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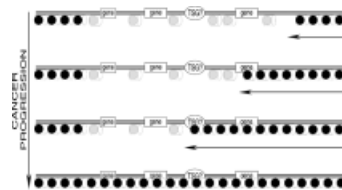
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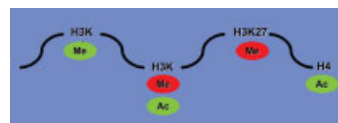
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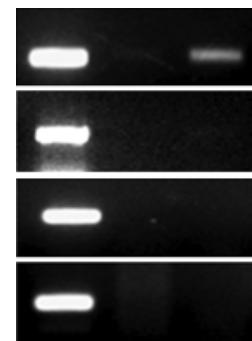
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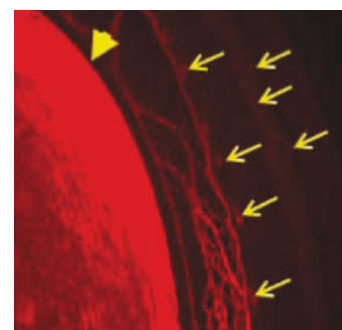


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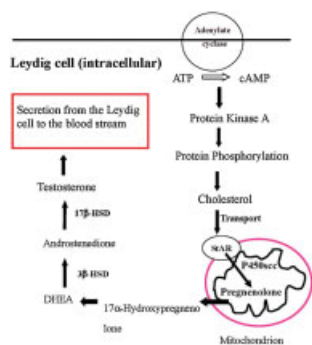
Cover: Cover shows the crystal structure of the *T. thermophilus* 30S ribosomal subunit (PDB accession 1JG0) revealing the relative positions of the Shine-Dalgarno sequence within the 5'-untranslated region of the mRNA (highlighted red), the G693 loop of the 16S rRNA (homologous to the G961 Loop of eukaryotic 18S rRNA, highlighted green), and the anti-Shine-Dalgarno segment of the 16S rRNA (highlighted blue, not present in eukaryotes). See article in this issue by Meng et al, pages 531–544.



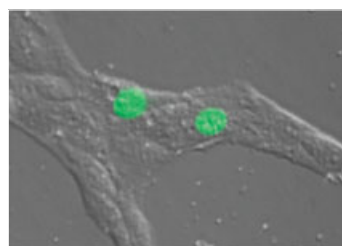
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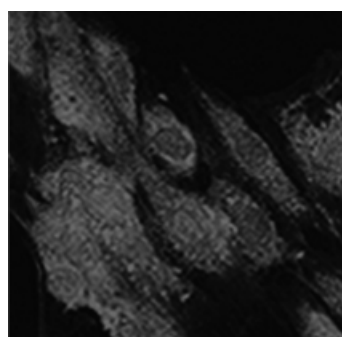
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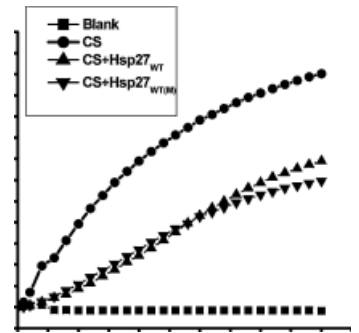
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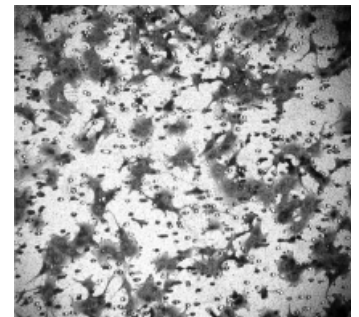
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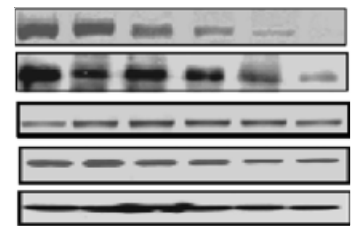
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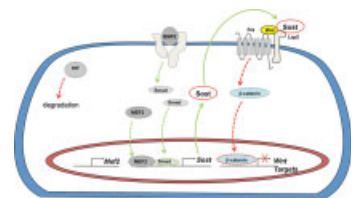
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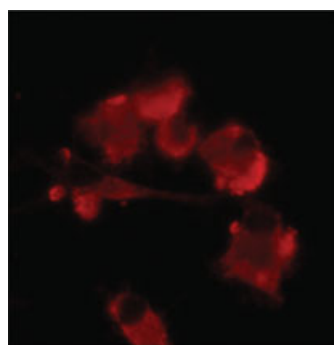


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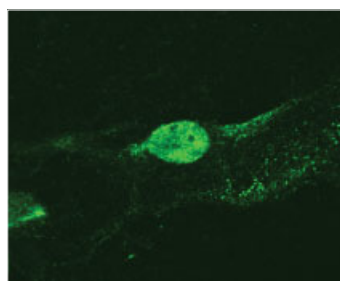
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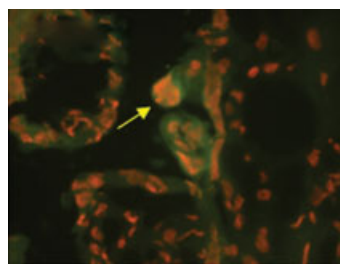
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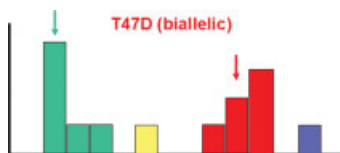
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