

제2형 당뇨병 조기발견을 위한 전략

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CONTENTS

- Basic epidemiology of T2DM* and Prediabetes
- Diabetes Screening Guidelines
- Natural Course of Diabetes Development
- Risk Factors of Diabetes: Genetic and Environmental
- Strategies for Early Diagnosis of T2DM

*T2DM: Type 2 Diabetes

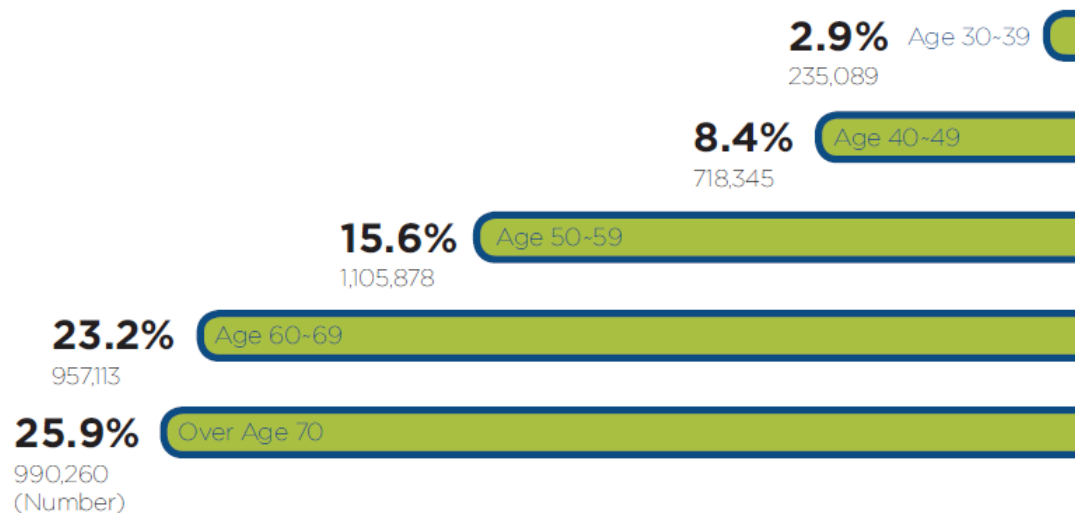
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PREVALENCE OF DIABETES 2011 (≥ 30 YRS OLD)

- > The prevalence of diabetes in adults 30 years and older is 12.4%.
- > As of 2011, an estimated 4.0 million people (about 1 every 8 adults) had diabetes.



IMPAIRED FASTING GLUCOSE



Total
19.3%
6,100,430

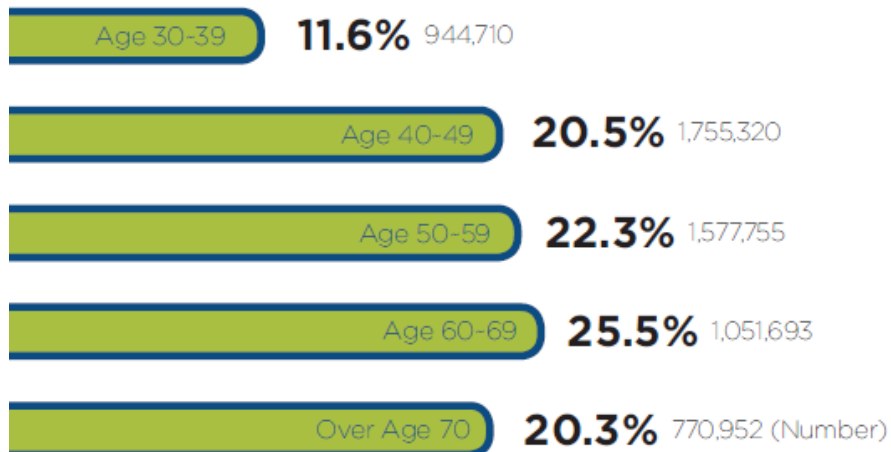


Men
23.8%
3,667,699



Women
14.9%
2,432,730

- > Approximately 20% of adults 30 years and older (6.1 million people) have impaired fasting glucose.
- > Therefore, about 1 in 3 adults was diabetes or had potential risk for diabetes.



AWARENESS OF DIABETES

3 out of 10

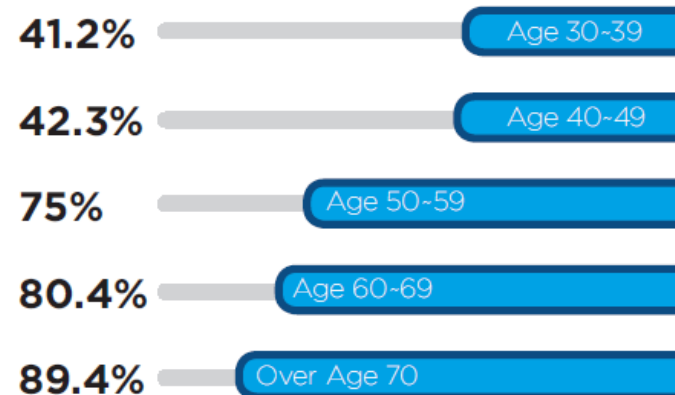
diabetic patients are not aware of their condition.
(diabetes awareness rate: 72%).



In a younger population of age < 50,

6 out of 10

are unaware of their diabetes.



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Diabetes Screening (ADA)

Table 2.2—Criteria for testing for diabetes or prediabetes in asymptomatic adults

1. Testing should be considered in all adults who are overweight (BMI ≥ 25 kg/m² or ≥ 23 kg/m² in Asian Americans) and have additional risk factors:
 - physical inactivity
 - first-degree relative with diabetes
 - high-risk race/ethnicity (e.g., African American, Latino, Native American, Asian American, Pacific Islander)
 - women who delivered a baby weighing >9 lb or were diagnosed with GDM
 - hypertension ($\geq 140/90$ mmHg or on therapy for hypertension)
 - HDL cholesterol level <35 mg/dL (0.90 mmol/L) and/or a triglyceride level >250 mg/dL (2.82 mmol/L)
 - women with polycystic ovary syndrome
 - A1C $\geq 5.7\%$, IGT, or IFG on previous testing
 - other clinical conditions associated with insulin resistance (e.g., severe obesity, acanthosis nigricans)
 - history of CVD
 2. For all patients, particularly those who are overweight or obese, testing should begin at age 45 years.
 3. If results are normal, testing should be repeated at a minimum of 3-year intervals, with consideration of more frequent testing depending on initial results (e.g., those with prediabetes should be tested yearly) and risk status.
-

Diabetes Screening (대한당뇨병학회)

- 40세 이상에서 선별 검사를 시행
- 아래 위험인자가 있는 경우 30세 이상

표 1. 제2형 당뇨병의 위험인자

- ▶ 과체중 (체질량지수 23 kg/m^2 이상)
- ▶ 직계 가족 (부모, 형제자매)에 당뇨병이 있는 경우
- ▶ 공복혈당장애나 내당능장애의 과거력
- ▶ 임신성 당뇨병이나 4 kg 이상의 거대아 출산력
- ▶ 고혈압 (140/90 mmHg 이상, 또는 약제 복용)
- ▶ HDL 콜레스테롤 35 mg/dL 미만 혹은 중성지방 250 mg/dL 이상
- ▶ 인슐린저항성 (다낭난소증후군, 흑색가지세포증 등)
- ▶ 심혈관질환 (뇌졸중, 관상동맥질환 등)

Diabetes Screening (USPSTF)

Annals of Internal Medicine

REVIEW

Screening for Type 2 Diabetes Mellitus: A Systematic Review for the U.S. Preventive Services Task Force

Shelley Selph, MD, MPH; Tracy Dana, MLS; Ian Blazina, MPH; Christina Bougatsos, MPH; Hetal Patel, MD; and Roger Chou, MD

- Screening for diabetes did not improve mortality rates after 10years follow-up in 2 trials
- But it was found to decrease mortality rate in a lifestyle intervention study with 23 years of follow up
- More evidence is needed to determine the effectiveness of treatments for screen detected diabetes
- Treatment of IFG or IGT was associated with delayed progression to diabetes

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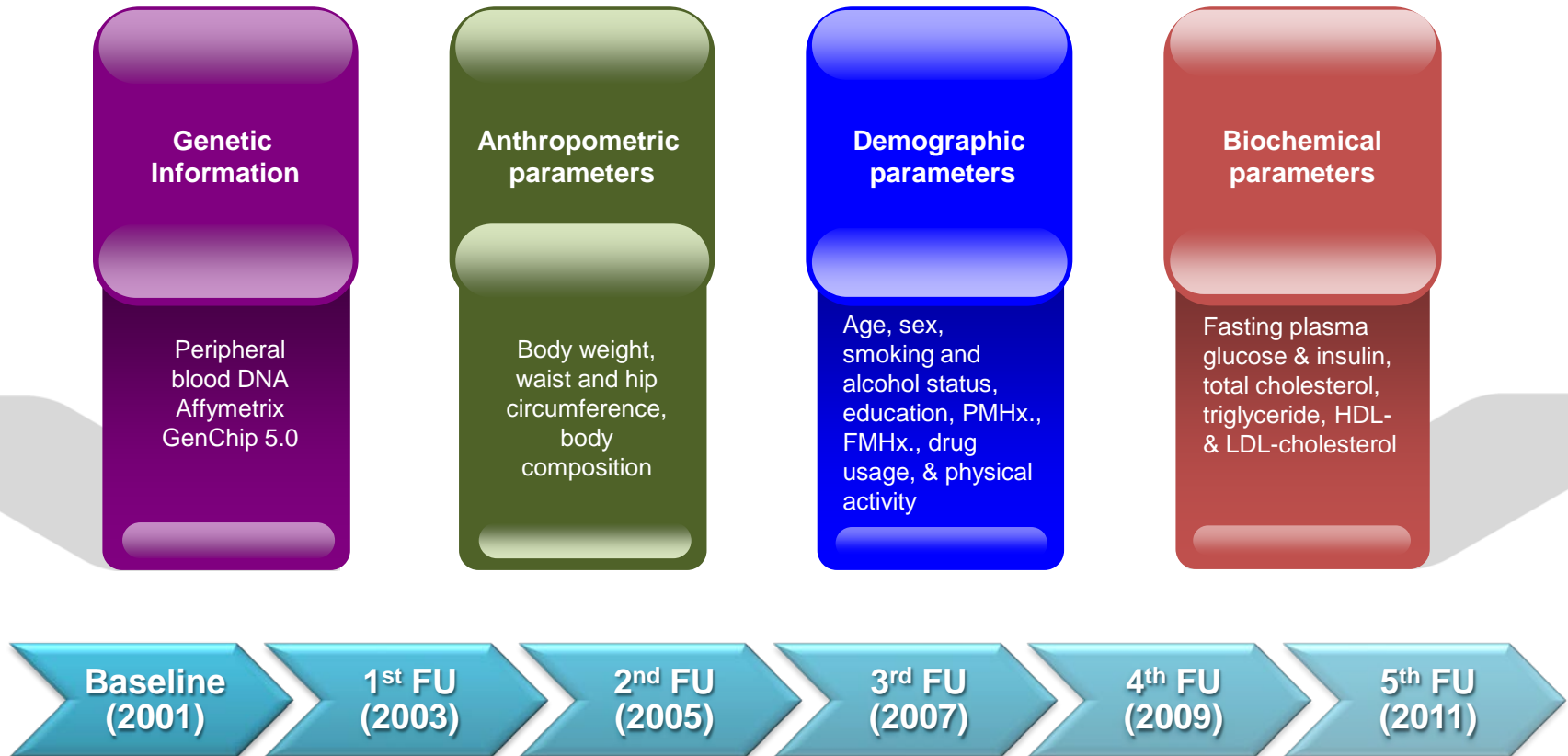
*T2DM: Type 2 Diabetes

Korea Human Genomic Study -Research Design and Methods

- Ansong cohort (Rural)
 - Population: 135,000
 - Farming area
 - Age: 40-69 yr
 - Subject: 5,024
 - Ansan cohort (Urban)
 - Population: 550,000
 - Industrial area
 - Age: 40-69 yr
 - Subject: 5,014
- * Eligibility criteria
- 40-69 years,
 - residence within the borders of the survey area for at least 6 months
 - mental and physical ability to participate.



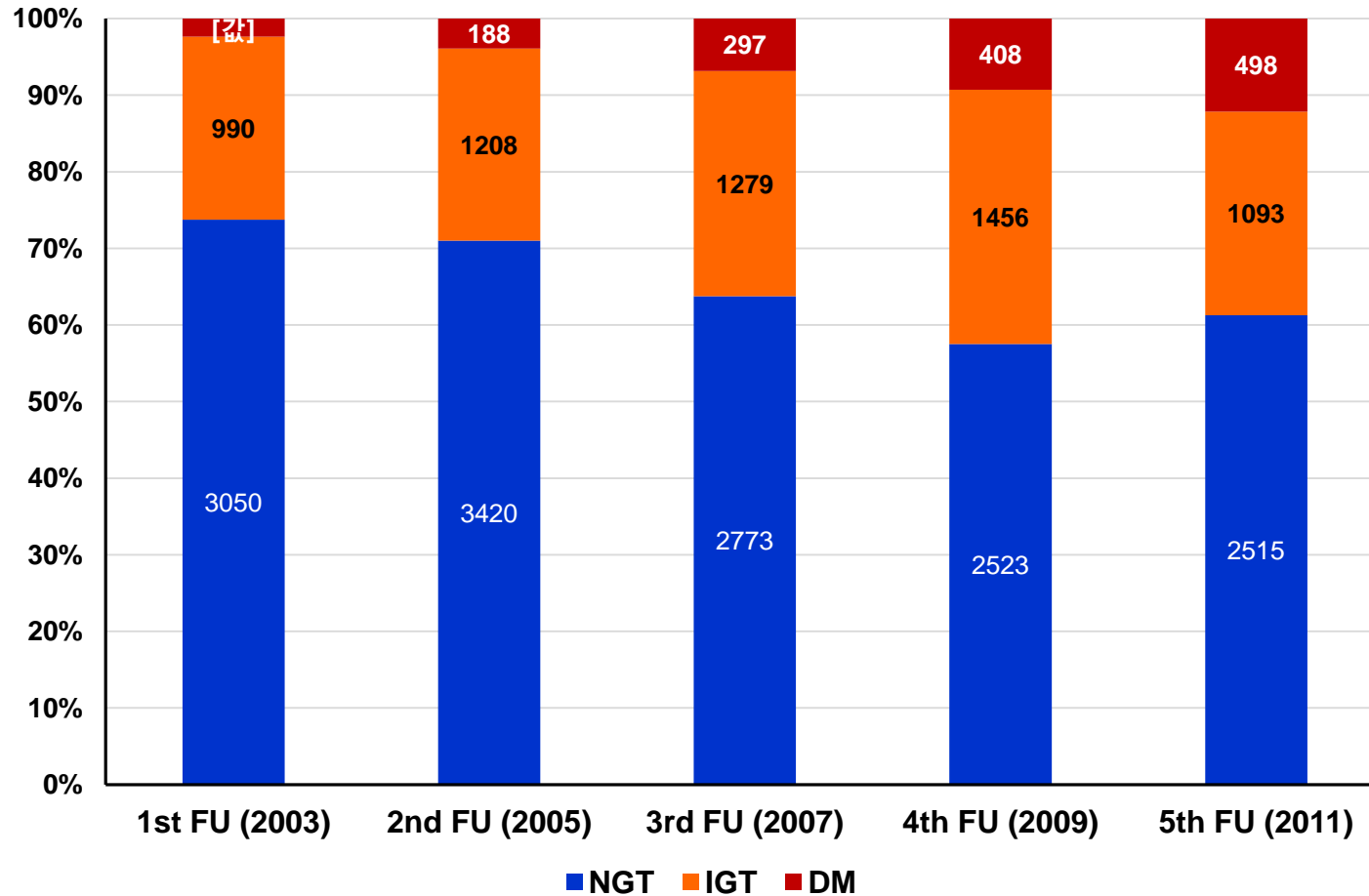
Biannual FU with Detailed Clinical and Biochemical Information



The follow-up is still ongoing

10 Year FU of Baseline NGT

Cumulative Incidence of IGT, and DM (Baseline NGT)



Measurement of beta cell function and insulin resistance

- 10,038 participant
- 75 g OGTT with 0, 60, 120 min Glucose and Insulin
- **Biennial follow-up (2001-2010)**

– Insulin secretion index

- Insulinogenic Index (60 min) = $\frac{(Insulin_{60} - Insulin_0, \mu U/ml)}{(Glucose_{60} - Glucose_0, mmol/l)}$
- HOMA-B = $20 \times (Insulin_0, \mu U/ml) \times (Glucose_0 - 3.5, mmol/l)$

– Insulin sensitivity index

- Matsuda Index =
$$\frac{10,000}{\sqrt{(Glucose_0, mg/dl) \times (Insulin_0, \mu U/ml) \times (mean\ Glucose, mg/dl) \times (mean\ Insulin, \mu U/ml)}}$$
- HOMA-IR = $\frac{(Glucose_0, mmol/l) \times (Insulin_0, \mu U/ml)}{22.5}$

– Disposition index

- (Insulinogenic Index) x (Matsuda Index)

NATURAL COURSE OF DIABETES

Baseline characteristics of study participants by the glycemic status at the end of 10-year follow-up

	Non-progressor (n=2,515)	Progressor to prediabetes (n=1,093)	Progressor to diabetes (n=498)	p value
Age (yr)	50.6±8.3	51.4±8.2	52.8±8.9	<0.001
Male (%)	1134(45.1)	526(48.1)	285(57.2)	<0.001
BMI (kg/m ²)	24.0±2.9	25.0±3.0	24.9±3.2	<0.001
Waist circumference (cm)	80.9±8.4	83.3±8.2	84.5±8.7	<0.001
Systolic BP (mmHg)	114±16	118±17	119±18	<0.001
Diastolic BP (mmHg)	73±11	76±12	77±11	<0.001
FPG (mmol/L)	4.5±0.4	4.7±0.4	4.8±0.4	<0.001
2-h glucose (mmol/L)	5.7±1.1	6.1±1.1	6.3±1.1	<0.001
HbA1c (%)	5.2±0.3	5.4±0.3	5.5±0.4	<0.001
Fasting insulin (pmol/L)*	37.8(36.9-38.7)	40.1(38.6-41.5)	39.0(36.9-41.2)	0.015
IGI ₆₀ *	7.9(7.4-8.3)	6.6(6.1-7.1)	5.1(4.5-5.7)	<0.001
HOMA-β*	131(128-135)	122(117-127)	110(103-117)	<0.001
Composite ISI*	10.0(9.8-10.2)	8.6(8.3-8.9)	8.2(7.8-8.7)	<0.001
HOMA-IR*	1.27(1.24-1.30)	1.38(1.33-1.43)	1.37(1.29-1.45)	<0.001
DI*	72.9(69.1-76.9)	52.9(49.3-56.6)	39.7(36.1-43.7)	<0.001
Total cholesterol (mmol/L)*	4.79(4.76-4.83)	4.89(4.84-4.94)	4.96(4.88-5.04)	<0.001
HDL cholesterol (mmol/L)*	1.20(1.19-1.21)	1.14(1.13-1.16)	1.11(1.09-1.13)	<0.001
Triglycerides (mmol/L)*	1.34(1.32-1.37)	1.58(1.54-1.63)	1.77(1.70-1.85)	<0.001
Hypertension (%)	169(6.7%)	129(11.8%)	80(16.1%)	<0.001
Family history of diabetes (%)	223(8.9%)	139(12.7%)	64(12.9%)	<0.001

P values are adjusted for age and sex. Data are means±SD, geometric means(95% CI), or n(percentage).

*Log transformed before statistical analysis.

BMI, body mass index; BP, blood pressure; FPG, fasting plasma glucose; IGI₆₀, insulinogenic index at 60 minute; HOMA-β, Homeostasis Model Assessment of β-cell function; ISI, insulin sensitivity index; HOMA-IR, Homeostasis Model Assessment of insulin resistance; DI, disposition index.

Impairment in β -cell function has more profound impact on incident diabetes compared to decreased insulin sensitivity

	High-IGI ₆₀ , High-ISI	High-IGI ₆₀ , Low-ISI		Low-IGI ₆₀ , High-ISI		Low-IGI ₆₀ , Low-ISI	
	Reference	HR (95% CI)	p value	HR (95% CI)	p value	HR (95% CI)	p value
Number of subjects	658	717		1,574		425	
Incident diabetes (%)	23(3.5%)	84(11.7%)		213(13.5%)		106(24.9%)	
Model A	1.00	3.4(2.2-5.4)	<0.001	4.1(2.7-6.3)	<0.001	8.0(5.1-12.6)	<0.001
Model B	1.00	3.5(2.2-5.5)	<0.001	4.0(2.6-6.1)	<0.001	8.2(5.2-12.8)	<0.001
Model C	1.00	3.3(2.1-5.3)	<0.001	3.8(2.5-5.9)	<0.001	7.9(5.0-12.5)	<0.001
Model D	1.00	2.6 (1.6-4.1)	<0.001	3.8 (2.4-5.8)	<0.001	6.2 (3.9-9.8)	<0.001
PAF(95% CI), %^a	-	12.1 (7.5-14.9)		36.7 (29.5-41.4)		20.9 (18.5-22.3)	

The cutoffs for dichotomizing IGI₆₀ and composite ISI into high vs. low groups are **10.5** and **6.9**, respectively.

Model A: unadjusted.

Model B: age, sex, and center adjusted.

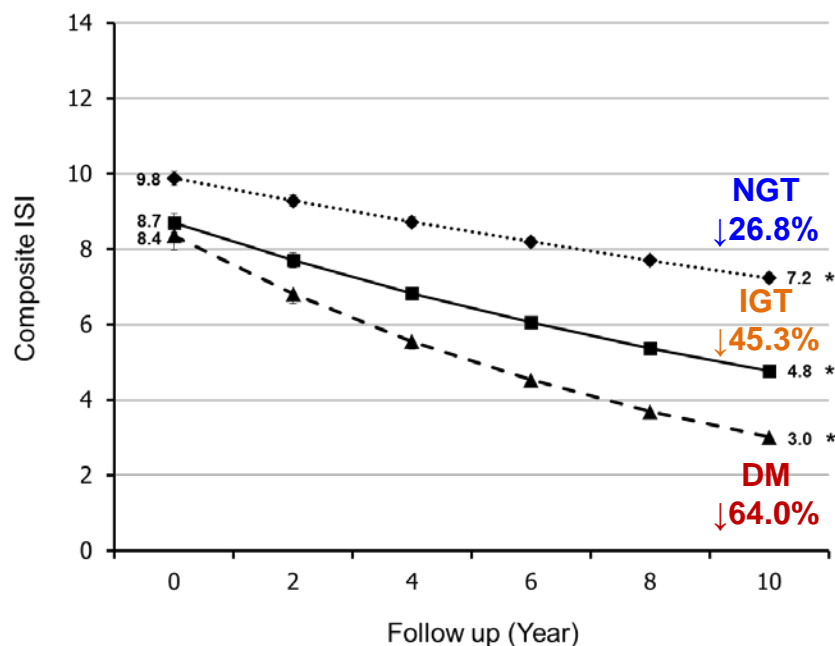
Model C: Model B + smoking, sporting activity, alcohol intake, and family history of diabetes adjusted.

Model D: Model C + systolic BP, waist circumference, ALT, total cholesterol, HDL cholesterol, and triglyceride adjusted.

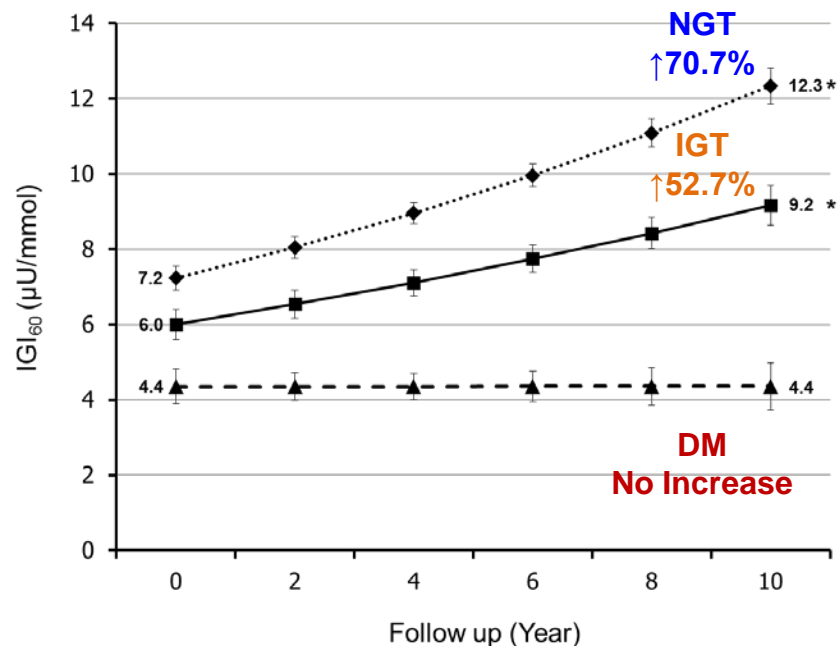
^aPopulation-attributable fraction(PAF, %) was calculated using multivariable-adjusted HR (Model D).

Trajectory of Insulin Sensitivity and Insulin Secretion During 10 Years Of Follow-up.

Change in Insulin Sensitivity Composite ISI (Matsuda Index)



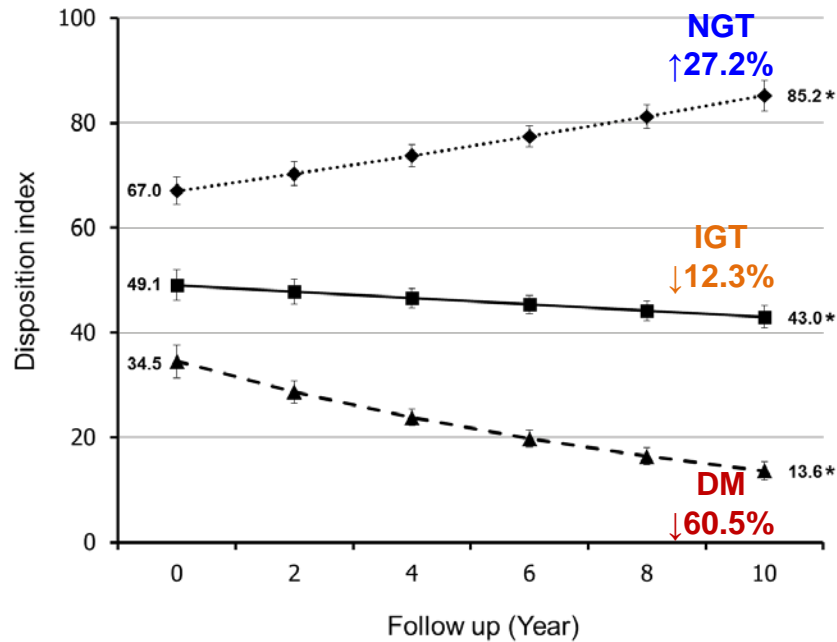
Change in Insulin Secretion Insulogenic Index (60min)



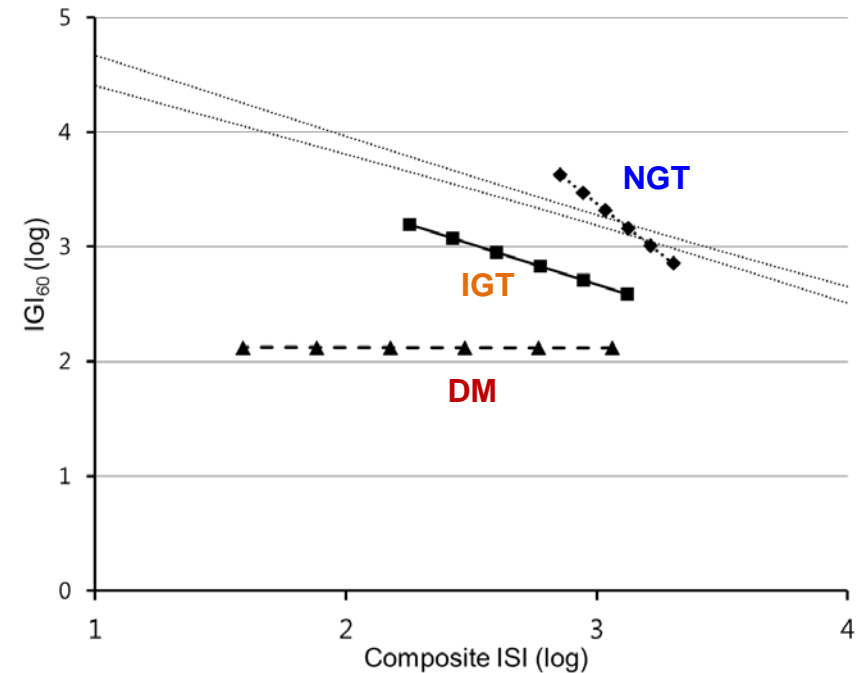
*p<0.01 for 10 vs. 0 years.

Overall Trajectory of Disposition Index, Insulin Sensitivity and Insulin Secretory Function

Change in Disposition Index



Change in Insulin Sensitivity and Insulin Secretion



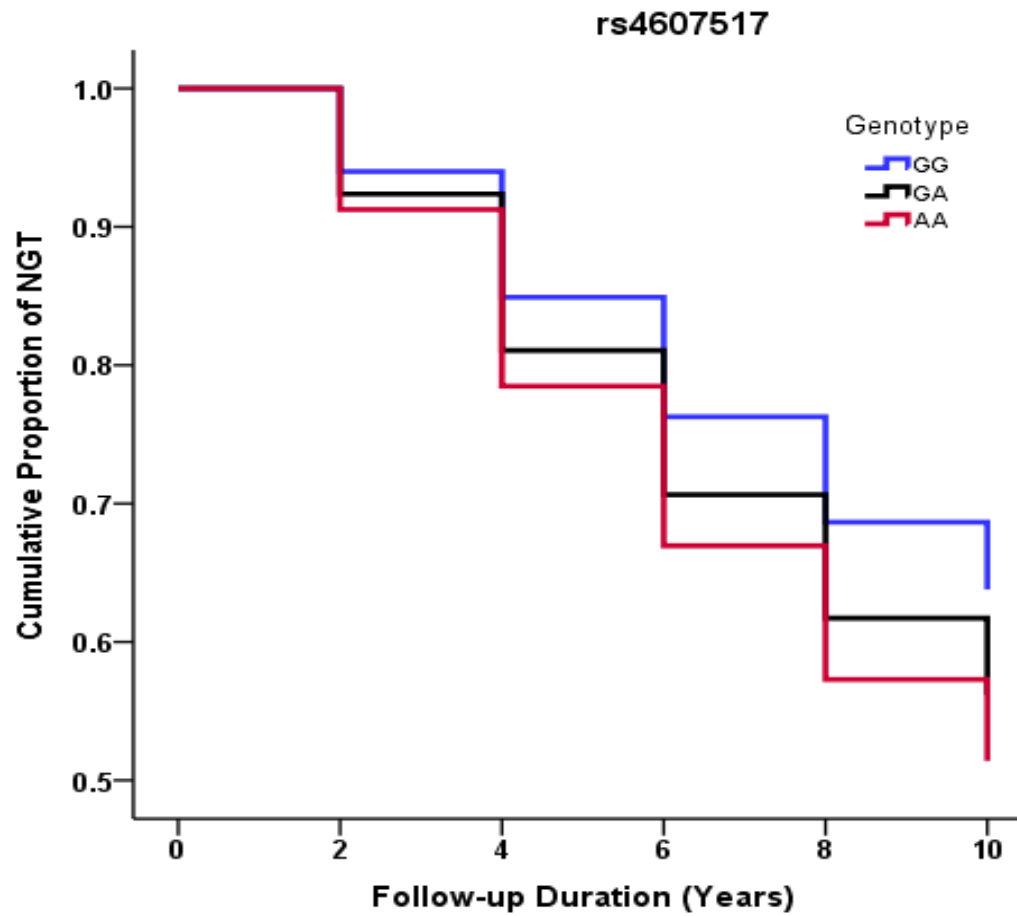
*p<0.01 for 10 vs. 0 years.

NATURAL COURSE OF DIABETES

Genetic Variants Associated With Progression To Diabetes

SNP	Chr	Position	Nearest Gene	Risk allele	Non-progressor vs. progressor to either prediabetes or diabetes					Non-progressor vs. progressor to diabetes			
					RAF in non-progressor (n=2,046)	RAF in prediabetes or diabetes (n=1,349)	HR	95% CI	p value	RAF in diabetes (n=431)	HR	95% CI	p value
rs17106184	1	50,682,573	<i>FAF1</i>	G	0.88	0.89	1.06	(0.94 to 1.19)	0.33	0.91	1.33	(1.06 to 1.67)	0.014
rs10923931	1	120,319,482	<i>NOTCH2</i>	T	0.03	0.03	1.10	(0.88 to 1.39)	0.40	0.03	1.21	(0.83 to 1.77)	0.33
rs2075423	1	212,221,342	<i>PROX1</i>	G	0.81	0.82	1.04	(0.94 to 1.15)	0.46	0.82	1.01	(0.85 to 1.21)	0.88
rs780094	2	27,594,741	<i>GCKR</i>	C	0.47	0.47	1.00	(0.93 to 1.08)	0.92	0.49	1.07	(0.93 to 1.22)	0.33
rs10203174	2	43,543,534	<i>THADA</i>	C	0.99	0.99	2.14	(0.96 to 4.77)	0.063	0.99	3.72	(0.52 to 26.49)	0.19
rs243088	2	60,422,249	<i>BCL11A</i>	T	0.64	0.65	1.05	(0.96 to 1.13)	0.29	0.65	1.03	(0.89 to 1.19)	0.66
rs6723108	2	135,196,450	<i>TMEM163</i>	T	0.98	0.98	0.86	(0.67 to 1.11)	0.24	0.98	0.79	(0.51 to 1.21)	0.28
rs7560163	2	151,346,182	<i>RND3</i>	C	0.90	0.90	0.93	(0.81 to 1.05)	0.23	0.88	0.73	(0.59 to 0.90)	0.0027
rs7593730	2	160,879,700	<i>RBMS1</i>	C	0.82	0.83	1.04	(0.94 to 1.16)	0.41	0.85	1.19	(0.99 to 1.44)	0.064
rs3923113	2	165,210,095	<i>GRB14</i>	A	0.88	0.88	1.09	(0.96 to 1.24)	0.20	0.88	1.03	(0.83 to 1.28)	0.81
rs2943640	2	226,801,829	<i>IRS1</i>	C	0.95	0.95	0.99	(0.84 to 1.18)	0.93	0.95	1.07	(0.78 to 1.46)	0.68
rs1801282	3	12,368,125	<i>PPARG</i>	C	0.94	0.95	1.16	(0.96 to 1.39)	0.12	0.95	1.08	(0.79 to 1.49)	0.62
rs7612463	3	23,311,454	<i>UBE2E2</i>	C	0.80	0.81	1.11	(1.00 to 1.22)	0.043	0.82	1.16	(0.98 to 1.38)	0.093
rs831571	3	64,023,337	<i>PSMD6</i>	C	0.62	0.63	1.02	(0.94 to 1.10)	0.67	0.64	1.07	(0.93 to 1.23)	0.37
rs6795735	3	64,680,405	<i>ADAMTS9</i>	C	0.20	0.21	1.03	(0.94 to 1.14)	0.49	0.21	1.02	(0.86 to 1.21)	0.82
rs4402960	3	186,994,381	<i>IGF2BP2</i>	T	0.28	0.29	1.04	(0.96 to 1.13)	0.31	0.29	1.06	(0.91 to 1.22)	0.46
rs16861329	3	188,149,155	<i>ST6GAL1</i>	C	0.85	0.84	0.88	(0.78 to 0.98)	0.025	0.85	0.96	(0.78 to 1.18)	0.70
rs6815464	4	1,299,901	<i>MAEA</i>	C	0.60	0.60	1.01	(0.93 to 1.09)	0.79	0.62	1.10	(0.96 to 1.26)	0.18
rs4458523	4	6,340,887	<i>WFS1</i>	G	0.98	0.97	0.97	(0.77 to 1.24)	0.83	0.97	0.96	(0.63 to 1.46)	0.84
rs6813195	4	153,739,925	<i>TMEM154</i>	C	0.51	0.54	1.10	(1.02 to 1.19)	0.011	0.55	1.12	(0.98 to 1.28)	0.091
rs702634	5	53,307,177	<i>ARL15</i>	A	0.89	0.90	1.08	(0.96 to 1.23)	0.21	0.90	1.17	(0.93 to 1.47)	0.17
rs459193	5	55,842,508	<i>ANKRD55</i>	G	0.51	0.53	1.04	(0.97 to 1.12)	0.29	0.53	1.04	(0.91 to 1.19)	0.57
rs9502570	6	7,235,436	<i>SSR1/RREB1</i>	T	0.59	0.59	1.05	(0.97 to 1.14)	0.26	0.59	1.04	(0.91 to 1.20)	0.56
rs7756992	6	20,787,688	<i>CDKAL1</i>	G	0.53	0.54	1.05	(0.97 to 1.13)	0.25	0.52	0.97	(0.84 to 1.11)	0.62
rs9470794	6	38,214,822	<i>ZFAND3</i>	C	0.25	0.24	0.97	(0.88 to 1.06)	0.47	0.23	0.93	(0.79 to 1.09)	0.36
rs1535500	6	39,392,028	<i>KCNK16</i>	T	0.41	0.42	1.01	(0.94 to 1.09)	0.77	0.41	0.98	(0.85 to 1.12)	0.74
rs17168486	7	14,864,807	<i>DGKB</i>	T	0.38	0.39	1.08	(0.99 to 1.16)	0.071	0.36	0.89	(0.77 to 1.03)	0.11
rs849135	7	28,162,938	<i>JAZF1</i>	G	0.99	0.99	0.95	(0.50 to 1.84)	0.89	0.99	0.91	(0.29 to 2.84)	0.87
rs4607517	7	44,223,721	GCK	A	0.16	0.21	1.27	(1.16 to 1.38)	1.70 × 10⁻⁷	0.21	1.20	(1.02 to 1.40)	0.025
rs6467136	7	126,952,194	<i>GCC1</i>	G	0.76	0.75	0.97	(0.89 to 1.07)	0.56	0.75	0.97	(0.83 to 1.14)	0.72
rs13233731	7	130,088,229	<i>KLF14</i>	G	0.70	0.69	0.99	(0.92 to 1.08)	0.89	0.69	1.00	(0.87 to 1.16)	1.00
rs516946	8	41,638,405	<i>ANK1</i>	C	0.86	0.88	1.13	(1.01 to 1.27)	0.038	0.89	1.19	(0.97 to 1.47)	0.10
rs7845219	8	96,006,678	<i>TP53INP1</i>	T	0.28	0.28	1.04	(0.96 to 1.13)	0.32	0.29	1.11	(0.96 to 1.29)	0.16

Genetic Variant in Glucokinase



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Optimal Cutoff for BMI

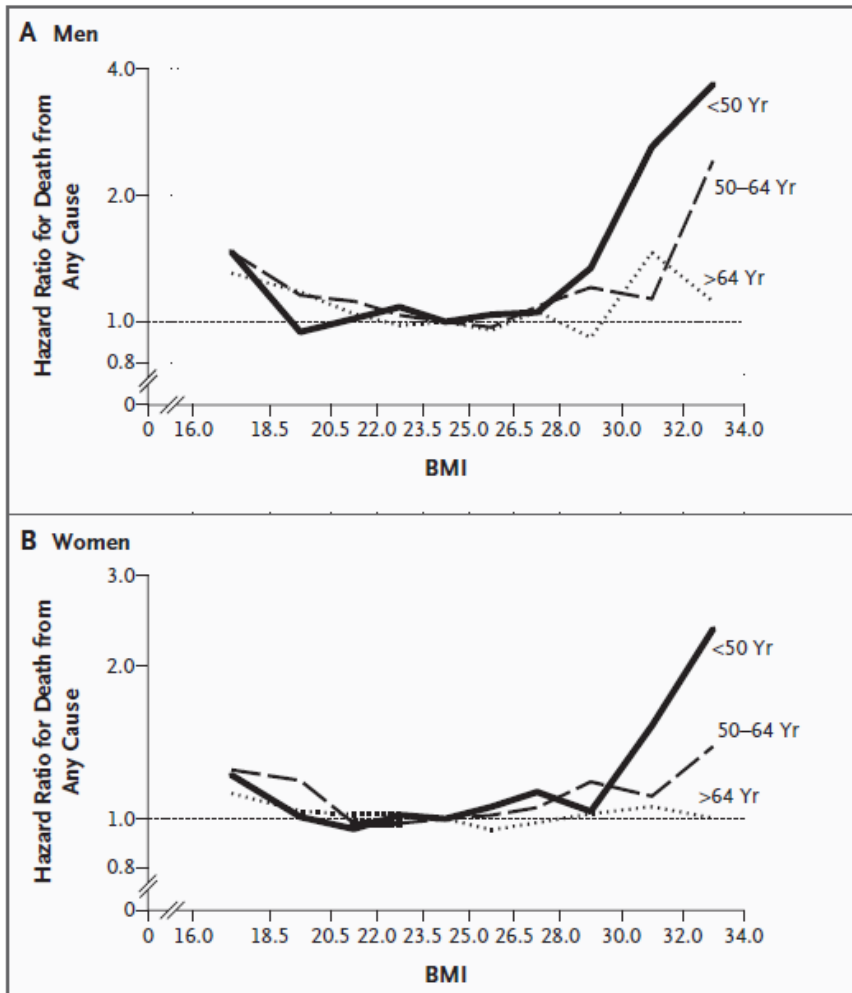


Figure 3. Hazard Ratios for Death from Any Cause among Men and Women with No History of Smoking, According to Age Group and BMI.

Data are from the KCPS, 1993–2004.^{12,13} The reference category was a BMI of 23.0 to 24.9. All hazard ratios were adjusted for age. Panel A and Panel B have different scales for hazard ratios.

OBESITY

- ✓ Based on Mortality
- ✓ BMI ≥ 25 kg/m²

OVERWEIGHT

- ✓ Based on Morbidity
- ✓ BMI ≥ 23 kg/m²

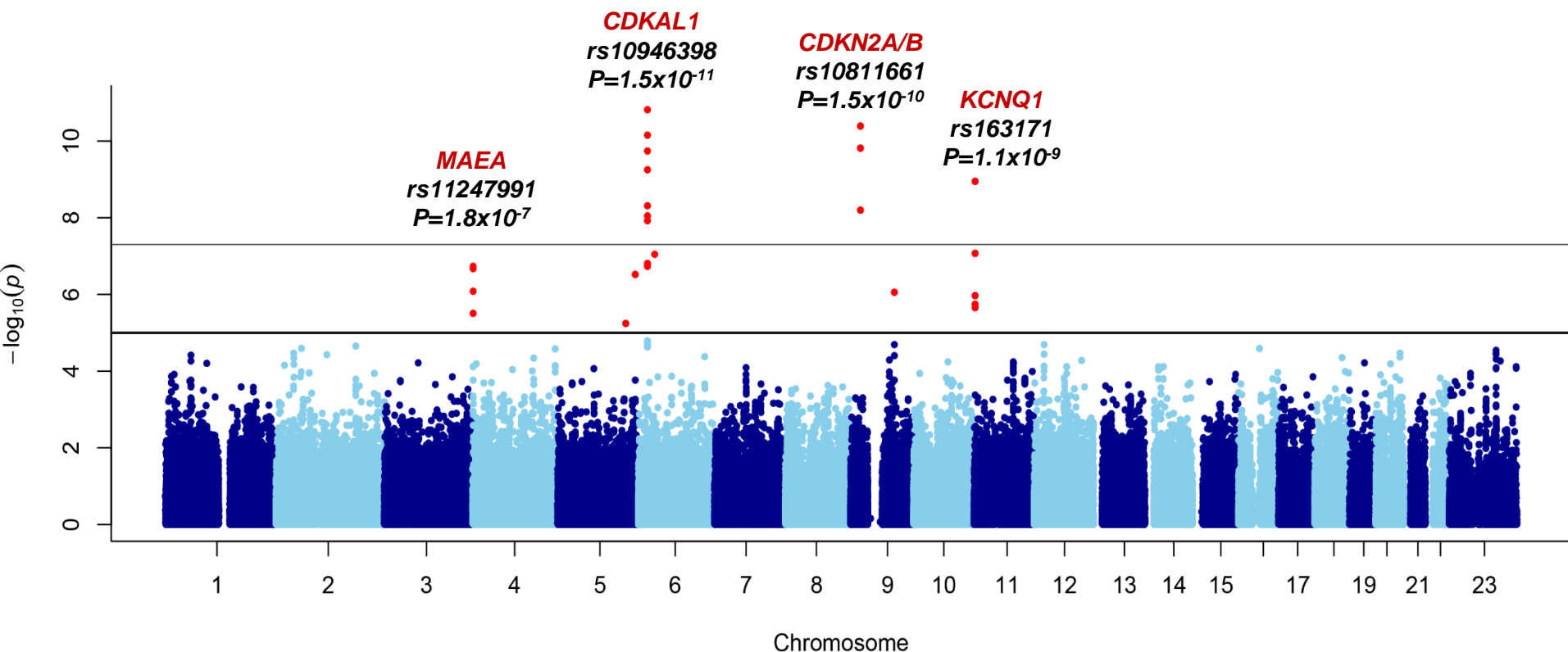
WEIGHT CIRCUMFERENCE

- ✓ Male ≥ 90 cm
- ✓ Female ≥ 85 cm

Evidence of Genetic Etiology in DM

1. Familial clustering:
 - “Diabetes runs in my family”
 - Life time risk of diabetes: one parent with T2DM – 35%
both parents with T2DM – 70%
2. Sibling recurrence risk ratio (λ_s): 2-3
3. Twin study:
 - Heritability (h^2) in monozygotic twins: 35-70%
 - Heritability (h^2) in dizygotic twins: 20-30%
4. Ethnic difference in prevalence of diabetes
5. Monogenic diabetes as specific type (MODY)

Genome-Wide Association Study in Korean T2DM vs Control Subjects



Subjects are from Ansung cohort and Seoul National University Hospital.

N = **3,596** (Cases/Controls = 2,601/995).

Logistic regression analysis adjusting for age, sex, and BMI.

Normal controls: Age > 60, NGT, No FHx of DM, No PHx of DM.

PAX4 (Paired Box 4)

Chr	rsID	Position	Non-Ref Allele		Non-Ref AF in T2DM	Non-Ref AF in Controls	No. Subjects (Case/Control)	OR	se	P
7	rs2233580	127,253,550	T	Stage 1	0.122	0.057	917 (619/298)	2.54	0.23	2.91x10⁻⁵
				Stage 2*	0.094	0.051	3,039 (2,021/1,018)	1.46	0.09	2.26x10⁻⁶
				Meta-Analysis**			3,956 (2,640/1,316)	1.81	0.10	5.48x10⁻⁹
7	rs3824004	127,253,551	T	Stage 1	0.045	0.037	917 (619/298)	1.06	0.31	0.858
				Stage 2*	0.054	0.029	3,036 (2,018/1,018)	1.81	0.15	5.75x10⁻⁵
				Meta-Analysis**			3,956 (2,640/1,316)	1.63	0.14	4.26x10⁻⁴

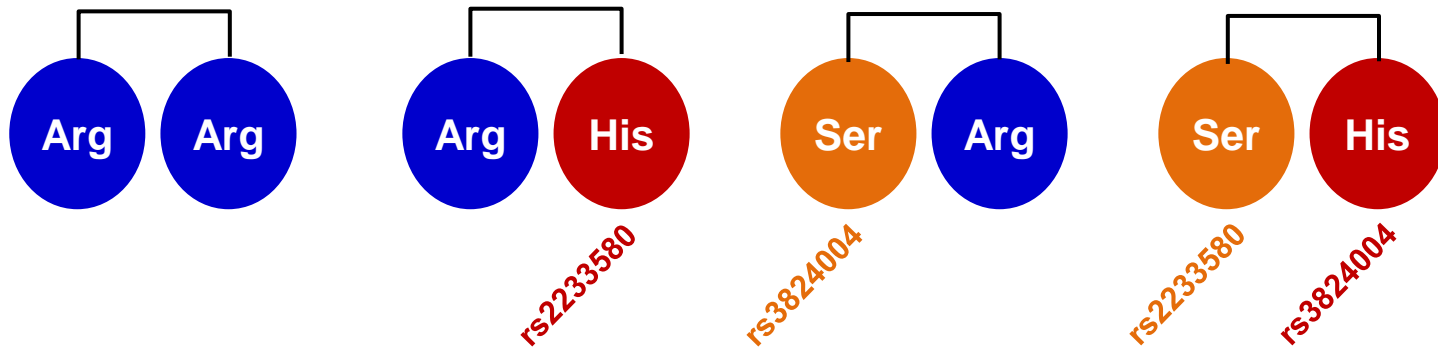
P values are from logistic regression adjusting for age and sex. ** Meta-analysis was done using METAL with inverse-variance weighted fixed effect model.

PAX4 Haplotype Association

CHR	SNP1	SNP2	HAPLOTYPE	AA	Group	Frequency	OR	P	Overall P
7	rs2233580	rs3824004	ACG	Arginine	Positive Charge	0.874	0.57	2.53E-09	1.81E-08
7	rs2233580	rs3824004	ATG	Histidine	Positive Charge	0.080	1.66	1.11E-05	
7	rs2233580	rs3824004	ACT	Serine	Polar Uncharged	0.044	1.80	0.000199	
7	rs2233580	rs3824004	ATT	Asparagine	Polar Uncharged	0	-	-	

LD measure for two SNPs: R-sq 0.002, D'=0.619

Possible Amino Acid Changes



Dosage

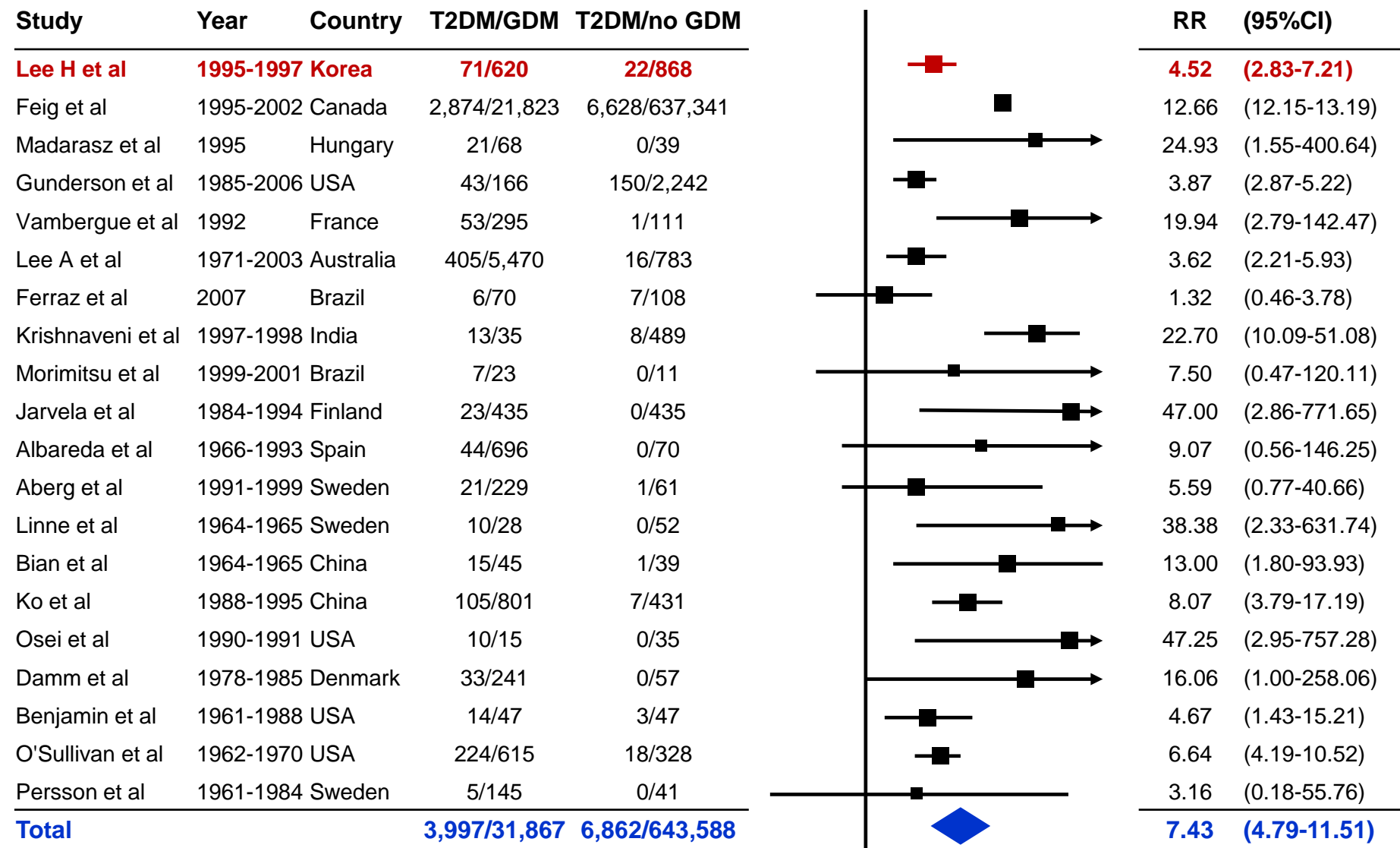
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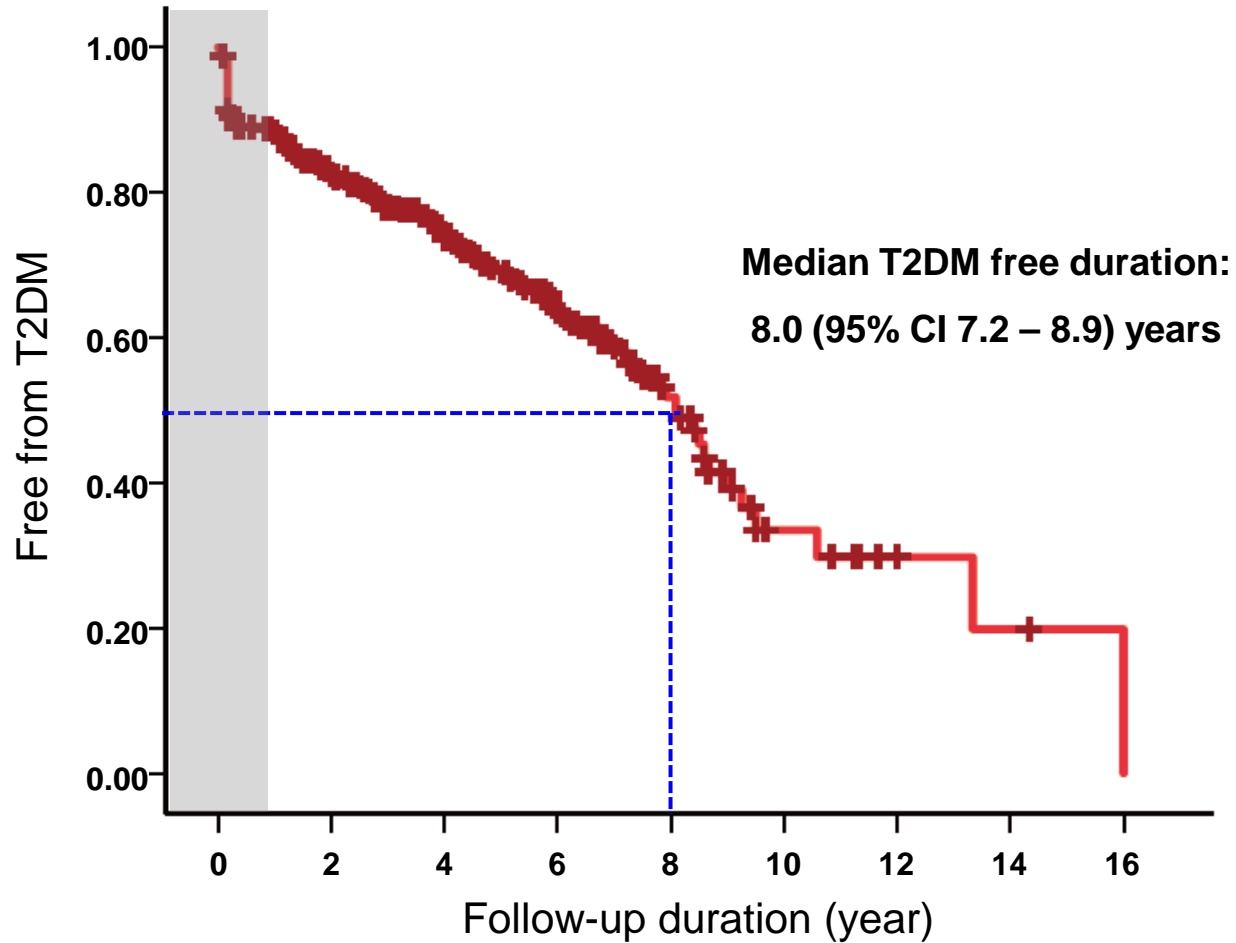
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2

Risk of T2DM after GDM



Incidence of T2DM after GDM in Koreans



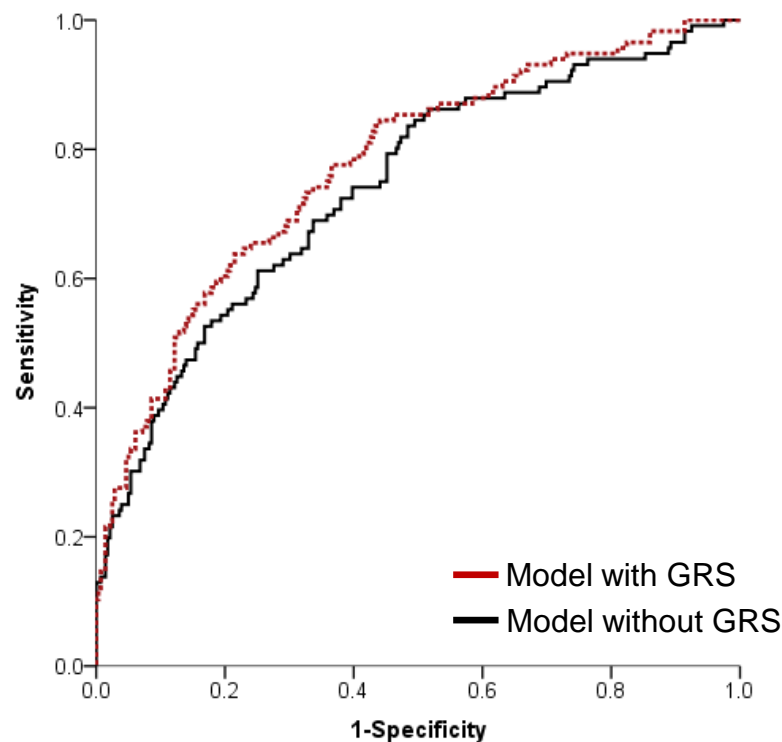
Early vs. Late Converters: Genetic Risk Factors

CHR	Nearby Gene	SNP	Risk Allele	Risk Allele Frequency			OR		P	
				Early Converters (N=64)	Late Converters (N=75)	Non-diabetic Controls (N=632)	Early Converters	Late Converters	Early Converters	Late Converters
3	<i>PPARG</i>	rs1801282	C	0.97	0.95	0.95	1.74	1.14	0.286	0.742
3	<i>PPARG</i>	rs3856806	C	0.84	0.85	0.82	1.09	1.24	0.731	0.366
3	<i>IGF2BP2</i>	rs4402960	T	0.36	0.42	0.30	1.34	1.71	0.136	0.00253
6	<i>CDKAL1</i>	rs7754840	C	0.57	0.65	0.46	1.54	2.18	0.02165	0.000012
6	<i>CDKAL1</i>	rs7756992	G	0.60	0.66	0.53	1.37	1.73	0.098	0.00256
8	<i>SLC30A8</i>	rs13266634	C	0.64	0.70	0.59	1.26	1.62	0.226	0.00947
9	<i>CDKN2A/2B</i>	rs564398	T	0.89	0.89	0.87	1.20	1.13	0.542	0.645
9	<i>CDKN2A/2B</i>	rs1333040	C	0.34	0.31	0.32	1.14	0.98	0.516	0.904
9	<i>CDKN2A/2B</i>	rs10757278	G	0.52	0.48	0.45	1.29	1.12	0.177	0.527
9	<i>CDKN2A/2B</i>	rs10811661	T	0.69	0.65	0.51	2.10	1.75	0.00015	0.00177
10	<i>HHEX</i>	rs1111875	C	0.44	0.37	0.30	1.77	1.35	0.00211	0.097
10	<i>HHEX</i>	rs5015480	C	0.28	0.24	0.19	1.71	1.38	0.00944	0.112
10	<i>HHEX</i>	rs7923837	G	0.33	0.27	0.21	1.88	1.40	0.00145	0.091
10	<i>TCF7L2</i>	rs7903146	T	0.04	0.06	0.02	1.60	2.52	0.332	0.014
10	<i>TCF7L2</i>	rs12255372	T	0.01	0.01	0.00	4.95	4.22	0.149	0.202
11	<i>KCNQ1</i>	rs2074196	G	0.63	0.61	0.58	1.22	1.14	0.308	0.458
11	<i>KCNQ1</i>	rs2237892	C	0.68	0.65	0.61	1.37	1.20	0.123	0.329
11	<i>KCNQ1</i>	rs2237895	A	0.72	0.35	0.70	1.10	1.27	0.643	0.189
11	<i>KCNJ11</i>	rs5215	G	0.41	0.65	0.38	1.13	1.11	0.523	0.565
11	<i>KCNJ11</i>	rs5219	A	0.42	0.64	0.38	1.18	1.10	0.399	0.587
16	<i>FTO</i>	rs8050136	C	0.88	0.91	0.88	1.05	1.36	0.863	0.299

T2DM Prediction Model Using GRS

- Model adjusted for Age, Prepregnancy BMI, Family History of DM, Mean Arterial Pressure, Fasting Glucose, Fasting Insulin

	Without wGRS	With wGRS
Age (years)		
OR (95% CI)	1.01 (0.95 - 1.07)	1.01 (0.95 - 1.07)
<i>P</i>	0.876	0.765
Family history of diabetes		
OR (95% CI)	1.49 (0.92 - 2.43)	1.53 (0.93 - 2.52)
<i>P</i>	0.107	0.094
Prepregnancy BMI		
OR (95% CI)	1.10 (1.02 - 1.19)	1.10 (1.02 - 1.19)
<i>P</i>	0.015	0.014
Mean arterial pressure		
OR (95% CI)	1.01 (0.98 - 1.04)	1.01 (0.98 - 1.04)
<i>P</i>	0.518	0.502
Fasting glucose		
OR (95% CI)	2.47 (1.79 - 3.40)	2.33 (1.69 - 3.20)
<i>P</i>	3.6×10^{-8}	2.4×10^{-7}
Log fasting insulin*		
OR (95% CI)	0.62 (0.19 - 2.02)	0.64 (0.18 - 2.21)
<i>P</i>	0.423	0.476
Weighted GRS		
OR (95% CI)	-	1.66 (1.30 - 2.13)
<i>P</i>	-	6.3×10^{-5}
Discriminatory capability		
C statistic (95% CI)	0.74 (0.69 - 0.80)	0.78 (0.72 - 0.83)
<i>P</i> for difference		0.015
Calibration accuracy		
χ^2	3.72	7.34
<i>P</i>	0.88	0.5
Reclassification improvement		
Continuous NRI (95% CI)	-	0.430 (0.218 - 0.642)
<i>P</i>	-	7.0×10^{-5}



CONTENTS

- Basic epidemiology of T2DM* and Prediabetes
- Diabetes Screening Guidelines
- Natural Course of Diabetes Development
- Risk Factors of Diabetes: Genetic and Environmental
- **Strategies for Early Diagnosis of T2DM**

*T2DM: Type 2 Diabetes

Strategies for Early Diagnosis of T2DM

1. Measurement of Beta Cell Dysfunction
2. Genetic Information (Family History)
3. Environmental Factors
 - Age
 - Obesity (BMI, Waist Circumference)
 - Diet, Physical Inactivity, Smoking, Alcohol
4. Simple Score Screening Tools
5. HbA1c

SIMPLE SCORE SCREENING TOOL

Question	Answer (Score)	Enter your Score (Enter 0 if you don't know)								
1. Your age group?	< 35 y (0 point) 35–44 y (2 points) ≥ 45 y (3 points)									
2. Have either of your parents or siblings been diagnosed with diabetes?	No (0 point) Yes (1 point)									
3. Are you currently taking medication for hypertension or do you have hypertension (i.e.g, blood pressures greater than 140/90 mmHg)?	No (0 point) Yes (1 point)									
4. What is your waist circumference (taken below the ribs, usually at the level of the navel) ?	<table border="0"> <tr> <td>Men</td> <td>Women</td> </tr> <tr> <td>< 84 cm (33 inch) (0 point)</td> <td>< 77 cm (30 inch) (0 point)</td> </tr> <tr> <td>84–89.9 cm (33–34.9 inch) (2 points)</td> <td>77–83.9 cm (30–32.9 inch) (2 points)</td> </tr> <tr> <td>≥ 90 cm (35 inch) (3 points)</td> <td>≥ 84 cm (33 inch) (3 points)</td> </tr> </table>	Men	Women	< 84 cm (33 inch) (0 point)	< 77 cm (30 inch) (0 point)	84–89.9 cm (33–34.9 inch) (2 points)	77–83.9 cm (30–32.9 inch) (2 points)	≥ 90 cm (35 inch) (3 points)	≥ 84 cm (33 inch) (3 points)	
Men	Women									
< 84 cm (33 inch) (0 point)	< 77 cm (30 inch) (0 point)									
84–89.9 cm (33–34.9 inch) (2 points)	77–83.9 cm (30–32.9 inch) (2 points)									
≥ 90 cm (35 inch) (3 points)	≥ 84 cm (33 inch) (3 points)									
5. Do you currently smoke cigarettes on a daily basis?	Never or Ex-smoker (0 point) Current smoker (1 point)									
6. How much alcohol do you drink on a daily basis? (regardless of types of alcohols)	Never or less than 1 drink / day (0 point) 1–4.9 drinks /day (1 point) ≥ 5 drinks /day (2 points)									
TOTAL SCORE (add points from questions 1–6)										
If the TOTAL SCORE is ≥5, you are at high risk for diabetes, so see your doctor for a blood test.										

Your total score from questionnaire	Probability of having diabetes now
≤ 4	2%
5~7	6%
8~9	12%
≥ 10	19%

Hemoglobin A_{1c} as a Diagnostic Tool for Diabetes Screening and New-Onset Diabetes Prediction

A 6-year community-based prospective study

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 HAK C. JANG, MD, PHD^{1,2}
 NAM H. CHO, MD, PHD³

indicated by the estimated 40% of people who have undiagnosed diabetes (1).
 The hemoglobin A_{1c} (A1C) level is

Table 2—Sensitivity, specificity, and positive and negative predictive value of increasing A1C cutoff levels for detecting undiagnosed diabetes and for predicting the incidence of type 2 diabetes at the 6-year follow-up

A1C cutoff (%)	Baseline undiagnosed diabetes				Incident diabetes after 6 years of follow-up			
	Predictive value				Predictive value			
	Sensitivity	Specificity	Positive	Negative	Sensitivity	Specificity	Positive	Negative
5.0 (−1.00 SDs above normal mean)	0.972	0.115	0.074	0.982	0.962	0.121	0.162	0.947
5.1 (−0.75 SDs above normal mean)	0.956	0.185	0.079	0.98	0.935	0.198	0.171	0.945
5.2 (−0.50 SDs above normal mean)	0.945	0.279	0.087	0.986	0.886	0.302	0.184	0.937
5.3 (−0.25 SDs above normal mean)	0.915	0.390	0.098	0.984	0.827	0.419	0.201	0.932
5.4 (0.00 SDs above normal mean)	0.887	0.506	0.115	0.984	0.768	0.547	0.231	0.930
5.5 (0.25 SDs above normal mean)	0.866	0.616	0.141	0.984	0.682	0.665	0.265	0.922
5.6 (0.50 SDs above normal mean)	0.822	0.717	0.174	0.982	0.594	0.769	0.313	0.914
5.7 (0.75 SDs above normal mean)	0.770	0.797	0.216	0.979	0.508	0.847	0.370	0.907
5.8 (1.00 SDs above normal mean)	0.720	0.862	0.274	0.977	0.420	0.908	0.448	0.898
5.9 (1.25 SDs above normal mean)	0.676	0.907	0.344	0.975	0.333	0.947	0.527	0.889
6.0 (1.50 SDs above normal mean)	0.619	0.935	0.411	0.971	0.263	0.967	0.586	0.881
6.2 (2.00 SDs above normal mean)	0.523	0.968	0.544	0.965	0.152	0.987	0.677	0.868
6.6 (3.00 SDs above normal mean)	0.372	0.992	0.771	0.956	0.051	0.999	0.885	0.856

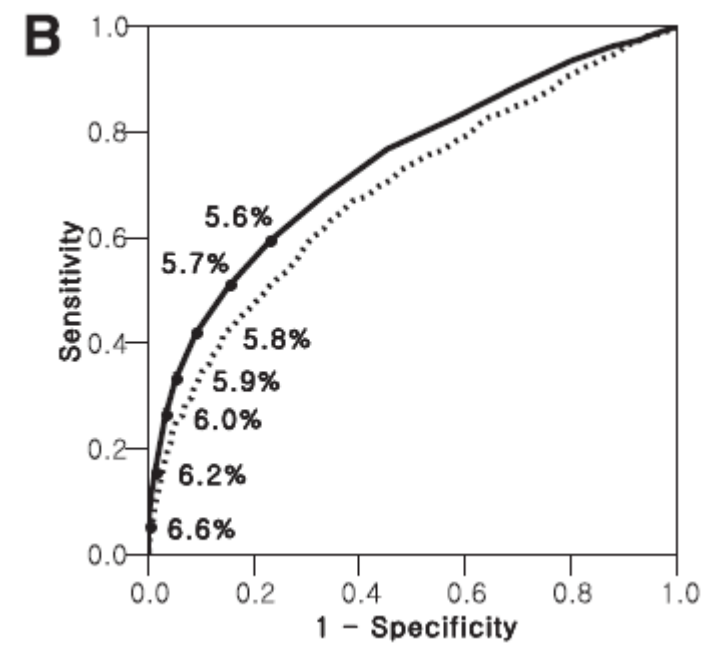
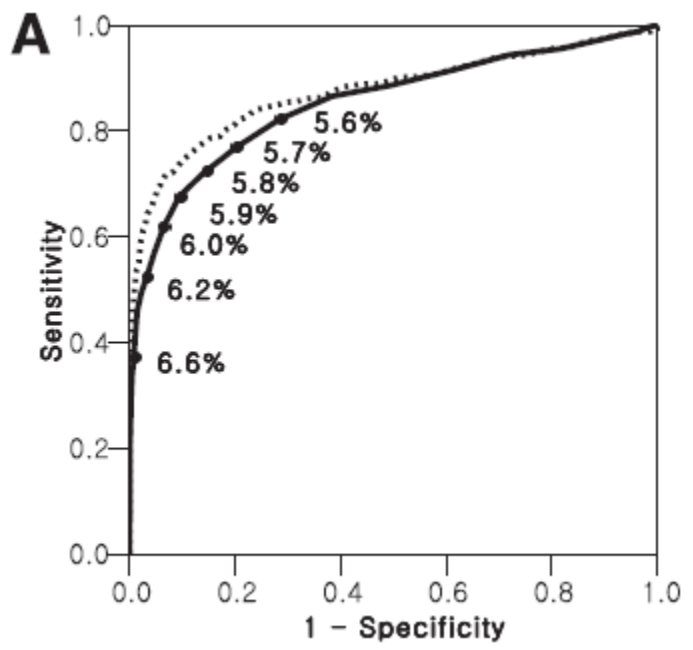
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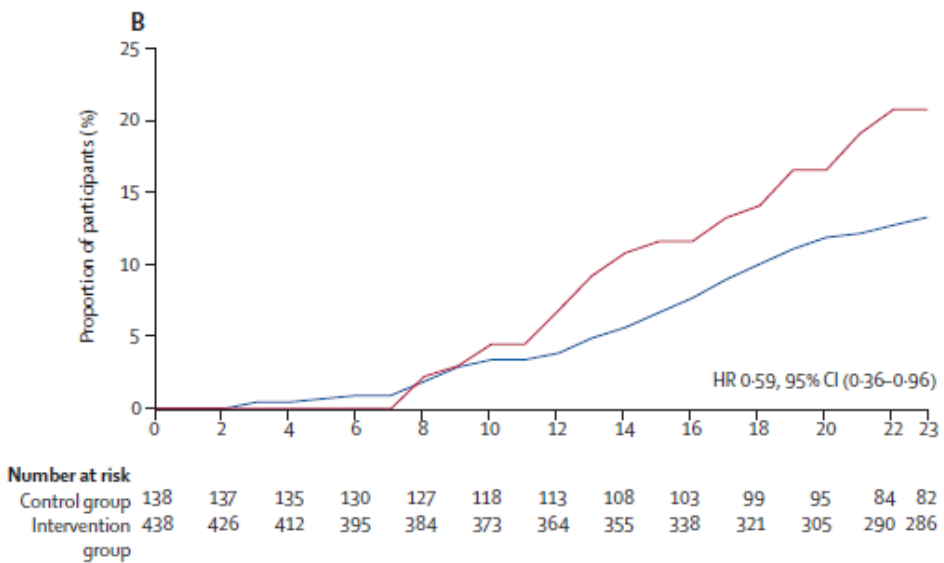
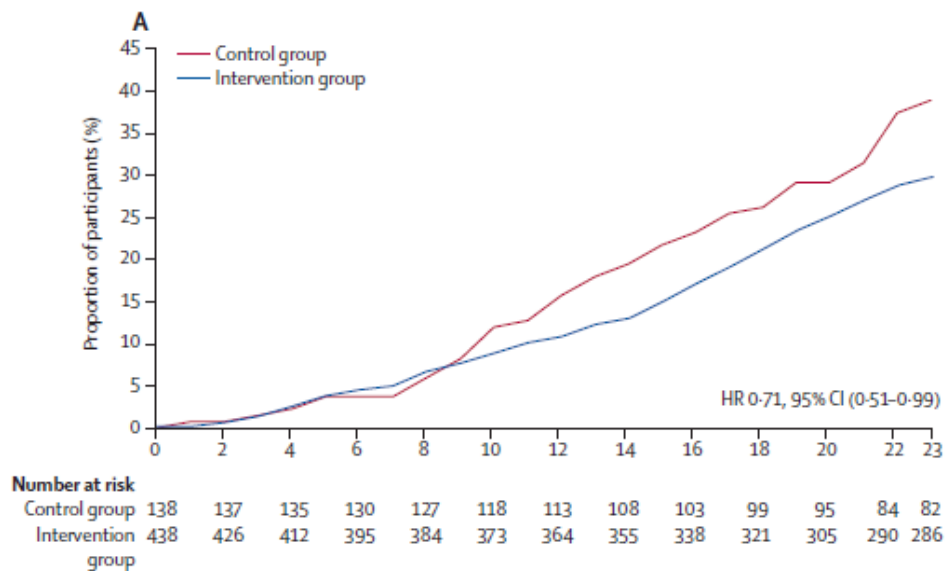
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OUTCOME OF DIABETES SCREENING?



SUMMARY

- 우리나라 당뇨병 유병율은 30세 이상 성인에서 12.4%에 이르며 공복혈당 장애의 유병율은 19.3%임. 당뇨병 환자 수는 2050년 경에는 약 591만명으로 현재 보다 약 1.9배 정도 증가할 것으로 추정됨
- 국내외 당뇨병 관련 선별 권고안이 있으나 목적 및 실정에 맞게 변형이 필요함
- 한국인의 전향적 코호트 연구에 의하면 나이가 증가함에 따라 인슐린 감수성이 감소하며 이를 보상하기 위한 베타세포 기능의 대상성 증가가 나타나지 않는 것이 당뇨병 발병의 중요한 병태 생리 기전임. 베타세포 관련 기능은 유전적 소인에 의하여 일부 결정됨.
- 당뇨병 발병의 위험 요소로 유전적 소인, 임신성 당뇨병의 과거력, 인슐린 분비 능력의 저하, 비만 등을 고려해야 함
- 당뇨병 선별을 위하여 고위험 요소를 파악하는 것이 중요하고 특히 간편선별 도구를 이용하는 것이 좋은 방법이 될 수 있음. 유전적 소인에 대한 연구는 아직 더 많은 연구가 필요하며 선별시 당화혈색소 등의 검사를 사용하는 것이 도움이 됨.