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It still says in blue- grey above the logo (in the Japanese version), in other countries the logo is normally white. It's being developed by a doujin circle known as Little Meteor, though I'm not sure if it's one of the doujin circles that made the other Muv-Luv fan games. The doujin circle's home page says they've released two products so far. but I'm not sure if those are the products mentioned in that link. They have been making Muv-Luv related fan games for quite some time, but only up to chapter 5. I don't know when they plan to release the final product of the fandisc - if it does ever come out. PS: I have a bit of a soft spot for the franchise because of my favorite episode of the anime is "The Deimos Duel", I guess. PS2: The official Muv-Luv site is currently down, so I'm linking to the fansite. A: They have a game called Little Meteor, it's an online multiplayer tactical shooter. And yes, they are related to MUV-Luv, I've been a player for a month now (I'm Japanese) and I've been playing it for a month. I wouldn't call myself a fan though. I just wanted to play more of it. UPDATE: Ah, here it is, and it looks pretty good. It doesn't seem like there's an english version, but that didn't bother me too much. TAS3: A comprehensive cancer genomics pipeline from DNA to androgen receptor signaling. The androgen receptor (AR) is a well known mediator of the cancer promoting activities of androgens in prostate and other cancers. Until recently, the analysis of AR signaling was largely restricted to 2D or 3D cell culture models, and also a limited number of patient samples. The TREAT-Androgen Receptor 3 (TAS3) pipeline enables the global characterization of AR signaling networks using high-throughput data from TCGA as a training set for a machine-learning algorithm that infers signaling maps from primary patient tumor transcriptome data. Gene expression and chromatin state at the genes encoding established AR target genes were inferred in 5,346 prostate cancer samples from The Cancer Genome Atlas (TCGA) and independently validated in 10,402 prostate cancer samples from the Gen

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