Arnaud Lagarde

List of Publications by Year
in descending order

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25 papers	2,199 citations	567281 15 h-index	610901 24 g-index
26 all docs	26 docs citations	26 times ranked	5133 citing authors

#	Article	IF	Citations
1	Gene Expression Classification of Colon Cancer into Molecular Subtypes: Characterization, Validation, and Prognostic Value. PLoS Medicine, 2013, 10, e1001453.	8.4	1,064
2	Mutations of polycombâ€associated gene ASXL1 in myelodysplastic syndromes and chronic myelomonocytic leukaemia. British Journal of Haematology, 2009, 145, 788-800.	2.5	537
3	Genome profiling of pancreatic adenocarcinoma. Genes Chromosomes and Cancer, 2011, 50, 456-465.	2.8	107
4	A Seven-Gene Signature Aggregates a Subgroup of Stage II Colon Cancers with Stage III. OMICS A Journal of Integrative Biology, 2012, 16, 560-565.	2.0	69
5	A polymorphism of EGFR extracellular domain is associated with progression free-survival in metastatic colorectal cancer patients receiving cetuximab-based treatment. BMC Cancer, 2008, 8, 169.	2.6	65
6	NCOA3, a new fusion partner for MOZ/MYST3 in M5 acute myeloid leukemia. Leukemia, 2008, 22, 663-665.	7.2	57
7	Molecular combing reveals complex 4q35 rearrangements in Facioscapulohumeral dystrophy. Human Mutation, 2017, 38, 1432-1441.	2.5	39
8	8q24 Cancer Risk Allele Associated with Major Metastatic Risk in Inflammatory Breast Cancer. PLoS ONE, 2012, 7, e37943.	2.5	34
9	Germline APC mutation spectrum derived from 863 genomic variations identified through a 15-year medical genetics service to French patients with FAP. Journal of Medical Genetics, 2010, 47, 721-722.	3.2	32
10	A Heterozygous ZMPSTE24 Mutation Associated with Severe Metabolic Syndrome, Ectopic Fat Accumulation, and Dilated Cardiomyopathy. Cells, 2016, 5, 21.	4.1	28
11	Vascular Endothelial Growth Factor A c.*237C>T polymorphism is associated with bevacizumab efficacy and related hypertension in metastatic colorectal cancer. Digestive and Liver Disease, 2015, 47, 331-337.	0.9	23
12	High Frequency of Chromosome 14 Deletion in Early-Onset Colon Cancer. Diseases of the Colon and Rectum, 2007, 50, 1881-1886.	1.3	22
13	Improving molecular diagnosis of distal myopathies by targeted next-generation sequencing: TableÂ1. Journal of Neurology, Neurosurgery and Psychiatry, 2016, 87, 340-342.	1.9	20
14	Acromegaly in Carney complex. Pituitary, 2019, 22, 456-466.	2.9	20
15	Expression Profiles in Stage II Colon Cancer According to APC Gene Status. Translational Oncology, 2012, 5, 72-76.	3.7	18
16	Methylation hotspots evidenced by deep sequencing in patients with facioscapulohumeral dystrophy and mosaicism. Neurology: Genetics, 2019, 5, e372.	1.9	16
17	Identification of a ERCC5 c.2333T>C (L778P) Variant in Two Tunisian Siblings With Mild Xeroderma Pigmentosum Phenotype. Frontiers in Genetics, 2019, 10, 111.	2.3	11
18	Germinal defects of SDHx genes in patients with isolated pituitary adenoma. European Journal of Endocrinology, 2020, 183, 369-379.	3.7	11

#	Article	IF	Citations
19	Performance of semiconductor sequencing platform for nonâ€invasive prenatal genetic screening for fetal aneuploidy: results from a multicenter prospective cohort study in a clinical setting. Ultrasound in Obstetrics and Gynecology, 2019, 54, 246-254.	1.7	9
20	Genomic profile concordance between pancreatic cyst fluid and neoplastic tissue. World Journal of Gastroenterology, 2019, 25, 5530-5542.	3.3	7
21	Identification of a CDH12 potential candidate genetic variant for an autosomal dominant form of transgrediens and progrediens palmoplantar keratoderma in a Tunisian family. Journal of Human Genetics, 2020, 65, 397-410.	2.3	4
22	Clinical profile of comorbidity of rare diseases in a Tunisian patient: a case report associating incontinentia pigmenti and Noonan syndrome. BMC Pediatrics, 2018, 18, 286.	1.7	3
23	Case Report: Identification of Novel Variants in ERCC4 and DDB2 Genes in Two Tunisian Patients With Atypical Xeroderma Pigmentosum Phenotype. Frontiers in Genetics, 2021, 12, 650639.	2.3	2
24	Next generation sequencing in non metastatic high grade pediatric osteosarcoma: A useful tool to identify new therapeutic targets.. Journal of Clinical Oncology, 2018, 36, e24235-e24235.	1.6	0
25	Genomic analysis of paired IDHwt glioblastoma (GB) to reveal recurrent alterations of MPDZ at relapse after radiotherapy and temozolomide (RTCT).. Journal of Clinical Oncology, 2019, 37, e13535-e13535.	1.6	0