

Wendy Wood

List of Publications by Year in descending order

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Version: 2024-02-01

17
papers

2,470
citations

840776
11
h-index

888059
17
g-index

18
all docs

18
docs citations

18
times ranked

3000
citing authors

#	ARTICLE	IF	CITATIONS
1	How can risk of COVID-19 transmission be minimised in domiciliary care for older people: development, parameterisation and initial results of a simple mathematical model. <i>Epidemiology and Infection</i> , 2022, 150, .	2.1	2
2	Conservative treatment for uncomplicated appendicitis in children: the CONTRACT feasibility study, including feasibility RCT. <i>Health Technology Assessment</i> , 2021, 25, 1-192.	2.8	10
3	CONTRACT Study - CONservative TRreatment of Appendicitis in Children (feasibility): study protocol for a randomised controlled Trial. <i>Trials</i> , 2018, 19, 153.	1.6	27
4	Emollient bath additives for the treatment of childhood eczema (BATHE): multicentre pragmatic parallel group randomised controlled trial of clinical and cost effectiveness. <i>BMJ: British Medical Journal</i> , 2018, 361, k1332.	2.3	50
5	Adding emollient bath additives to standard eczema management for children with eczema: the BATHE RCT. <i>Health Technology Assessment</i> , 2018, 22, 1-116.	2.8	14
6	Timing of surgical intervention for developmental dysplasia of the hip: a randomised controlled trial (Hip â€™Op). <i>Health Technology Assessment</i> , 2017, 21, 1-84.	2.8	6
7	Wilmsâ€™ tumour antigen 1 Immunity via DNA fusion gene vaccination in haematological malignancies by intramuscular injection followed by intramuscular electroporation: a Phase II non-randomised clinical trial (WIN). <i>Efficacy and Mechanism Evaluation</i> , 2016, 3, 1-80.	0.7	7
8	Bath additives for the treatment of childhood eczema (BATHE): protocol for multicentre parallel group randomised trial. <i>BMJ Open</i> , 2015, 5, e009575.	1.9	10
9	A genome-wide association study identifies colorectal cancer susceptibility loci on chromosomes 10p14 and 8q23.3. <i>Nature Genetics</i> , 2008, 40, 623-630.	21.4	514
10	Common genetic variants at the CRAC1 (HMPS) locus on chromosome 15q13.3 influence colorectal cancer risk. <i>Nature Genetics</i> , 2008, 40, 26-28.	21.4	277
11	Refinement of the basis and impact of common 11q23.1 variation to the risk of developing colorectal cancer. <i>Human Molecular Genetics</i> , 2008, 17, 3720-3727.	2.9	61
12	A genome-wide association study shows that common alleles of SMAD7 influence colorectal cancer risk. <i>Nature Genetics</i> , 2007, 39, 1315-1317.	21.4	463
13	A genome-wide association scan of tag SNPs identifies a susceptibility variant for colorectal cancer at 8q24.21. <i>Nature Genetics</i> , 2007, 39, 984-988.	21.4	754
14	Evidence for a colorectal cancer susceptibility locus on chromosome 3q21-q24 from a high-density SNP genome-wide linkage scan. <i>Human Molecular Genetics</i> , 2006, 15, 2903-2910.	2.9	52
15	Evidence of Linkage to Chromosome 9q22.33 in Colorectal Cancer Kindreds from the United Kingdom. <i>Cancer Research</i> , 2006, 66, 5003-5006.	0.9	51
16	Disruption of the Interaction of Mammalian Protein Synthesis Eukaryotic Initiation Factor 4B with the Poly(A)-binding Protein by Caspase- and Viral Protease-mediated Cleavages. <i>Journal of Biological Chemistry</i> , 2001, 276, 23922-23928.	3.4	91
17	Changes in integrity and association of eukaryotic protein synthesis initiation factors during apoptosis. <i>FEBS Journal</i> , 2000, 267, 1083-1091.	0.2	80