess whether the association of high pulse pressure with proteinuria ($\geq 1+$) was attenuated by the potential confounding effects of glucose and lipid parameters (Model 2) and systolic BP (Model 3). In addition, to minimize the influence of systolic BP in the association between pulse pressure and proteinuria, we examined the association only in patients with diabetes whose systolic BP was within the normal BP range (i.e., < 130 mmHg) (22). Finally, we examined the association of a +1 SD increase of pulse pressure (+13 mmHg), rather than pulse pressure as a dichotomous variable, with proteinuria in patients with diabetes by a multivariable logistic regression analysis. Statistical significance was defined as $P < 0.05$.

RESULTS

Clinical characteristics of the study population

The mean age \pm SD of the 228,778 subjects was 63.2 ± 8.9 years, and 89,877 of

the subjects (39.3%) were men. There were 27,913 subjects (12.2% of the total subject population) with diabetes, of whom 10,980 subjects (39.1%) were taking antihyperglycemic medications. There were 100,214 subjects (43.8%) with prediabetes. The clinical characteristics according to the presence of diabetes or prediabetes are shown in Table 1. Compared with subjects with normal glucose tolerance (as a reference), the odds ratio (OR) for the increased risk of proteinuria ($\geq 1+$) in diabetes itself was 2.14 (95% CI 2.03–2.25), and that in prediabetes was 1.10 (1.05–1.14), even after adjustment for significant covariates, such as age, sex, BMI, current smoking and daily drinking, the presence of antihypertensive medications, and systolic BP level (both $P < 0.001$).

Pulse pressure and proteinuria

Clinical characteristics and metabolic or BP parameters according to the quintile of pulse pressure are shown in Supplementary Table 2. The increasing prevalence of proteinuria ($\geq 1+$) in accordance with the increasing pulse pressure was more prominent in subjects with diabetes than those without diabetes (Fig. 1). Supplementary Table 3 shows the prevalence of proteinuria subdivided by the dipstick positive scale according to the quintile of pulse pressure with or without diabetes.

Next, a multivariable logistic regression analysis was performed to examine the independent association between the highest quintile of pulse pressure and proteinuria, separately in subjects with diabetes, prediabetes, and normal glucose tolerance. In patients with diabetes, the highest quintile of pulse pressure (≥ 63 mmHg) was positively associated with proteinuria, independently of significant covariates, including systolic BP (Models 1–3 in Table 2). When we examined the association between pulse pressure and proteinuria only in patients with diabetes whose systolic BP was within the normal range (i.e., < 130 mmHg, $n = 11,074$ [39.7%]), the highest quintile of pulse pressure still remained significantly associated with proteinuria (OR 1.46 [95% CI 1.03–2.08]; $P = 0.04$, respectively), even after adjustment for significant covariates, as shown in Model 2 in Table 2. When diastolic BP or mean BP was entered into Model 3 in Table 3 in place of systolic BP, the association between the highest quintile of pulse pressure and proteinuria still remained significant (1.61 [1.49–1.75] and 1.42 [1.31–1.55]; both $P < 0.001$, respectively). In contrast, the highest quintile of pulse pressure in subjects with prediabetes or normal glucose tolerance was not associated with proteinuria independently of systolic BP (Model 3 in Table 2). When

Table 1—Characteristics of the study population according to the presence of diabetes or prediabetes

	Diabetes (n = 27,913)	Prediabetes (n = 100,214)	Normal glucose tolerance (n = 100,651)	P value
Age (years)	65.2 \pm 7.3	64.2 \pm 7.9	61.6 \pm 9.8	<0.001
Men, n (%)	14,626 (52.4)	40,077 (40.0)	35,174 (34.9)	<0.001
BMI (kg/m ²)	24.1 \pm 3.7	23.3 \pm 3.3	22.5 \pm 3.1	<0.001
Current smoker, n (%)	4,846 (17.4)	12,960 (12.9)	13,971 (13.9)	<0.001
Daily drinker, n (%)	7,162 (25.7)	22,825 (22.8)	21,521 (21.4)	<0.001
eGFR (mL/min/1.73 m ²)	76.2 \pm 17.8	74.7 \pm 15.6	76.1 \pm 15.9	<0.001
Proteinuria ($\geq 1+$), n (%)	3,164 (11.3)	5,013 (5.0)	3,913 (3.9)	<0.001
Glucose and lipid parameters				
Fasting glucose (mg/dL)*	125.0 (100.0–143.0)	98.0 (90.0–105.0)	89.0 (84.0–93.0)	<0.001
HbA _{1c} (%)*	6.2 (5.6–6.9)	5.4 (5.3–5.6)	5.0 (4.8–5.1)	<0.001
Triglycerides (mg/dL)*	112.0 (79.0–162.0)	101.0 (74.0–142.0)	91.0 (67.0–127.0)	<0.001
LDL (mg/dL)*	123.0 (104.0–145.0)	127.0 (108.0–148.0)	124.0 (105.0–144.0)	<0.001
HDL (mg/dL)*	57.0 (48.0–68.0)	60.0 (51.0–72.0)	63.0 (53.0–75.0)	<0.001
Antihypertensive drugs, n (%)	11,101 (39.8)	29,157 (29.1)	21,410 (21.3)	<0.001
Antihyperlipidemic drugs, n (%)	6,823 (24.4)	17,440 (17.4)	12,233 (12.2)	<0.001
Antihyperglycemic drugs, n (%)	10,980 (39.1)	0 (0)	0 (0)	<0.001
BP parameters				
Systolic BP (mmHg)	133.4 \pm 17.5	129.7 \pm 17.0	125.7 \pm 17.2	<0.001
Diastolic BP (mmHg)	77.1 \pm 10.8	76.8 \pm 10.5	75.1 \pm 10.7	<0.001
Pulse pressure (mmHg)	56.2 \pm 13.4	52.9 \pm 12.4	50.6 \pm 12.2	<0.001

Data are expressed as the means \pm SD or percentage. P values were obtained by ANOVA or χ^2 test. *Variables with skewed distribution are expressed as median (interquartile range).

Pulse pressure and proteinuria

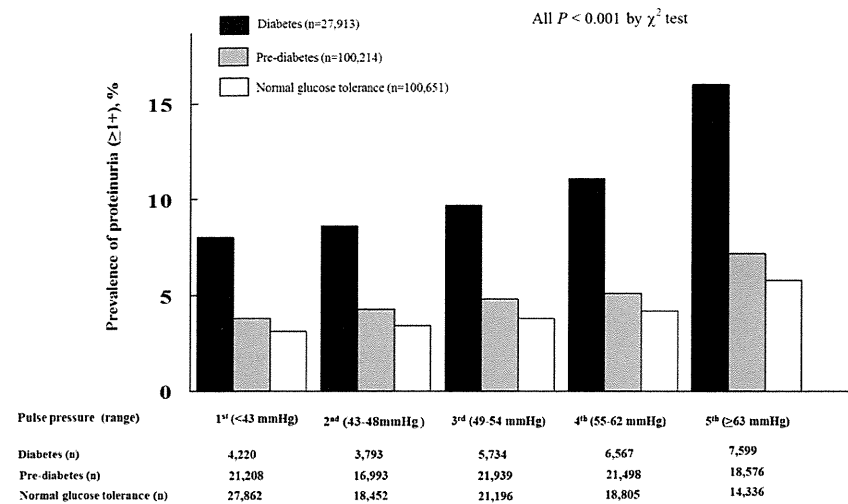


Figure 1—Prevalence of proteinuria according to the quintile of pulse pressure in subjects with diabetes, prediabetes, or normal glucose tolerance. The prevalence of proteinuria ($\geq 1+$) was calculated among each group of the quintiles of pulse pressure separately in subjects with diabetes, prediabetes, or normal glucose tolerance, respectively. The P value was obtained by a χ^2 test among each group of the quintiles of pulse pressure.

we examined the risk of the highest quintile of pulse pressure on proteinuria among subjects without antihypertensive medications ($n = 167,110$), the conclusion remained unchanged (Model 4 in Table 2). Use of antihyperglycemic or antihyperlipidemic drugs did not influence any of the above results (data not shown). In contrast, systolic BP, used as an adjusted factor in Model 3 in Table 2, showed significant associations with proteinuria in subjects with diabetes, prediabetes, and normal glucose tolerance (data not shown).

Finally, we analyzed the association of a +1 SD increase of pulse pressure (+13

mmHg), rather than pulse pressure as a dichotomous variable, with proteinuria in patients with diabetes. We found that a +1 SD increase of pulse pressure was associated with proteinuria independently of significant covariates, including systolic BP (Table 3), diastolic BP, or mean BP (data not shown).

CONCLUSIONS—In this nationwide study of 228,778 Japanese people (mean age 63.2 years) who had no known cardiovascular disease, we demonstrated for the first time that there was a significant difference in the association between the highest

quintile of pulse pressure (≥ 63 mmHg) and proteinuria ($\geq 1+$ on dipstick) among subjects with diabetes, prediabetes, and normal glucose tolerance. The cross-sectional design of the current study did not allow us to elucidate the pathophysiological pathway linking high pulse pressure and proteinuria ($\geq 1+$). However, there are some possible explanations for the observed association.

Pulse pressure, proteinuria, and patients with diabetes

Since the glomerular afferent arterioles provide relatively low resistance, the glomerulus is susceptible to barotrauma if the pulse pressure is elevated (1–6). In fact, prior studies have demonstrated an association of high pulse pressure with microalbuminuria even in subjects without diabetes (7,8). In the current study, we examined the possible association of high pulse pressure and proteinuria ($\geq 1+$), i.e., macroalbuminuria, and found that this association was not significant independently of systolic BP in subjects without diabetes. In contrast, systolic BP was significantly associated with proteinuria in these subjects. Although the usefulness of the urine dipstick test for risk stratification of renal and cardiovascular disease has been recognized, this method is a less sensitive measure of albuminuria compared with the measurement of urinary albumin excretion (23–26). Accordingly, we cannot deny the possibility of an association between high pulse pressure and microalbuminuria in subjects without diabetes.

Table 2—OR for the highest quintile of pulse pressure in the association of proteinuria ($\geq 1+$) according to the presence of diabetes or prediabetes

Model	Adjusted covariates	OR (95% CI)		
Overall ($n = 228,778$)		Diabetes ($n = 27,913$)	Prediabetes ($n = 100,214$)	Normal glucose tolerance ($n = 100,651$)
Model 1	Age + sex + BMI + current-smoking + daily drinking + antihypertensive medications + eGFR	1.72 (1.59–1.87)‡	1.45 (1.35–1.55)‡	1.48 (1.37–1.61)‡
Model 2	Model 1 + fasting glucose + triglycerides + HDL + LDL	1.63 (1.50–1.77)‡	1.41 (1.31–1.50)‡	1.48 (1.36–1.60)‡
Model 3	Model 2 + systolic BP	1.16 (1.05–1.29)†	0.97 (0.89–1.05)	1.08 (0.98–1.20)
Subjects without antihypertensive medications ($n = 167,110$)		Diabetes ($n = 16,812$)	Prediabetes ($n = 71,057$)	Normal glucose tolerance ($n = 79,241$)
Model 4	Age + sex + BMI + current smoking + daily drinking + eGFR + fasting glucose + triglycerides + HDL + LDL + systolic BP	1.21 (1.03–1.43)*	1.09 (0.97–1.23)	1.13 (0.98–1.29)

OR (95% CI) of proteinuria ($\geq 1+$) was calculated for highest quintile of pulse pressure (≥ 63 mmHg, $n = 40,511$) vs. lower quintiles of pulse pressure (< 63 mmHg) in each model. Statistical significance was defined as $P < 0.05$. * $P < 0.05$. † $P < 0.01$. ‡ $P < 0.001$.

Table 3—OR (95% CI) for proteinuria in diabetes (n = 27,913)

Model	OR (95% CI)	P value
Age (+9 years)*	0.94 (0.89–1.00)	0.04
Sex (0, men; 1, women)	0.55 (0.50–0.60)	<0.001
BMI (+3 kg/m ²)*	1.18 (1.14–1.22)	<0.001
Current smoking (0, no; 1, yes)	1.49 (1.35–1.65)	<0.001
Daily drinking (0, no; 1, yes)	0.90 (0.82–0.99)	0.04
Antihypertensive medications (0, no; 1, yes)	0.59 (0.54–0.64)	<0.001
eGFR (+16 mL/min/1.73 m ²)*	0.76 (0.73–0.79)	<0.001
Fasting glucose (+21 mg/dL)*	1.20 (1.18–1.22)	<0.001
Triglycerides (+78 mg/dL)*	1.06 (1.03–1.09)	<0.001
LDL (+30 mg/dL)*	1.07 (1.03–1.11)	<0.001
HDL (+16 mg/dL)*	1.02 (0.98–1.07)	0.39
Systolic BP (+17 mmHg)*	1.27 (1.20–1.36)	<0.001
Pulse pressure (+13 mmHg)*	1.08 (1.01–1.14)	0.02

Statistical significance was defined as $P < 0.05$. *The OR (95% CI) of proteinuria ($\geq 1+$) was calculated for a +1 SD increase of each indicated variable as well as dichromatic variables.

In spite of the strict collinearity between systolic BP and pulse pressure, the OR of high pulse pressure to proteinuria was reduced but remained significant even after adjustment for systolic BP in patients with diabetes (Table 2). Table 3 also shows that a +1 SD increase of systolic BP and a +1 SD increase of pulse pressure were associated with proteinuria independently of each other, with the OR of the systolic BP increase on proteinuria being higher than that of the pulse pressure increase. These findings indicate that high systolic BP showed a confirmed association with proteinuria and is an important confounder explaining the association between high pulse pressure and proteinuria; however, even after adjustment for systolic BP, the pulsatile component of BP itself was still significantly associated with proteinuria in patients with diabetes. Intriguingly, even in the patients with diabetes who were within the normal range of systolic BP values, high pulse pressure was associated with proteinuria. Some possible explanations for these findings exist. First, since renal autoregulation is impaired in diabetes (1–3,11–13), it may be possible that when pulse pressure is elevated, more barotrauma-induced glomerular ultrastructural changes leading to albuminuria occur in subjects with diabetes than in those without diabetes (1–5). Second, much as in the previous reports (27,28), higher pulse pressure was observed in diabetes than nondiabetes (Table 1), suggesting the possibility that diabetes accelerates aortic and large arterial stiffness (29). Aortic stiffness itself has a potential etiologic role in the causation and progression of renal dysfunction (30–32), because loss of the

damping of ventricular ejection in the stiffened aortae could lead to an increase in the transmission of these pressure changes to the renal microcirculation. In the current study, however, we did not use any measure of vascular stiffness more direct than pulse pressure, such as pulse wave velocity, and thus the potential efficacy of such measures will need to be investigated in the future. Third, overt proteinuria in patients with diabetes, which is observed in long-standing diabetes, together with hypertension and increased arterial stiffness, is a surrogate marker not only for renal structural damages but also generalized vascular damages (3,6,24,25). Therefore, we speculate that patients with diabetes with proteinuria are likely to have systemic vasculopathy, and as a consequence, they have high pulse pressure. Lastly, since the current study is a cross-sectional analysis, we have to pay attention to another possibility that diabetic renal disease indicated by greater proteinuria raises systolic BP as well as pulse pressure rather than the reverse in patients with diabetes.

Pulse pressure, proteinuria, and prediabetes

The current study provided the first examination of the association of pulse pressure with proteinuria in prediabetes using a large sample size. Understanding such risk estimates is important, given the increases in the prevalence of prediabetes that have occurred in many populations in conjunction with the increasing prevalence of obesity, particularly in Asian populations (33,34). In the current study, the prevalence of prediabetes was substantially high (44%). Another Japanese study performed in healthy Japanese

people ($n = 6,636$, mean age 50 years) demonstrated that the prevalence of prediabetes was 32% (35). This survey was performed between 1997 and 2003, and since the prevalence of diabetes in Asian populations has increased rapidly in recent years (33,34), the high prevalence of prediabetes in the current study was not entirely unexpected.

Several limitations of our study should be mentioned. First, single-measurement readings of BP, fasting glucose or HbA_{1c}, and proteinuria cannot be considered fully accurate. In particular, some of the dipstick-positive proteinuria could have been transient, and thus could not be taken as definitive evidence of the presence of persisting proteinuria. These factors may introduce a source of variability that could have led to a tendency to underestimate the true association between pulse pressure and proteinuria. Second, we could not separate diabetes into type 1 or type 2 diabetes. However, the incidence of type 1 diabetes is extremely low (approximately two cases/year/100,000 individuals), and Japan has one of the lowest incidence rates of type 1 diabetes in the world (36). Third, we could not assess the diabetes- and atherosclerosis-related information, such as the duration of diabetes and the presence of diabetes complications (e.g., neuropathy), which would be informative and extend the knowledge achieved in the current study. Lastly, we could not assess what kinds of antihypertensive drugs had been prescribed in treated hypertensive subjects. Some antihypertensive drugs (e.g., angiotensin receptor blockers or angiotensin enzyme-converting inhibitors) have more favorable effects on vascular and renal protection (37). Therefore, their use was potentially confounding, although our conclusions remained unchanged when we analyzed our data while excluding the subjects with antihypertensive medications.

In conclusion, among the Japanese general population, high pulse pressure, particularly in individuals with diabetes, was associated with proteinuria, and this information has the potential to supplement other BP indices. To confirm our findings, a prospective study as well as interventions that examine whether or not reduction of pulse pressure can enhance nephron-protective benefits in diabetes will be required.

Acknowledgments—This work was supported by Health and Labor Sciences Research grants for “Research on the Positioning of Chronic

Kidney Disease in Specific Health Check and Guidance in Japan” (20230601) from the Ministry of Health, Labor, and Welfare of Japan.

No potential conflicts of interest relevant to this article were reported.

Y.Y. and Y.S. analyzed the data. S.F. designed the study, collected data, and wrote the paper. T.K. and K.I. designed the study and collected data. T.M., K.Y., K.T., H.Y., K.A., I.K., Y.O., and T.W. designed the study, collected data, supervised the study, and revised the manuscript. S.F. is the guarantor of this work and, as such, had full access to all the data in the study and takes responsibility for the integrity of the data and the accuracy of the data analysis.

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Received for publication: 18.12.2011; Accepted in revised form: 20.5.2012

Nephrol Dial Transplant (2012) 27: 3862–3868

doi: 10.1093/ndt/gfs324

Advance Access publication 1 August 2012

Glycohemoglobin not as predictive as fasting glucose as a measure of prediabetes in predicting proteinuria

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Abstract

Background. There is little data on the assessment of prediabetes with proteinuria.

Methods. This is a cross-sectional cohort study assessing prediabetes with proteinuria in a large Japanese population. Using a nationwide health checkup database of 228 778 Japanese aged ≥ 20 years (median 66 years; 39.3% were men; none had pre-existing cardiovascular disease), we examined the association between prediabetes and proteinuria ($\geq 1+$ on dipstick) separately in prediabetes subjects diagnosed with the new hemoglobin A1c (HbA1c) criterion only (PD-A1c), the impaired fasting plasma glucose only (PD-IFG) and fulfilling both criteria (PD-Both).

Results. According to the American Diabetes Association’s (ADA’s) criterion of 5.7–6.4% HbA1c and/or 100–125 mg/dL fasting plasma glucose, 43.8% of the subjects were judged as having prediabetes. Prediabetes subjects were divided into subclasses of PD-A1c (53.7%), PD-IFG (21.7%) and PD-Both (24.5%), respectively. Therefore, 21.7% of prediabetes subjects were missed using the new

HbA1c criterion only. Compared with subjects with normal glucose tolerance (as a reference), the odds ratio (OR) [95% confidence interval (95% CI)] for the increased risk of proteinuria ($\geq 1+$) in diabetes itself was 2.191 (2.081–2.307) and in whole prediabetes was 1.093 (1.046–1.142); when prediabetes was subdivided, the OR for proteinuria in PD-IFG was 1.217 (1.140–1.300) and that in PD-Both was 1.249 (1.174–1.329), but that in PD-A1c was not significant, even after adjustment for significant covariates, such as age, sex, body mass index, systolic blood pressure, antihypertensive medication, eGFR, lifestyle and lipid profile.

Conclusions. Prediabetes is a significant risk factor for proteinuria compared with completely normal glucose level, and subjects with prediabetes defined using IFG are at significantly higher risk for proteinuria than those defined by HbA1c only.

Keywords: odds ratio; prediabetes; proteinuria