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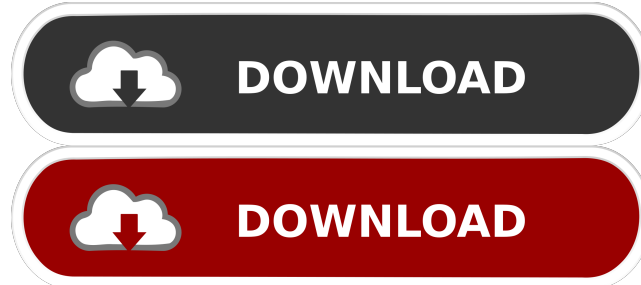
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ovulation-inducing follicles, adeno-hypophyseal and secondary sex-steroid-producing corpus luteum and Leydig cells. All of these sub-sites may be expected to exhibit tissue-specific genomic imprinting, however, the extent to which this occurs remains undetermined. Using a panel of 12 highly polymorphic microsatellite markers, the parental origin of these tissues was determined in 39 F1 hybrid females, 6 of which were progeny of matings between Wistar rats and Fischer 344/Brown Norway rats. The follicle population contained both oocyte and granulosa cells. A strong bias for the maternal origin of both components of the follicle was observed, indicating that, within the follicle, only the oocyte-expressed X-chromosome is involved in genomic imprinting. No imprinting of the Z-chromosome was observed. The corpus luteum contained both oocyte and granulosa cells and also displayed very strong maternal bias in the imprinting of both components, indicating that it also exhibits tissue-specific imprinting. Thus, in the adult female rat, both granulosa and thecal cells display maternal imprinting. In contrast, no parental bias was observed in the imprinting of Leydig cells, indicating that these

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cells do not exhibit tissue-specific imprinting. These data indicate that each of the major sub-sites of the female rat ovary is associated with the same level of tissue-specific imprint

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