Supplementary Table 1S . Standard gene name tags used in CandidaDB

| Tag | Interpretation |
|-------|------------------------------------------------------------------------------------|
| .5 | On the basis of the homology to other proteins, the protein coding sequence only |
| | corresponds to the 5'-end of the gene and the cognate 3'-end has not been |
| | identified. |
| .3 | The protein coding sequence only corresponds to the 3'-end of the gene and the |
| | cognate 5'-end has not been identified. This tag is frequently assigned to protein |
| | coding sequences where a start codon has not been identified. |
| .5f | The protein coding sequence only corresponds to the 5'-end of the gene and the |
| | corresponding 3'-end is found contiguously on the genome. |
| .3f | The protein coding sequence only corresponds to the 3'-end of the gene and the |
| | corresponding 5'-end is found contiguously on the genome. |
| .5eoc | The protein coding sequence only corresponds to the 5'-end of a gene and is |
| | located at the end of a Contig. |
| .3eoc | The protein coding sequence only corresponds to the 3'-end of a gene and is |
| | located at the end of a Contig. |