SUPPLEMENTARY DATA

Box S1

Operational definition of clinical trial and types of studies

Consistent with the World Health Organization (WHO) and the International Committee of Medical Journal Editors (ICMJE) for purposes of registration, ^{1,2} a clinical trial is defined here as any experimental or interventional study that prospectively assigns subjects (individually or clusters) to an intervention(s), with or without concurrent comparison or control groups, to evaluate the effects on biomedical outcomes (inflammatory biomarkers). The term "prospectively assigns" refers to a pre-defined process specified stipulating the assignment of research subjects to one or more arms of the clinical trial. To be prospectively assigned, there need not be randomisation or a control group. The term "intervention" is defined as a manipulation of the individual's home built environment (BE) for modifying the health-related endpoint (see Box S2).

Therefore, we will include all types of clinical trials, experimental studies (including randomised and pseudo-randomised, controlled and uncontrolled trials, etc.) We adhere to the Cochrane Handbook,³ and we refer to non-randomised trials as a quantitative experimental study design in which individuals (or households) are prospectively assigned to an intervention(s) using methods that differ from the randomisation procedure. Randomised controlled trials (RCTs) are prospective, experimental study design involving random allocation of participants to interventions; that includes two-arm parallel, cluster, crossover, stepped-wedge RCTs, etc. The term "controlled" refers to the presence of a concurrent control or comparator group. We will include any clinical trial, regardless of whether the blind method is used or not (e.g., open-label, single-, double-blind, etc.). Mixed-methods studies employing quantitative data will be included if meeting the inclusion criteria.

Observational studies, that means non-experimental or non-interventional studies, including case reports, case series, cross-sectional, case—control, cohort, ecological, etc., will be excluded.⁵ Other studies in which the intervention is not manipulated experimentally to change the home built environment (i.e., policy intervention) and provided or assigned by a research team will be excluded.⁵ Pilot and feasibility studies of clinical trials will be considered if meeting eligibility criteria.

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Box S2

Home built environment definition and specification for eligibility criteria.

The umbrella term Home Built Environment and its acronym (H-BE) is used in this study protocol to refer to the BE of homes.

The H-BE as place of usual residency is defined as a housing facility for a person living on their own or with a family group or non-relative members –regardless of the dwelling type, tenure status and housing conditions– for an indefinite period. This broad definition will embrace private dwellings, public housing facilities, sheltered housing, retirement homes and recreated naturalistic home settings.

A residential healthcare facility provides a home-like atmosphere, spaces, and recreational activities, facilitate communal living spaces for long-term care, and provide overnight permanent accommodation. However, these facilities will not be considered in this study as they do not reflect typical domestic microenvironments in terms of resident's behaviour, activity-time-space patterns and spatial configuration. Health care environments which offer rehabilitations interventions in which the patient spent less than 24-hour period or short stays (i.e., day hospitals, day centres) will not be considered as H-BE and will be excluded. The same applies to studies carried out in institutional settings of medium- and long-term residence (i.e., hospitals, prisons). A BE that provides accommodation for a finite period (i.e., college student settings, campus dormitories) will be also excluded.

Studies carried out in experimental settings will be left out as we seek investigations evaluating household environmental conditions in naturalistic home settings. But those conducted in recreated naturalistic settings that simulate a conventional home space(s) will be considered for inclusion. Research carried out across several settings (i.e., college student settings and private homes) will be included if quantitative data is clearly differentiated between the settings.

Box S3

Strategy for involvement of non-academic GP and eligibility criteria.

A group of non-academic, English or Spanish speaking, practising GPs will be actively involved in conducting the screening process for study selection in cooperation with the first reviewer (EHG). Non-academic GPs have showed willingness to participate in research studies if these address topics of high relevance to their daily practice, specially caring for multimorbid patients at home. The effectiveness of H-BE interventions on inflammatory status in community-dwelling adults is a topic of particular interest of GPs for caring patients with long-term and chronic conditions, and consequently, we expect a high level of GP's willingness to participate in this research study.

Involvement of GPs will be sought through different strategies. In a first round, they will be invited to participate through online cancer community and primary care networks. The use of Primary Care Networks has showed the highest recruitment rate (41%) with no financial and human resources, compared with published ads, personal visits in practice and calls. Our involvement activity will be disseminated in regular newsletters and/or social media platforms from different cancer and primary care networks- e.g., Cancer and Primary Care Research International Network (Ca-PRI), Royal College of General Practitioners (RCGP), Doctors.net, Cancer Research UK (CRUK), Spanish Society of Primary Care Physicians (SEMERGEN), Spanish Society of Family and Community Medicine (semFYC), etc. If there is low initial response, we will use snowball sampling methods that are successful used to recruit hard-to-reach subjects in research studies. Initial GPs involved in the first round that meet eligibility criteria will recruit others from their surgeries or social network who also meet the study inclusion criteria. This strategy will be supplemented by a third round, in which a convenience sample of GPs will be first approached and recruited if meeting the eligibility criteria. Primary care settings will be asked to gain permission to mail invitations to their clinician staff.

Once both GPs show their willingness to participate, a cooperation agreement accompanied by further material about the study and involvement activity will be provided. They will be provided with a complete list of the 287 biomarkers and H-BE interventions, including variants and abbreviations. The signed cooperation agreement will be considered as a successful inclusion.

Each GP group member involved in the research will screen no more than 300 titles and abstracts of the total number of identified papers in a first phase, in duplicate with the first reviewer (EHG). Then, they will be asked for supporting the screening process of the remaining studies.

Inclusion criteria

We will include a primary care physician defined as (1) currently practising as a GP, family practitioner or general internist, (2) being able to read, write and understand English language, (3) meeting technical requirements (internet connection), (4) willing to participate in a short session about the H-BE interventions and screening process in Rayyan software, and (5) signed cooperation agreement.

Quality control mechanisms into the screening process

The following mechanisms for quality controls into the screening process conducted by non-experienced GPs will be implemented:

- Before screening begins, GPs will (1) be provided of an abstract screening tool
 consisting of the set of eligibility criteria organised hierarchically across the PICO
 question; and (2) conduct a pilot screening 10 abstracts together with the first
 reviewer (EHG) in a training phase.⁹
- During abstract screening, (3) GPs will be monitored continuously as the agreement rates at individual level are visible in the Rayyan screening software by the first reviewer; this allows determining if any GP with high levels of disagreement (agreement rates < 75%) may require booster training;⁹ (4) the text mining function in Rayyan screening tool will successfully assist GP reviewers to identify the relevant studies early in the process.¹⁰
- After screening ends, (5) a second experienced reviewer will independently be acting as a resolver to screen all records that received discordant assessments between the first reviewer and the crowd of GPs.¹¹ Following this partial replication of citation-screening task, the remaining disagreements between the first and second reviewers will be resolved by consensus. The same approach will be applied in the full-text article screening.

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Table S1 Environmental exposure categories within home built environments.

Category	Terms for environmental attributes ¹							
	Bioaerosols	Aerosols	Carbon monoxide					
	Volatile organic compounds	Particulate matter	Nitrogen dioxide					
	Fungi	Fine/ulftrafine particles	Nitrous acid					
Air quality	Mold	Formaldehyde	Sulfur dioxide					
All quality	Ventilation	Polycyclic aromatic hydrocarbons	Ozone					
	Air-conditioning	Phthalic acids						
	Air filtration	High-efficiency particulat air	e					
Thermal comfort	Temperature	Thermal insulation	warmth					
	Heating							
	Natural lighting	Artificial lighting	Illumination					
	Daylight	Night lighting	Illuminance					
Non-ionising	Sunlight	Light at night	Photoperiod					
radiation	Electromagnetic fields	Microwaves	VLF-EMFs					
	Electric / Magnetic field	Radiofrequency	ELF-EMFs					
	Electricity							
Noise	Sound	Acoustic	Noise					
INOISE	Soundscape							
	Green	Natural	Houseplant					
Nature	Greenspace	Garden	Vegetation					
	Blue space	Plants						
\\/	Drinking water	Water filtration	Water supply					
Water	Water quality	Water purification						

VLF, very low frequency; ELF, extremely low frequency; EMFs, electromagnetic fields.

¹ A full list of variants terminology, synonyms, alternative topics, or spelling variations for each term is provided by requesting to the corresponding author. These variants are shown in the search strategy (Table S4).

Table S2

Potential home built environment interventions for improving or modifying quality of household environmental exposure.

Environmental attribute	Description of some home-built environment interventions ^{1, 2}
Warmth, indoor air and energy efficiency improvements	Heating installation/repair, electric storage heater, thermal insulation, reproofing, window replacement, provision of ventilation, air filter installations, delivery of portable ventilation devices, use of low volatile organic compound products and materials, provision of elements/devices for air ionisation, improvements for avoiding moisture exposure.
Artificial lighting and sunlighting ³	Replacement of luminaries, space layout and re-configurations of surface area of window or internal space, terrace, patios. Home-based bright light therapy with devices will be included if these are integrated into domestic luminaries or other built environment elements.
Non-ionising radiation ³	Installing protective panels, unplugging devices not in use to reduce field strengths in the home, increasing the distance or changing phase and neutral line (system upgrade), relocation from apartment building with new electric system.
Noise, soundscape	Installing soundproof windows/panels, acoustic insulation in party walls or building facades to block external noises, ventilation device replacement to mask unwanted sounds, supplying phono-acoustic material, indoor technological systems to release wanted soundscape.
Nature	Provision and improvement (e.g., accessibility and/or visibility) of natural elements to individual/community private gardens, incorporating greater number of houseplants, spatial re-design to provide domestic greenspaces, supplying devices to recreate digitally nature views.
Water	Installation of sustainable water systems, home treatment devices for production of water with ionisation characteristics and specific potential of hydrogen (pH).

¹The proposed home built environment interventions do not pretend to represent a comprehensive list, but to serve as potential examples.

²Those studies where residents were rehoused or received housing refurbishment will be included if enough data about the specific home built environment intervention(s) are available.

³Although the artificial and natural lighting belong to the non-ionising radiation group, the former is dealt with separately because of the large number of research studies examining the health effects from these environmental attributes.

Table S3
Cancer-associated systemic inflammation biomarkers

			Circulating individual inflammator	y marker	S ^{1, 2}
Group	$[ID_i-ID_f]$	ID	Single marker	ID	Single marker
G1-10	Inflammato	ry med	iators		
G1	[1-99]	Cyto	kines		
	[1-41]	Inter	leukins		
	[1-9]	Inter	leukin-1 family		
		1	Interleukin-1 alpha	6	Interleukin-36 gamma
		2	Interleukin-1 beta	7	Interleukin-36 alpha
		3	Interleukin-1Ra	8	Interleukin-37
		4	Interleukin-18	9	Interleukin-38
		5	Interleukin-33		
	[10-15]		leukin-2 family		
		10	Interleukin-2	13	Interleukin-9
		11	Interleukin-4	14	Interleukin-15
	[40.04]	12	Interleukin-7	15	Interleukin-21
	[16-24]		leukin-6 family		
		16	Interleukin-6	21	Cardiotrophin-like cytokine factor 1
		17	Interleukin-11	22	Ciliary neurotrophic factor
		18	Leukemia inhibitory factor	23	Neuropoietin
		19	Oncostatin	24	Interleukin-31
	[05 04]	20	Cardiotrophin-1		
	[25-31]		leukin-10 family		lataria di 20
		25 26	Interleukin-10 Interleukin-19	29 30	Interleukin-24 Interleukin-26
		27	Interleukin-19 Interleukin-20	31	Interleukin-29
		28	Interleukin-22	31	IIIteriedKiii-29
	[32-35]		leukin-12 family		
	[02-00]	32	Interleukin-12	34	Interleukin-27
		33	Interleukin-23	35	Interleukin-35
	[36]	36	Interleukin-13		
	[37, 38]	37	Interleukin-17	38	Interleukin-25 (/-17E)
	[39, 40]	39	Interleukin-32	40	Interleukin-32 gamma
	[41]	41	Interleukin-34		<u> </u>
	[42-46]	Colo	ony-stimulating factor		
		42	Granulocyte colony stimulating factor	45	Multipotential colony stimulating
		43	Granulocyte-macrophage colony		factor
		. •	stimulating factor		Interleukin-3
		44	Macrophage colony stimulating	46	Erythropoietin
			factor		Hematopoietin
	[47-50]	Adip	okines		
	-	47	Adiponectin	49	Resistin
		48	Leptin	50	Visfatin
	[51-86]		mokines		
			family		
	[51-58]		Receptors		
		51	CC Receptor 1	55	CC Receptor 6
		52	CC Receptor 2	56	CC Receptor 7
		53	CC Receptor 4	57	CC Receptor 9
	[50 70]	54	CC Receptor 5	58	CC Receptor 10
	[59-73]		Ligand		001:140
		59	CC Ligand 2	66 67	CC Ligand 19
		60	CC Ligand 4	67	CC Ligand 20
		61 62	CC Ligand 5	68 60	CC Ligand 22
		62	CC Ligand 5 RANTES	69 70	CC Ligand 22 CC Ligand 24
		63	CC Ligand 7	70 71	CC Ligand 24 CC Ligand 25
		00	CO Ligaria /	, ,	OO Ligaria 20

		64 CC	Ligand 8	72	CC Ligand 27
			Ligand 18	73	CC Ligand 28
		CXC family			_
	[74-79]	CXC Rece	,		
	[•]		C Receptor 1	77	CXC Receptor 4
			C Receptor 2	78	CXC Receptor 6
			C Receptor 3	79	CXC Receptor 7
	[80-92]	CXC Ligar			
	[00 02]		C Ligand 1	87	CXC Ligand 9
			C Ligand 2	88	CXC Ligand 10
			C Ligand 3	89	CXC Ligand 11
			C Ligand 4	90	CXC Ligand 12
		84 CXC	C Ligand 5	91	CXC Ligand 14
			C Ligand 6	92	CXC Ligand 16
			rleukin-8	02	ono Ligaria 10
			C Ligand 8		
	[93-95]	Interferons			
	[50 50]		rferon gamma	95	Interferon alpha
			rferon beta	50	interior diplia
	[96-98]		rosis factor		
	[96]		nor necrosis factor alpha		
	[97, 98]		crosis factor receptor	00	Tumor pocresis factor recentor 2
	[00]		nor necrosis factor receptor 1	98	Tumor necrosis factor receptor 2
	[99]		crophage migration inhibitory		
	[400,400]	facto			
G2	[100-129]	Growth fac			
	[100]		nsforming growth factor-beta		
	[101-103]		ndothelial growth factor		
		••••••	cular endothelial growth factor		
	[102,104]		ndothelial growth factor receptor	······ - ········	
			cular endothelial growth factor	104	Vascular endothelial growth factor
			eptor 1		receptor 3
			cular endothelial growth factor		
			eptor 2		
	[105]		elet-derived growth factor		
	[106-108]		growth factor		
			oblast growth factor 1	108	Fibroblast growth factor receptor
			ic fibroblast growth factor		
	[109, 110]		growth factor		
		109 Epic			
			dermal growth factor	110	Epidermal growth factor receptor
	[111]		dermal growth factor cental growth factor	110	Epidermal growth factor receptor
	[111] [112]	111 Plac		110	Epidermal growth factor receptor
	[112] _[113]	111 Plac 112 Hep	cental growth factor	110	Epidermal growth factor receptor
	[112]	111 Plac 112 Hep	cental growth factor atocyte growth factor	110	Epidermal growth factor receptor
	[112] _[113]	111 Plac 112 Hep 113 Ner	cental growth factor atocyte growth factor ve growth factor	110	Epidermal growth factor receptor Insulin-like growth factor
	[112] _[113]	111 Plac 112 Hep 113 Ner Insulin	cental growth factor latocyte growth factor live growth factor		
	[112] [113] [114, 115]	111 Place 112 Hep 113 Nerr Insulin 114 Insu	cental growth factor latocyte growth factor live growth factor		
	[112] [113] [114, 115]	111 Plac 112 Hep 113 Ner Insulin 114 Insu Endothelin 116 End	cental growth factor atocyte growth factor we growth factor allin	115	Insulin-like growth factor
	[112] [113] [114, 115] [116-118]	111 Place 112 Hep 113 Nerr Insulin 114 Insu Endothelin 116 End 117 End	cental growth factor vatocyte growth factor ve growth factor llin s othelin 1 othelin 2	115	Insulin-like growth factor
	[112] [113] [114, 115]	111 Place 112 Hep 113 Ner Insulin 114 Insu Endothelin 116 End 117 End Renin-ang	cental growth factor latocyte growth factor lines southlin 1 othelin 2 iotensin system	115	Insulin-like growth factor Endothelin 3
	[112] [113] [114, 115] [116-118] [119, 120]	111 Place 112 Hep 113 Ner Insulin 114 Insu Endothelin 116 End 117 End Renin-ang 119 Ang	cental growth factor vatocyte growth factor ve growth factor llin s othelin 1 othelin 2 iotensin system iotensin 1	115	Insulin-like growth factor
	[112] [113] [114, 115] [116-118]	111 Plac 112 Hep 113 Ner Insulin 114 Insu Endothelin 116 End 117 End Renin-ang 119 Ang	cental growth factor latocyte growth factor line line line lise lothelin 1 lothelin 2 liotensin system liotensin 1 line	115 118 120	Insulin-like growth factor Endothelin 3 Angiotensin 2
	[112] [113] [114, 115] [116-118] [119, 120] [121, 122]	111 Plac 112 Hep 113 Ner Insulin 114 Insu Endothelin 116 End 117 End Renin-ang 119 Ang Angiopoiet 121 Ang	cental growth factor vatocyte growth factor ve growth factor velin s othelin 1 othelin 2 iotensin system iotensin 1 iins iopoietin 1	115	Insulin-like growth factor Endothelin 3
	[112] [113] [114, 115] [116-118] [119, 120]	111 Plac 112 Hep 113 Ner Insulin 114 Insu Endothelin 116 End 117 End Renin-ang 119 Ang Angiopoiet 121 Ang Angiopoiet	cental growth factor catocyte growth factor we growth factor llin ss othelin 1 othelin 2 iotensin system iotensin 1 iins iopoietin 1 iinslike protein family	115 118 120 122	Insulin-like growth factor Endothelin 3 Angiotensin 2 Angiopoietin 2
	[112] [113] [114, 115] [116-118] [119, 120] [121, 122]	111 Place 112 Hep 113 Ner Insulin 114 Insu Endothelin 116 End 117 End Renin-ang 119 Ang Angiopoiet 121 Ang Angiopoiet 123 Ang	cental growth factor latocyte growth factor we growth factor llin lis othelin 1 othelin 2 iotensin system iotensin 1 lins iopoietin 1 lin-like protein family iopoietin-like protein 1	115 118 120	Insulin-like growth factor Endothelin 3 Angiotensin 2 Angiopoietin 2 Angiopoietin related growth factor
	[112] [113] [114, 115] [116-118] [119, 120] [121, 122]	111 Plac 112 Hep 113 Ner Insulin 114 Insu Endothelin 116 End 117 End Renin-ang 119 Ang Angiopoiet 121 Ang Angiopoiet 123 Ang 124 Ang	cental growth factor vatocyte growth factor ve growth factor llin s othelin 1 othelin 2 iotensin system iotensin 1 iins iopoietin 1 iin-like protein family iopoietin-like protein 1 iopoietin-like protein 2	115 118 120 122 128	Insulin-like growth factor Endothelin 3 Angiotensin 2 Angiopoietin 2 Angiopoietin related growth factor Angiopoietin-like protein 6
	[112] [113] [114, 115] [116-118] [119, 120] [121, 122]	111 Plac 112 Hep 113 Ner Insulin 114 Insu Endothelin 116 End 117 End Renin-ang 119 Ang Angiopoiet 121 Ang Angiopoiet 123 Ang 124 Ang 125 Ang	cental growth factor latocyte growth factor we growth factor llin s othelin 1 othelin 2 iotensin system iotensin 1 iins iopoietin 1 iin-like protein family iopoietin-like protein 1 iopoietin-like protein 2 iopoietin-like protein 3	115 118 120 122	Insulin-like growth factor Endothelin 3 Angiotensin 2 Angiopoietin 2 Angiopoietin related growth factor Angiopoietin-like protein 6 Cornea-derived transcript 6
	[112] [113] [114, 115] [116-118] [119, 120] [121, 122]	111 Plac 112 Hep 113 Ner Insulin 114 Insu Endothelin 116 End 117 End Renin-ang 119 Ang Angiopoiet 121 Ang Angiopoiet 123 Ang 124 Ang 125 Ang 126 Ang	cental growth factor ratocyte growth factor we growth factor llin s othelin 1 othelin 2 iotensin system iotensin 1 cins iopoietin 1 cin-like protein family iopoietin-like protein 1 iopoietin-like protein 2 iopoietin-like protein 3 iopoietin-like protein 4	115 118 120 122 128	Insulin-like growth factor Endothelin 3 Angiotensin 2 Angiopoietin 2 Angiopoietin related growth factor Angiopoietin-like protein 6
G2	[112] [113] [114, 115] [116-118] [119, 120] [121, 122] [123-129]	111 Plac 112 Hep 113 Ner Insulin 114 Insu Endothelin 116 End 117 End Renin-ang 119 Ang Angiopoiet 121 Ang Angiopoiet 123 Ang 124 Ang 125 Ang 126 Ang 127 Ang	cental growth factor latocyte growth factor line growth factor lline s othelin 1 othelin 2 iotensin system iotensin 1 lins iopoietin 1 lin-like protein family iopoietin-like protein 1 iopoietin-like protein 2 iopoietin-like protein 3 iopoietin-like protein 4 iopoietin-like protein 5	115 118 120 122 128	Insulin-like growth factor Endothelin 3 Angiotensin 2 Angiopoietin 2 Angiopoietin related growth factor Angiopoietin-like protein 6 Cornea-derived transcript 6
G 3	[112] [113] [114, 115] [116-118] [119, 120] [121, 122] [123-129]	111 Place 112 Hep 113 Ner Insulin 114 Insulin 116 End 117 End Renin-ang 119 Ang Angiopoiet 121 Ang Angiopoiet 123 Ang 124 Ang 125 Ang 126 Ang 127 Ang Transcripti	cental growth factor latocyte growth factor we growth factor llin lin lis othelin 1 othelin 2 liotensin system liotensin 1 lins liopoletin 1 lin-like protein family liopoletin-like protein 1 liopoletin-like protein 2 liopoletin-like protein 3 liopoletin-like protein 4 liopoletin-like protein 5 on factor family	115 118 120 122 128	Insulin-like growth factor Endothelin 3 Angiotensin 2 Angiopoietin 2 Angiopoietin related growth factor Angiopoietin-like protein 6 Cornea-derived transcript 6
G3	[112] [113] [114, 115] [116-118] [119, 120] [121, 122] [123-129]	111 Place 112 Hep 113 Ner Insulin 114 Insu Endothelin 116 End 117 End Renin-ang 119 Ang Angiopoiet 121 Ang Angiopoiet 123 Ang 124 Ang 125 Ang 126 Ang 127 Ang Transcripti 130 Nuc	cental growth factor latocyte growth factor line growth factor lline s othelin 1 othelin 2 iotensin system iotensin 1 lins iopoietin 1 lin-like protein family iopoietin-like protein 1 iopoietin-like protein 2 iopoietin-like protein 3 iopoietin-like protein 4 iopoietin-like protein 5	115 118 120 122 128	Insulin-like growth factor Endothelin 3 Angiotensin 2 Angiopoietin 2 Angiopoietin related growth factor Angiopoietin-like protein 6 Cornea-derived transcript 6

		factor 1	r erythroid-2 related	132	Nuclear factor erythroid-2 related factor 2
	[133-138]		and activators of transci		
			ed- Signal transducer of transcription 1	136	phosphorylated- Signal transducer and activator of transcription 4
			ed- Signal transducer of transcription 2	137	phosphorylated- Signal transducer and activator of transcription 5
			ed- Signal transducer of transcription 3	138	phosphorylated- Signal transducer and activator of transcription 6
	[139-141]	Hypoxia-inducible fa	actors		transonption o
			cible factor-1 alpha	141	Hypoxia-inducible factor-2α
		140 Hypoxia-indu	cible factor-1β		
G4	[142-152]	Immunoglobulins. G			
	[142-148]	Cell-adhesion mole			
		142 Intercellular of molecules		145	Epithelial cell-adhesion molecules
			adhesion molecules thelial cell-adhesion	146	Neural cell-adhesion molecules
	[147-149]	Selectin family			
		147 E-selectin		148	P-selectin
			eukocyte adhesion	149	L-selectin
		molecule 1	1.4. 2.1. 11		
		Leukocyte-er			
	[450 450]	adhesion mo			
	[150-152] [150]	Programmed cell de 150 Programmed			
		Programmed cell de	cell death protein 1		
	[151, 152]		cell death protein	152	Programmed cell death protein
		ligand 1	con acam protein	102	ligand 2
G5	[153-165]	Eicosanoids			
	[153-159]	Cyclooxygenase pa	thwavs		
	[153, 154]	Cyclooxygenases			
	. , .	153 Cyclooxygen	ase 1 n-endoperoxide	154	Cyclooxygenase 2 Prostaglandin-endoperoxide synthase 2
	[155-158]	Prostaglandins			
		155 Prostaglandir	n D2	150	
		156 Prostaglandir	n E2	158	Prostaglandin I2 Prostacyclin
	[450]	156 Prostaglandir157 Prostaglandir	n E2	158	
	[159]	156 Prostaglandir157 ProstaglandirTromboxanes	n E2 n F2 alpha	158	
		156 Prostaglandir 157 Prostaglandir Tromboxanes 159 Tromboxane	n E2 n F2 alpha A2	156	
	[159]	156 Prostaglandir 157 Prostaglandir Tromboxanes 159 Tromboxane Lipoxygenase pathy	n E2 n F2 alpha A2	156	
	[160-165]	156 Prostaglandir 157 Prostaglandir Tromboxanes 159 Tromboxane Lipoxygenase pathy Lipoxygenases	n E2 n F2 alpha A2 vays	136	
	[160-165] [160]	156 Prostaglandin 157 Prostaglandin Tromboxanes 159 Tromboxane Lipoxygenase pathy Lipoxygenases 160 5-lipoxygena	n E2 n F2 alpha A2 vays	136	
	[160-165]	156 Prostaglandin 157 Prostaglandin Tromboxanes 159 Tromboxane Lipoxygenase patho Lipoxygenases 160 5-lipoxygenase Leukotrienes	n E2 n F2 alpha A2 ways		Prostacyclin
	[160-165] [160]	156 Prostaglandin 157 Prostaglandin Tromboxanes 159 Tromboxane Lipoxygenase pathv Lipoxygenases 160 5-lipoxygena Leukotrienes 161 Leukotriene	n E2 n F2 alpha A2 ways se	163	Prostacyclin Leukotriene C4
	[160-165] [160] [161-164]	156 Prostaglandin 157 Prostaglandin 157 Prostaglandin 159 Tromboxane 159 Tromboxane Lipoxygenase patho Lipoxygenases 160 5-lipoxygena Leukotrienes 161 Leukotriene 6 162 Leukotriene 6	n E2 n F2 alpha A2 ways se		Prostacyclin
	[160-165] [160]	156 Prostaglandin 157 Prostaglandin Tromboxanes 159 Tromboxane Lipoxygenase pathy Lipoxygenases 160 5-lipoxygenas Leukotrienes 161 Leukotriene Marches Lipoxines	n E2 n F2 alpha A2 ways se	163	Prostacyclin Leukotriene C4
G6	[160-165] [160] [161-164] [165]	156 Prostaglandin 157 Prostaglandin 157 Prostaglandin 159 Tromboxane Lipoxygenase patho Lipoxygenases 160 5-lipoxygenase Leukotrienes 161 Leukotriene March 162 Leukotriene E Lipoxines 165 Lipoxine A4	n E2 n F2 alpha A2 vays se	163	Prostacyclin Leukotriene C4
G6	[160-165] [160] [161-164] [165] [166-209]	156 Prostaglandin 157 Prostaglandin 157 Prostaglandin 159 Tromboxane 159 Tromboxane Lipoxygenase patho Lipoxygenases 160 5-lipoxygena Leukotrienes 161 Leukotriene R 162 Leukotriene R Lipoxines 165 Lipoxine A4 Acute phase protein	n E2 n F2 alpha A2 vays se	163	Prostacyclin Leukotriene C4
G 6	[160-165] [160] [161-164] [165] [166-209] [166-205]	156 Prostaglandin 157 Prostaglandin 157 Prostaglandin 159 Tromboxanes 159 Tromboxane Lipoxygenase patho Lipoxygenases 160 5-lipoxygena Leukotrienes 161 Leukotriene R 162 Leukotriene R Lipoxines 165 Lipoxine A4 Acute phase protein Up-regulation	n E2 n F2 alpha A2 vays se	163	Prostacyclin Leukotriene C4
G6	[160-165] [160] [161-164] [165] [166-209] [166-205] [166-172]	156 Prostaglandin 157 Prostaglandin 157 Prostaglandin 159 Tromboxanes 159 Tromboxane Lipoxygenase patho Lipoxygenases 160 5-lipoxygena Leukotrienes 161 Leukotriene E 162 Leukotriene E Lipoxines 165 Lipoxine A4 Acute phase protein Up-regulation Pentraxins family	n E2 n F2 alpha A2 vays se	163	Prostacyclin Leukotriene C4
G6	[160-165] [160] [161-164] [165] [166-209] [166-205]	156 Prostaglandin 157 Prostaglandin 157 Prostaglandin 159 Tromboxanes 159 Tromboxane Lipoxygenase patho Lipoxygenases 160 5-lipoxygena Leukotrienes 161 Leukotriene R 162 Leukotriene R Lipoxines 165 Lipoxine A4 Acute phase protein Up-regulation	n E2 n F2 alpha A2 ways see A4 34	163	Prostacyclin Leukotriene C4

	[169-172]		raxins		
	[170 171]	169	Pentraxin-3		
	[170,171]	170	onal pentraxin Neuronal pentraxin 1	171	Neuronal pentraxin 2
	[172]	172	Serum amyloid P	171	Neuronal pentraxiii 2
	[173-175]		m amyloid A		
	[170 170]	173	Serum amyloid A1	175	Serum amyloid A3
		174	Serum amyloid A2		Beta globulin
	[176-178]		a 1 globulins		3
	[176,177]		a 1acid-glycoprotein		
		176	Orosomucoid 1	177	Orosomucoid 2
			Alpha 1acid-glycoprotein 1		Alpha 1acid-glycoprotein 2
	[178]	178	Alpha 1 antitrypsin		
	[179-184]		cellular matrix proteins		
		179	Fibronectin	183	Vitronectin
		180	Ferritin (L-ferrint, H-ferritin)	184	Lipopolysaccharide binding
		404	11.6		protein
		181	H-ferritin		
		182	Osteopontin		
			Early T lymphocyte activation gene	•	
	[185-195]	Coar	gulation and fibrinolytic system		
	[185-187]	185	Fibrinogen	187	Thrombin
	[100 107]	186	D-dimer	101	
	[188-195]		minogen activation system		
		188	Plasmin	192	Plasminogen activator inhibitory-1
		189	Plasminogen	193	Plasminogen activator inhibitory-2
		190	Urokinase-plasminogen activator	194	Neuroserpin
		191	Urokinase-plasminogen activator	195	Tissue-type plasminogen activator
			receptor		
	[196-197]		oglobulins		
		196	Beta 2 microglobulin	197	Alpha 1 microglobulin
	[198-201]		sport proteins		
		198	Ceruloplasmin	200	Hemopexin
		199	Haptoglobin	201	Alpha 2 macroglobulin
	[000 005]	0	Alpha 2 globulin		Alpha 2 globulin
	[202-205]		plement system plement 3		
	[202, 203]		Complement 3a	203	Complement 2h
	[204,205]		plement 5	203	Complement 3b
	[204,205]	204	Complement 5a	205	Complement 5b-9
	[206-209]		n-regulation	200	Complement 30-8
	[206, 207]	206	Pre-albumin	207	Albumin
	[200, 207]	200	Transthyretin	201	AIRMITIIT
	[208, 209]	Trans	sferrin		
	[200, 200]	208	Transferrin receptor 1	209	Transferrin receptor 2
			Beta globulin		Beta globulin
G7	[210-225]	Matri	x metalloproteinases		<u> </u>
		210	Matrix metalloprotenase 1	218	Matrix metalloprotenase 12
		211	Matrix metalloprotenase 2	219	Matrix metalloprotenase 13
		212	Matrix metalloprotenase 3	220	Matrix metalloprotenase 14
		213	Matrix metalloprotenase 7	221	Matrix metalloprotenase 16
		214	Matrix metalloprotenase 8	222	Matrix metalloprotenase 17
		215	Matrix metalloprotenase 9	223	Matrix metalloprotenase 19
		216	Matrix metalloprotenase 10	224	Matrix metalloprotenase 26
		217	Matrix metalloprotenase 11	225	Matrix metalloprotenase 28
G8	[226-235]		ox active mediators		
	[226,227]		lloproteins		
		226	Hemoglobin	227	Cell-free heme
			Cell-free hemoglobin		
	[228-233]	Vitan	nin D		

		228	25-hydroxyvitamin D	231	hydroxycholecalciferol
		229	25-hydroxyvitamin D3 1-alpha- hydroxylase	232	Calcidiol, 25-hydroxyvitamin D ₃
		230	25-hydroxyvitamin D3 1-α- hydroxylase	233	Calcitriol, 1,25-dihydroxyvitamin D3
	[234-235]	Mela	tonin		
	-	234	Melatonin	235	6-sulfatoxymelatonin
G9	[236-242]	Lipor	oroteins		
		236	Very low density lipoprotein	240	Apolipoprotein
		237	Low density lipoprotein	241	Total cholesterol
		238	High density lipoprotein	242	Triglycerides
		239	Oxidized low density lipoprotein		
G10	[243-252]	Adre	nal cortex hormones and neurotransr	nitters	
			ocorticoids		
		243	Cortisone	244	Cortisol
		245	Corticotropin-releasing hormone	248	Serotonin
		246	Adrenocorticotropic hormone	249	5-hydroxytryptamine
		247	Gamma aminobutyric acid		
			cholamines		
		250	Epinephrine, adrenaline	252	Dopamine
		251	Norepinephrine		
G11-13	[253-264]	Inflar	mmatory effector cells		
G11	[253]		elets parameters		
	•	253	Platelets		
			Platelet count		
G12	[254-257]	Eryth	nrocytes parameters		
		254	Red blood cell count	256	Erythrocyte sedimentation rate*
		255	Red blood cell distribution width*	257	Plasma viscosity
G13	[258-264]	Leuk	ocytes		•
	[258-260]	258	White blood cell count	260	Neutrophil count
		259	Monocyte count Macrophages		
	[261-264]	261	Total lymphocyte count	263	Regulatory T lymphocytes
		262	Absolute lymphocyte count	264	Circulating plasma cells
		Cor	mbining multiple inflammatory marker	s (into a	score) ^{1, 2}
Group	[ID _i -ID _f]	ID	Multiple markers		[calculated]
-	-		-	ID	Combined single and/or multi-
					markers
cG11.13	[265-271]				
	[=== =: :]	Leuk	ocytes-platelets parameters		
	[265-267]		sical scores		
					[259:261]
		Class	sical scores	259	Monocyte count
		Class 265	sical scores Lymphocyte-to-monocyte ratio	259 261	Monocyte count Total lymphocyte count
		Class	sical scores	261	Monocyte count Total lymphocyte count [260:261]
		Class 265	sical scores Lymphocyte-to-monocyte ratio	261 260	Monocyte count Total lymphocyte count [260:261] Neutrophil count
		265 266	sical scores Lymphocyte-to-monocyte ratio Neutrophil-to-lymphocyte ratio	261	Monocyte count Total lymphocyte count [260:261] Neutrophil count Total lymphocyte count
		Class 265	sical scores Lymphocyte-to-monocyte ratio	261 260 261	Monocyte count Total lymphocyte count [260:261] Neutrophil count Total lymphocyte count [253:261]
		265 266	sical scores Lymphocyte-to-monocyte ratio Neutrophil-to-lymphocyte ratio	261 260 261 253	Monocyte count Total lymphocyte count [260:261] Neutrophil count Total lymphocyte count [253:261] Platelet count
	[265-267]	Class 265 266 267	Lymphocyte-to-monocyte ratio Neutrophil-to-lymphocyte ratio Platelet-to-lymphocyte ratio	261 260 261	Monocyte count Total lymphocyte count [260:261] Neutrophil count Total lymphocyte count [253:261]
		265 266 267 Nove	Lymphocyte-to-monocyte ratio Neutrophil-to-lymphocyte ratio Platelet-to-lymphocyte ratio el combining scores ³	261 260 261 253	Monocyte count Total lymphocyte count [260:261] Neutrophil count Total lymphocyte count [253:261] Platelet count Total lymphocyte count
	[265-267]	Class 265 266 267	Lymphocyte-to-monocyte ratio Neutrophil-to-lymphocyte ratio Platelet-to-lymphocyte ratio el combining scores ³ Derived neutrophil-to-lymphocyte	261 260 261 253 261	Monocyte count Total lymphocyte count [260:261] Neutrophil count Total lymphocyte count [253:261] Platelet count Total lymphocyte count [260:(258-262)]
	[265-267]	265 266 267 Nove	Lymphocyte-to-monocyte ratio Neutrophil-to-lymphocyte ratio Platelet-to-lymphocyte ratio el combining scores ³	261 260 261 253 261 258	Monocyte count Total lymphocyte count [260:261] Neutrophil count Total lymphocyte count [253:261] Platelet count Total lymphocyte count [260:(258-262)] White blood cell count
	[265-267]	265 266 267 Nove	Lymphocyte-to-monocyte ratio Neutrophil-to-lymphocyte ratio Platelet-to-lymphocyte ratio el combining scores ³ Derived neutrophil-to-lymphocyte	261 260 261 253 261 258 260	Monocyte count Total lymphocyte count [260:261] Neutrophil count Total lymphocyte count [253:261] Platelet count Total lymphocyte count [260:(258-262)] White blood cell count Neutrophil count
	[265-267]	265 266 267 Nove	Lymphocyte-to-monocyte ratio Neutrophil-to-lymphocyte ratio Platelet-to-lymphocyte ratio el combining scores ³ Derived neutrophil-to-lymphocyte ratio	261 260 261 253 261 258	Monocyte count Total lymphocyte count [260:261] Neutrophil count Total lymphocyte count [253:261] Platelet count Total lymphocyte count [260:(258-262)] White blood cell count Neutrophil count Absolute lymphocyte count
	[265-267]	265 266 267 Nove	Lymphocyte-to-monocyte ratio Neutrophil-to-lymphocyte ratio Platelet-to-lymphocyte ratio Platelet-to-lymphocyte ratio combining scores ³ Derived neutrophil-to-lymphocyte ratio Systemic immune-inflammation	261 260 261 253 261 258 260 262	Monocyte count Total lymphocyte count [260:261] Neutrophil count Total lymphocyte count [253:261] Platelet count Total lymphocyte count [260:(258-262)] White blood cell count Neutrophil count Absolute lymphocyte count [253x260:261]
	[265-267]	265 266 267 Nove	Lymphocyte-to-monocyte ratio Neutrophil-to-lymphocyte ratio Platelet-to-lymphocyte ratio el combining scores ³ Derived neutrophil-to-lymphocyte ratio	261 260 261 253 261 258 260	Monocyte count Total lymphocyte count [260:261] Neutrophil count Total lymphocyte count [253:261] Platelet count Total lymphocyte count [260:(258-262)] White blood cell count Neutrophil count Absolute lymphocyte count

				261	Total lymphocyte count
		270	Combination of platelet count and		[253; (260:261)]
			neutrophil-to-lymphocyte ratio	253	Platelet count
				260	Neutrophil count and
				261	Total lymphocyte count
		271	Combined neutrophil/platelet/		[tumor differentiation score x
			lymphocyte / differentiation score		(260:261)-(253:261) score)]
				253	Platelet count
				260	Neutrophil count
				261	Total lymphocyte count
cG6	[272-279]		e phase proteins, combinations		
	[272-275]	Class	sical scores		
		272	Glasgow prognostic score		[165;206]
				165	C-reactive protein
				206	Albumin
		273	Modified glasgow prognostic score		[165;206]
				165	C-reactive protein
				206	Albumin
		274	Glasgow prognostic score:modified		[165;206]
			glasgow prognostic score	165	C-reactive protein
			5 5 1 5	206	Albumin
		275	Prognostic inflammatory and		[165x(175,176)x206x205]
		•	nutritional index	165	C-reactive protein
				175	Alpha 1 acid-glycoprotein
				176	Alpha 1 acid-glycoprotein
				205	Transthyretin
				206	Albumin
	[276-279]	Nove	l combined scoring system ³		
		276	C-reactive protein-to-albumin ratio		[165:206]
		2.0	o reactive protein to albamin ratio	165	C-reactive protein
				206	Albumin
		277	Pre-albumin-to-C-reactive protein		[205:165]
		211	prognostic score	165	C-reactive protein
			progressio score	205	Pre-albumin
		278	Albumin-to-globulin ratio		[206: <i>globulins</i> (total proteins-206)]
		210	Albaniin-to-globalin ratio	206	Albumin
				200	Globulins. <i>Alpha 1, alpha 2, beta</i>
					and gamma globulins
		279	Inflammation based index		[(253x260):261]
		219	illiailillation based ilidex	252	
				253	Platelet count
				260	Neutrophil count
				261	Total lymphocyte count
cG11.6	[280-284]	Lauk	ocytes-acute phase proteins, combina	tions ³	
7011.0	[200-204]	280	c-reactive protein and neutrophil-	110113	[165;(260:261)]
		200	lymphocyte ratio score	165	C-reactive protein
			iyiripilooyte tallo soole		
				260	Neutrophil count
				260 261	Neutrophil count
		7 21	Dariyad nautrophil lymphosyto	260 261	Total lymphocyte count
		281	Derived neutrophil-lymphocyte	261	Total lymphocyte count [260:(258-262);206+0.005x261]
		281	ratio and prognostic nutritional	261 206	Total lymphocyte count [260:(258-262);206+0.005x261] Albumin
		281		261 206 258	Total lymphocyte count [260:(258-262);206+0.005x261] Albumin White blood cell count
		281	ratio and prognostic nutritional	261 206 258 260	Total lymphocyte count [260:(258-262);206+0.005x261] Albumin White blood cell count Neutrophil count
		281	ratio and prognostic nutritional	261 206 258 260 261	Total lymphocyte count [260:(258-262);206+0.005x261] Albumin White blood cell count Neutrophil count Total lymphocyte count
			ratio and prognostic nutritional index	261 206 258 260	Total lymphocyte count [260:(258-262);206+0.005x261] Albumin White blood cell count Neutrophil count Total lymphocyte count Absolute lymphocyte count
		281	ratio and prognostic nutritional index Combination of neutrophil-	261 206 258 260 261 262	Total lymphocyte count [260:(258-262);206+0.005x261] Albumin White blood cell count Neutrophil count Total lymphocyte count Absolute lymphocyte count [(260:261); 165, 206]
			ratio and prognostic nutritional index Combination of neutrophillymphocyte ratio and glasgow	261 206 258 260 261 262	Total lymphocyte count [260:(258-262);206+0.005x261] Albumin White blood cell count Neutrophil count Total lymphocyte count Absolute lymphocyte count [(260:261); 165, 206] C-reactive protein
			ratio and prognostic nutritional index Combination of neutrophil-	261 206 258 260 261 262 165 206	Total lymphocyte count [260:(258-262);206+0.005x261] Albumin White blood cell count Neutrophil count Total lymphocyte count Absolute lymphocyte count [(260:261); 165, 206] C-reactive protein Albumin
			ratio and prognostic nutritional index Combination of neutrophillymphocyte ratio and glasgow	261 206 258 260 261 262 165 206 260	Total lymphocyte count [260:(258-262);206+0.005x261] Albumin White blood cell count Neutrophil count Total lymphocyte count Absolute lymphocyte count [(260:261); 165, 206] C-reactive protein Albumin Neutrophil count
		282	ratio and prognostic nutritional index Combination of neutrophillymphocyte ratio and glasgow prognostic score	261 206 258 260 261 262 165 206	Total lymphocyte count [260:(258-262);206+0.005x261] Albumin White blood cell count Neutrophil count Total lymphocyte count Absolute lymphocyte count [(260:261); 165, 206] C-reactive protein Albumin Neutrophil count Total lymphocyte count
			ratio and prognostic nutritional index Combination of neutrophillymphocyte ratio and glasgow prognostic score Combination of fibrinogen and	261 206 258 260 261 262 165 206 260 261	Total lymphocyte count [260:(258-262);206+0.005x261] Albumin White blood cell count Neutrophil count Total lymphocyte count Absolute lymphocyte count [(260:261); 165, 206] C-reactive protein Albumin Neutrophil count Total lymphocyte count [184; (260:261)]
		282	ratio and prognostic nutritional index Combination of neutrophillymphocyte ratio and glasgow prognostic score	261 206 258 260 261 262 165 206 260	Total lymphocyte count [260:(258-262);206+0.005x261] Albumin White blood cell count Neutrophil count Total lymphocyte count Absolute lymphocyte count [(260:261); 165, 206] C-reactive protein Albumin Neutrophil count Total lymphocyte count

				261	Total lymphocyte count
		284	Inflammation-based cumulative		[165;206;(259:261);(260:261);
			prognostic score system		(253:258);184]
				165	C-reactive protein
				184	Fibrinogen
				206	Albumin
				253	Platelet count
				259	Monocyte count
				260	Neutrophil count
				261	Total lymphocyte count
cG11.6.8	[285]	Leuk	ocytes-acute phase proteins-redox ac	tive med	diators, combinations ³
		285	Combined hemoglobin, albumin,		[206x225x253x261]
			lymphocyte, platelet	206	Albumin
				225	Hemoglobin
				253	Platelet count
				261	Total lymphocyte count
cG9.6	[286]	Lipop	protein particle-derived measure of ins	sulin res	istance
		286	Lipoprotein insulin resistance score		[L235sz+L235p+237sz+237p+S2 36sz+S236p]
				235	Very low-density lipoprotein
				236	Low-density lipoprotein
				237	High-density lipoprotein
		287	AC score		[165;240]
		_0,	7.0 000.0	240	Apolipoprotein A-1
				165	C-reactive protein

G, group; cG, combined group; ID, identifier; L, large; P, particle concentration; S, small; Sz, particle size. Cursive, globulin group to which the marker belongs.
*Method of measuring the marker, clinical test.

¹The proposed panel of 287 cancer-associated inflammatory biomarkers could be modified and upgraded over time in

accordance with clinical efficacy tested and promising clinical results of novel candidates.

² Novel combined systemic inflammation-based scoring systems proposed in further research will be incorporated into the panel.

Table S4

Search strategy for Pubmed-Medline

Key:

mh = Medical Subject Heading (MeSH) terms
mh:noexp = MeSH terms:no explode
tiab = title/abstract
* = truncation

Search	Query
Home b	uilt environment
#1	((((((((((((((((((((((((((((((((((((((
Environ	mental parameters
#2	((((((air pollution, indoor[mh]) OR (air pollutants[mh])) OR (air pollut*[tiab])) OR (air
	quality[tiab])) OR (air contamin*[tiab])) OR (aerosols[mh])) OR (aerosol*[tiab])
#3	((particulate matter[mh]) OR (particulate matter[tiab])) OR (particle*[tiab])
#4	((((((((((((((((((((((((((((((((((((((
#5	((((((((((((((((((((((((((((((((((((((
#6	((((((((((((((((((((((((((((((((((((((
#7	((((((((((((((((((((((((((((((((((((((
#8	(((((((temperature[mh]) OR (temperature*[tiab])) OR (thermal comfort[tiab])) OR (thermal condition*[tiab])) OR (thermal insulation[tiab])) OR (warmth[tiab])) OR (heat*[tiab])
#9	((((((((((((((((((((((((((((((((((((((
#10	((((((((((((((((((((((((((((((((((((
#11	((((acoustics[mh])) OR (acoustic*[tiab])) OR (sound[mh])) OR (sound*[tiab])) OR (noise[tiab])
#12	(((((((((nature[tiab])) OR (natural[tiab])) OR (green*[tiab])) OR (greenspace*[tiab])) OR ("blue space*"[tiab])) OR (garden*[tiab])) OR (plant*[tiab])) OR (houseplant*[tiab])) OR (vegetation[tiab])
#13	(((water[mh]) OR (water[tiab])) OR (drinking water[mh])) OR (water quality[mh])
#14	#2 OR #3 OR #4 OR #5 OR #6 OR#13
	ers of systemic inflammatory response
#15	(((((((biomarker*[tiab])) OR (marker*[tiab])) OR (indicator*[tiab])) OR (mediator*[tiab])) OR (profile*[tiab])) OR (score*[tiab])) OR (index*[tiab])) AND (((((inflammat*[tiab])) OR (immun*[tiab])) OR (lipid*[tiab])) OR (metabolic[tiab])) OR (endocrine[tiab])) OR (biomarkers[mh:noexp])) OR (inflammation[mh:noexp])

#16	((((((((cytokines[mh]) OR (cytokine*[tiab]) OR (interleukin*[tiab])) OR (leukemia inhibitory factor[tiab])) OR (oncostatin[tiab])) OR (cardiotrophin[tiab])) OR (ciliary neurotrophic factor[tiab])) OR (colony stimulating factor*[tiab])) OR (erythropoietin[tiab])) OR (hematopoietin[tiab])
#17	((((((((((((((((((((((((((((((((((((((
#18	(((((((chemokine*[tiab])) OR (receptors, chemokine[mh])) OR (interferon*[tiab])) OR (tumor necrosis factor*[tiab])) OR (macrophage migration-inhibitory factor*[tiab])) OR (glycosylation-inhibiting factor[tiab])
#19	((((((((((((((((((((((((((((((((((((((
#20	((((insulin[mh]) OR (insulin[tiab])) OR (endothelins[mh]) OR (endothelin*[tiab]) OR (angiotensins[mh])) OR (angiotensin*[tiab])) OR (angiopoietin*[tiab])
#21	(((((((NF-kappa B[mh])) OR (NF-kappa B[tiab])) OR (nuclear factor kappa B[tiab])) OR (nuclear factor erythroid[tiab])) OR (STAT transcription factors[mh])) OR (STAT[tiab]) OR(hypoxia-inducible factor[mh])) OR(hypoxia-inducible factor[tiab])
#22	((((((((serum globulins[MeSH Terms]) OR (globulin*[Title/Abstract])) OR (immunoglobulin*[Title/Abstract])) OR (cell adhesion molecules[MeSH Terms:noexp])) OR (cell adhesion molecule*[Title/Abstract])) OR (selectins[MeSH Terms])) (selectin*[Title/Abstract])) OR (endothelial-leukocyte adhesion molecule[Title/Abstract])) OR (programmed cell death protein*[Title/Abstract])
#23	((((((((((((((((((((((((((((((((((((((
#24	((((((((((((((((((((((((((((((((((((((
#25	(((((((((((((((((((((((((((((((((((((
#26	((((((lipoproteins[mh]) OR (lipoprotein*[tiab])) OR (apolipoprotein*[tiab])) OR (cholesterol[tiab])) OR (triglycerides[mh])) OR (triglyceride*[tiab])
#27	((((((((((((((((((((((((((((((((((((((
#28	((((((((((((((((((((((((((((((((((((((

	width[tiab])) OR (blood sedimentation[tiab])) OR (erythrocyte sedimentation rate[tiab])) OR(hemorheology[mh])) OR (blood viscosity[tiab])) OR (plasma viscosity[tiab])
#29	((((((((((((((((((((((((((((((((((((((
#30	#15 OR #16 OR #17 OR #18 OR #19 OR#29
#31	#1 AND #14 AND #30
#32	#31 NOT (animals[mh] NOT humans[mh])

Table S5

Variables collected in pre-established data extraction template.

Construct	Variables
(A) Study information	First author's last name, publication year
. ,	Study design
	Journal name
	 Geographical region in which the study was conducted
	Duration of follow-up or study period
	Aims and/or purpose of the study
(B) participant, patient	Sample size
characteristics	Gender composition
	Mean or median age
	Ethnicity
	Socio-economic status (if available)
(C) clinicopathological	Comorbidities: medical classification topography code,
characteristics (if	collection method, measure of association 1
available)	Baseline clinical parameters
	Body mass index
	Prescribed medications
(D) Home built	Device(s), technology and/or architecture element(s)
environment	information
intervention	 Technical specifications
	Duration of treatment period
	Timing, delivery
	Co-interventions
	Economic information
(E) improved/modified	Parameter type within categories (i.e., pollutants)
environmental factor	• Sources
	Exposure levels
	Method(s) of measurement
	Equipment
	 Measurements: dose, duration, intensity, frequency
(F) Housing	Study setting
characteristics and	Housing typology
design (if reported)	Residence type
	Building age
	Material specification
	Spatial characteristics
	Engineering systems
(G) Inflammatory	Sample types
biomarker data	Laboratory assay methods
	Cut-offs, midpoints to report up- or down-regulation and to
	distinguish marker levels from unresolved to resolving
	outcomes
	 Concentration level for each marker (pre- and post-
	intervention; repeated over intervention period).
(H) Outcomes	Analysis models
	Effect estimates
	Adjusted potential confounders
(I) Plausible	Initiating event(s)
mechanistic pathways ²	Key events (e.g., cellular/tissue/system responses)
	Adverse outcome(s)
1 International Classification	

¹ International Classification of Diseases (ICD-11) to group reported medical conditions. However, adjust of terms to the International Classification of Primary Care, 3rd edition (ICPC-3) into WHO Family of Internationals Classifications (FIC) [https://browser.icpc-3.info/] will be performed to classify the clinicopathological characteristics and patient and clinical data in the domains of General/Family Practice and Primary Care. ² The plausible mechanistic pathways will be organised and structured according to the Adverse Outcome Pathway (AOP) framework.