![A close up of a piece of paper

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Eggert, S., Huyck, K., Somasundaram, P., Kavalla, R., Stewart, E., Lu, A., … Morton, C. (2012). Genome-wide linkage and association analyses implicate FASN in predisposition to uterine leiomyomata. *American Journal of Human Genetics*, 91(4), 621–628. DOI: 10.1016/j.ajhg.2012.08.009

![A screenshot of a cell phone

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Hellwege, J. N., Jeff, J. M., Wise, L. A., Gallagher, C. S., Wellons, M., Hartmann, K. E., … Velez Edwards, D. R. (2017). A multi-stage genome-wide association study of uterine fibroids in African Americans. *Human Genetics*, 136(10), 1363–1373. DOI:10.1007/s00439-017-1836-1

![A picture containing text, map

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Rafnar, T., Gunnarsson, B., Stefansson, O.A., Sulem, P., Ingason, A., Frigge, M.L., … Stefansson, K. (2018). Variants associating with uterine leiomyoma highlight genetic background shared by various cancers and hormone-related traits. *Nature Communications*, 9:3636. DOI:10.1038/s41467-018-05428-6

![A picture containing screenshot

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Cha, P, Takahashi, A., Hosono, N., Low, S., Kamatani, N., Kubo, M., & Nakamura, Y. (2011). A genome-wide association study identifies three loci associated with susceptibility to uterine fibroids. *Nature Genetics*, 43(5).