

| Location (nt) | AA change | Frequency | Source |
|---------------|-----------|---------------|----------------|
| 252C>T | Silent | 2 samples | this study |
| 678C>T | Pro>Ser | not validated | dbSNP:11548847 |
| 1061A>C | Pro>Gln | not validated | dbSNP:11548850 |
| 1108 to 1122 | Deletion | 2 samples | this study |

Supporting Information Figure S2. SNPs (single nucleotide polymorphisms) in the hnRNP K gene were extracted from Ensembl (www.ensembl.org) and Celera databases. In addition, we cloned and sequenced the hnRNP K full length cDNA from 18 normal volunteers. Only 2 SNPs were reported from the dbSNP and both have not been validated. We have found a novel SNP at nt252 that involves a C>T change, but is non-synonymous. An interesting deletion was observed in 2 different samples in which a 15 base deletion was found just upstream of the KH3 domain (nt1108 to nt1122, nt: ATGATTATTCCTATG). There are a number of SNPs in the UTR (untranslated region) and intronic regions reported in the public databases, but none have been validated (results not shown).