

Table S1. The search strategy used in different databases.

Medline via OVID	<pre> #1 exp dietary fiber/ #2 dietary fiber*.tw #3 dietary fibre*.tw #4 Roughage*.tw #5 wheat bran*.tw #6. whole grain*.tw #7 alimentary fiber*.tw #8 alimentary fibre*.tw #9 biomarker*.tw #10 biologic marker*.tw #11 bioindicator*.tw #12 biological marker*.tw #13 exp biological marker/ #14 biological indicator*.tw #15 1 or 2 or 3 or 4 or 6 or 7 or 8 #16 9 or 10 or 11 or 12 or 13 or 14 #17 15 and 16 #18 17 not (exp animals/ not humans.sh.) </pre>
EMBASE via OVID	<pre> #1 exp dietary fiber/ #2 dietary fiber*.tw #3 dietary fibre*.tw #4 Roughage*.tw #5 wheat bran*.tw #6. whole grain*.tw #7 alimentary fiber*.tw #8 alimentary fibre*.tw #9 biomarker*.tw #10 biologic marker*.tw #11 bioindicator*.tw #12 biological marker*.tw #13 exp biological marker/ #14 biological indicator*.tw #15 1 or 2 or 3 or 4 or 6 or 7 or 8 #16 9 or 10 or 11 or 12 or 13 or 14 #17 15 and 16 #18 17 not ((exp animal/ or nonhuman/) not exp human/) </pre>
Web of Science	<pre> # 1 TS = ("dietary fiber**" or "dietary fibre**" or roughage* or "whole grain**" or "alimentary fiber**" or "alimentary fibre**" or "wheat bran**") #2 TI = ("Biological marker**" or "Biomarker**" or "Biologic marker**" or "Biological indicator**" or Bioindicator* or "biologic marker**") #3 #2 AND #1 </pre>
The Cochrane database	<pre> #1 "dietary fiber**":ti,ab,kw #2 "dietary fibre":ti,ab,kw #3 "roughage":ti,ab,kw #4 "wheat bran":ti,ab,kw #5 "whole grain":ti,ab,kw #6 alimentary fiber*:ti,ab,kw #7 alimentary fibre:ti,ab,kw #8 MeSH descriptor: [Dietary Fiber] explode all trees #9 #1 or #2 or #3 or #4 or #5 or #6 or #7 or #8 #10 MeSH descriptor: [Biomarkers] this term only </pre>

	<p>#11 biomarker*:ti,ab,kw #12 biologic marker*:ti,ab,kw #13 bioindicator*:ti,ab,kw #14 biological marker*:ti,ab,kw #15 biological indicator*:ti,ab,kw #16 #10 or #11 or #12 or #13 or #14 or #15 #17 #9 and #16</p>
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Table 2. Quality assessment and the assessment of the risk of bias in the included studies.

Author	Study design	Study		Methods			Assessment					Analyses				Fast ing blo od sam ples ?	Typ e of uri ne coll ecti on
		Total Sam ple size	Reported inclusion and exclusion criteria	Reported health status	Well-de scribed interven tion/s et-up	Qua lifie d asse ssor	Bl in de as se ss or	Adjusted for potential confoun ders	Same apparatu s (calibrati on) for all participa nts	Valid questio nnaire for the study	Recall period for grain intake in questio nnaire	Time period assessed by questio nnaire	Blood samples within 24 h since last interventio n day?				
Ampatzoglou, 2015 [36]	RCT X-over non-blinded	33	Y	Y	Y	Y	-	N	Y	-	<3 d	6 w	Y	Y	#		
Biltoft-Jensen, 2016 [37]	RCT X-over non-blinded	750	Y	Y	Y	Y	-	Y	Y	-	<3 d	3 m	N	Y	#		
Landberg, 2008 [38]	RCT X-over non-blinded	30	N	N	Y	-	-	Y	Y	-	1 d	6 w	Y	Y	#		
Landberg, 2009 [39]	RCT X-over n/blinded	16	N	Y	Y	-	N	Y	Y	-	7 d	1 d	Y	Y	24 h		
Linko, 2005 [40]	RCT X-over non-blinded	39	Y	Y	Y	-	-	N	-	-	-	8 w	-	-	#		
Hanhineva, 2014 [62]	RCT X-over non-blinded	12	Y	Y	N	Y	-	N	Y	#	#	#	Y	Y	#		
Magnusdottir, 2013 [41]	RCT parallel non-blinded	158	Y	Y	Y	Y	-	Y	N	-	1 d	18-24 w	-	-	#		
McKeown, 2016 [42]	RCT X-over non-blinded	19	Y	Y	Y	-	N	Y	-	-	3 d	6 d	Y	Y	24 h		
Wu, 2015 [43]	RCT parallel non-blinded	16	Y	Y	Y	Y	-	-	N	Y	-	1 d	<4 w	-	Y	#	
Landberg, 2009 [44]	RCT X-over non-blinded	17	Y	Y	Y	Y	Y	N	Y	Y	-	1 d	6 w	#	Y	#	
Ross, 2012 [45]	RCT parallel non-blinded	266	Y	Y	Y	Y	Y	N	Y	-	Y	1 d	16 w	-	Y	#	
Knudsen, 2014 [46]	Case-Control Nested	900	Y	Y	Y	-	-	Y	Y	Y	-	-	-	-	Y/N	#	
Drake, 2014 [60]	Case-Control Nested	2927	Y	Y	N	Y	-	Y	Y	-	<1 d	1 d	-	N	#		

Meija, 2015 [56]	Case–Control Unmatched	31	Y	Y	Y	Y	-	N	Y	-	<1 d	1 d	Y	Y	12 h
Aubertin- Leheudre, 2008 and 2010 [34,59]	Cohort	56	Y	Y	Y	Y	-	Y	Y	Y	1 d	1 d	Y	Y	72 h
Aubertin- Leheudre, 2010 [52]	Cohort	60	Y	Y	Y	Y	-	Y	Y	Y	1 d	1 d	Y	Y	72 h
Andersson, 2011 [35]	Cohort	72	Y	Y	Y	-	-	N	-	-	1 d	1 d	N	Y/N	#
Linko, 2005 [51]	Cohort	9	N	N	N	N	-	N	-	/	-	1-2 w	y	Y	#
Landberg, 2012 [54]	Cohort	104	Y	N	Y	Y	N	Y	-	-	1 yr	1 yr	#	#	spot
Marklund, 2013 [55]	Cohort	66	Y	N	Y	Y	-	Y	Y	-	<1 d	<1 d	#	#	24 h
Soderholm, 2009 [61]	Cohort	15	Y	Y	Y	-	-	N	Y	#	#	#	Y	Y	#
Ross, 2004 [57]	Cohort	1	N	N	N	Y	-	#	N	Y	#	#	#	#	12 h
Landberg, 2018 [33]	Cohort	40	Y	N	Y	Y	-	Y	Y	Y	<1 d	<1 d	#	#	spot
Wierzbicka, 2017 [32]	Cohort	69	Y	N	Y	-	-	N	-	Y	<1 d	<1 d	Y	-	24 h
Wang, 2017 [63]	Cohort	12	Y	Y	Y	-	-	#	Y	#	#	#	#	#	48 h at 8 interval s
Zhu, 2014 [58]	Cohort	12	Y	Y	Y	-	-	N	Y	#	#	#	#	#	24– 36 h
Wu, 2018 [47]	Cohort	258	Y	N	Y	-	-	Y	-	-	1 yr	1 yr	N	Y	#
Landberg, 2011 [48]	Cross- sectional	360	N	N	N	-	-	Y	Y	-	1 yr	1 yr	Y	N	#

Guyman, 2008 [53]	Cross-sectional	99	Y	Y	Y	Y	-	Y	Y	-	<3 m	<3 m	#	#	12 h
McKeown, 2016 [49]	Cross-sectional	190	Y	Y	Y	-	-	Y	Y	-	<1 d	1 yr	-	-	#
Jansson, 2010 [50]	Cross-sectional	20	N	N	N	-	N	N	Y	-	1 yr	1 yr	-	F*	#
Bondia-Pons, 2013 [64]	Untargeted metabolomic RCT-X-over n/blinded	20	Y	Y	Y	Y	-	-	Y	-	1 d	4 w	#	#	24 h
Johansson- Persson, 2013 [65]	Untargeted metabolomic RCT-X-over n/blinded	25	Y	Y	Y	-	Y	N	Y	-	1 d	5 w	Y	Y	#
Hanhineva, 2015 [66]	Untargeted metabolomic RCT- Parallel n/blinded	106	Y	Y	Y	-	-	-	Y	-	1 d	3 w	y	y	#
Zhu, 2016 [67]	Untargeted metabolomic Cohort	12	Y	Y	Y	-	-	N	Y	#	#	#	#	#	2 × 24- h urin e at 6 tim e poi nts
Coulomb, 2015 [68]	Untargeted metabolomic Cohort	1	N	N	Y	-	-	N	Y	#	#	#	#	#	24 h
Garcia-Aloy, 2014 [69]	Untargeted metabolomic Cross-sectional	155	Y	N	Y	-	-	N	Y	Y	1 m	1 m	#	#	Spo t urin e
Hanhineva, 2015 [70]	Untargeted metabolomic Cross-sectional	66			Y	-	-	N	Y	-	1 d	1 d	#	#	24 h

Note: Abbreviation: (-), not reported; (#), not applicable; d, day; h, hours; RCT, randomized controlled trial; x-over, cross over design; N, no; w, week; Y, yes; yr, year;

