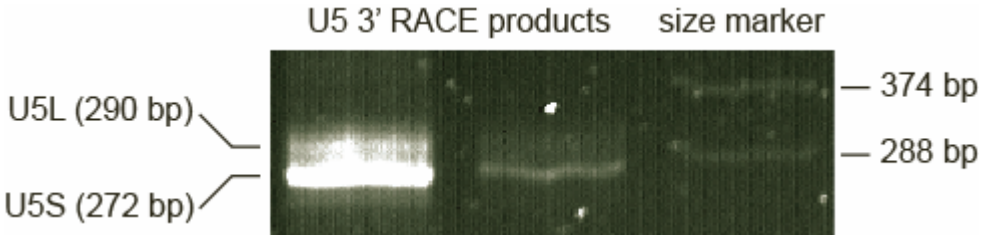

Supplemental Figure 1. Analysis of the short (U5S) and long (U5L) forms of *C. albicans* U5 snRNA. 3' RACE products were resolved on 1X TBE, 3% NuSieveGTG agarose gels (Lonza) and stained with ethidium bromide. Product sizes represent fusions of both the 3' ends of U5 snRNAs and the 5' end of 18S rRNA (see Materials and Methods). The first two lanes contain different loadings of the same sample; the third lane contains unrelated restriction digest products of known size.

Sequence information

I. Sequences of oligonucleotides used for analysis of snRNA 3' ends.

Reverse-transcription of snRNA/18S rRNA fusions

5'-CCACGGTTATCCAAGTAG-3'

Primary PCR

18S rRNA: 5'-TAGCGGCCGCACGGTTATCCAAGTAGTAAGG-3'
U1 snRNA: 5'-TATCTAGAATTTGATGGAGGCTCTGGTG-3'
U2 snRNA: 5'-TATCTAGAACATAGTGTCGTTGCCACTG-3'
U4 snRNA: 5'-GGGCATTACGTGTATCAGTG-3'
U5 snRNA: 5'-CGCCAGAAGATCAATTTCAG-3'
U6 snRNA: 5'-CATTTCCCCTGCATAAGGAC-3'

Nested PCR and sequencing

18S rRNA: 5'-TATCTAGATCCAAGTAGTAAGGTACTATC-3'
U1 snRNA: 5'-TAGCGGCCGCGGAGGCTCTGGTGATGTTC-3'
U2 snRNA: 5'-TAGCGGCCGCTGTCGTTGCCACTGTACTTT-3'
U4 snRNA: 5'-TAGCGGCCGCCGTGTATCAGTGAGGATTCG-3'
U5 snRNA: 5'-TAGCGGCCGCAGAAGATCAATTTCAGTTTACC-3'
U6 snRNA: 5'-TAGCGGCCGCCCTGCATAAGGACGGAACG-3'

II. Yeast snRNA sequences.

With the exception of the *S. cerevisiae* snRNAs and the *C. albicans* snRNA 3' ends, the reported sequences are based on computational rather than experimental analyses, and the precise sequence termini are therefore predictions. In addition to hemiascomycete snRNAs, we include U1 snRNAs from more distantly related yeasts, as their identification required searching by methods other than BLAST (see Results and Discussion). *Y. lipolytica*, *A. nidulans* and *N. crassa* each have more than one gene encoding U1 snRNA; as the U1 genes within these species are highly similar to each other, we report only one representative for each.

Predicted snRNAs of hemiascomycetes listed in Figure 2

>*Saccharomyces_cerevisiae*_U1_snRNA
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>*Candida_glabrata*_U1_snRNA
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>*Saccharomyces_cerevisiae*_U2_snRNA
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ACTTTTTGGAAAATTGTTTCAAGACTTTGTCTTGT
>*Candida_dubliniensis*_U5_snRNA
ATCGCCAGAAGATCAATTTCAATTTACCTACGTACATGGTTCTTGTCTTTTCCCAGAACCATCCGGGAATTGTCTTCATTCAATGGTGTATCAAAATTTT
TACTTTTTGGAAACCTGTTTCAAGACTTTGTCTTGT
>*Candida_tropicalis*_U5_snRNA
ATCGCCAGAAGATCAATTTCGTTCAATCGTACATGGTTCTTGTCTTTTCCCAGAACCATCCGGGAATTGTCTTCATTCAATGGTGTATCAAATTTTAATT
TTTTGGAACTCCCCCTTCATTATGATATAAAAAGGACTTTGTCTTTTTA
>*Lodderomyces_elongisporus*_U5_snRNA
ATCGCCAGGAGATCAATTCCAACGAACATGGTTCTTGCCTTTTCCCAGAACCATCCGGGAATTGTCTCCCGTTAGTCAAATGGTGTATCAATTATCTAAT
TTTTGGAATATTTATCATTTAGTTTTGATT
>*Candida_parapsilosis*_U5_snRNA
CACACCAGAAGATCAATTTCATCAACGAACATGGTTCTTGTCTTTTCCCAGAACCATCCGGGAATTGTCTTCCGTTTTAAGTCAAATGGTGTTTTAATTG
TTTTAATTTTTGGAAACCTTTTTCAACTTTTGAGTTT
>*Clavispora_lusitaniae*_U5_snRNA
ATGACGCCAGAAGATCAGTTTCTTAAGCGTACATGGTTCTTGCCTTTTACCAGAACCATCCGGGAACTGTCTTTGCTAATGGTGTTTTTTTTCATTTTTT
GGAACTCTCCCATGTGCTAAATCATTTATGATTAGCCCATT

>*Pichia_guilliermondii*_U5_snRNA
CTGGCGCCAGAAGATCAATATCTATCGTACATGGTTCTTGCCTTTTACCAGAACCATCCGGGTATTGTCTTTATATGGTGCTCGCTATTTTTTGGAACCT
TCTCATGATGCTTTGCATCAGT
>*Debaryomyces_hansenii*_U5_snRNA
ATCACCAGAAGATCAATTTCAAAATCGTACATGGTTCTTGCCTTTTACCAGAACCATCCGGGAATTGTCTTCACTACTGGTGTTTTCACTATTTTTTGGA
ACCTTTTTCTTTATATTAAGATTTATTCTAGTATAT
>*Yarrowia_lipolytica*_U5_snRNA
CAACAACGGAGGTCGACATCGACGTACATGGCTCTTGCCTTTTACCAGAGCCATCCGGGTGTCGTCTCTACAAGTTGTTTTTCTAGTTTTTGGAAACTTC
TTCTTATGGAGACAAGT

>*Saccharomyces_cerevisiae*_U6_snRNA
GTTCGCGAAGTAACCCTTCGTGGACATTTGGTCAATTTGAAACAATACAGAGATGATCAGCAGTTCCCCTGCATAAGGATGAACCGTTTTACAAAGAGAT
TTATTTCGTTTT
>*Candida_glabrata*_U6_snRNA
GTCCTTGAAGTAACCCTTCTTGGACATTTGGTCAATTTGAAACAATACAGAGATGATCAGCAGTTCCCCTGCATAAGGATGAACCGTTTTACAAAGAGAT
TTATTCGTTTT
>*Kluyveromyces_lactis*_U6_snRNA
GTCCATATTACCCTCCGTGGTTTTATGGACATTTGGTCAATTTGAAACAATACAGAGATGATCAGCAGTTCCCCTGCATAAGGATGAACCGTTTTACAAA
GAGATTTAAGATTTT
>*Kluyveromyces_waltii*_U6_snRNA
GTCCCGCTCATTTCGGTGGGCGTGGACATTTGGTCAATTTGAAACAATACAGAGATGATCAGCAGTTCCCCTGCATAAGGATGAACCGTTTTACAAAGAG
ATTTACAAGTTTT
>*Saccharomyces_kluyveri*_U6_snRNA
GTCCAAGACATTTCGGTGTTTTGGACATTTGGTCAATTTGAAACAATACAGAGATGATCAGCAGTTCCCCTGCATAAGGATGAACCGTTTTACAAAGAGA
TTTACTTCATTTT
>*Candida_albicans*_U6_snRNA
GTTGCGCGTTCGGAACTTGTCAACTTTAAAATTACAGAGAAAATTAGCATTTCCCCTGCATAAGGACGGAACGCTTTACAAAGAGATTTACCATTTTT
>*Candida_dubliniensis*_U6_snRNA
GTTGCGCGTTCGGAACTTGTCAACTTTAAAATTACAGAGAAAATTAGCATTTCCCCTGCATAAGGACGGAACGCTTTACAAAGAGATTTAACATTTTT
>*Candida_tropicalis*_U6_snRNA
GTCGCCGGTTCGGAACTTGTCAAATTTAAAATTACAGAGAAAATTAGCATTTCCCCTGCATAAGGACGGAACGCTTTACAAAGAGATTTACAACATTTTT
>*Lodderomyces_elongisporus*_U6_snRNA
GTACTATCTTCGGATAGTACATTTGTCAATTTTAACAATACAGAGAAGATTAGCATTGTCCTTGCATAAAGATGACACGCTTTACAAAGAGATTTACAGT
TTTT
>*Candida_parapsilosis*_U6_snRNA
GTATTATCTTCGGATAGTACATTTGTCAAATTTAACAATACAGAGAAGATTAGCATTGTCCTTGCATAAAGATGACACGCTTTACAAAGAGATTTACCAT
TTTT
>*Clavispora_lusitaniae*_U6_snRNA
GTACTATTGTTCGCAATAGTACATCTGTCAATTTTAACAATACAGAGAAGATTAGCATTGTCCTTGCATAAAGATGACACGCTATACAAAGAGATTTACC
ATTTTT
>*Pichia_guilliermondii*_U6_snRNA
GTACTATCGTTCGCGATAGTACATTTGTCAATTTTAACAATACAGAGAAGATTAGCATTGTCCTTGCATAAAGATGACACGCTTTACAAAGAGATTTACC
ATTTTT
>*Debaryomyces_hansenii*_U6_snRNA
GTACTATCCTTCTGGATAGTACATTTGTCAATTTTAACAATACAGAGAAGATTAGCATTGTCCTTGCATAAAGATGACACGCTTTACAAAGAGATTTAAC
CATTTTT
>*Yarrowia_lipolytica*_U6_snRNA
GTCCTTCGGGACATATGGTCAATTTGAAAAAATACAGAGAAGATTAGCATGGCCCCTGCATAAGGATGACACGCTTTACAAAGAGATTTAAACCGTTTTT

Other yeast U1 snRNAs

>*Aspergillus_nidulans*_U1_snRNA
ATACTTACCTTGTTTGGGTGGCTGCGATCAAGAAGGCGGCCCACCACAGACTGGTCCCTGCATTGCACAGCGCGGCGGAACGGTCTGTTCAACGGCCTTC
GGGTCATTGACAGGCGTATTCTTTGGCACTTTCC
>*Coccidioides_immitis*_U1_snRNA
ATACTTACCTTGTCTCGGGAACCGCGATCATGAAGGCGGGCTCCCATGAGATGGTCCCTGCATTGCACAGCGCGGCGGACGCGTCTTGTTCATCGGCCTT
CGGGTCTTTGAGAGGCGTAATTTTTGACTCCCTTC
>*Histoplasma_capsulatum*_U1_snRNA
ATACTTACCTTGTTTGGGCATCCGCGATCAAGAAGGCGGTCTGCCGTGGAGGGGCTCCTACATTGCACATCGTGGCGAGCCTTGCCACTCAACATCCTTC
GGGGTTTTGACAGGCATATTCTTTGGCTCCCTTCT
>*Magnaporthe_grisea*_U1_snRNA
ATACTTACCTTGTTGGCCGAGTTCCGTGATCAAGAAGACGGAGCCGGAGAACGGGAGACCTGCATTGCACACCGTGGCGTCGACCGTTCTGGTTTGTCCT
TCGGGGCCAACCCTGGCGCAATTTTTGATAACCCACCTCTTTTC
>*Neurospora_crassa*_U1_snRNA
ATACTTACCTTGGTCGATTGCGCGGAGATCAAGAACACCGCGTAGTATGGCGGAGGGCCTGCATTGCACATCGTGGTGCCAGCCGCCATGAATTGTCCTT
CGGGGCCGTTCCGGCCGTAATTTTTGATGTCAATCC
>*Uncinocarpus_reesii*_U1_snRNA
ATACTTACCTTGTCTCGGGAACCGCGATCATGAAGGCGGGCTCCCATGAGATGGTCCCTGCATTGCACAGCGCGGCGGACTCGTCTTGTTCATCGGCCTT
CGGGTCTTTGAGAGGCGCAATTTTTGACTCTTTACTT