



67° CONGRESSO NAZIONALE SIGG

LA LONGEVITÀ DECLINATA AL FEMMINILE

Diabete e Gender

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SOCIETÀ ITALIANA
DI GERONTOLOGIA
E GERIATRIA

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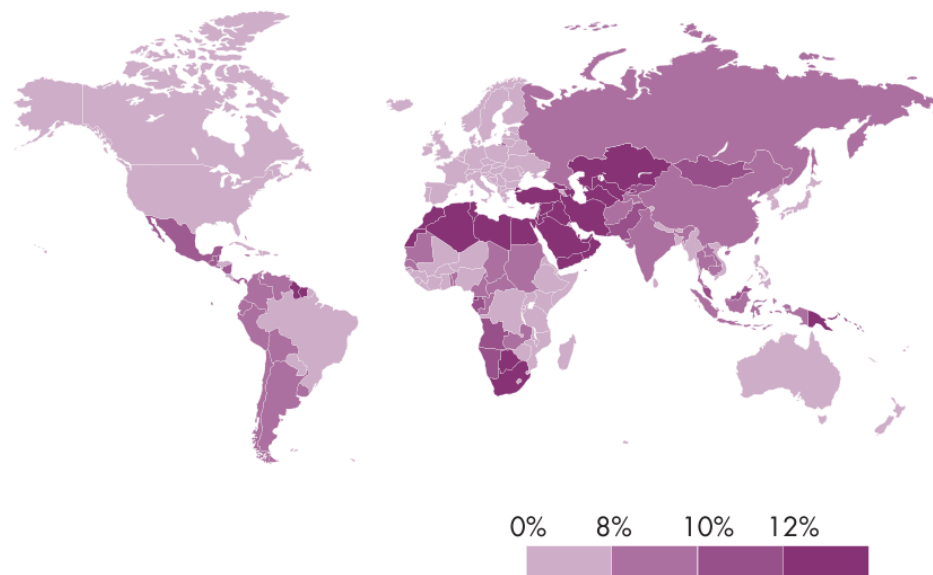
Diabetes around the world in 2021



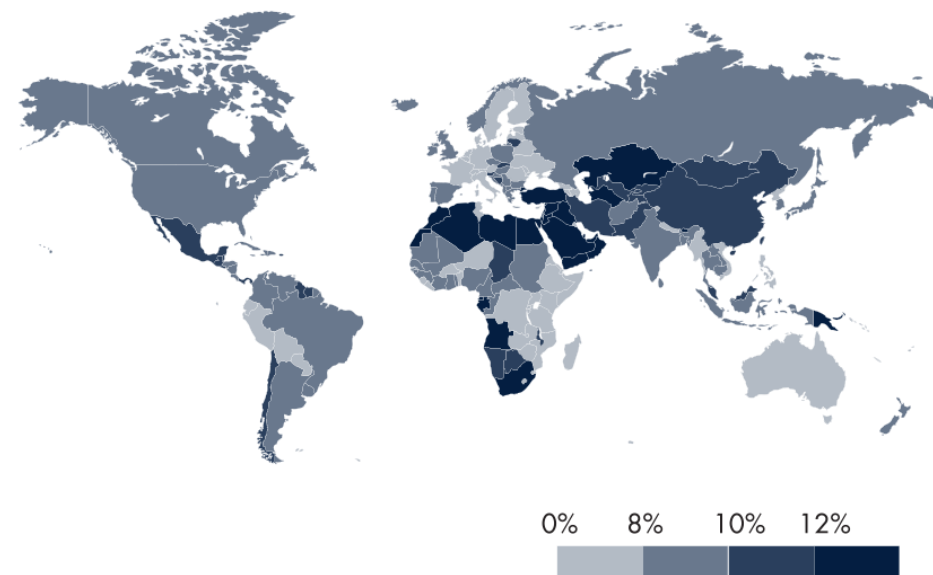


Diabetes prevalence in woman and men

Women

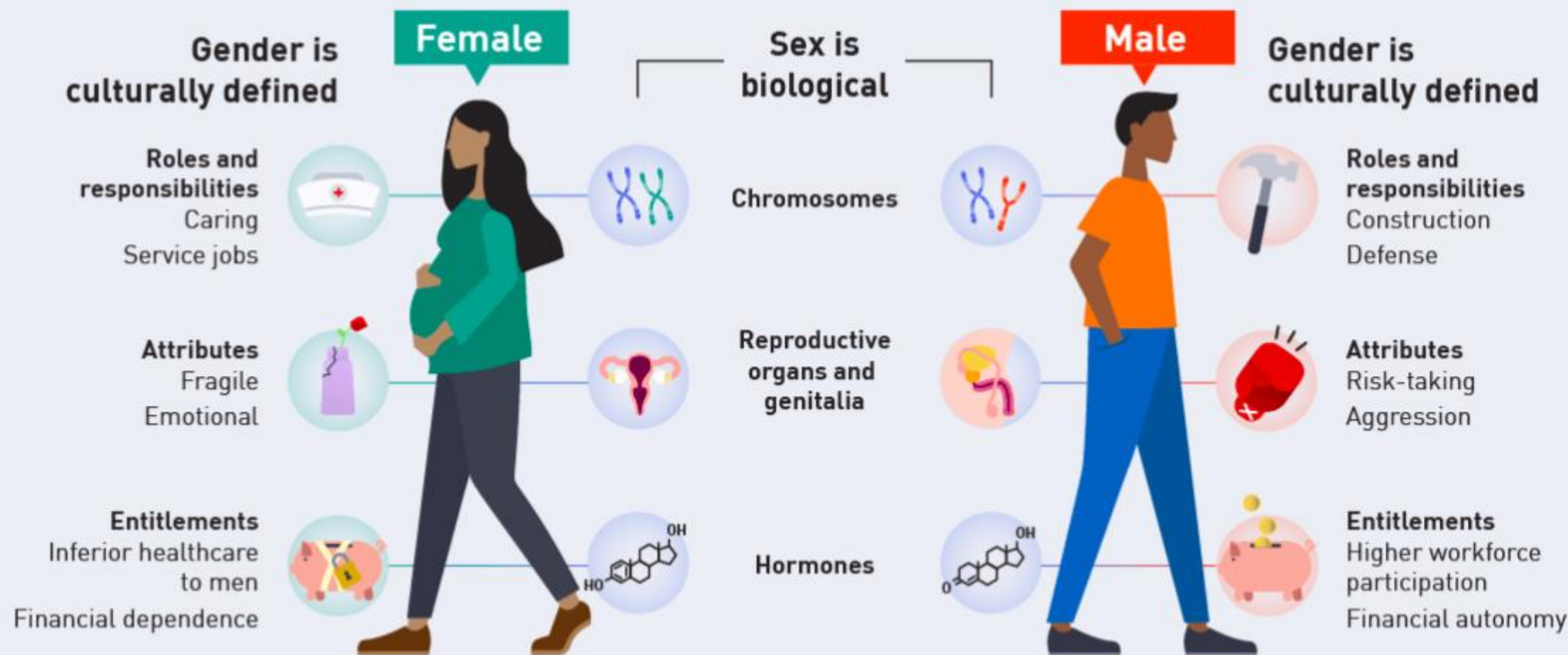


Men



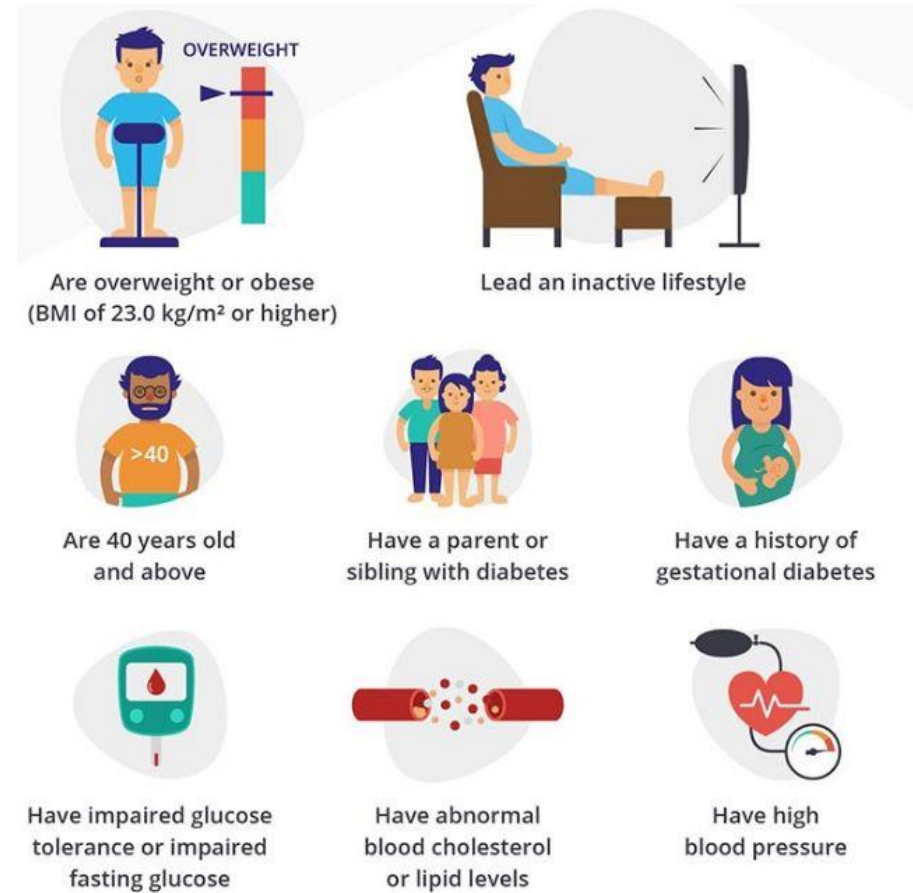


Sex vs Gender



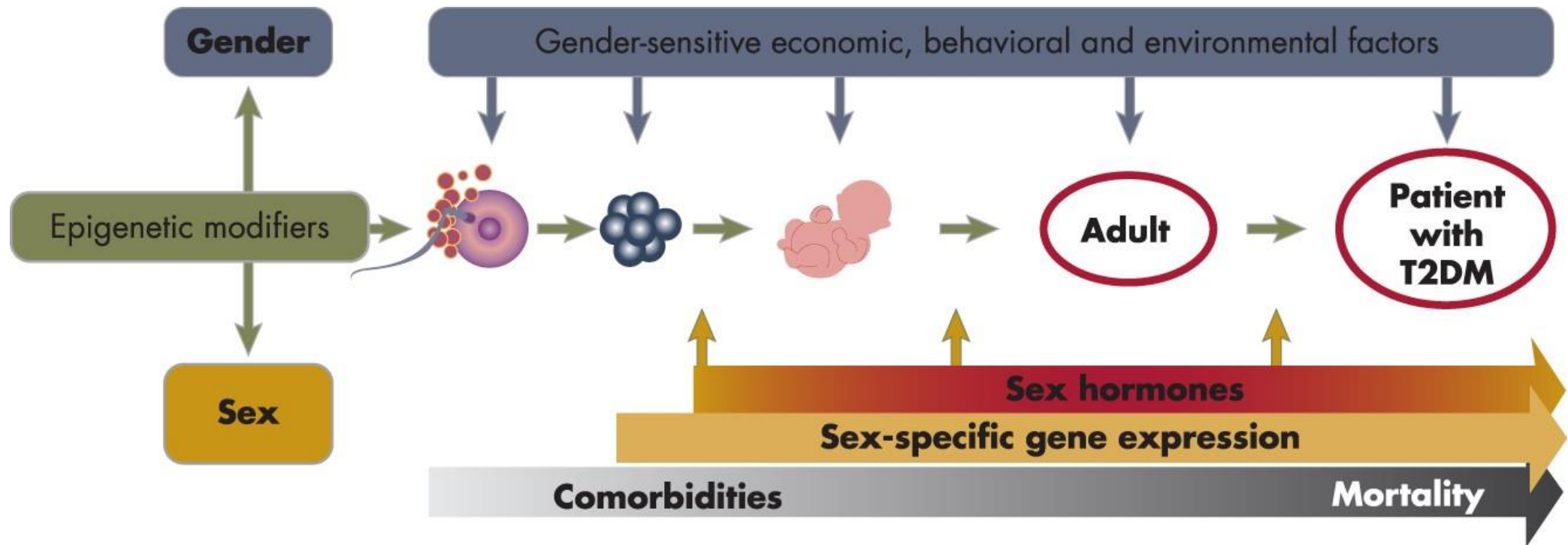


- **Biological risk factors**
- Psychological risk factors
- Genetic predisposition
- Diabetes complications
- Antidiabetic Agent





Lifelong impact and interaction between sex and gender on development and outcomes of Type 2 Diabetes Mellitus (T2D)



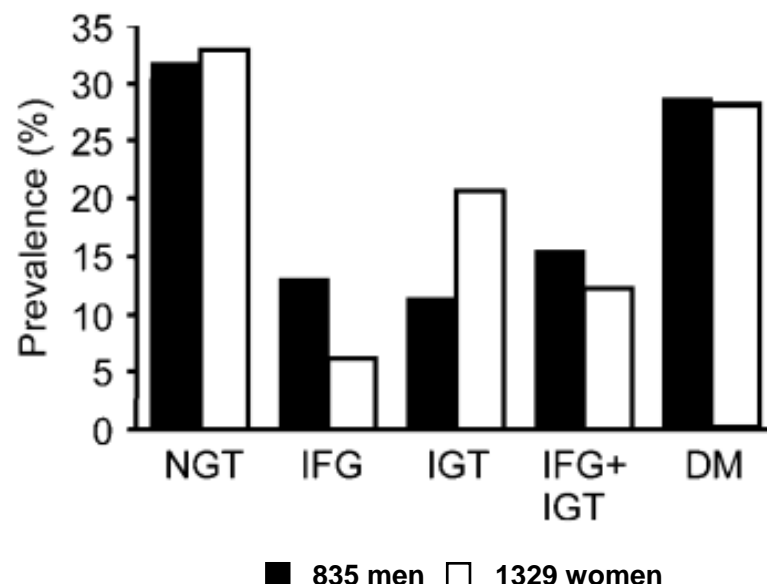


Sex and prevalence of impaired fasting glucose, impaired glucose tolerance and T2DM

- AusDiabStudy
- Inter99 Study
- GENNID Study Group

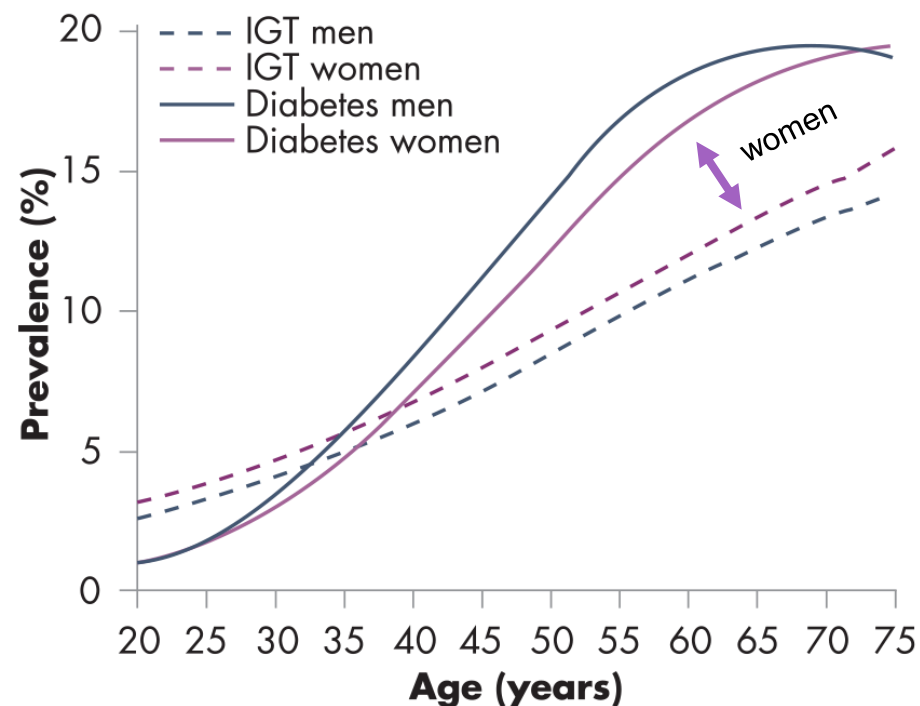


Isolated IFG was more frequent in men (13.1 vs. 6.2%), whereas **isolated IGT** was more frequent in women (20.7 vs. 11.3%).





Prevalence of prediabetes and diabetes in women



Women show an acceleration of developing diabetes after menopause



The associations of menopausal age and reproductive life span (menopausal age minus menarcheal age) with diabetes risk: results from the EPIC-INTERACT STUDY

		HR (95% CI)			
	N total/cases	Crude	Model 1	Model 2	Model 3
Menopausal age (years)					
<40	419/220	1.50 (1.22–1.85)	1.50 (1.22–1.85)	1.28 (1.00–1.64)	1.32 (1.04–1.69)
40–44	887/424	1.19 (1.02–1.38)	1.18 (1.02–1.37)	1.08 (0.89–1.30)	1.09 (0.90–1.31)
45–49	2,570/1,186	1.05 (0.95–1.17)	1.05 (0.95–1.17)	0.97 (0.86–1.10)	0.97 (0.86–1.10)
50–54	3,333/1,554	Ref. (1.00)	Ref. (1.00)	Ref. (1.00)	Ref. (1.00)
≥55	655/307	0.91 (0.77–1.08)	0.92 (0.77–1.08)	0.84 (0.69–1.02)	0.85 (0.70–1.03)
Menopausal age (per SD decrease)	7,864/3,691	1.11 (1.06–1.16)	1.11 (1.06–1.16)	1.07 (1.01–1.13)	1.08 (1.02–1.14)
Reproductive life span (years)					
Quartile 1 (<33)	1,982/959	1.11 (0.97–1.28)	1.11 (0.97–1.28)	1.16 (0.97–1.38)	1.17 (0.98–1.39)
Quartile 2 (33–36)	2,364/1,077	0.94 (0.82–1.08)	0.94 (0.82–1.08)	1.00 (0.85–1.19)	1.00 (0.85–1.19)
Quartile 3 (37–39)	1,979/897	0.91 (0.79–1.04)	0.91 (0.79–1.04)	0.97 (0.82–1.14)	0.96 (0.82–1.14)
Quartile 4 (≥40)	1,443/710	Ref. (1.00)	Ref. (1.00)	Ref. (1.00)	Ref. (1.00)
Reproductive life span (per SD decrease)	7,768/3,643	1.07 (1.02–1.12)	1.07 (1.02–1.12)	1.06 (1.00–1.12)	1.06 (1.01–1.12)



Testosterone deficiency and T2DM in men

Endogenous Sex Hormones and the Development of Type 2 Diabetes in Older Men and Women: the RANCHO BERNARDO STUDY

Independent variables†	Covariates‡	Men		Women	
		β	P	β	P
Testosterone	Model 1	-0.37	0.001	-0.023	0.71
	Model 2	-0.24	0.02	0.020	0.76
	Model 3	-0.31	0.02	0.011	0.88
Bioavailable testosterone	Model 1	-0.10	0.39	0.15	0.003
	Model 2	-0.13	0.23	0.15	0.01
	Model 3	-0.15	0.28	0.15	0.02
Estradiol	Model 1	-0.13	0.24	0.035	0.61
	Model 2	-0.08	0.47	0.054	0.49
	Model 3	-0.05	0.64	0.025	0.77
Bioavailable estradiol	Model 1	-0.03	0.79	0.14	0.01
	Model 2	-0.07	0.49	0.15	0.02
	Model 3	-0.05	0.66	0.14	0.055



Testosterone excess and T2DM in women

Diabetologia

Plasma sex steroid hormones and risk of developing type 2 diabetes in women: a prospective study

Analysis	Oestradiol ^a		Free oestradiol ^b		Testosterone ^c		Free testosterone ^d	
	RR (95% CI)	<i>p</i> value for trend	RR (95% CI)	<i>p</i> value for trend	RR (95% CI)	<i>p</i> value for trend	RR (95% CI)	<i>p</i> value for trend
Multivariate	1.97 (1.28–3.04)	0.002	2.78 (1.89–4.10)	<0.001	1.48 (1.01–2.17)	0.05	2.30 (1.60–3.30)	<0.001
Multivariate (excluding first 3 years follow-up)	1.88 (1.09–3.25)	0.02	2.97 (1.82–4.85)	<0.001	1.46 (0.92–2.34)	0.11	2.60 (1.63–4.14)	<0.001
Multivariate +HDL + TG ^e	2.08 (0.93–4.65)	0.07	2.91 (1.47–5.73)	0.002	1.72 (0.92–3.20)	0.09	2.38 (1.14–4.97)	0.02
Multivariate + waist circumference	1.80 (1.15–2.83)	0.01	2.55 (1.72–3.78)	<0.001	1.43 (0.93–2.20)	0.11	2.14 (1.48–3.10)	<0.001
Multivariate + C-reactive protein	2.25 (1.39–3.66)	0.001	2.70 (1.78–4.10)	<0.001	1.33 (0.88–2.02)	0.18	1.99 (1.31–3.03)	0.001
Multivariate (among those with HbA _{1c} <6%)	2.00 (1.18–3.36)	0.009	3.07 (1.85–5.09)	<0.001	1.39 (0.86–2.24)	0.17	2.40 (1.49–3.88)	<0.001

Results

In PostMenopausal women, higher plasma levels of oestradiol and testosterone were strongly and prospectively related to increased risk of developing type 2 diabetes.



Gestational Diabetes and T2DM

Persistence of Risk for Type 2 Diabetes After Gestational Diabetes Mellitus

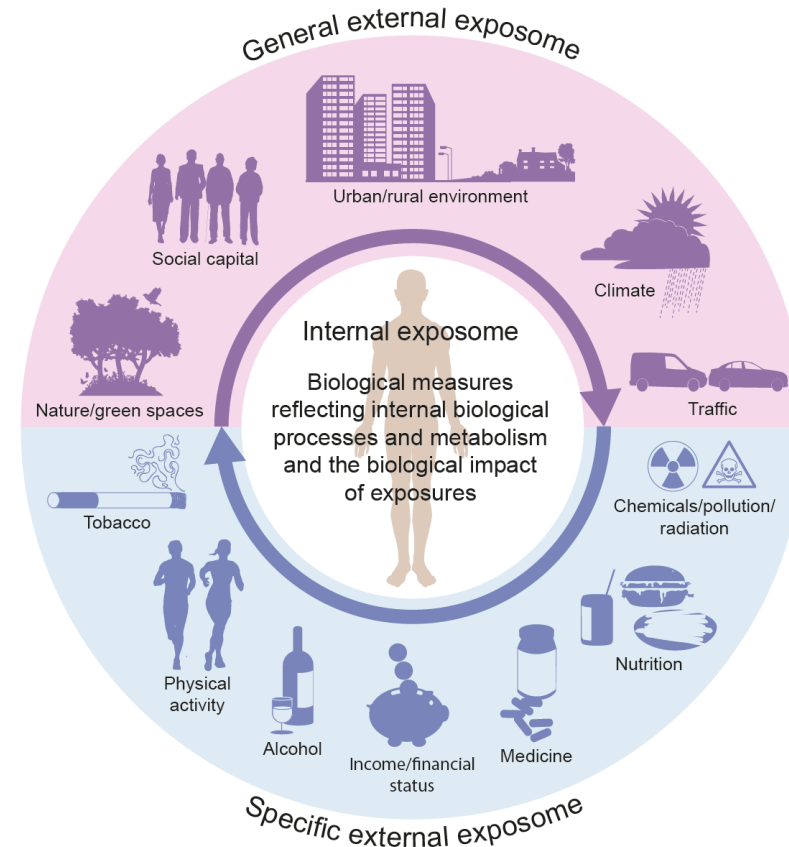
Conclusions

Gestational diabetes mellitus predicted markedly increased rates of type 2 diabetes.

	Ever having pregnancy with GDM, HR (95% CI)		Time since last pregnancy with GDM, HR (95% CI)
	No GDM	GDM	
Person-years	462,042	10,958	—
Participants, <i>n</i>	45,357	1,172	—
Model 1: model with no time since GDM interaction	1 (ref)	2.50 (2.15–2.91)	—
Model 2: model with time since GDM interaction term† GDM effect†	1 (ref)	5.07 (3.36–7.65)	0.76 (0.66–0.88)
Model 3: model stratified by time since last pregnancy with GDM, years			
6–15	1 (ref)	3.87 (2.60–5.75)	—
16–25	1 (ref)	3.50 (2.79–4.40)	—
26–35	1 (ref)	1.95 (1.46–2.61)	—
>35	1 (ref)	1.62 (1.12–2.33)	—



- Biological risk factors
- **Psychological risk factors**
- Genetic predisposition
- Diabetes complications
- Antidiabetic Agent





Work stress and T2DM

Work Stress, Obesity and the Risk of Type 2 Diabetes: Gender-Specific Bidirectional Effect in the Whitehall II Study

Among women, work stress was associated with higher risk of T2DM in the obese (HR 2.01: 1.06; 3.92) but not in the nonobese ($P_{\text{INTERACTION}} = 0.005$)

All men		Nonobese men (BMI <30 kg/m ²)		Obese men (BMI ≥30 kg/m ²)		P for interaction ^a
Cases/total	HR (95% CI)	Cases/total	HR (95% CI)	Cases/total	HR (95% CI)	
389/3,689	0.80 (0.63; 1.02)	310/3,429	0.70 (0.53; 0.93)	79/260	1.05 (0.63; 1.75)	0.17
All women		Nonobese women (BMI <30 kg/m ²)		Obese women (BMI ≥30 kg/m ²)		P for interaction ^a
Cases/total	HR (95% CI)	Cases/total	HR (95% CI)	Cases/total	HR (95% CI)	
151/1,449	1.37 (0.98; 1.92)	104/1,248	1.18 (0.63; 2.10)	47/201	2.01 (1.06; 3.82)	0.005



- Biological risk factors
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**GENETIC
PREDISPOSITION**



Sexual dimorphism in genetic predisposition

Fetal sex influences gene expression and produces functional differences in the human placentas that increase the susceptibility to developing type 2 diabetes in the offspring.

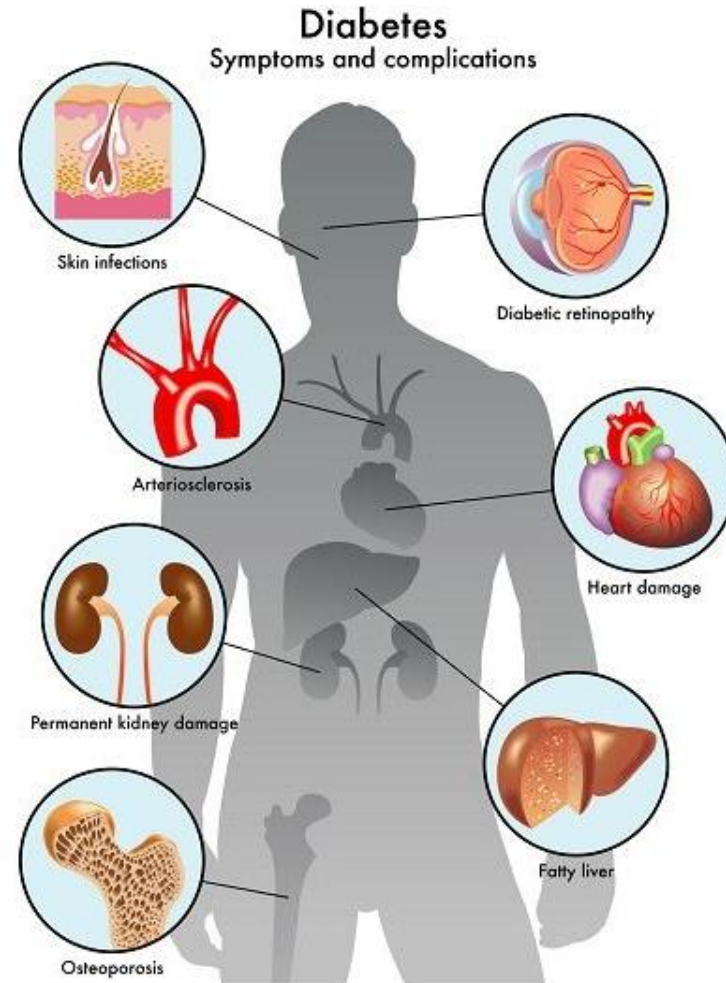
SNP	Trait	Chr	Locus	EA*	EAF	Sex-combined			Women			Men			Sex diff.
						β	P	N	β	P	N	β	P	N	
Loci achieving genome-wide significance in European-ancestry meta-analyses															
rs10925060	WCadjBMI	1	OR2W5- NLRP3	T	0.03	0.017	2.2×10^{-5}	140,515	0.002	6.8×10^{-1}	85,186	0.045	9.1×10^{-13}	55,522	1.7×10^{-8}
rs10929925	HIP	2	SOX11	C	0.55	0.020	4.5×10^{-8}	207,648	0.021	9.0×10^{-6}	115,428	0.018	3.2×10^{-4}	92,499	6.1×10^{-1}
rs2124969	WCadjBMI	2	ITGB6	C	0.42	0.020	7.1×10^{-9}	231,284	0.016	3.5×10^{-4}	127,437	0.025	2.3×10^{-7}	104,039	1.4×10^{-1}
rs17472426	WCadjBMI	5	CCNJL	T	0.92	0.014	3.1×10^{-2}	217,564	-0.014	1.0×10^{-1}	119,804	0.052	4.3×10^{-8}	97,954	3.9×10^{-8}
rs7739232	HIPadjBMI	6	KLHL31	A	0.07	0.037	5.4×10^{-5}	131,877	0.063	1.0×10^{-8}	80,475	-0.004	7.5×10^{-1}	51,589	2.9×10^{-5}
rs13241538	HIPadjBMI	7	KLF14	C	0.48	0.017	1.6×10^{-6}	210,935	0.033	9.9×10^{-14}	117,210	-0.003	5.0×10^{-1}	93,911	2.0×10^{-9}
rs7044106	HIPadjBMI	9	C5	C	0.24	0.023	4.1×10^{-5}	143,412	0.039	5.7×10^{-9}	86,733	-0.003	6.9×10^{-1}	56,865	1.3×10^{-5}
rs11607976	HIP	11	MYEOV	C	0.70	0.022	4.2×10^{-8}	212,815	0.019	1.9×10^{-4}	118,391	0.024	7.7×10^{-6}	94,701	4.4×10^{-1}
rs1784203	WCadjBMI	11	KIAA1737	A	0.01	0.031	1.3×10^{-8}	63,892	0.000	9.9×10^{-1}	35,539	0.075	1.0×10^{-19}	28,353	1.2×10^{-1}
rs1394461	WHR	11	CNTN5	C	0.25	0.017	4.7×10^{-4}	144,349	0.035	3.6×10^{-8}	87,441	-0.011	1.6×10^{-1}	57,094	1.1×10^{-6}
rs319564	WHR	13	GPC6	C	0.45	0.014	3.4×10^{-5}	212,137	0.003	5.3E-01	117,970	0.027	1.6×10^{-8}	94,350	6.0×10^{-5}
rs2047937	WCadjBMI	16	ZNF423	C	0.50	0.019	4.7×10^{-8}	231,009	0.022	5.5×10^{-7}	127,288	0.014	3.6×10^{-3}	103,914	2.0×10^{-1}
rs2034088	HIPadjBMI	17	VPS53	T	0.53	0.021	4.8×10^{-9}	210,737	0.028	9.6×10^{-10}	117,142	0.014	6.5×10^{-3}	93,781	2.5×10^{-2}
rs1053593	HIPadjBMI	22	HMGXB4	T	0.65	0.021	3.9×10^{-8}	202,070	0.029	1.8×10^{-9}	114,347	0.011	5.1×10^{-2}	87,908	6.2×10^{-3}
Loci achieving genome-wide significance in all-ancestry meta-analyses															
rs1664789	WCadjBMI	5	ARL15	C	0.41	0.014	2.6×10^{-5}	244,110	0.005	2.8×10^{-1}	133,052	0.026	3.6×10^{-8}	109,025	4.4×10^{-4}
rs722585	HIPadjBMI	6	GMDS	G	0.68	0.015	2.1×10^{-4}	205,815	-0.001	8.8×10^{-1}	113,965	0.032	9.2×10^{-9}	89,831	4.3×10^{-6}
rs1144	WCadjBMI	7	SRPK2	C	0.34	0.019	3.1×10^{-8}	239,342	0.020	1.2×10^{-5}	131,398	0.018	4.1×10^{-4}	105,911	7.8×10^{-1}
rs2398893	WHR	9	PTPDC1	A	0.71	0.020	4.0×10^{-8}	226,572	0.019	5.1×10^{-5}	124,577	0.019	2.7×10^{-4}	99,968	9.5×10^{-1}
rs4985155‡	HIP	16	PDXDC1	A	0.66	0.018	4.5×10^{-7}	227,296	0.011	1.6×10^{-2}	125,048	0.029	9.7×10^{-9}	100,313	6.3×10^{-3}

Results:

Sex-specific differences were found in genetic loci, which are involved in regulatory functions of adipose tissue and insulin biology

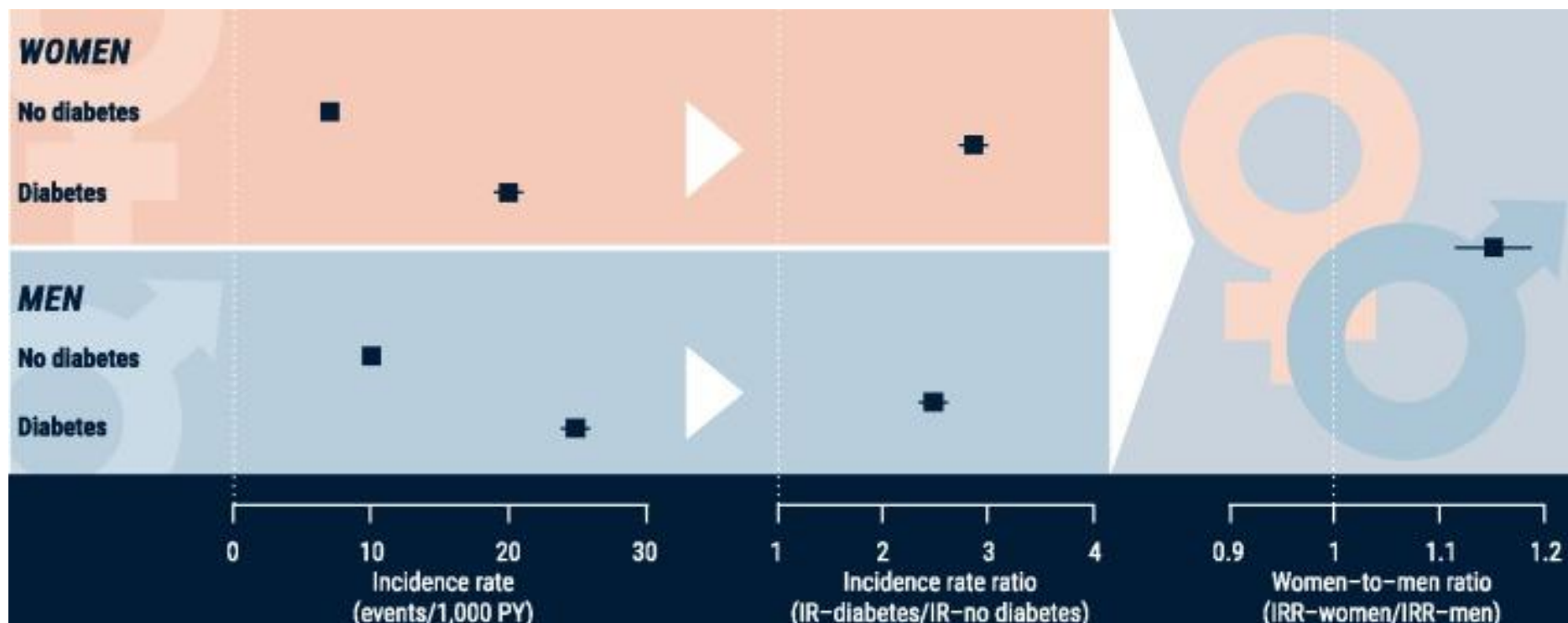


- Biological risk factors
- Psychological risk factors
- Genetic predisposition
- **Diabetes complications**
- Antidiabetic Agent



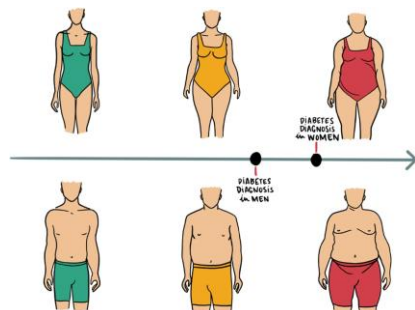


Women has major risk to develop cardiovascular diseases

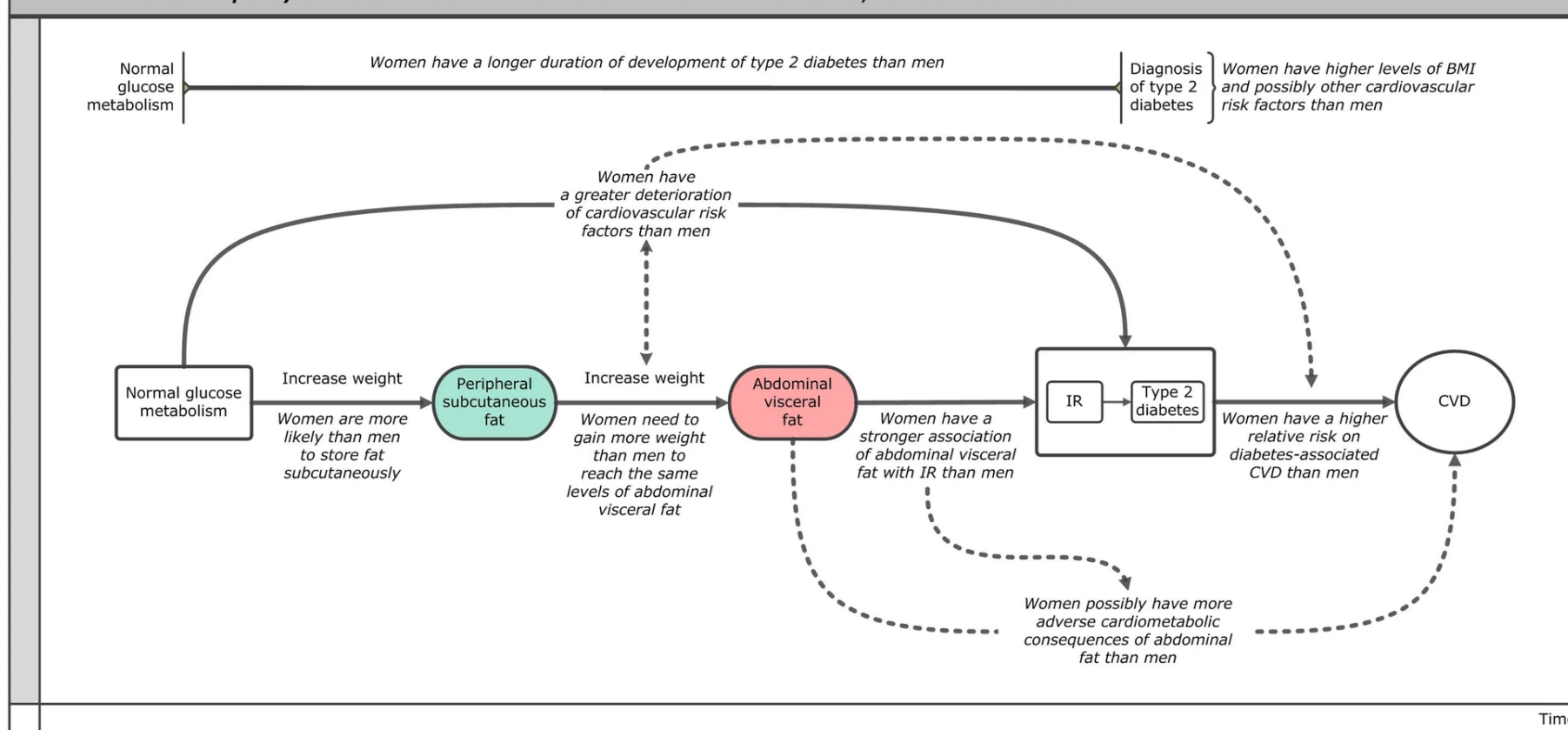




Sex differences in the risk of vascular disease associated with diabetes

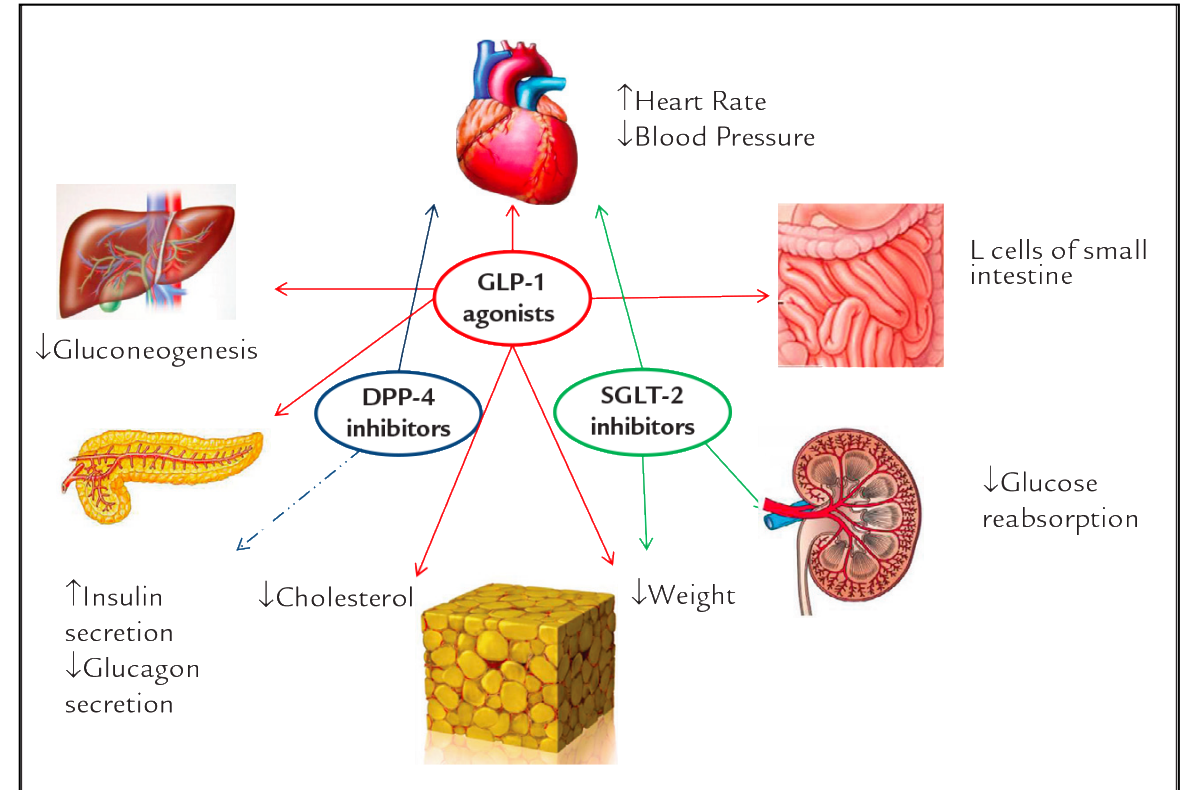


Sex differences in adiposity in association with diabetes and cardiovascular disease; women versus men





- Biological risk factors
- Psychological risk factors
- Genetic predisposition
- Diabetes complications
- **Antidiabetic Agent**





Long-Term Effectiveness of Liraglutide for Weight Management and Glycemic Control in Type 2 Diabetes

	B	Beta	T	<i>p</i> -Value
BMI	0.387	0.380	2.533	0.016
Female gender	5.086	0.365	2.420	0.020
* Female gender	6.459	0.464	2.975	0.005

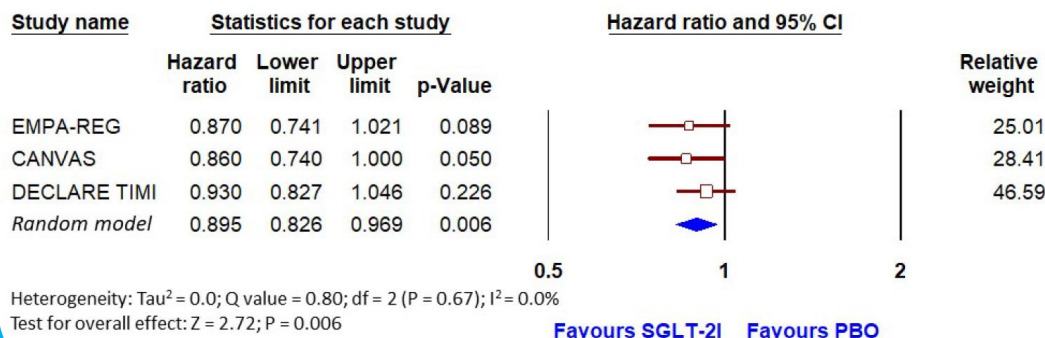
Conclusions:

Prolonging treatment with Liraglutide can lead to durable benefits in relation to weight and glycemic control, with a greater impact on women.

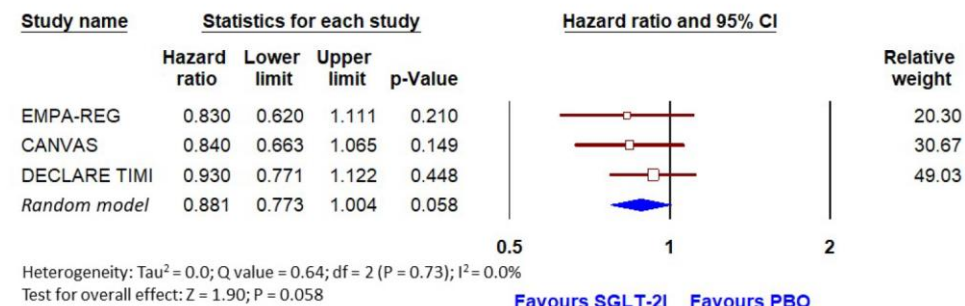


Gender difference in Cardiovascular Outcomes with SGLT-2 inhibitors and GLP-1 receptor agonist in Type 2 Diabetes: A Systematic Review and Meta-analysis of Cardio-Vascular Outcome Trials

MACE outcome in Male on SGLT-2i: A Meta-analysis of CVOTs



MACE outcome in Female on SGLT-2i: A Meta-analysis of CVOTs





Conclusions

- Sex is a fundamental biological factor, which plays a key role in regulation of homeostasis in health and causes vulnerability to cardiometabolic risk factors, as well as manifestation, clinical picture, and management of T2DM;
- The care of diabetic pregnancy demands special attention, because this vulnerable phase programs health of offspring even in a sex-specific way;
- Psychosocial factors also impact development and progression of diabetes and coping in a gender-dimorphic way.



Modern personalized treatment has to consider differences in biological factors, like genetic predisposition, sex hormones, and neurohumoral pathways, as well as behavioral and environmental differences between men and women