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## EST Profile

[Mm.339215](#) - Gm4340: Predicted gene 4340

### Breakdown by Body Sites

[Mm.339215](#)

adipose tissue	0	0 / 1564
adrenal gland	0	0 / 2044
bladder	0	0 / 13461
blood	0	0 / 16832
bone	0	0 / 32327
bone marrow	0	0 / 137491
brain	0	0 / 459991
connective tissue	0	0 / 19790
dorsal root ganglion	0	0 / 10186
embryonic tissue	7	5 / 664846
epididymis	0	0 / 2652
extraembryonic tissue	0	0 / 73200
eye	0	0 / 184900
fertilized ovum	0	0 / 26353
heart	0	0 / 52131
inner ear	0	0 / 37947
intestine	0	0 / 84236
joint	0	0 / 17136
kidney	0	0 / 122453
liver	0	0 / 108957
lung	0	0 / 96442
lymph node	0	0 / 15170
mammary gland	0	0 / 302624
molar	0	0 / 3601
muscle	0	0 / 27256
nasopharynx	0	0 / 8067
olfactory mucosa	0	0 / 3281
ovary	0	0 / 53434
oviduct	0	0 / 3201
pancreas	0	0 / 105756
pineal gland	0	0 / 3914
pituitary gland	0	0 / 16965
prostate	0	0 / 29888
salivary gland	0	0 / 19345
skin	0	0 / 114950
spinal cord	0	0 / 21805
spleen	0	0 / 94430
stomach	0	0 / 29435
sympathetic ganglion	0	0 / 9045
testis	0	0 / 113759
thymus	0	0 / 116168
thyroid	0	0 / 8859
tongue	0	0 / 9861
turbinate	0	0 / 1383

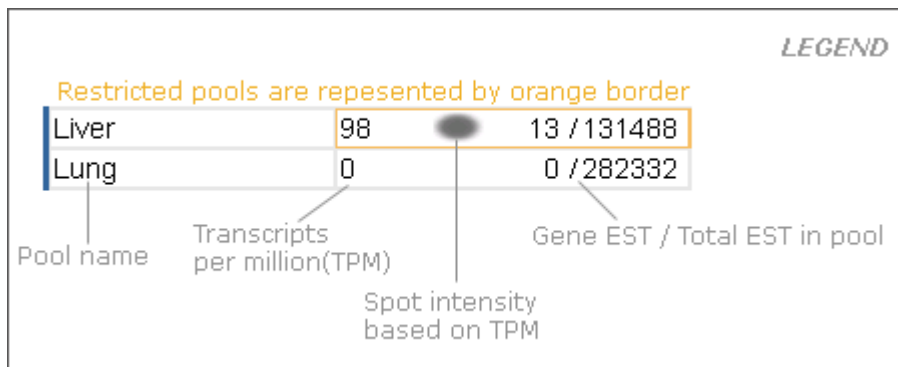
uterus	0	0 / 6842
vagina	0	0 / 5171
vesicular gland	0	0 / 2187

**Breakdown by Developmental Stage**

Mm.339215

oocyte	0	0 / 19423
unfertilized ovum	0	0 / 19710
zygote	0	0 / 27241
cleavage	36	1 / 27056
morula	0	0 / 37647
blastocyst	0	0 / 69305
egg cylinder	0	0 / 12083
gastrula	0	0 / 27624
organogenesis	0	0 / 125372
fetus	0	0 / 653835
neonate	0	0 / 103512
juvenile	0	0 / 291299
adult	0	0 / 995094

EST profiles show **approximate** gene expression patterns as inferred from EST counts and the cDNA library sources (as reported by sequence submitters). Libraries known to be normalized, subtracted, or otherwise biased have been removed, but for a variety of reasons, EST counts may not be a true indication of gene activity.



**Questions or Comments?**

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