## Duo Xu

## List of Publications by Year in descending order

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Abstract 235: Identifying potential drug targets using patient-derived, tissue specific, gene regulatory networks., 2021, , .

Abstract 2034: Patient-specific enhancer-gene networks for hundreds of primary cancer samples. , 2021, , .

4 Loss-of-function tolerance of enhancers in the human genome. PLoS Genetics, 2020, 16, el008663.
3.5

12

| 5 | Structural Variants in Ancient Cenomes. Population Genomics, 2018, , 375-391. | 0.5 |
| :--- | :--- | :--- |

6 Archaic Hominin Introgression in Africa Contributes to Functional Salivary MUC7 Genetic Variation. Molecular Biology and Evolution, 2017, 34, 2704-2715.
8.9

57

VCFtoTree: a user-friendly tool to construct locus-specific alignments and phylogenies from
thousands of anthropologically relevant genome sequences. BMC Bioinformatics, 2017, 18, 426.
2.6

14

8
Evolution and protein interactions of AP2 proteins in Brassicaceae: Evidence linking development and environmental responses. Journal of Integrative Plant Biology, 2016, 58, 549-563.

Atopic Dermatitis Susceptibility Variants in Filaggrin<i>Hitchhike</i>Hornerin Selective Sweep.
Genome Biology and Evolution, 2016, 8, 3240-3255.

10 Recent evolution of the salivary mucin MUC7. Scientific Reports, 2016, 6, 31791.
3.3

30

The psoriasis-associated deletion of late cornified envelope genes LCE3B and LCE3C has been
11 maintained under balancing selection since Human Denisovan divergence. BMC Evolutionary Bi
11 maintained under balancing selection since Human Denisovan divergence. BMC Evolutionary Biology,
3.2

35
2016, 16, 265.
Functional Specialization of Human Salivary Glands and Origins of Proteins Intrinsic to Human Saliva. SSRN Electronic Journal, 0, , .

