Arnaud Lagarde

List of Publications by Year in descending order

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25 papers

2,199 citations

567281 15 h-index 24 g-index

26 all docs

26 does citations

times ranked

26

5133 citing authors

#	Article	IF	CITATIONS
1	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
2	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
3	Germinal defects of SDHx genes in patients with isolated pituitary adenoma. European Journal of Endocrinology, 2020, 183, 369-379.	3.7	11
4	Acromegaly in Carney complex. Pituitary, 2019, 22, 456-466.	2.9	20
5	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
6	Methylation hotspots evidenced by deep sequencing in patients with facioscapulohumeral dystrophy and mosaicism. Neurology: Genetics, 2019, 5, e372.	1.9	16
7	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
8	Genomic profile concordance between pancreatic cyst fluid and neoplastic tissue. World Journal of Gastroenterology, 2019, 25, 5530-5542.	3.3	7
9	Genomic analysis of paired <i>IDHwt</i> glioblastoma (GB) to reveal recurrent alterations of <i>MPDZ</i> at relapse after radiotherapy and temozolomide (RTCT) Journal of Clinical Oncology, 2019, 37, e13535-e13535.	1.6	0
10	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
11	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	O
12	Molecular combing reveals complex 4q35 rearrangements in Facioscapulohumeral dystrophy. Human Mutation, 2017, 38, 1432-1441.	2.5	39
13	A Heterozygous ZMPSTE24 Mutation Associated with Severe Metabolic Syndrome, Ectopic Fat Accumulation, and Dilated Cardiomyopathy. Cells, 2016, 5, 21.	4.1	28
14	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
15	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
16	Gene Expression Classification of Colon Cancer into Molecular Subtypes: Characterization, Validation, and Prognostic Value. PLoS Medicine, 2013, 10, e1001453.	8.4	1,064
17	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
18	Expression Profiles in Stage II Colon Cancer According to APC Gene Status. Translational Oncology, 2012, 5, 72-76.	3.7	18

#	Article	IF	CITATION
19	8q24 Cancer Risk Allele Associated with Major Metastatic Risk in Inflammatory Breast Cancer. PLoS ONE, 2012, 7, e37943.	2.5	34
20	Genome profiling of pancreatic adenocarcinoma. Genes Chromosomes and Cancer, 2011, 50, 456-465.	2.8	107
21	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
22	Mutations of polycombâ€essociated gene <i>ASXL1</i> in myelodysplastic syndromes and chronic myelomonocytic leukaemia. British Journal of Haematology, 2009, 145, 788-800.	2.5	537
23	NCOA3, a new fusion partner for MOZ/MYST3 in M5 acute myeloid leukemia. Leukemia, 2008, 22, 663-665.	7.2	57
24	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
25	High Frequency of Chromosome 14 Deletion in Early-Onset Colon Cancer. Diseases of the Colon and Rectum, 2007, 50, 1881-1886.	1.3	22